

A BIOLOGICAL CONDITION GRADIENT (BCG) ASSESSMENT MODEL FOR STREAM FISH COMMUNITIES OF CONNECTICUT

FINAL REPORT



Prepared for:

**U.S. EPA Office of Science and Technology
Susan K. Jackson, Work Assignment Manager**

**USEPA Region 1
David McDonald, Work Assignment Manager**

and

**Prepared for CT DEEP Bureau of Water Protection and Land Reuse
Michael Beauchene* and Chris Bellucci**

Prepared by:

**Jennifer Stamp
Jeroen Gerritsen**

**Tetra Tech, Inc.
400 Red Brook Blvd., Suite 200
Owings Mills, MD 21117**

March 31, 2013

*at the initiation of this project affiliated with WPLR Monitoring and Assessment and is currently in the Inland Fisheries Division

EXECUTIVE SUMMARY

Since 2007, Connecticut Department of Energy and Environmental Protection (CT DEEP) Bureau of Water Protection and Land Reuse (WPLR) has been using a macroinvertebrate multi-metric index (MMI) and a Biological Condition Gradient (BCG) assessment model for determination of aquatic life use support (ALUS) assessments in high gradient streams (Gerritsen and Jessup 2007). The BCG supports development of biological criteria in a state's water quality standards that can protect the best quality waters, that can be used as a tool to prevent or remediate cumulative, incremental degradation, and that can help to establish realistic management goals for impaired waters. The basis of the framework is recognition that biological condition of water bodies responds to human-caused disturbance and stress, and that the biological condition can be measured reliably.

The BCG is a universal measurement system or yardstick that is calibrated on a common scale for all states and regions. It is divided into biologically recognizable categories of condition. The BCG is not a management system, nor does it describe management goals. However, biological information as measured by the BCG can tell us if criteria are being met.

WPLR has long recognized the need to obtain a broader perspective of biological integrity through incorporation of fish community assessment data into the biological monitoring process. This document describes the development of BCG assessment models for fish assemblages in freshwater small-cold and medium-large wadeable streams of Connecticut. The BCG model incorporates multiple attribute decision criteria to assign streams to levels of the BCG, and it can be directly applied to designation of multiple aquatic life uses in Connecticut's Criteria and Standards. These fish BCG models will complement Connecticut's existing macroinvertebrate assessment tools (MMI and BCG) along with the recently developed cold and mixed water fish MMIs, and could potentially serve as a starting point for a regional fish BCG model for New England.

ACKNOWLEDGEMENTS

The participants in this effort invested significant time and commitment in the process. We are grateful for their hard work and enthusiasm.

| Organization | Name |
|---|----------------|
| Connecticut Department of Energy and Environmental Protection (CT DEEP), Bureau of Water Protection and Land Reuse (WPLR) | Chris Bellucci |
| | Guy Hoffman |
| CT DEEP, Bureau of Bureau of Natural Resources (BNR) Inland Fisheries Division | Mike Beauchene |
| | Neal Hagstrom |
| | Brian Murphy |
| | Mike Humphreys |
| Silvio O. Conte Anadromous Fish Research Center, U.S. Geological Survey | Yoichiro Kanno |
| Vermont Department of Environmental Conservation (VT DEC) | Rich Langdon |
| | Aaron Moore |
| Maine Department of Environmental Protection (Maine DEP) | Dave Halliwell |
| U.S. EPA Region 1 | Hilary Snook |
| Midwest Biodiversity Institute (MBI) | Chris Yoder |

ACRONYMS

| | |
|---------|---|
| ALUS | Aquatic Life Use Support |
| BCG | Biological Condition Gradient |
| CT | Connecticut |
| CT DEEP | Connecticut Department of Energy and Environmental Protection |
| EPA | United States Environmental Protection Agency |
| MBI | Midwest Biodiversity Institute |
| ME DEP | Maine Department of Environmental Protection |
| MMI | Multi-metric Index |
| NEIWPC | New England Interstate Water Pollution Control Commission |
| OTU | Operational Taxonomic Unit |
| SOP | Standard Operating Procedure |
| TALU | Tiered Aquatic Life Use |
| TNC | The Nature Conservancy |
| VT DEC | Vermont Department of Environmental Conservation |
| WPLR | Bureau of Water Protection and Land Reuse |

TABLE OF CONTENTS

| | |
|--|-----|
| ACKNOWLEDGEMENTS..... | ii |
| ACRONYMS..... | iii |
| | |
| 1 INTRODUCTION..... | 1 |
| 2 METHODS..... | 1 |
| 2.1 Study Design..... | 1 |
| 2.2 Sampling Methods..... | 3 |
| 2.3 BCG Exercise..... | 3 |
| 2.4 Quantitative Description..... | 6 |
| 2.5 Develop Decision Criteria Model..... | 7 |
| 2.6 Development of a BCG Model Using a Decision Criteria Approach..... | 7 |
| 3 COMPREHENSIVE DECISION RULES AND BCG MODEL – SMALL-COLD..... | 9 |
| 3.1. Site Assignments and BCG Level Descriptions..... | 9 |
| 3.2 BCG Attribute Metrics..... | 10 |
| 3.3 BCG Rule Development..... | 16 |
| 3.4 Model Performance..... | 20 |
| 4 COMPREHENSIVE DECISION RULES AND BCG MODEL – MEDIUM-LARGE..... | 21 |
| 4.1 Site Assignments and BCG Level Descriptions..... | 21 |
| 4.2 BCG Attribute Metrics..... | 22 |
| 4.3 BCG Rule Development..... | 27 |
| 4.4 Model Performance..... | 29 |
| 5 MMI PERFORMANCE..... | 30 |
| 6 DISCUSSION..... | 32 |
| 7 LITERATURE CITED..... | 34 |

APPENDIXES

- A Distribution Maps
- B Additional BCG Background Information
- C Fish BCG Attribute Assignments
- D Capture Probability Modeled vs. Disturbance Gradient
- E Sample Worksheet
- F Box plots – all metrics – small-cold samples
- G Small-cold BCG level Assignments
- H Box plots – all metrics – medium-large samples
- I Medium-large BCG Level Assignments

LIST OF TABLES

Table 2-1. Descriptions of the BCG attributes assigned to fish taxa for this exercise..... 5

Table 3-1. BCG decision rules for fish assemblages in small-cold (<6 m²) and medium-large (≥6 m²) streams..... 17

Table 3-2. Summary of differences between model and panelist BCG level assignments for small-cold water samples..... 21

Table 4-1. Summary of differences between model and panelist BCG level assignments for medium-large samples. 30

LIST OF FIGURES

| | |
|---|----|
| Figure 2-1. Relationship between watershed size (mi ²) and % most sensitive (attribute 2) individuals, fit with a logarithmic trend line. | 2 |
| Figure 2-2. The Biological Condition Gradient (BCG) (modified from Davies and Jackson 2006)..... | 4 |
| Figure 2-3. Fuzzy set membership functions assigning linguistic values of Total Taxa to defined quantitative ranges..... | 9 |
| Figure 3-1. Locations of assessed small-cold samples (sites with drainage areas <6 mi ²), coded by panelist BCG level assignment. | 10 |
| Figure 3-2. Box plots of total taxa metric values for small-cold samples, grouped by nominal BCG level (group majority choice). | 12 |
| Figure 3-3. Relationship between total taxa metric values and watershed area for small-cold samples (r=0.53, p<0.01). | 12 |
| Figure 3-4. Box plots for a subset of BCG attribute percent individual metrics for the 54 small-cold samples that were assessed (this includes both calibration and validation samples), grouped by nominal BCG level (group majority choice)..... | 13 |
| Figure 3-5. Box plots for a subset of BCG attribute percent taxa metrics for the 54 small-cold samples that were assessed (this includes both calibration and validation samples), grouped by nominal BCG level (group majority choice)..... | 14 |
| Figure 3-6. Box plots of native brook trout and brown trout percent individual metrics for the 54 small-cold samples that were assessed (this includes both calibration and validation samples), grouped by nominal BCG level (group majority choice)..... | 15 |
| Figure 4-1. Locations of assessed medium-large (drainage areas ≥6 mi ²) sites, coded by panelist BCG level assignment. | 22 |
| Figure 4-2. Box plots of total taxa metric values for medium-large samples, grouped by nominal BCG level (group majority choice). | 24 |
| Figure 4-3. Relationship between total taxa metric values and watershed area for medium-large samples (r=0.37, p<0.01). Samples are coded by nominal BCG level (group majority choice). | 24 |
| Figure 4-4. Box plots for a subset of BCG attribute percent taxa metrics for 70 medium-large samples that were assessed (this includes both calibration and validation samples), grouped by nominal BCG level (group majority choice)..... | 25 |
| Figure 4-5. Box plots for a subset of BCG attribute percent taxa metrics for 70 medium-large samples that were assessed (this includes both calibration and validation samples), grouped by nominal BCG level (group majority choice). | 26 |
| Figure 4-6. Box plots of the percent Centrarchidae individuals and percent Cyprinid taxa metrics for 70 medium-large samples that were assessed (this includes both calibration and validation samples), grouped by nominal BCG level (group majority choice)..... | 27 |
| Figure 5-1. Comparison of panelist BCG level consensus assignments for small-cold samples with cold water MMI scores (top), and BCG level calls on medium-large samples with mixed water MMI scores (bottom). | 31 |

1 INTRODUCTION

The Connecticut Department of Energy and Environmental Protection (CT DEEP) Bureau of Water Protection and Land Reuse (WPLR) routinely samples macroinvertebrates and fish as part of its comprehensive ambient water quality monitoring strategy (CT DEP 2005). Since 2007, WPLR has been using a macroinvertebrate multi-metric index (MMI) and a Biological Condition Gradient (BCG) assessment model for determination of aquatic life use support (ALUS) assessments in high gradient streams (Gerritsen and Jessup 2007). WPLR has long recognized the need to obtain a broader perspective of biological integrity through incorporation of fish community assessment data into the biological monitoring process. The fish assessments would complement the existing macroinvertebrate tools to ultimately produce the most accurate and appropriate aquatic life use support assessments.

There have been recent advancements in fish community assessments in Connecticut. Since 2009, CT DEEP has collected continuous water temperature data from hundreds of sites in Connecticut's inland flowing waters. From these data, CT DEEP has defined 3 major thermal habitat types (cold, cool, warm) and has conducted analyses to identify which fish species are most strongly associated with each habitat type (Beauchene et al. 2012). Patterns of fish distributions in Connecticut's wadeable streams were documented in a publication by Kanno and Vokoun (2008). WPLR fish community data were used to develop cold and mixed water MMIs specific to Connecticut streams (Kanno et al. 2010a) as well as to examine the effects of water withdrawals and impoundments on fish assemblages in southern New England streams (Kanno and Vokoun 2010b). The MMIs are currently being incorporated into WPLR's ALUS. In developing the methods and indexes, WPLR has incorporated the regional New England context, as well as general knowledge on coldwater fish community assessments (e.g., Vermont's Index of Biological Integrity (IBI) (VT DEC 2004); Halliwell et al. 1999; Lyons et al. 2009).

This document describes the development of BCG assessment models for small, cold and medium –large cool wadeable streams of Connecticut. The BCG model incorporates multiple attribute decision criteria to assign streams to levels of the BCG, and it can be directly applied to designation of multiple aquatic life uses in Connecticut's Criteria and Standards, to improve the precision of aquatic life use criteria and assessments. The fish BCG models are intended to support development of fish community structure metrics that will provide a more quantitative approach to WPLR's assessment process. The fish BCG models and MMIs will also supplement Connecticut's macroinvertebrate MMI and BCG model.

2 METHODS

2.1 Study Design

The WPLR fish dataset consisted of 967 samples from 676 unique stations, with sample dates ranging from 1999-2010. The data used passed WPLR's quality assurance procedures. WPLR also provided attribute data for 68 species of fish, along with data on fish size class, water chemistry and land use. Distribution maps for 62 of the fish species are shown in Appendix A.

Samples were initially grouped into temperature subclasses (cold, transitional cool, transitional warm) based on The Nature Conservancy's (TNC) Northeast Aquatic Habitat Classification (Olivero and Anderson 2008). The TNC designations are based on stream size, air temperature, gradient, and groundwater inputs, and are intended to represent natural flowing-water aquatic habitat types across the region. However, the TNC designations were found to be a poor predictor of actual temperature regime in small streams (Mike Beauchene, unpublished data).

Watershed size was deemed to be a more consistent and accurate indicator of temperature class than TNC designations. Stream size exerts a major influence on the longitudinal shift in fish assemblages (Vannote et al., 1980; Kanno and Vokoun, 2008). CT DEEP's continuous temperature sensor data supports the assertion that most of Connecticut's small streams are coldwater, while medium to large-sized streams provide mixed water habitats.

Accordingly, we changed the stream temperature classification to agree with the above findings. We used 6 square miles (=15 km²) as the threshold for separating small from medium-large streams (Kanno et al. 2010a). This threshold also corresponds with a fairly distinct drop in percent sensitive individuals such as brook trout and slimy sculpin (Figure 2-1). There are obviously exceptions to our broad groupings (e.g., naturally occurring small cool, small warm and medium-large coldwater streams in Connecticut), but these occur in low numbers. While not applicable to all streams, the models we have developed should cover the majority of Connecticut streams.

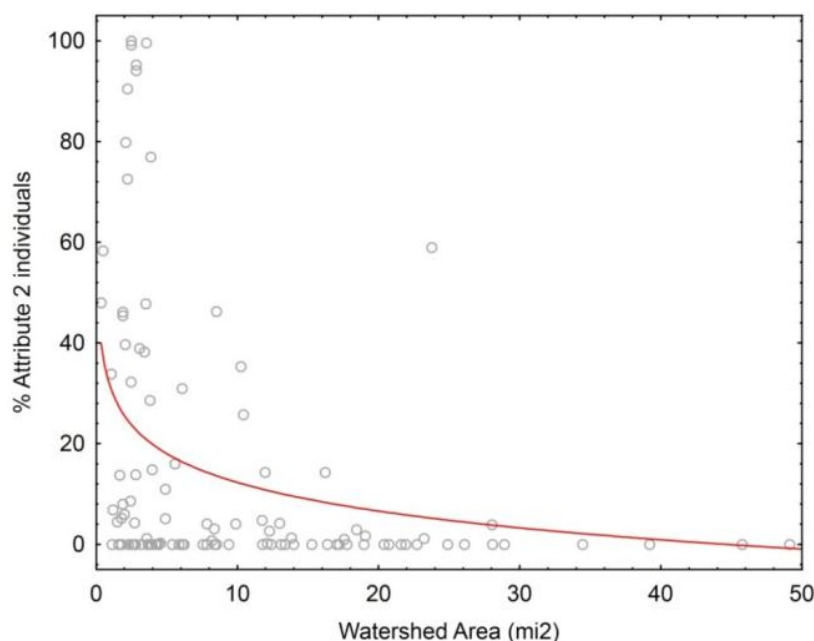


Figure 2-1. Relationship between watershed size (mi²) and % most sensitive (attribute 2) individuals, fit with a logarithmic trend line. In this figure, the scale of the x-axis has been limited to a maximum value of 50.

2.2 Sampling Methods

Samples in the WPLR fish dataset were collected using comparable single pass methods (CT DEEP 2013). Crews sample approximately 100-150 meter reaches or 10-20 mean stream widths. In larger streams, they may sample 200-300 meters. The larger the river, the fewer mean stream widths are typically sampled. The type of gear that a crew uses depends on the stream width. In small streams, crews typically sample with one backpack shocker. In medium-sized streams, they use 2 backpack shockers or 1 tote barge, and in large streams, crews sample with multi-tote barges. All captured individuals are measured to the nearest centimeter and are identified to the species level. Samples are collected during a June-September index period (Kanno et al. 2010a; M. Beauchene and Y Kanno unpublished data).

2.3 BCG Exercise

Biological condition levels and associated attributes are narrative statements on presence, absence, abundance, and relative abundance of several groups of taxa that have been empirically observed to have differing responses to stressors caused by human disturbance, as well as statements on system connectivity and ecosystem attributes (e.g., production, material cycling). The USEPA Tiered Aquatic Life Uses (TALU) national workgroup developed the statements out of consensus best professional judgments (Davies and Jackson 2006). The attributes and transitions between the levels that are described in the BCG model are based on years of biologists' field experience in a given region and reflect accumulated biological knowledge. The current generalized BCG model evolved from a prototype model that was adjusted following a series of exercises, conducted in several different regions of the United States, in which biologists attempted to place actual biomonitoring data into BCG levels (Figure 2-2). Greater detail about the BCG conceptual model may be found in Davies and Jackson (2006).

The BCG is presented as a 6 by 10 matrix of levels and attributes that describe differences in the relative condition of the levels (Appendix B). The attributes are:

- I. Historically documented, sensitive, long-lived or regionally endemic taxa
- II. Highly sensitive taxa, often at low abundance
- III. Sensitive but ubiquitous taxa
- IV. Taxa of intermediate tolerance
- V. Tolerant taxa
- VI. Non-native taxa
- VII. Organism condition
- VIII. Ecosystem functions
- IX. Spatial and temporal extent of detrimental effects
- X. Ecosystem connectivity

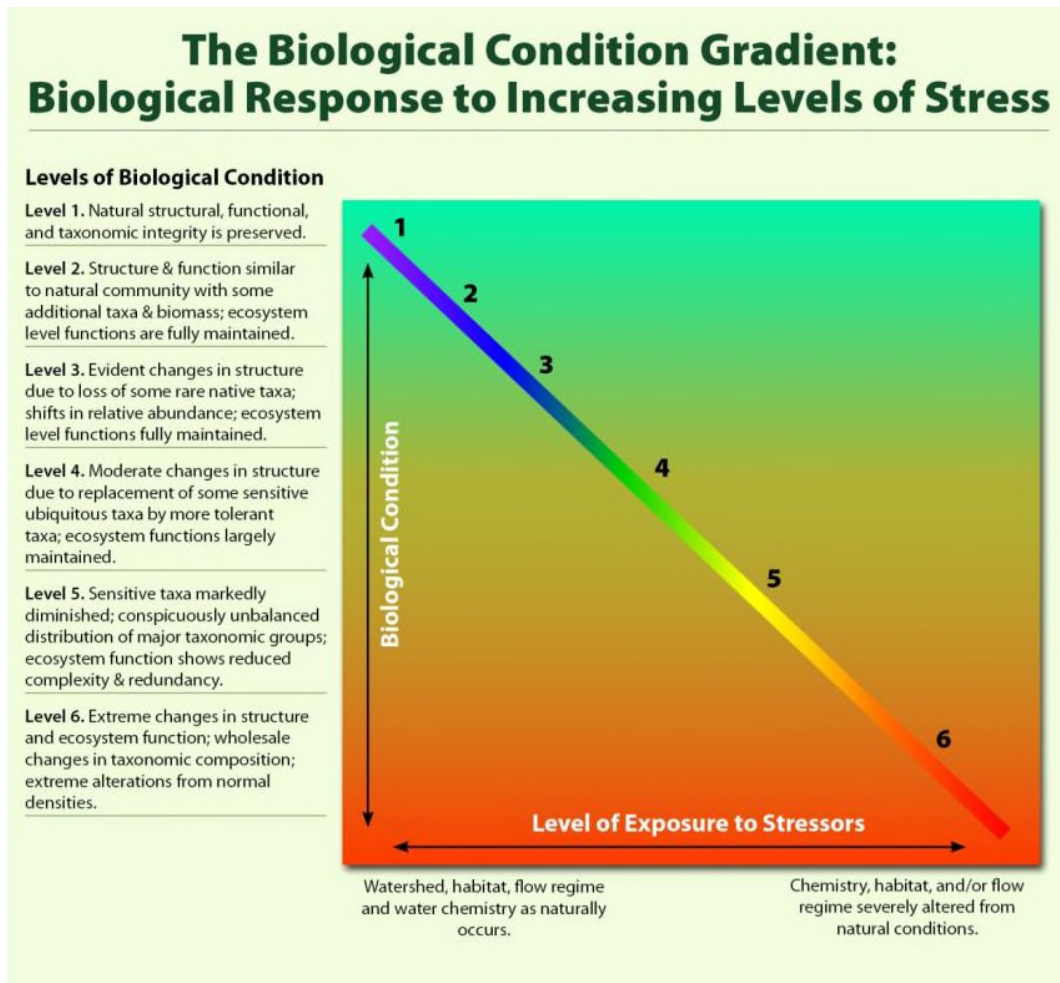


Figure 2-2. The Biological Condition Gradient (BCG) (modified from Davies and Jackson 2006). The BCG was developed to serve as a scientific framework to synthesize expert knowledge with empirical observations and develop testable hypotheses on the response of aquatic biota to increasing levels of stress. It is intended to help support more consistent interpretations of the response of aquatic biota to stressors and to clearly communicate this information to the public, and it is being evaluated and piloted in several regions and states.

The ten attributes presented in the BCG describe multiple aspects of ecological condition, including taxonomic and structural information at the site scale (Attributes I-VI), organism and system performance at the site scale (Attributes VII and VIII), and physical-biotic interactions at broader temporal and spatial scales (Attributes IX and X). Some of the attributes in the BCG represent core data elements that are commonly measured in most state/tribal biological monitoring programs (e.g., Attributes II, III, IV, V, VI, VII) while others, though recognized as very important (e.g., Attributes I, VIII, IX and X), are not commonly measured due to resource limitations and technical complexity.

Development of the BCG is a collective exercise among regional biologists to develop consensus assessments of sites, and then to elicit the rules that the biologists use to assess the sites (Davies and Jackson 2006, US EPA 2007). The goal of this project was to develop a set of decision criteria rules for assigning sites to the BCG levels for small-cold and medium-large wadeable streams based on fish assemblages.

As part of this process, panelists first assigned BCG attributes to fish taxa (attribute assignments can be found in Appendix C). Attribute assignments were initially made during a November 8-9, 2010 workshop in Hartford, CT, and some were later revised following further examination of data and assessments. To help inform the attribute assignments, capture probability was plotted against disturbance gradient (plots can be found in Appendix D).

The panel created new sub-attributes from attribute 6, non-native taxa, to distinguish sensitive intermediate and tolerant non-native taxa (brown trout) and also to identify species that are technically native but have been locally extirpated and exist in some streams only through annual stocking of fry (Atlantic salmon). The sensitive nonnative category included naturalized nonnative salmonids and fry-stocked brown trout and Atlantic salmon. This distinction allowed the assessments to take into account that naturalized nonnative salmonids are highly valued, and are indicators of good water quality, good habitat, and naturally cold or cool temperature.

Table 2-1. Descriptions of the BCG attributes assigned to fish taxa for this exercise.

| BCG | Description |
|-----|---|
| 1 | Historically documented, sensitive, long-lived or regionally endemic taxa |
| 2 | Highly sensitive taxa, often occur in low abundance |
| 3 | Intermediate sensitive taxa |
| 4 | Taxa of intermediate tolerance |
| 5 | Tolerant native taxa |
| 6 | Non-native taxa of intermediate tolerance |
| 6a | Highly tolerant non-native taxa |
| 6b | Sensitive non-native salmonids (=highly valued recreational taxa) |
| 10 | Catadromous fish, indicating ecosystem connectivity |
| x | No attribute assignment (insufficient information) |

Next the panelists examined biological data from individual sites and assigned those samples to levels 1 to 6 of the BCG. The intent was to achieve consensus and to identify rules that panelists were using to make their assignments. Sometimes questions arose regarding the classification of samples (e.g., if an obvious coldwater assemblage was being assessed with the group of coolwater samples). In these situations, we asked CTDEEP personnel to verify (or nullify) the classification based on their knowledge. When there was not an obvious error and when we could not verify or nullify a classification based on site knowledge, panelists made BCG level assignments under the assumption that the site classification was correct.

The panel met in person and per teleconference several times in the period November 2010 to October 2012. During the work, the panelists' thinking evolved on attribute assignments and some of the rules. Accordingly, updates were made to some of the species attribute assignments (as described above), a revised classification scheme was put into place (small-cold and medium-large vs. the original size-TNC temperature subclasses, as described in Section 2.1) and changes were made to the BCG rules table.

In October 2012, the panelists made BCG level assignments on 50 additional samples (25 small-cold and 25 medium-large samples). Some of these were repeat samples, meaning that they had also been assessed during a previous calibration round. If there were discrepancies in the BCG level consensus calls for the repeat samples (e.g., in the first round a sample was assigned to BCG level 3 but in the second round, it was assigned to BCG level 4) we used the consensus call from the second round, since this captured the panelists' most recent thinking, and reflected the evolution in classification scheme, attribute assignments and BCG rules that occurred over the course of the exercise. The repeat samples were included in the calibration dataset. Samples that had not been assessed before were placed in a validation dataset and were used to confirm the models. A total of 124 samples were assessed, of which 94 were included in the calibration dataset, and 30 were placed in a validation dataset and were used to confirm the models.

The data that the experts examined when making BCG level assignments were provided in worksheets. The worksheets contained lists of taxa, taxa abundances, BCG attribute levels assigned to the taxa, BCG attribute metrics and limited site information, such as watershed area, TNC temperature/geology/gradient classifications, average July temperature (if available), and % forest. Participants were not allowed to view StationIDs or waterbody names when making BCG level assignments, as this might bias their assignments. A sample worksheet can be found in Appendix E. Other information that was gathered but not included in the worksheets was latitude and longitude, chemical water quality data and land use information. Site information data were not gathered with the intent of developing causal relationships; rather the intent was to define a stress gradient (mainly from land use data) and to learn more about the full range of anthropogenic disturbances that may be occurring in these streams.

2.4 Quantitative Description

Level descriptions in the conceptual model tend to be rather general (e.g., "reduced richness"). To allow for consistent assignments of sites to levels, it is necessary to formalize the expert knowledge by codifying level descriptions into a set of rules (e.g., Droesen 1996). If formalized properly, any person (with data) can follow the rules to obtain the same level assignments as the group of experts. This makes the actual decision criteria transparent to stakeholders.

Rules are logic statements that experts use to make their decisions; for example, "If taxon richness is high, then biological condition is high." Rules on attributes can be combined, for example: "If the number of highly sensitive taxa (Attribute II) is high, and the number of tolerant individuals (Attribute V) is low, then assignment is Level 2." In questioning individuals on how decisions are made in assigning sites to levels, people generally do not use inflexible, "crisp" rules, for example, the following rule is unlikely to be adopted:

"Level 2 always has 10 or more Attribute II taxa; 9 Attribute II taxa is always Level 3."

Rather, people use strength of evidence in allowing some deviation from their ideal for any individual attributes, as long as most attributes are in or near the desired range. Clearly, the definitions of "high," "moderate," "low," etc., are fuzzy. These rules preserve the collective professional judgment of the expert group and set the stage for the development of models that

reliably assign sites to levels without having to reconvene the same group. In essence, the rules and the models capture the panel's collective decision criteria.

Rule development requires discussion and documentation of BCG level assignment decisions and the reasoning behind the decisions. During our discussions, facilitators recorded:

- Each participant's decision ("vote") for the site
- The critical or most important information for the decision—for example, the number of taxa of a certain attribute, the abundance of an attribute, the presence of indicator taxa, etc.
- Any confounding or conflicting information and how this was resolved for the eventual decision

The criteria that panelists use to make their decisions are captured in preliminary, narrative rules. For example, "For BCG Level 2, sensitive taxa must make up half or more of all taxa in a sample." The decision rule for a single level of the BCG does not always rest on a single attribute (e.g., highly sensitive taxa) but may include other attributes as well (intermediate sensitive taxa, tolerant taxa, indicator species), so these are termed "Multiple Attribute Decision Rules." With data from the sites, the rules can be checked and quantified. Quantification of rules will allow the agency to consistently assess sites according to the same rules used by the expert panel, and will allow a computer algorithm, or other persons, to obtain the same level assignments as the panel.

Following the initial site assignment and rule development, we developed descriptive statistics of the attributes and other biological indicators for each BCG level determined by the panel. These descriptions assisted in review of the rules and their iteration for testing and refinement.

2.5 Develop Decision Criteria Model

Consensus professional judgment used to describe the BCG levels can take into account nonlinear responses, uncommon stressors, masking of responses, and unequal weighting of attributes. This is in contrast to the commonly used biological indexes, which are typically unweighted sums of attributes (e.g., multimetric indexes; Barbour et al. 1999; Karr and Chu 1999), or a single attribute, such as observed to expected taxa (e.g., Simpson and Norris 2000; Wright 2000). Consensus assessments built from the professional judgment of many experts result in a high degree of confidence in the assessments, but the assessments are labor-intensive (several experts must rate each site). It is also not practical to reconvene the same group of experts for every site that is monitored in the long term. Since experts may be replaced on a panel over time, assessments may in turn "drift" due to individual differences of new panelists. Management and regulation, however, require clear and consistent methods and rules for assessment, which do not change unless deliberately reset.

2.6 Development of a BCG Model Using a Decision Criteria Approach

A BCG-based index for use in routine monitoring and assessment thus requires a way to automate the consensus expert judgment so that the assessments are consistent. We incorporated

the decision criteria into a decision model, which has the advantage that the criteria are visible and transparent. The model replicates the decision criteria of the expert panel by assembling the decision rules using logic and set theory, in the same way the experts used the rules.

Instead of a statistical prediction of expert judgment, this approach directly and transparently converts the expert consensus to automated site assessment. The method uses modern mathematical set theory and logic (called “fuzzy set theory”) applied to rules developed by the group of experts. Fuzzy set theory is directly applicable to environmental assessment, and has been used extensively in engineering applications worldwide (e.g., Demicco and Klir 2004) and environmental applications have been explored in Europe and Asia (e.g., Castella and Speight 1996; Ibelings et al. 2003).

Mathematical fuzzy set theory allows degrees of membership in sets, and degrees of truth in logic, compared to all-or-nothing in classical set theory and logic. Membership of an object in a set is defined by its membership function, a function that varies between 0 and 1. As an example, we compare how classical set theory and fuzzy set theory treat the common classification of sediment, where sand is defined as particles less than or equal to 2.0 mm diameter, and gravel is greater than 2.0 mm (Demicco and Klir 2004). In classical “crisp” set theory, a particle with diameter of 1.999 mm is classified as “sand”, and one with 2.001 mm diameter is classified as “gravel.” In fuzzy set theory, both particles have nearly equal membership in both classes (Demicco 2004). Measurement error of 0.005 mm in particle diameter greatly increases the uncertainty of classification in classical set theory, but not in fuzzy set theory (Demicco and Klir 2004). Demicco and Klir (2004) proposed four reasons why fuzzy sets and fuzzy logic enhance scientific methodology:

- Fuzzy set theory has greater capability to deal with “irreducible measurement uncertainty,” as in the sand/gravel example above.
- Fuzzy set theory captures vagueness of linguistic terms, such as “many,” “large” or “few.”
- Fuzzy set theory and logic can be used to manage complexity and computational costs of control and decision systems.
- Fuzzy set theory enhances the ability to model human reasoning and decision-making, which is critically important for defining thresholds and decision levels for environmental management.

In order to develop the decision criteria inference model, each attribute variable (e.g., “high taxon richness”) was defined quantitatively as a fuzzy set (e.g., Klir, 2004). A fuzzy set has a membership function. An example of membership functions of different classes of taxon richness are shown in Figure 2-3. We used piecewise linear functions to assign membership of a sample to the fuzzy sets. Numbers below a lower threshold have membership of 0, and numbers above an upper threshold have membership of one, and membership is a straight line between the lower and upper thresholds. For example, in Figure 2-3, a sample with 20 taxa would have a membership of 0.50 in the set “Low-moderate Taxa” and a membership of 0.50 in the set “Moderate Taxa.”

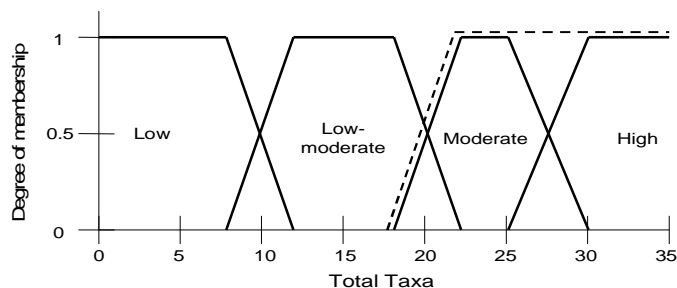


Figure 2-3. Fuzzy set membership functions assigning linguistic values of Total Taxa to defined quantitative ranges. Heavy dashed line shows membership of fuzzy set defined by “Total taxa are moderate to high.”

Inference uses the logic statements developed by expert consensus. In “crisp” logic, an AND statement is the same as “Intersection” in crisp set theory, and logical OR is equivalent to set theory “Union”. These are the same in fuzzy logic, however, a fuzzy AND uses the minimum membership of the two sets, and a fuzzy OR uses the maximum (Klir, 2004). For example, suppose a sample has membership of 0.25 in the set “Highly Sensitive taxa are Moderate” and membership of 0.75 in “Sensitive Taxa are High.” If the rule is a fuzzy AND statement (e.g., Highly Sensitive taxa are Moderate AND Sensitive Taxa are High), then its membership in level 2 is $\min(0.75, 0.25) = 0.25$. If the rule is a fuzzy OR statement, then its membership in level 2 equals $\max(0.75, 0.25) = 0.75$. Output of the inference model may include membership of a sample in a single level only, ties between levels, and varying memberships among two or more levels. The level with the highest membership value is taken as the nominal level.

3 COMPREHENSIVE DECISION RULES AND BCG MODEL – SMALL-COLD

3.1. Site Assignments and BCG Level Descriptions

During the calibration exercise, participants made BCG level assignments on 40 small-cold calibration samples and 14 validation samples. Locations of the assessed small-cold sites are shown in Figure 3-1. These samples were assigned to 5 BCG levels (BCG levels 1-5)¹. Five samples were assigned to BCG level 1, which consists of nearly pristine sites (Davies and Jackson 2006). Designating BCG level 1 samples is challenging because we lack sufficient information to know what the historical undisturbed assemblage in this region looked like.

¹ There was one majority opinion for a BCG Level 6 assignment, which is the most disturbed condition, but this was a questionable sample (Gulf Brook, StationID 5923 – it contained 1 American eel) so we did not include it in our calibration dataset.

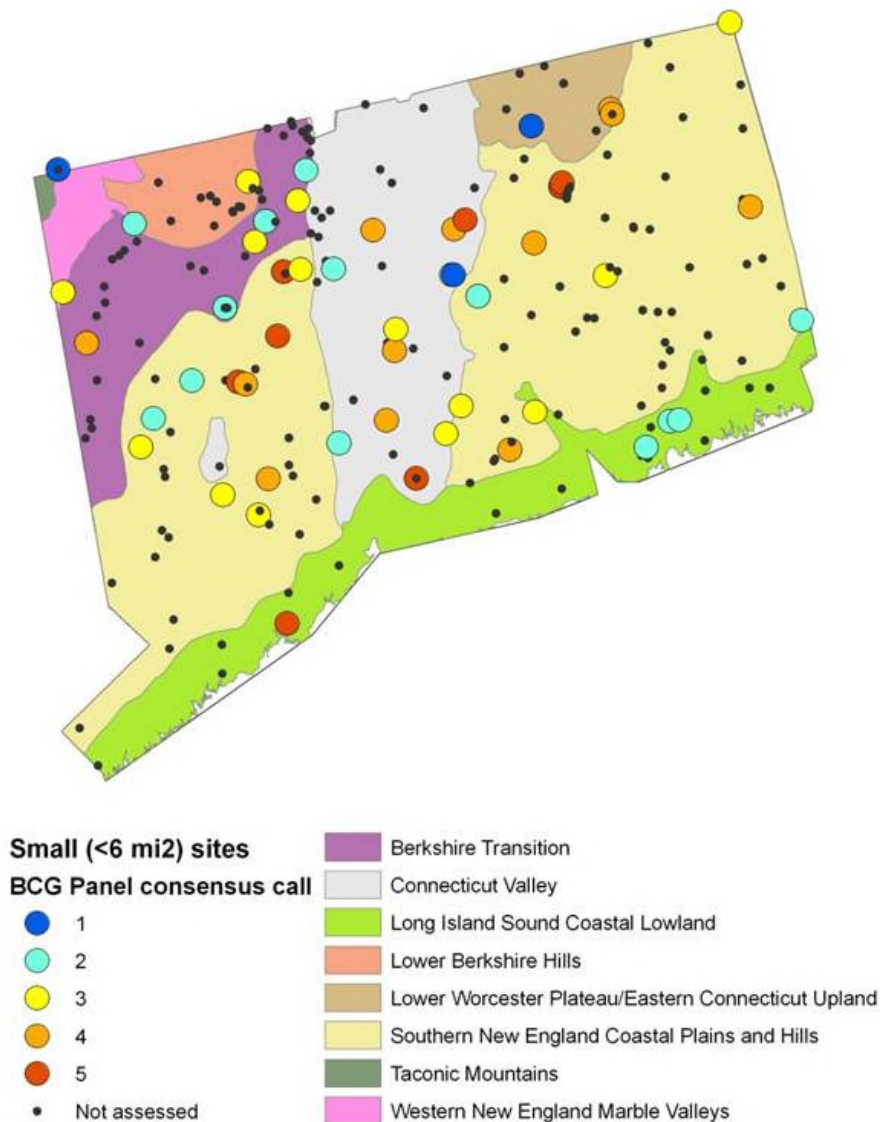


Figure 3-1. Locations of assessed small-cold samples (sites with drainage areas <6 mi²), coded by panelist BCG level assignment. This map also shows U.S. EPA Level 4 ecoregions. Ecoregions are delineated based on similarities in characteristics such as geology, physiography, vegetation, climate, soils, land use, wildlife, and hydrology (Omernik 1987, Omernik 1995).

3.2 BCG Attribute Metrics

We considered over 90 different metrics when calibrating the BCG model. Examinations of taxonomic attributes among the BCG levels determined by the panel showed that several of the attributes are useful in distinguishing levels, and indeed, were used by the panelists for decision criteria. The most important considerations were number of total taxa, percent native brook trout individuals and percent individuals and percent taxa metrics for sensitive (Attribute II+III), tolerant (Attribute V+VIa), non-native-non-salmonid (VI+VIa) and Attribute I-IV+VIb taxa.

Total richness showed a distinctly modal pattern, increasing as the assigned BCG level went from 1 to 4 (Figure 3-2), and then sharply fewer taxa in BCG Level 5. Watershed size was significantly and positively correlated with the total number of taxa ($r=0.53$, $p<0.01$) (Figure 3-3). Expectations of the panelists were in keeping with this relationship. In small, high quality coldwater streams (BCG levels 1-3), panelists expected the assemblage to be comprised of 6 or fewer species (the threshold of 6 is based on best professional judgment). As the streams increase in size, panelists expected more species to naturally be present. In BCG level 1 and 2 samples, the panelists expected to see high densities of native brook trout. If slimy sculpin and/or American Brook lamprey were also present, the panelists viewed this favorably, but since these species have limited spatial distributions, their presence was not required.

For the BCG attribute metrics, the percent individuals and percent taxa metrics were generally more effective at discriminating between BCG levels than absolute richness metrics. The Attribute II, II+III and IV metrics show relatively monotonic patterns, with Attribute IV metrics increasing and Attribute II+III metrics decreasing as the assigned BCG level goes from 1 to 5. The total taxa, Attribute II, II+III, percent wild brook individuals, percent tolerant (Attribute V+VIa) and percent non-native (Attribute VI+VIa+VIb) metrics were most effective at discriminating between BCG levels 1 and 2. All but one of these metrics (the percent non-native (Attribute VI+VIa+VIb) metric) also effectively captured the transition from BCG level 2 to 3. The percent non-native individuals (Attribute VI+VIa) was effective at distinguishing between BCG level 2 and 3 when nonnative salmonids were excluded (Attribute VIb). The transition from BCG level 3 to 4 was best captured by the Attribute II, Attribute II+III, Attribute II+III+non-native salmonid (VIb), Attribute II+III+IV+non-native salmonid (VIb) and tolerant (Attribute V+VIa) percent individuals and taxa metrics. BCG level 5 was discriminated from other BCG levels by the complete loss of Attribute II taxa, a decrease in Attribute II+III taxa and the concomitant increase in Attribute IV and percent tolerant (Attribute V + VIa) individuals. Distributions of various percent individuals and percent taxa metrics across BCG levels are shown graphically in Figures 3-4 through 3-6. Box plots for additional metrics can be found in Appendix F.

Presence and relative abundance of non-native taxa, in particular non-native trout, was another important consideration when panelists made BCG level assignments. Non-native trout were regarded as indicators of good water quality and coldwater habitat, but they also represent an altered fish assemblage. Panelists had different opinions on whether non-native trout could be present in BCG level 1 samples. In the end, a rule was established that requires all non-native taxa (including sensitive trout species) to be absent from BCG level 1 samples. In the original BCG level descriptions (Davies and Jackson 2006; Appendix B), the definition of BCG Level 1 does not explicitly state that non-natives cannot be present; however it does state that native structure must be preserved, so if non-natives are present, they cannot be displacing natives. The difference between BCG level 1 and 2 is subtle and comprises small changes in taxonomic composition versus functional degradation.

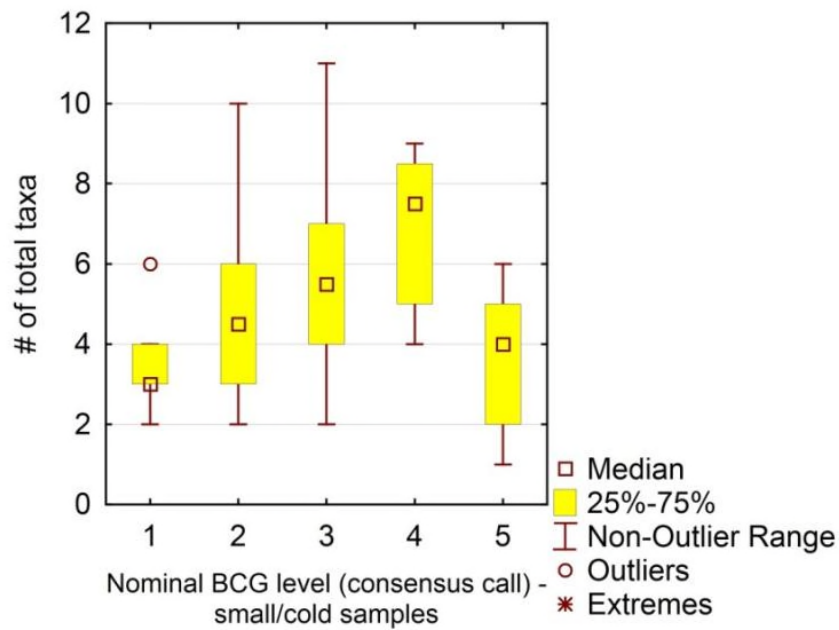


Figure 3-2. Box plots of total taxa metric values for small-cold samples, grouped by nominal BCG level (group majority choice). Sample size (which includes both calibration and validation samples) for BCG level 1 = 5, level 2 = 14, level 3 = 14, level 4 = 12, and level 5 = 9. The total taxa metric counts *all* taxa (even singletons), and counts native and non-native brook trout as separate species.

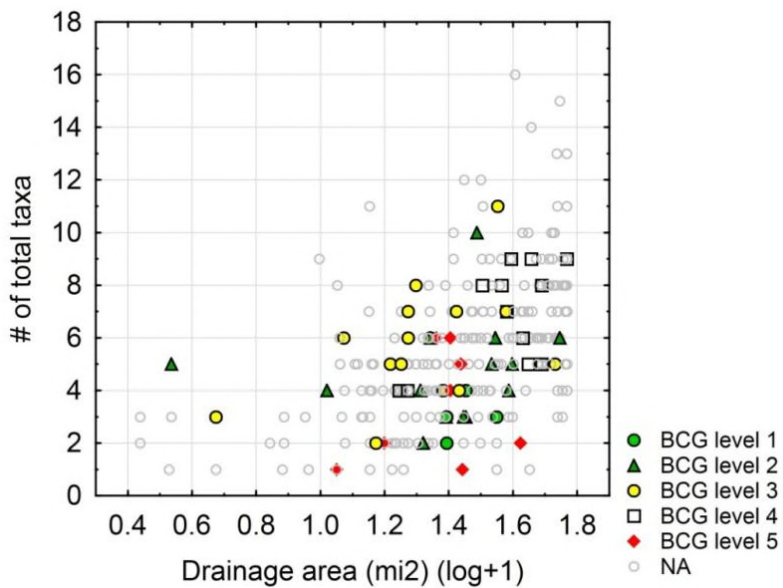


Figure 3-3. Relationship between total taxa metric values and watershed area for small-cold samples ($r=0.53$, $p<0.01$). Samples are coded by nominal BCG level (group majority choice).

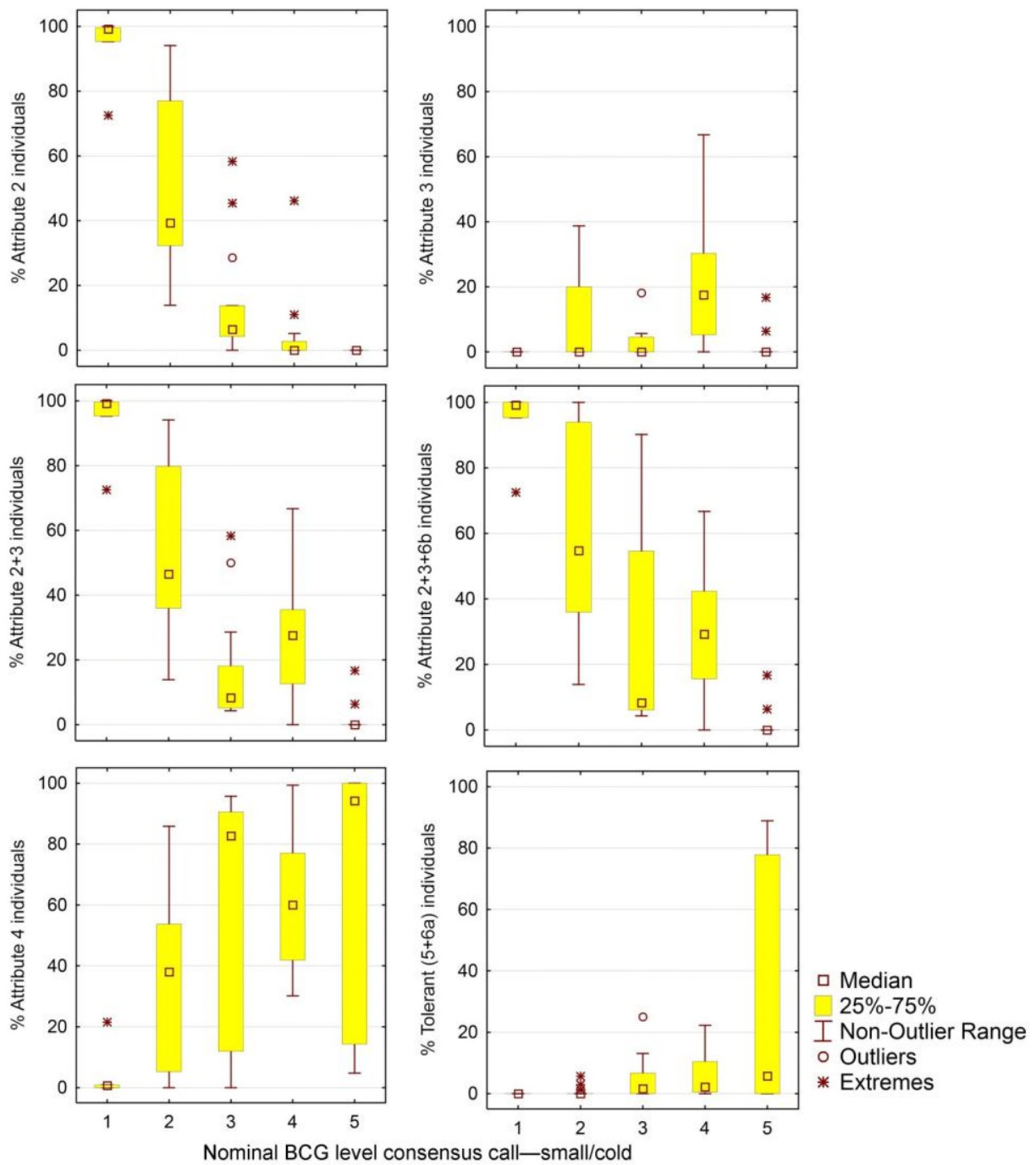


Figure 3-4. Box plots for a subset of BCG attribute percent individual metrics for the 54 small-cold samples that were assessed (this includes both calibration and validation samples), grouped by nominal BCG level (group majority choice). Sample size for BCG level 1 = 5, level 2 = 14, level 3 = 14, level 4 = 12, and level 5 = 9.

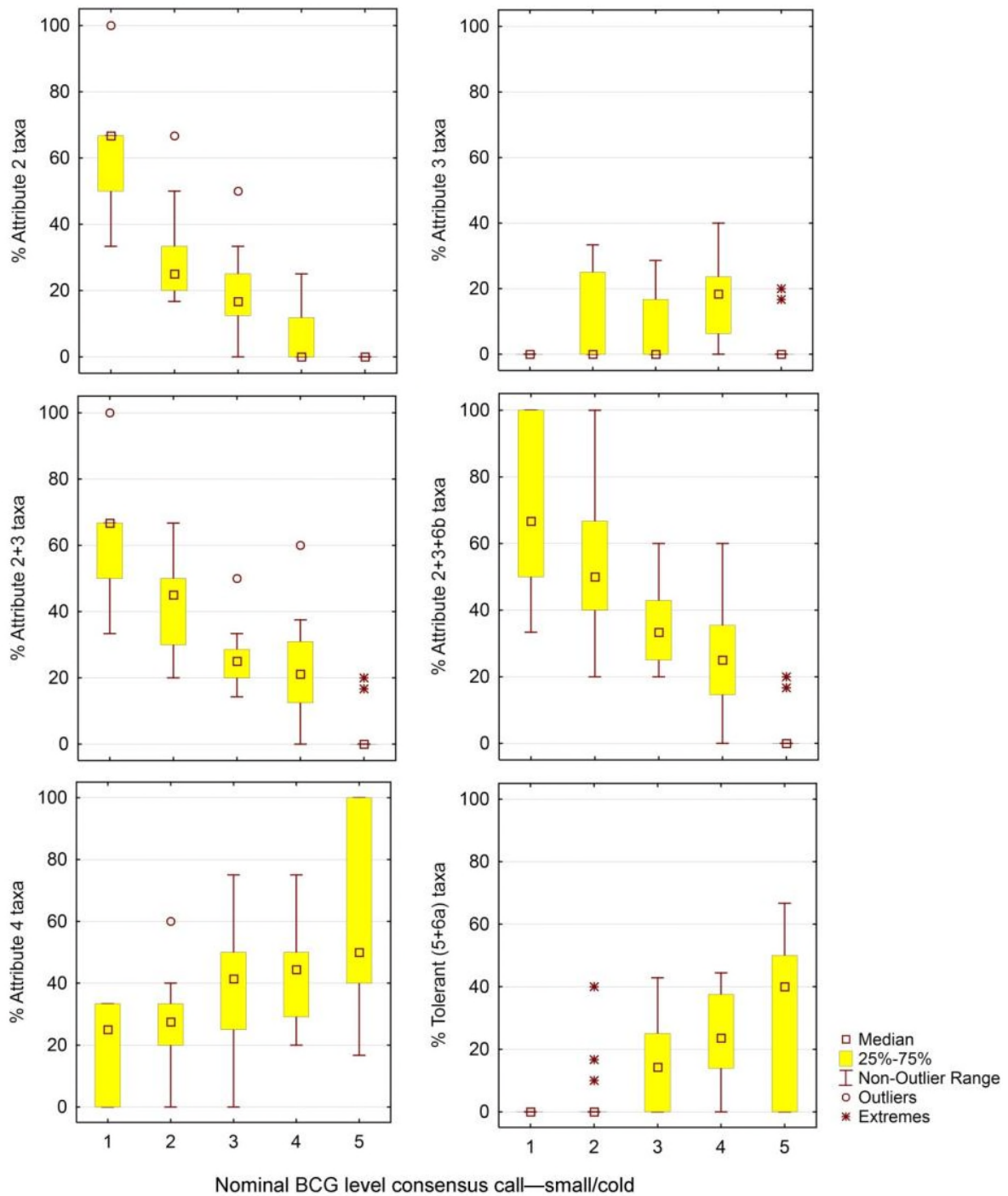


Figure 3-5. Box plots for a subset of BCG attribute percent taxa metrics for the 54 small-cold samples that were assessed (this includes both calibration and validation samples), grouped by nominal BCG level (group majority choice). Sample size for BCG level 1 = 5, level 2 = 14, level 3 = 14, level 4 = 12, and level 5 = 9.

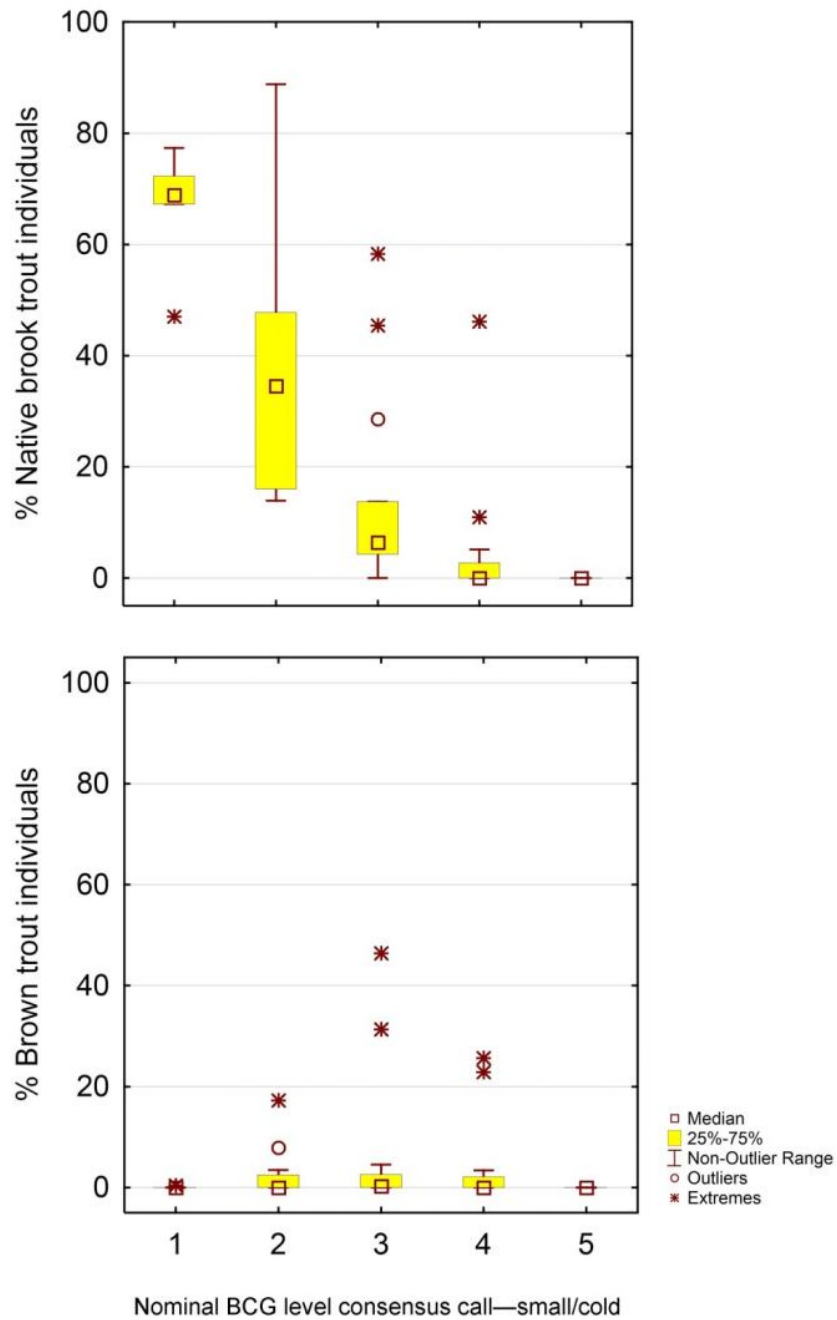


Figure 3-6. Box plots of native brook trout and brown trout percent individual metrics for the 54 small-cold samples that were assessed (this includes both calibration and validation samples), grouped by nominal BCG level (group majority choice). Sample size for BCG level 1 = 5, level 2 = 14, level 3 = 14, level 4 = 12, and level 5 = 9.

3.3 BCG Rule Development

The small-cold rules, which are shown in Table 3-1, were derived from discussions with the panelists on why individual sites were assessed at a certain level. They follow the observations shown in Figures 3-2 through 3-6. The rules were calibrated with 40 small-cold fish samples rated by the group, and were adjusted so that the model would replicate the panel's decisions as closely as possible. Inevitably, there were some places where the panel may have used different, unstated rules, or where rules were inconsistently applied. Panelist and model BCG level assignments for these 40 samples, along with site information, are summarized in Appendix G. Appendix G also contains panelist and model BCG level assignments for the 14 samples that were assessed during the validation round.

In the model, rules work as a logical cascade from BCG Level 1 to Level 6. A sample is first tested against the Level 1 rules; if a single rule fails, then the Level fails, and the assessment moves down to Level 2, and so on. All required rules must be true for a site to be assigned to a level. As described in Section 2.6, membership functions had to be defined for the richness and percent individual metrics.

The rules shown in Table 3-1 have been developed for distinguishing BCG levels for small-cold fish samples. They follow a general pattern of decreasing richness of sensitive taxa and increasing relative abundance of tolerant individuals as biological condition degrades. Small-cold BCG Level 1 rules require that fewer than 6 total taxa be present, that at least 70% of the assemblage be comprised of sensitive (Attribute I + II + III) individuals (at least 60% of which must be native brook trout), and that non-native (Attribute VI+VIa+VIb) taxa are absent.

In BCG Level 2 samples, fewer than 7 total taxa must be present. At least 30% of a BCG level 2 sample must be comprised of sensitive (Attribute I + II + III) individuals, at least 10% of the individuals must be native brook trout, and there must be less than 6 and 12% percent tolerant (Attribute V + VIa) and non-native, non-salmonid (Attribute VI + VIa) individuals, respectively.

BCG level 3 samples must have fewer than 9 total taxa. In addition, the percent Attribute II and percent sensitive (Attribute I + II + III) taxa metrics must exceed thresholds of 5 and 15%, respectively, there must be more than 5% percent sensitive (Attribute I + II + III) individuals, the most dominant Attribute V, VI or VIa taxon must comprise less than 50% of the assemblage and percent non-native, non-salmonid (Attribute VI + VIa) individuals must be less than 20%.

BCG Level 4 is characterized by decreased richness and abundance of sensitive (Attribute I + II + III) taxa. More than 3 total taxa must be present, or, alternately, if fewer than 3 are present, at least 1 of the taxa must be an Attribute II taxa. Sensitive taxa (Attribute II+III) must be present, and the assemblage must be comprised of more than 40% Attribute I + II + III + IV + non-native salmonid (VIb) individuals and taxa. There also must be less than 20% tolerant (Attribute V + VIa) individuals. BCG Level 5 rules require that Attribute I + II + III + IV + non-native salmonid (VIb) individuals comprise at least 30% of the taxa and at least 10% of the individuals

Table 3-1. BCG decision rules for fish assemblages in small-cold (<6 m²) and medium-large (≥6 m²) streams. Samples are flagged for professional assessment (outside experience of model) if < 20 total individuals are present (per Kanno et al. 2010a), if > 400 total individuals are present in streams < 6 mi² or if > 500 total individuals are present in streams ≥ 6 mi².

| BCG Level 1 | Small-Cold (n=5) | | | Medium-Large (n=1) | | |
|--|--------------------------|-------------|-----------------|---------------------------|---|-----------------|
| Metrics | data | rule | alt rule | data | rule | alt rule |
| # Total taxa | 2-6 | ≤ 5 | | 7 | | ≥ 6 |
| # Attribute I + II taxa | -- | -- | | 1 | | present |
| % Native brook trout individuals | 47-77% | > 60% | | -- | | -- |
| % Attribute I + II + III taxa | 33-100% | | | 28% | | > 25% |
| % Attribute I + II + III individuals | 72-100% | > 70% | | 79% | | > 25% |
| % Tolerant (V + VI a) individuals | -- | -- | | 0% | | < 5% |
| % Non-native (VI + VI a + VIb) individuals | 0-0.4% | absent | | 0.4% | | absent |
| BCG Level 2 | Small-Cold (n=14) | | | Medium-Large (n=6) | | |
| Metrics | data | rule | alt rule | data | rule | alt rule |
| # Total taxa | 2-10 | ≤ 6 | | 4-12 | area <20 mi ² , taxa ≥ 2; area >20 mi ² , taxa ≥ 8 | |
| % Native brook trout individuals | 14-89% | > 10% | | -- | | -- |
| % Attribute I + II + III taxa | -- | -- | | 9-44% | | > 20% |
| % Attribute I + II + III individuals | 14-94% | > 30% | | 1-64% | | > 20% |
| % Tolerant (V + VIa) individuals | 0-6% | < 6% | | -- | | -- |
| % Non-native, non-salmonid (VI + VIa) individuals | 0-5% | < 12% | | -- | | -- |
| % Most dominant intermediate tolerant (Att IV) taxon | -- | -- | | 14-52% | | < 40% |
| # Salmonidae taxa | -- | -- | | 1-4 | | present |
| % Centrarchidae individuals | -- | -- | | 0-14% | < 2 % OR | |
| % Attribute II individuals | -- | -- | | 0-59% | | > 5% OR |

Continued...

| BCG Level 3 | Small-Cold (n=14) | | | Medium-Large (n=24) | | |
|---|--------------------------|-------------|-----------------|----------------------------|---|-----------------|
| Metrics | data | rule | alt rule | data | rule | alt rule |
| # Total taxa | 2-11 | ≤ 8 | | 4-14 | area <20 mi ² , taxa ≥ 2; area >20 mi ² , taxa ≥ 8 | |
| % Attribute I + II taxa | 0-50% | > 5% | | -- | -- | |
| % Attribute I + II + III taxa | 14-50% | > 10% | | 11-67% | > 10% | |
| % Attribute I + II + III individuals | 4-58% | > 5% | | 3-64% | > 3% | |
| % Attribute I + II + III + non-native salmonid (VIb) taxa | -- | -- | | 14-67% | > 20% | |
| % Attribute I + II + III + non-native salmonid (VIb) individuals | -- | -- | | 11-65% | > 10% | |
| % Most dominant tolerant (Att V, VI or VIa) taxon | 0-25% | < 50% | | -- | -- | |
| % Most dominant tolerant (Att V) taxon | -- | -- | | 0-4% | < 5% | |
| % Centrarchidae individuals | -- | -- | | 0-25% | < 10% | |
| % Non-native, non-salmonid (VI + VIa) individuals | 0-14% | < 20% | | -- | -- | |
| % Cyprinid taxa | 0-67% | -- | | 22-67% | > 10% | |
| BCG Level 4 | Small-Cold (n=12) | | | Medium-Large (n=27) | | |
| Metrics | data | rule | alt rule | data | rule | alt rule |
| # Total taxa | 4-9 | > 3 | ≥ 1 | -- | -- | |
| # Attribute I + II taxa | 0-1 | -- | present | -- | -- | |
| # Attribute I + II + III taxa | 0-3 | present | | -- | -- | |
| % Attribute I + II + III + IV + non-native salmonid (VIb) taxa | 50-89% | > 40% | | 50-91% | > 40% | |
| % Attribute I + II + III + IV + non-native salmonid (VIb) individuals | 74-100% | > 40% | | 49-99% | > 40% | |
| % Centrarchidae individuals | -- | -- | | 0-53% | < 40% | |
| % Tolerant (V + VIa) individuals | 0-22% | < 20% | | -- | -- | |
| % Cyprinid taxa | -- | -- | | 9-50% | present | |
| % Non-native, non-salmonid (VI + VIa) individuals | -- | -- | | -- | < 50% | |

Continued...

| BCG Level 5 | Small-Cold (n=9) | | | Medium-Large (n=12) | | |
|---|------------------|------|----------|---------------------|------|----------|
| Metrics | data | rule | alt rule | data | rule | alt rule |
| # Total taxa | -- | | -- | 5-16 | | > 3 |
| % Attribute I + II + III + IV + non-native salmonid (VIb) taxa | 33-100% | | > 30% | 38-86% | | > 20% |
| % Attribute I + II + III + IV + non-native salmonid (VIb) individuals | 11-100% | | > 10% | 14-100% | | > 10% |

As mentioned in Section 3.2, a rule was also developed for all BCG levels to flag samples with low and high densities. These rules are as follows: low density < 20 total individuals; high density >400 individuals in small (<6 mi²) streams and >500 individuals in medium-large (≥ 6 mi²) streams. Failure of the density rule causes the sample to be flagged for professional assessment. It should be noted that the model still makes BCG level assignments for samples that are flagged, and that density is not a consideration in these BCG level assignments.

3.4 Model Performance

To evaluate the performance of the 41-sample small-cold calibration dataset and the 14-sample validation dataset, we considered two matches in BCG Level choice: an exact match, where the BCG decision model's nominal level matched the panel's majority choice; and a "near match", where the model predicted a BCG level within one level of the majority expert opinion. When model performance was evaluated, the small-cold fish model matched exactly with the regional biologists' BCG level assignments on 68% of the calibration samples (Table 3-2). Eleven (27%) of the model assignments were within one level of the majority expert opinion, and two (5%) were off by two BCG levels. Where there were differences, the tendency was for the model to rate samples worse than the panel; the model assigned 9 samples to a BCG level that was 1 level worse than the panelists, 1 sample to a BCG level that was 2 levels worse, 2 samples to a BCG level that was 1 level better than the panelists' assignment and 1 sample to a BCG level that was 2 levels better than the panelists' assignment. These results should be interpreted with caution because some of the calibration consensus calls were made early on in the process (e.g. at the November 2010 workshop) and may not reflect changes that occurred in the group's thinking.

In order to confirm the model, panelists made BCG level assignments on 14 additional small-cold samples. When nominal level assignments from the BCG decision model were compared to the panelists' nominal level assignments², the small-cold fish model matched exactly with the regional biologists' BCG level assignments on 100% of the confirmation samples (Table 3-2). We believe the model performance is better in the validation dataset than in the calibration dataset because the calibration dataset reflects the twists and turns that occurred in the group's thinking during the model development process; the results from the validation dataset suggest that the group has indeed converged on this latest set of rules, and that these rules appear to have been successfully captured in the model.

It is possible that model performance in the calibration dataset could improve if the group reassesses some of the original calibration samples. In addition to potentially revising some of the consensus calls, closer examination of the anomalous samples may reveal issues that make some samples fall outside the realm of experience for the BCG model. For example, in this case, the two samples that differed the most from BCG model assignments both had such issues: one (Menunketesuck River, StationID 1976) was flagged for low density; and the other (East Mountain Brook, StationID 2714) was located near a large river, which likely resulted in an

² For most small-cold validation samples, panelists worked independently to rate the samples and did not discuss these ratings as a group.

inflated species count (which caused the sample to exceed the total taxa richness threshold for small streams).

Table 3-2. Summary of differences between model and panelist BCG level assignments for small-cold water samples.

| Difference (model minus panel consensus call)* | Calibration | | Confirmation |
|--|-------------|---|--------------|
| | # samples | notes | # samples |
| 2 better** | 1 | this sample was flagged for low density | 0 |
| 1 better | 2 | | 0 |
| same | 28 | | 14 |
| 1 worse | 9 | 2 of these samples were flagged for high density; 5 of these had differences between BCG levels 3 & 4 | 0 |
| 2 worse | 1 | site located near large river, likely resulting in inflated species count | 0 |
| Total # Samples | 41 | | 14 |
| % Correct | 70% | | 100% |

* In some instances, the model output was a tie between two BCG levels. We considered these to be matches if the range of model assignments matched with the range of panelist calls. For example, if the model output was a tie between BCG levels 1 and 2 and the panelist calls ranged from 1 to 2, we called this a match. If the model output was a tie between BCG levels 1 and 2 and the panelist calls ranged from 2 to 3, we considered this to be a difference of 1 BCG level. If there were ties between panelist calls, we used the lower score (e.g., BCG level 2 vs. 3) as the consensus call.

** this means that the model score was 2 levels better than the panelist score; for example, the model assigned it to BCG level 2 and the panel assigned it to BCG level 4.

4 COMPREHENSIVE DECISION RULES AND BCG MODEL – MEDIUM-LARGE

4.1 Site Assignments and BCG Level Descriptions

Participants made BCG level assignments on 54 medium-large calibration samples and 16 validation samples. Locations of the assessed medium-large sites are shown in Figure 4-1. These samples were assigned to BCG levels 1-5. One sample (Green Fall River, StationID 606) was assigned to BCG level 1. Designating BCG level 1 samples was challenging because there is not enough information to know what the historical undisturbed fish assemblage in medium-large streams in this region looked like. Most panelists said that they had greater difficulty making BCG level assignments on samples from medium-large streams versus small-cold streams.

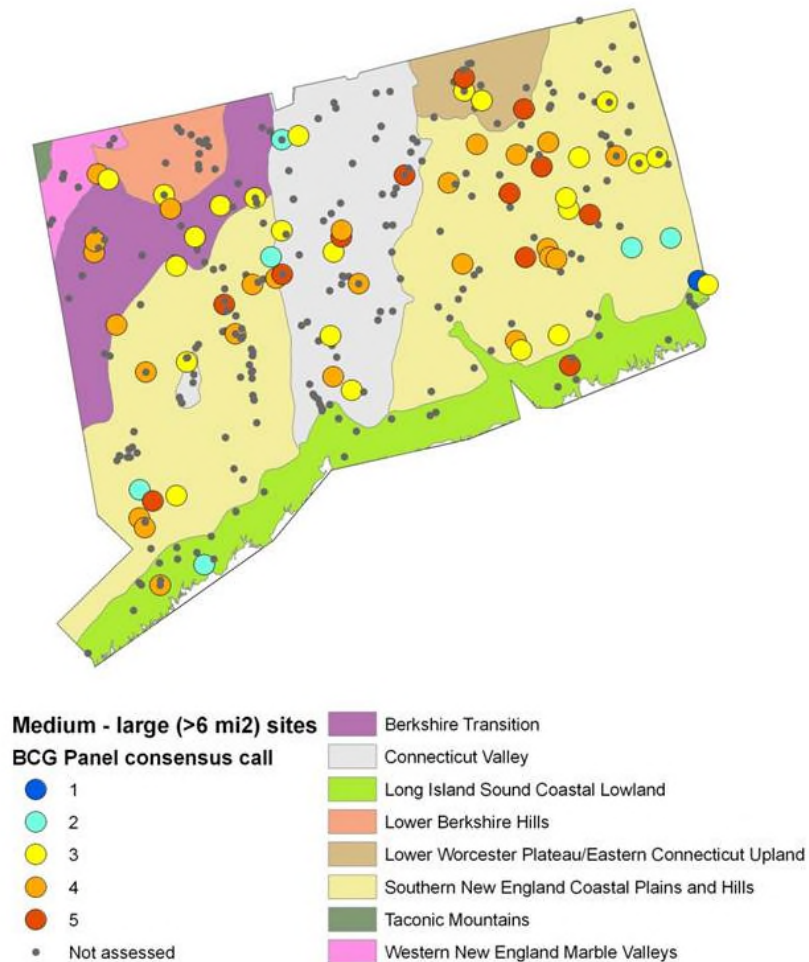


Figure 4-1. Locations of assessed medium-large (drainage areas ≥ 6 mi²) sites, coded by panelist BCG level assignment. This map also shows U.S. EPA Level 4 ecoregions. Ecoregions are delineated based on similarities in characteristics such as geology, physiography, vegetation, climate, soils, land use, wildlife, and hydrology (Omernik 1987, Omernik 1995).

4.2 BCG Attribute Metrics

The same metrics that were evaluated for coldwater samples (Section 3.2) were evaluated for samples from medium-large streams. Panelists considered an assortment of metrics when distinguishing between BCG levels for these samples. The most important considerations were number of total taxa, percent individuals and percent taxa metrics for sensitive (Attribute II+III), tolerant (Attribute V+VIa), non-native-non-salmonid (VI+VIa) taxa, Attribute I-IV+non-native salmonid (VIb) taxa, number of salmonid taxa, percent Cyprinid taxa and percent Centrarchidae individuals.

Total taxa richness did not show a clear pattern across BCG levels (Figure 4-2). As expected, watershed size was significantly and positively correlated with total fish species richness ($r=0.37$, $p<0.01$) (Figure 4-2). In high quality medium-large streams (BCG levels 1-3), panelists expected the assemblage to be comprised of at least 2 species. As the streams increase in size ($>20 \text{ mi}^2$), they expected more species (≥ 8) to naturally be present. In high quality, medium-large streams, panelists expected to see a balanced assemblage of cool water species, with at least 2 species of sensitive (Attribute II + III) taxa like longnose dace, common shiner and fallfish, and no hyperabundance of blacknose dace and/or white suckers. Panelists said they would consider samples with high proportions of native coldwater species to be BCG Level 1.

Overall, patterns in the medium-large BCG attribute metrics were less evident than those in the small-cold plots. However, the sensitive (Attribute II+III & Attribute II+III+VIb) and tolerant (Attribute VI + VIa) metrics did show fairly clear monotonic patterns, with sensitive metrics decreasing and tolerant metrics generally increasing as the assigned BCG level increased from 1 to 5 (Figures 4-3 & 4-4). Percent Cyprinid taxa and percent Centrarchidae individuals also show relatively monotonic patterns, with percent Cyprinid taxa decreasing and percent Centrarchidae individuals increasing with BCG level 1 to 5.

The total taxa, Attribute II+III and percent tolerant (Attribute V+VIa) are most effective at discriminating between BCG levels 1 and 2, and Attribute II taxa must be present and non-native taxa (Attribute VI+VIa+VIb) must be absent in BCG level 1 samples. Salmonids must be present in BCG level 2 samples, and the transition from BCG level 2 to 3 is captured by total taxa, Attribute II+III metrics, percent most dominant Attribute IV taxon and an alternate rule related to percent Centrarchidae individuals and percent Attribute II individuals. Metrics effective at discriminating between BCG levels 3 and 4 include Attribute II+III, Attribute II+III+ VIb and Attribute II+III+ IV+VIb metrics, as well as metrics related to Centrarchids and Cyprinids. The transition from BCG level 4 to 5 is captured by decreases in Attribute II+III+ IV+VIb metrics and increases in percent tolerant (Attribute V+VIa) individuals. Distributions of percent individuals and percent taxa metrics across BCG levels are shown graphically in Figures 4-2 through 4-6. Box plots for additional metrics can be found in Appendix H.

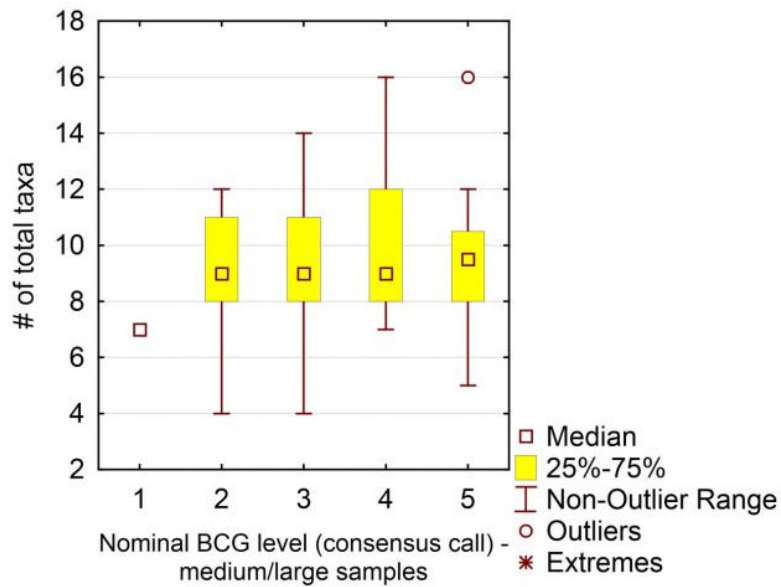


Figure 4-2. Box plots of total taxa metric values for medium-large samples, grouped by nominal BCG level (group majority choice). Sample size (which includes both calibration and validation samples) for BCG level 1 = 5, level 2 = 6, level 3 = 24, level 4 = 27, and level 5 = 12. The total taxa metric counts *all* taxa (even singletons), and counts native and non-native brook trout as separate species.

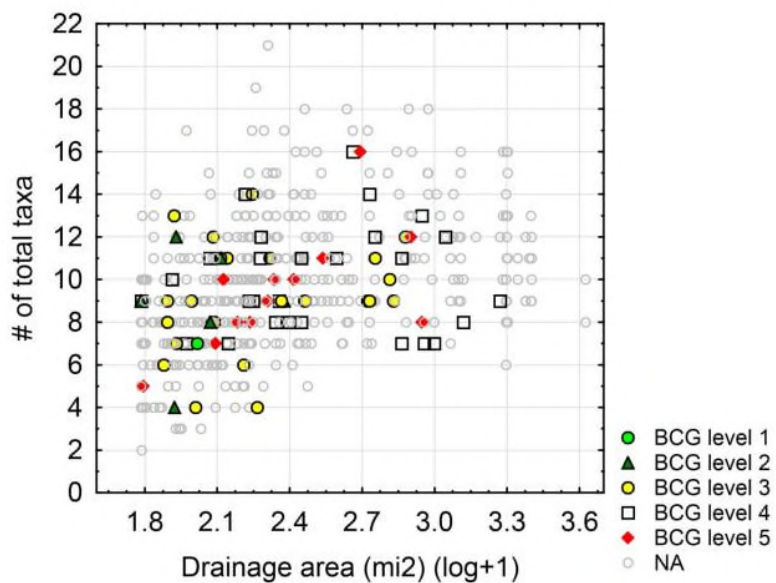


Figure 4-3. Relationship between total taxa metric values and watershed area for medium-large samples ($r=0.37$, $p<0.01$). Samples are coded by nominal BCG level (group majority choice).

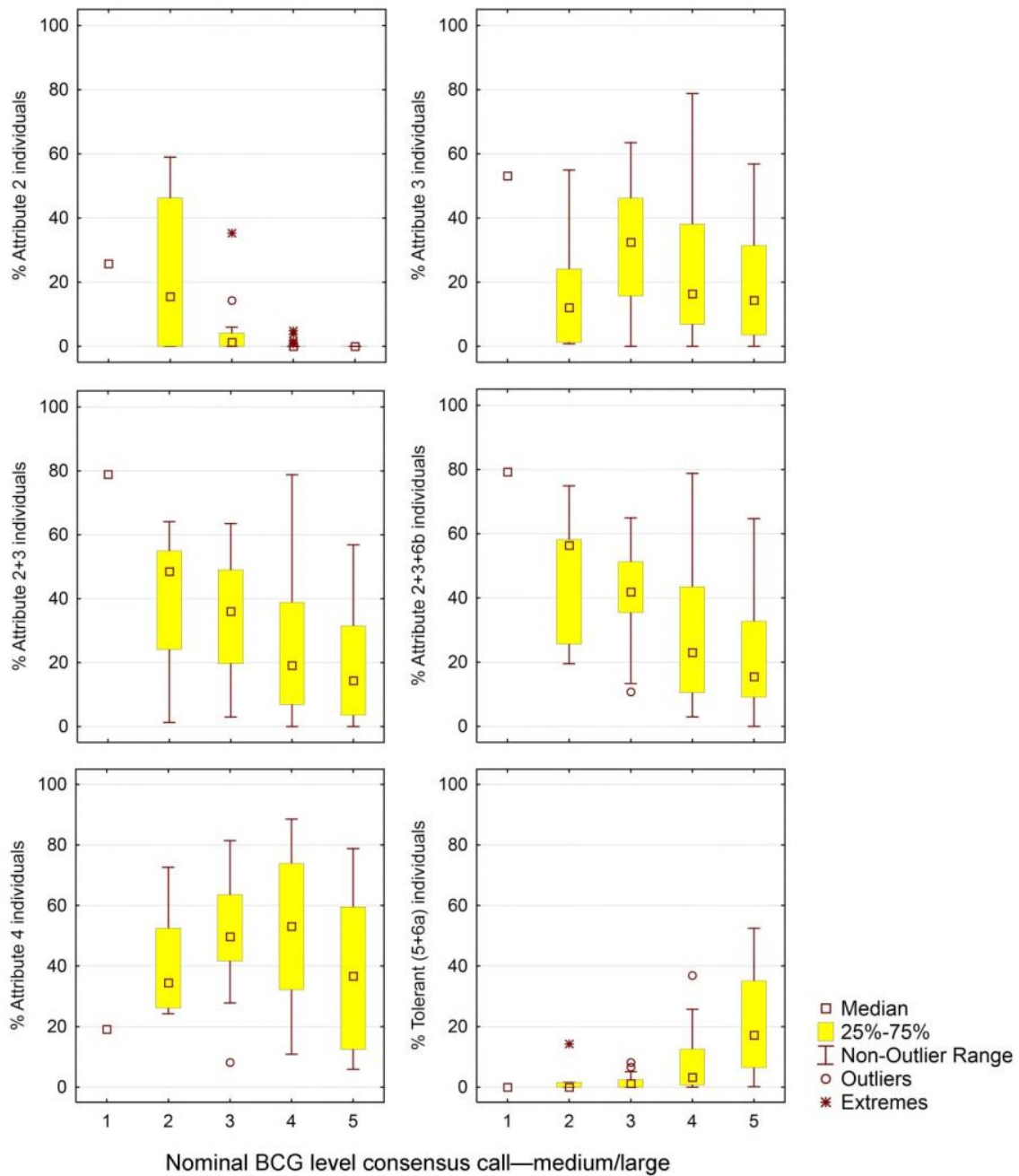


Figure 4-4. Box plots for a subset of BCG attribute percent taxa metrics for 70 medium-large samples that were assessed (this includes both calibration and validation samples), grouped by nominal BCG level (group majority choice). Sample size (which includes both calibration and validation samples) for BCG level 1 = 5, level 2 = 6, level 3 = 24, level 4 = 27, and level 5 = 12.

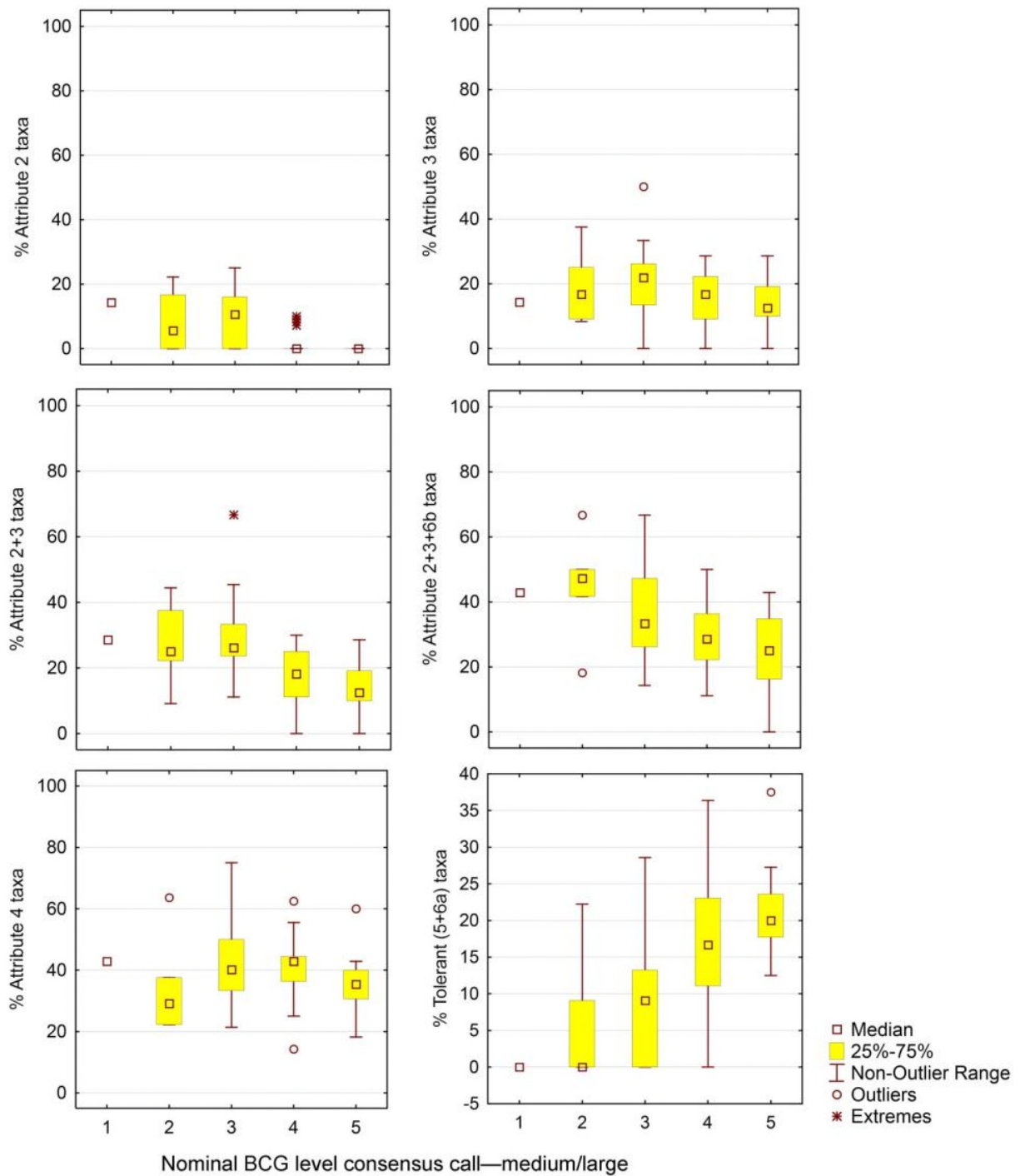


Figure 4-5. Box plots for a subset of BCG attribute percent taxa metrics for 70 medium-large samples that were assessed (this includes both calibration and validation samples), grouped by nominal BCG level (group majority choice). Sample size (which includes both calibration and validation samples) for BCG level 1 = 5, level 2 = 6, level 3 = 24, level 4 = 27, and level 5 = 12.

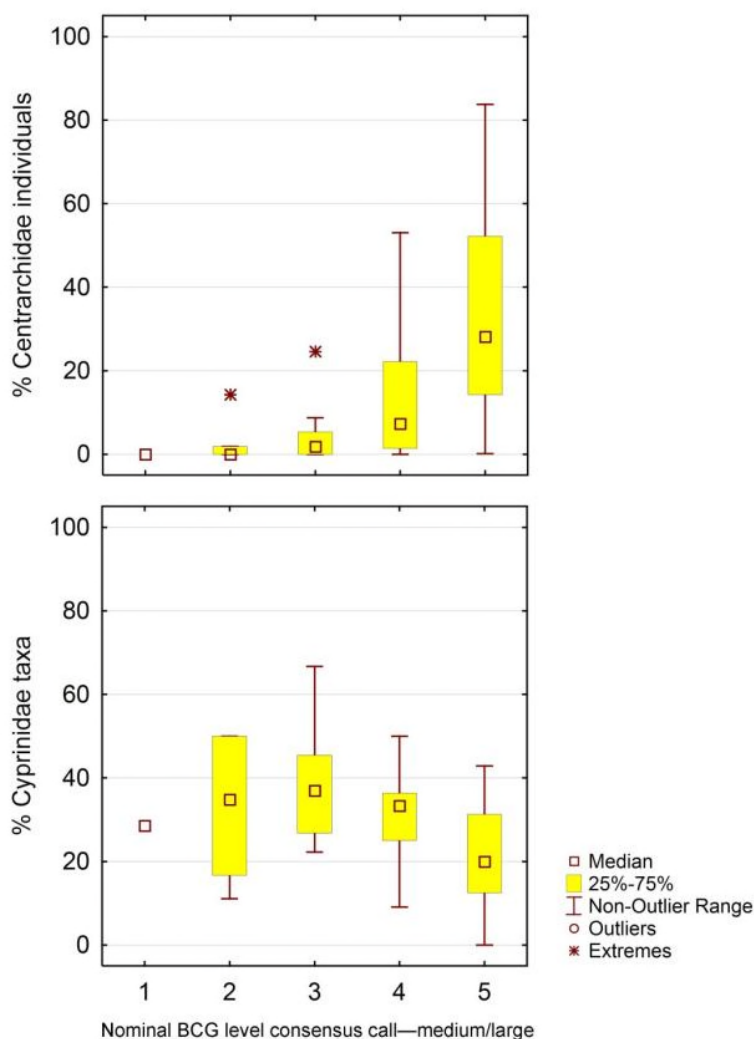


Figure 4-6. Box plots of the percent Centrarchidae individuals and percent Cyprinid taxa metrics for 70 medium-large samples that were assessed (this includes both calibration and validation samples), grouped by nominal BCG level (group majority choice). Sample size (which includes both calibration and validation samples) for BCG level 1 = 5, level 2 = 6, level 3 = 24, level 4 = 27, and level 5 = 12.

4.3 BCG Rule Development

Rules for medium-large ($\geq 6 \text{ mi}^2$) streams, which are shown in Table 3-1, were derived from discussions with the panelists on why individual sites were assessed at a certain level. They follow the observations shown in Figures 4-2 through 4-6. The rules were calibrated with the 54 medium-large samples rated by the group, and were adjusted so that the model would replicate the panel's decisions as closely as possible. Panelist and model BCG level assignments for these 54 samples, along with site information, are summarized in Appendix I. Appendix I also contains

panelist and model BCG level assignments for the 16 samples that were assessed during the validation round.

The rules for medium-large streams are shown in Table 3-1. They follow a general pattern of decreasing richness of sensitive taxa and increasing relative abundance of tolerant individuals as biological condition degrades. BCG Level 1 rules for medium-large streams require that at least 6 total taxa must be present, that at least 25% of the assemblage be comprised of sensitive (Attribute I + II + III) taxa and individuals, that less than 5% of the assemblage be comprised of tolerant (Attribute V+VIa) individuals and that non-native (Attribute VI+VIa+VIb) taxa be absent.

In BCG Level 2 samples, at least 2 total taxa must be present in streams less than 20 mi², and 8 or more taxa must be present in streams greater than 20 mi². At least 20% of a BCG level 2 sample must be comprised of sensitive (Attribute I + II + III) taxa and individuals and at least one species of salmonid must be present. The most dominant Attribute IV taxon must comprise less than 40% of the assemblage and there is an alternate rule in which percent Centrarchidae individuals must comprise less than 2% of the assemblage or percent Attribute II individuals must make up more than 5% of the assemblage.

The same total taxa richness rule holds true for BCG level 3 samples, with thresholds of 2 in smaller (<20 mi²) streams and 8 in larger (>20 mi²) streams. In addition, the percent sensitive (Attribute I + II + III) taxa and individuals metrics must exceed thresholds of 10 and 3%, respectively, the percent Attribute I + II + III + IV + non-native salmonid (VIb) taxa and individuals metrics must exceed 20 and 10%, respectively, the most dominant Attribute V taxon must not comprise more than 5% of the assemblage, percent Centrarchidae individuals must be less than 10% and percent Cyprinid taxa must be greater than 10%.

BCG Level 4 is characterized by decreased richness and abundance of sensitive (Attribute I + II + III) taxa. There is no rule for total taxa richness. Cyprinid taxa must be present, and percent Attribute I + II + III + IV + non-native salmonid (VIb) taxa and individuals must both exceed 40%. Percent Centrarchidae individuals must comprise less than 40% of the assemblage and non-native, non-salmonid (Attribute VI+VIa) must be less than 50%. BCG Level 5 rules require that more than 3 taxa be present, and that the percent Attribute I + II + III + IV + non-native salmonid (VIb) taxa and individuals metrics must exceed thresholds of 20 and 10%, respectively.

The rules for flagging high and low density samples (see Section 3.2) also apply to samples from medium-large streams.

4.4 Model Performance

To evaluate the performance of the 54-sample medium-large calibration dataset and the 16-sample validation dataset, we considered two matches in BCG Level choice: an exact match, where the BCG decision model's nominal level matched the panel's majority choice; and a "near match", where the model predicted a BCG level within one level of the majority expert opinion. When model performance was evaluated, the medium-large fish model matched exactly with the regional biologists' BCG level assignments on 61% of the calibration samples (Table 4-1). Eighteen of the model assignments were within one level of the majority expert opinion, one was off by two BCG levels and two were off by a half level (these were ties). Where there were differences, there wasn't a clear tendency for the model to rate samples better or worse than the panel; the model assigned 10 samples to a BCG level that was 1 level better than the panelists, 8 were assigned to a BCG level that was 1 level worse, and 1 was assigned to a BCG level that was 2 levels worse than the panelists' assignment. The results from the calibration dataset should be interpreted with caution because some of these BCG level assignments were made early in the process (e.g. at the November 2010 workshop) and may not reflect changes that have occurred in the group's thinking since that time.

In order to confirm the model, panelists made BCG level assignments on 16 additional medium-large samples. When nominal level assignments from the BCG decision model were compared to the panelists' nominal level assignments³, the model performed better on the validation dataset than on the calibration dataset. The medium-large fish model matched exactly with the regional biologists' BCG level assignments on 75% of the confirmation samples (Table 4-1). Where there were differences, there was a tendency for the model to rate samples worse than the panel; the model assigned 1 samples to a BCG level that was 1 level better than the panelists, 2 were assigned to a BCG level that was 1 level worse (one of these samples was flagged for high density), and 1 was assigned to a BCG level that was 2 levels worse than the panelists' assignment.

As with the small-cold samples, we believe the model performance is better in the validation dataset than in the calibration dataset because the calibration dataset reflects evolution that occurred in the group's thinking during the model development process; the improved results in the validation dataset suggest that the group has converged more on this latest set of rules, and the model appears to have captured this.

It is possible that model performance in the calibration dataset could improve if the group reassesses some of the original calibration samples. In addition to potentially revising some of the consensus calls, closer examination of the anomalous samples may reveal issues that make some samples fall outside the realm of experience for the BCG model. For example, of the 4 anomalous samples in the validation dataset, two (Pease Brook, StationID 1482 and Mount Hope River, StationID 1671) were flagged for high density.

³ For the medium/large validation samples, panelists first assessed samples independently but then discussed them as a group to reach a consensus call.

Table 4-1. Summary of differences between model and panelist BCG level assignments for medium-large samples.

| Difference (model minus panel consensus call) | Calibration | | Confirmation |
|---|-------------|---|----------------------------------|
| | # samples | notes | # samples |
| 1 better | 10 | 5 of these had differences between BCG levels 4 & 5 | 1 |
| 0.5 better* | 2 | | 0 |
| same | 33 | | 12 |
| 1 worse | 8 | 1 sample was flagged for high density; 4 had differences between BCG levels 3 & 4 | 2 (one flagged for high density) |
| 2 worse | 1 | | 1 |
| Total # Samples | 54 | | 16 |
| % Correct | 61% | | 75% |

* the model assignments were ties between BCG levels 3 & 4, and panelist scores were solid 4s; we considered this to be a difference of 0.5.

5 MMI PERFORMANCE

We examined how the panelist BCG level assignments compared to Connecticut MMI scores (Kanno et al. 2010a) for samples for which MMI scores were available. Figure 5-1 shows box plots of cold water MMI scores grouped by nominal BCG level (panelist consensus) for small-cold samples and mixed water MMI scores grouped by nominal BCG level for medium-large samples. Overall, there is good agreement between the two. In both plots, the MMI scores do not discriminate well between BCG levels 4 and 5, and for the small-cold streams, MMI does not discriminate well between BCG levels 3 and 4.

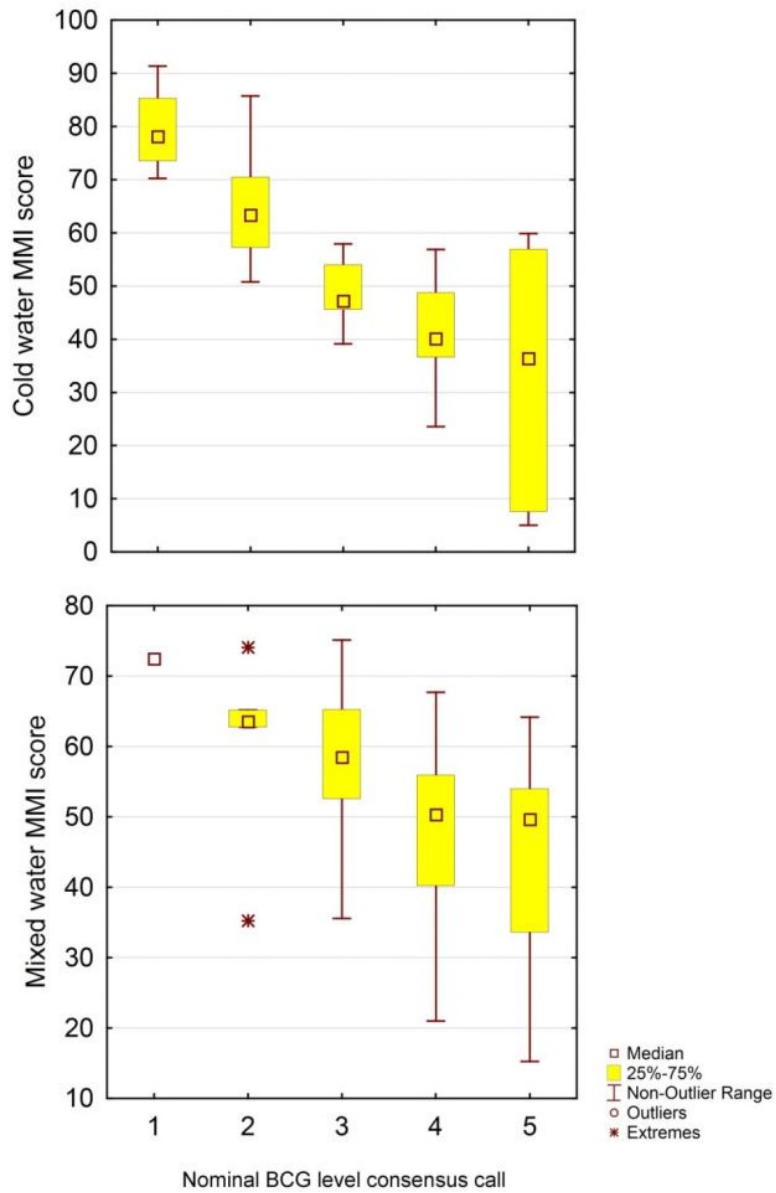


Figure 5-1. Comparison of panelist BCG level consensus assignments for small-cold samples with cold water MMI scores (top), and BCG level calls on medium-large samples with mixed water MMI scores (bottom).

6 DISCUSSION

The development of fish BCG models for small-cold and medium-large freshwater streams in Connecticut marks an important step towards the development of fish community structure metrics that will provide a more quantitative approach to WPLR's assessment process. This was a collective exercise among biologists to develop consensus on assessments of samples. We elicited the rules that the biologists used to assess the samples, and developed a set of quantitative decision criteria rules for assigning fish samples to BCG levels. The regional biologists were able to establish and quantify their differing expectations for small-cold and medium-large streams. These fish BCG models will complement Connecticut's existing macroinvertebrate assessment tools (MMI and BCG) along with the recently developed cold and mixed water fish MMIs, and could potentially serve as a starting point for a regional fish BCG model for New England (e.g., similar to the regional model that was developed for benthic macroinvertebrates (Stamp and Gerritsen 2009). The calibration exercise suggests that the BCG discriminates fair and poor levels of condition (BCG levels 4 and 5) more consistently than a fish MMI (Figure 5-1).

During this process, biologists were faced with a number of challenging questions, including:

- **What is the best classification scheme?** Samples were initially grouped into temperature subclasses (cold, transitional cool, transitional warm) based on TNC's Northeast Aquatic Habitat Classification (Olivero and Anderson 2008). During phase 2 of our work, watershed size was deemed to be a more consistent indicator of temperature class than the TNC designations, and we used 6 square miles as the threshold for separating small-cold from medium-large streams.
- **Do BCG level 1 communities exist in Connecticut streams?** There is not enough information to know what the historical undisturbed assemblage in this region looked like. The biologists felt that additional information, particularly on genetics (stocked vs. native) and age/size class, could help further refine the models (e.g., with active stocking of trout, and without reliable data on genetics and size/class, it is difficult to differentiate between a fish community in BCG level 2 versus 3).
- **How should samples with native vs. non-native trout be assessed?** Non-native trout are regarded as indicators of good water quality and coldwater habitat, but they also represent an altered fish assemblage. For purposes of this exercise, we developed the models in keeping with the traditional BCG levels. The consensus was that BCG level 1 required absence of non-native trout, and that non-native trout could not outnumber natives in BCG Level 2 (this is also consistent with cold and coolwater fish BCG models that have been developed in the Midwest (Gerritsen and Stamp 2012)). During these discussions, differences sometimes arose between those managing for recreational purposes and those managing for native assemblages. Recreational management of highly valued species such as brook, brown, and rainbow trout has a long history in Connecticut. CT DEEP Inland Fisheries Division has a trout management plan with a diverse set of

management objectives and actively stocks these fish into many waters of the state (Hyatt et al. 2000). Stocking occurs for a variety of reasons including: natural reproduction is not adequate to support harvest pressure; streams are capable of supporting trout seasonally but not year round; proximity of the waterbody to population centers; to increase the size and probability of capture; and to supplement populations in streams with marginal habitat.

There were also some lessons learned and blind alleys that we went down, including:

- **Not excluding singletons and doubletons.** We explored using richness metrics in which certain species were excluded if they occurred as singletons (only 1 individual of a species occurs in a sample) or doubletons (2 individuals of a species occurs in a sample). We did this because participants were consciously (or subconsciously) screening samples to exclude what they thought were transient taxa (versus residents). After exploratory analyses and much deliberation, we decided not to use the exclusion metrics because:
 - We lacked the data necessary to draw a clear line between transient vs. resident taxa
 - If rules were being applied, it was not being done in a way that was consistent across participants.
 - The exclusion metrics did not clearly show better performance/discriminatory ability across BCG levels than the original metrics.
- **Not establishing a rule based on an abundance threshold.** We initially tried to make a rule based on total number of individuals. However, we decided against this because it was difficult to set an abundance threshold, especially since expectations were influenced by factors such as watershed size and productivity. Also, panelists did not consistently apply rules related to abundance and sometimes placed more emphasis on the presence of particular species, such as brook trout. In the end, we flagged samples with high and low numbers of individuals since they are considered to be unusual samples that are outside the realm of the BCG model. We also considered including a rule for situations in which there is a hyperabundance of blacknose dace, creek chub, cutlips minnow and/or white suckers (60% was proposed as a threshold). However, this metric did not have enough discriminatory power to include in the final models.

While we were able to accomplish a great deal through this exercise, further work could be done. We conclude by making the following recommendations:

- **Work towards developing models for additional stream types.** There are exceptions to the broad groupings (small-cold and medium-large) that we used for this exercise. Examples include naturally occurring small cool, small warm and medium-large coldwater streams. In the future, if CT DEEP is able to identify, describe and classify these 'exceptions' and has sufficient samples to assess them, models could be developed for these additional stream types. CT DEEP could also explore developing a warm-water

model, although the scarcity of high quality warm-water streams in Connecticut would make this challenging.

- **Gather additional information on genetics (stocked vs. native) and age/size class.**
The biologists felt that this type of information could help further refine the models (e.g., with active stocking of trout, and without reliable data on genetics and size/class, it is difficult to differentiate between a fish community in BCG level 2 versus 3). If resources permit, we recommend that CT DEEP start to consistently gather information on genetics (stocked vs. native) and age/size class to help inform future assessments.

7 LITERATURE CITED

Barbour, M.T., J. Gerritsen, B.D. Snyder, and J.B. Stribling. 1999. Rapid Bioassessment Protocols for Use in Streams and Wadeable Rivers: Periphyton, Benthic Macroinvertebrates and Fish. Second Edition. EPA/841-B-99-002. U.S. Environmental Protection Agency, Office of Water, Washington, D.C.

Beauchene, M., Becker, M., Belluci, C., Hagstrom, N. and Y. Kanno. 2012. Defining Water Temperature Habitat Types; Connecticut Inland Streams and Rivers – Draft. Connecticut Department of Energy and Environmental Protection.

Castella, E. and M.C.D. Speight. 1996. Knowledge representation using fuzzy coded variables: an example based on the use of Syrphidae (Insecta, Diptera) in the assessment of riverine wetlands. *Ecological Modelling* 85:13-25.

Connecticut DEP. 2005. Connecticut Comprehensive Ambient Water Quality Monitoring Strategy. Connecticut Department of Environmental Protection, Bureau of Water Management, Hartford, CT.

Connecticut DEEP. 2013. Standard Operating Procedures For the Collection of Fish Community Data From Wadeable Streams For Aquatic Life Assessment. Connecticut Department of Energy and Environmental Protection, Bureau of Water Protection and Land Resuse, Hartford, CT.

Davies, S.B., and S. K. Jackson. 2006. The Biological Condition Gradient: A descriptive model for interpreting change in aquatic ecosystems. *Ecological Applications* 16(4):1251–1266.

Demicco, R.V. 2004. Fuzzy Logic and Earth Science: An Overview. In *Fuzzy Logic in Geology*, R.V. Demicco and G.J. Klir (eds.), pp. 11-61. Elsevier Academic Press, San Diego, CA.

Demicco, R.V. and G.J. Klir. 2004. Introduction. In *Fuzzy Logic in Geology*, R.V. Demicco and G.J. Klir (eds.), pp. 1-10. Elsevier Academic Press, San Diego, CA.

Droesen, W.J. 1996. Formalisation of ecohydrological expert knowledge applying fuzzy techniques. *Ecological Modelling* 85:75-81.

Gerritsen, J. and B. Jessup. 2007. Calibration of the Biological Condition Gradient for High Gradient Streams of Connecticut. Prepared for the U.S. EPA Office of Science and Technology and Connecticut Department of Environmental Protection. Prepared by Tetra Tech, Inc., Owings Mills, MD. 86 pp.

Gerritsen, J. and J. Stamp. 2012. Calibration of the Biological Condition Gradient (BCG) in Cold and Cool Waters of the Upper Midwest: Fish and Benthic Macroinvertebrate Assemblages. Prepared for U.S. EPA Office of Science and Technology and U.S. EPA Region 5; prepared by Tetra Tech, Inc. 400 Red Brook Blvd., Suite 200, Owings Mills, MD 21117.

Halliwell, D.B., Langdon, R.W., Daniels, R.A., Kurtenbach, J.P. and R.A. Jacobson. 1999. Classification of freshwater fish species of the Northeastern United States for use in the development of indices of biological integrity, with regional applications. In *Assessing the Sustainability and Biological Integrity of Water Resources Using Fish Communities*. CRC Press, LLC. Pp. 301-337.

Hyatt, W., M. Humphreys, and N. T. Hagstrom, 2000. A Trout Management Plan for Connecticut Rivers and Streams. Conn. D.E.P. Bureau of Natural Resources, Inland Fisheries Division, Hartford, CT. 50p.

Ibelings, B.W., M Vonk, H.F.J. Los, D.T. Van Der Molen, and W.M. Mooij. 2003. Fuzzy modeling of Cyanobacterial surface waterblooms: validation with NOAA-AVHRR satellite images. *Ecological Applications* 13:1456-1472.

Kanno, Y., Vokoun, J.C. and M. Beauchene. 2010a. Development of dual fish multi-metric indices of biological condition for streams with characteristic thermal gradients and low species richness. *Ecological Indicators* 10(3): 565-571.

Kanno, Y., and J.C. Vokoun. 2010b. Evaluating effects of water withdrawals and impoundments on fish assemblages in southern New England streams, USA. *Fisheries Management and Ecology* 17: 272–283.

Kanno, Y., and J.C. Vokoun. 2008. Biogeography of Stream Fishes in Connecticut: Defining Faunal Regions and Assemblage Types. *Northeastern Naturalist* 15(4):557–576.

Karr, J.R. and E.W. Chu. 1999. Restoring Life. In *Running Waters: Better Biological Monitoring*. Island Press, Washington, DC.

Klir, G.J. 2004. Fuzzy Logic: A Specialized Tutorial. In *Fuzzy Logic in Geology*, R.V. Demicco and G.J. Klir (eds.), pp. 11-61. Elsevier Academic Press, San Diego, CA.

Lyons, J., Zorn, T., Stewart, J., Seelbach, P., Wehrly, K. and L. Wang. 2009. Defining and Characterizing Coolwater Streams and Their Fish Assemblages in Michigan and Wisconsin, USA. *North American Journal of Fisheries Management* 29:1130–1151

Olivero, A.P. and Anderson, M.G. 2008. Northeast Aquatic Habitat Classification. The Nature Conservancy in Collaboration with the Northeast Association of Fish and Wildlife Agencies. 86 pp.

Omernik, J.M. 1987. Ecoregions of the conterminous United States. Map (scale 1:7,500,000). *Annals of the Association of American Geographers* 77(1):118-125.

Omernik, J.M. 1995. Ecoregions: A spatial framework for environmental management. In: *Biological Assessment and Criteria: Tools for Water Resource Planning and Decision Making*. Davis, W.S. and T.P. Simon (eds.), Lewis Publishers, Boca Raton, FL. p. 49-62.

Simpson, J.C. and R.H. Norris. 2000. Biological assessment of river quality: development of AusRivAS models and outputs. In *Assessing the Biological Quality of Fresh Waters: RIVPACS and Other Techniques*. J.F. Wright, D.W. Sutcliffe and M.T. Furse (eds.), pp. 125-142. Freshwater Biological Association, Ambleside, UK.

Stamp, J. and J. Gerritsen 2009. Methods and Assessment Comparability Among State and Federal Biological Monitoring Protocols. Prepared for New England Interstate Water Pollution Control Commission (NEIWPCC), Boott Mills South, 100 Foot of John Street, Lowell, MA 01852.

U.S. EPA. 2005. Use of biological information to tier designated aquatic life uses in state and tribal water quality standards. EPA-822-R-05-001.

U.S. EPA. 2007. The New England Wadeable Stream Survey (NEWS): Development of Common Assessments in the Framework of the Biological Condition Gradient. Prepared for the US EPA Office of Science and Technology and the US EPA Office of Watersheds.

Vannote, R. L., Minshall, G. W., Cummins, K. W., Sedell, J.R., and Cushing, C. E. 1980. The River Continuum Concept. *Canadian Journal of Fisheries and Aquatic Sciences* 37:130-137.

Vermont DEC. 2004. Wadeable Stream Biocriteria Development and Implementation Methods for Fish and Macroinvertebrate Assemblages in Vermont Wadeable Streams and Rivers. Water Quality Division, Biomonitoring and Aquatic Studies Section. 83 pp.

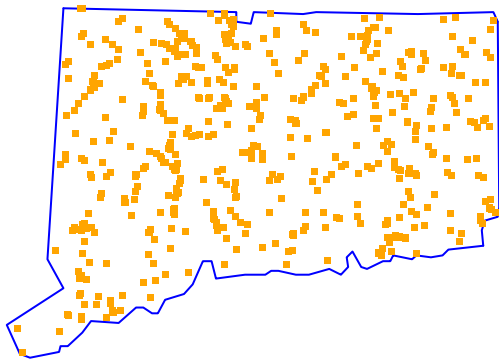
Whittier, T.R., Hughes, R.M., Stoddard, J.L., Lomnický, G.A., Peck, D.V., Herlihy, A.T., 2007. A structured approach for developing indices of biotic integrity: three examples from streams and rivers in the western USA. *Trans. Am. Fish. Soc.* 136, 718–735.

Wright, J.F. 2000. An introduction to RIVPACS. In *Assessing the Biological Quality of Fresh Waters: RIVPACS and Other Techniques*. J.F. Wright, D.W. Sutcliffe and M.T. Furse (eds.), pp. 1-24. Freshwater Biological Association, Ambleside, UK.

APPENDIX A

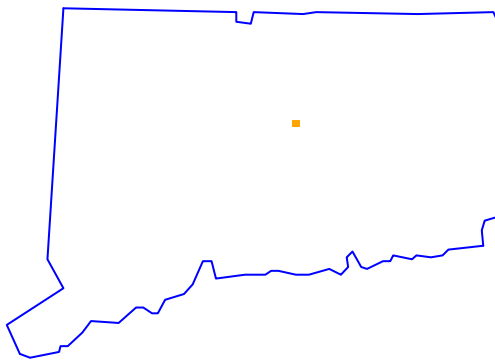
Distribution maps

Connecticut



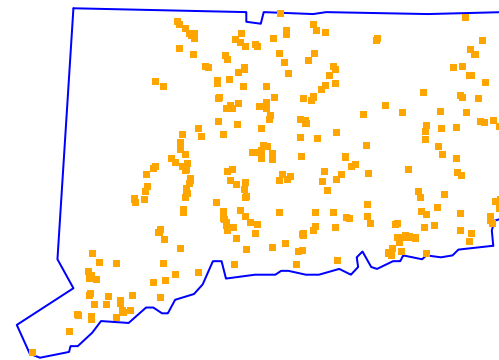
N = 790

American Brook lamprey



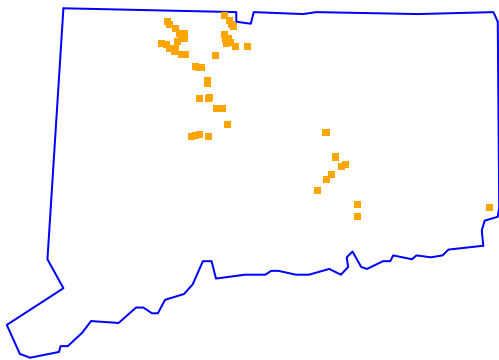
N = 2

American eel



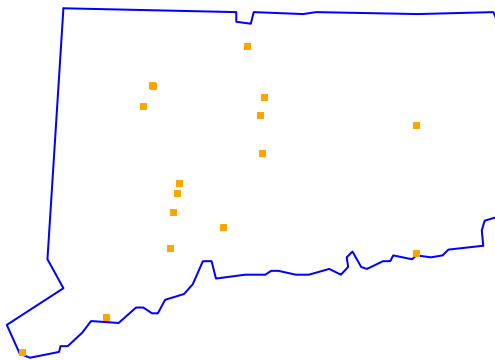
N = 427

Atlantic salmon, fry stocked



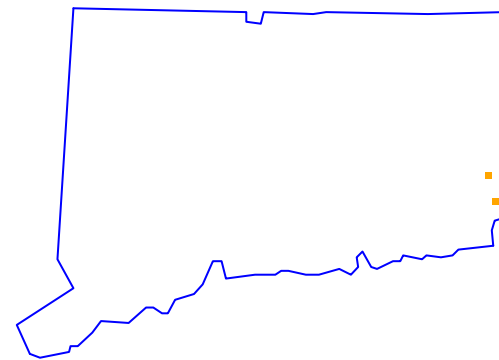
N = 81

Banded killifish



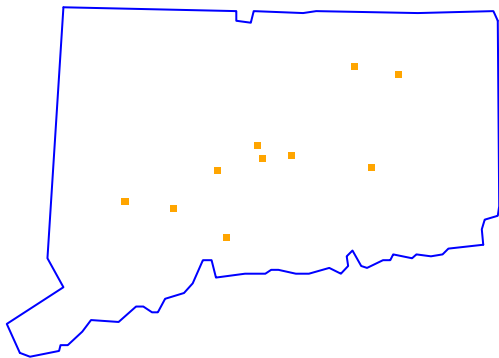
N = 19

Banded Sunfish



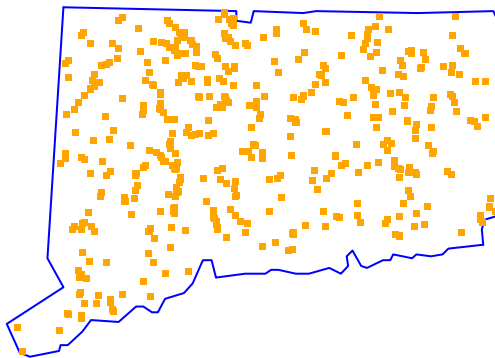
N = 3

Black Crappie



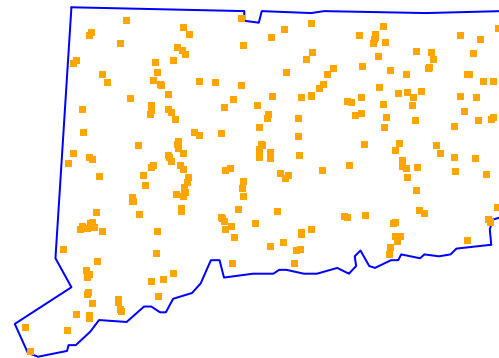
N = 12

Blacknose dace



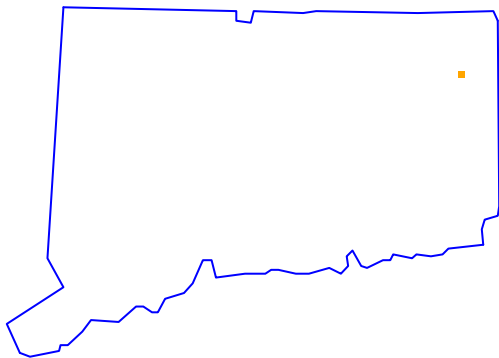
N = 648

Bluegill sunfish



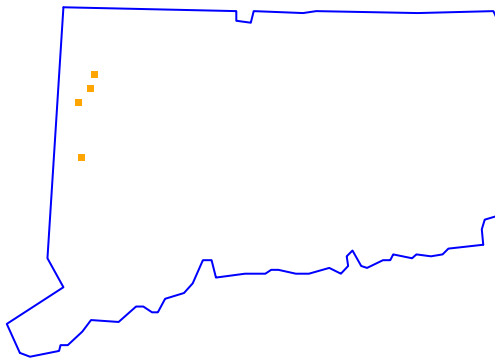
N = 320

Bluegill X Pumpkinseed



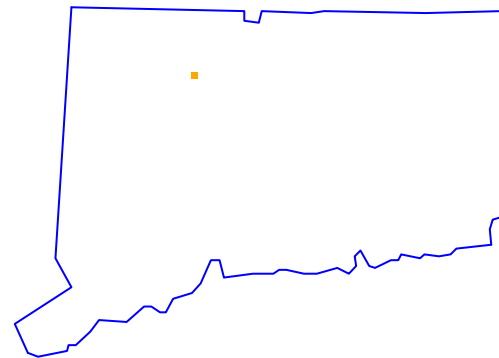
N = 1

Bluntnose minnow



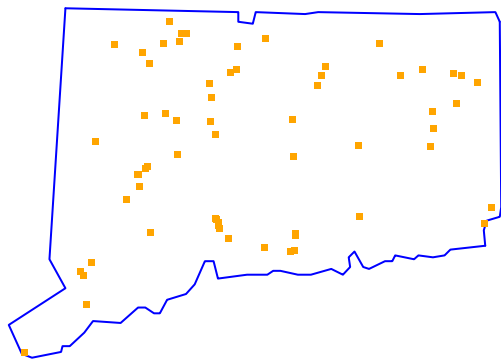
N = 4

Bridled shiner



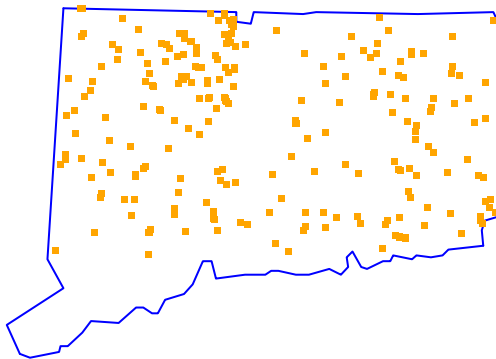
N = 1

Brook trout, stocked



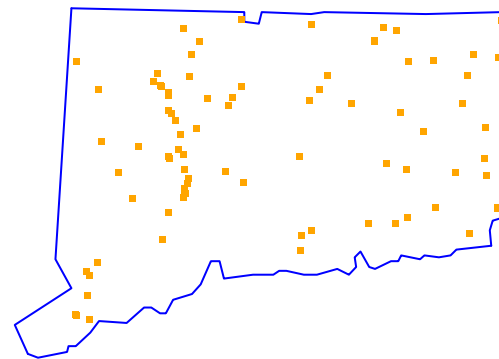
N = 73

Brook trout, wild



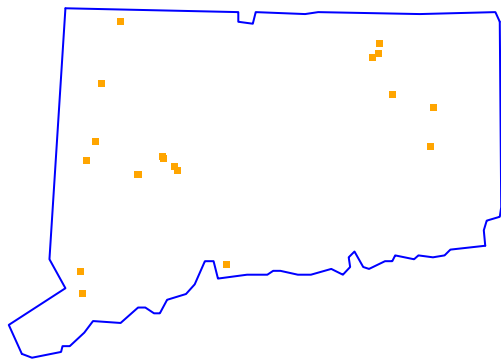
N = 316

Brown bullhead



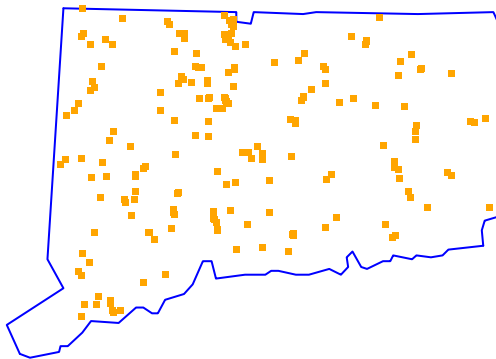
N = 98

Brown trout, Fry Stocked



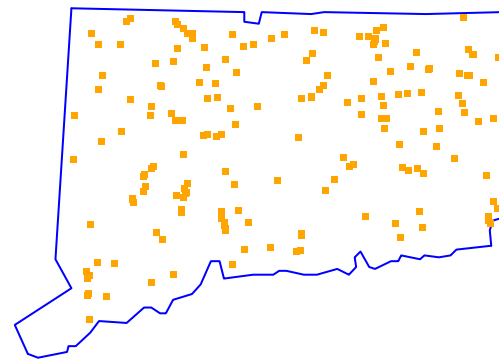
N = 28

Brown trout, naturalized



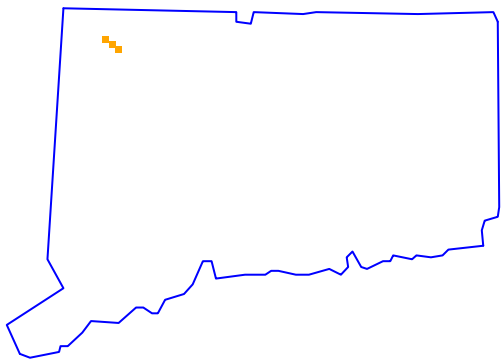
N = 235

Brown trout, stocked



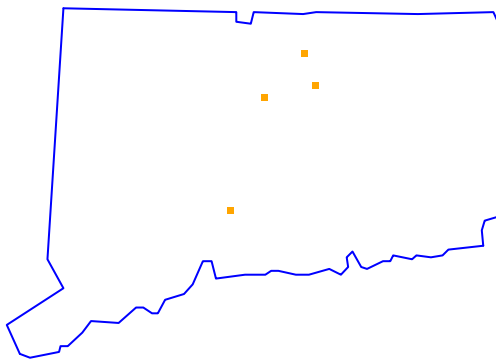
N = 233

Burbot



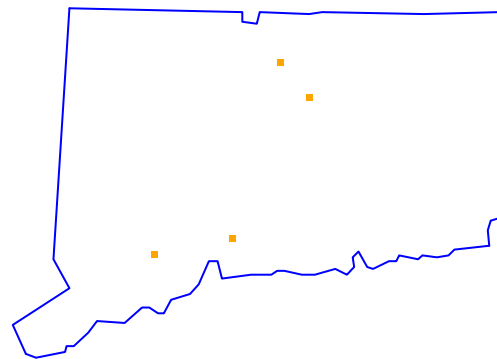
N = 4

Carp



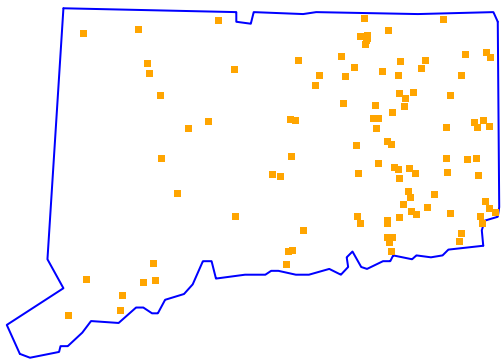
N = 4

Central mudminnow



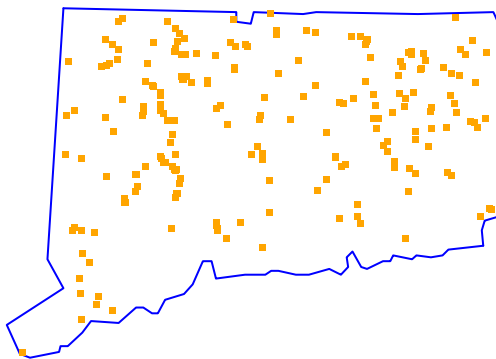
N = 4

Chain pickerel



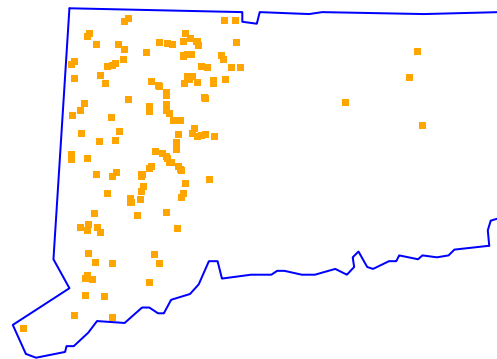
N = 124

Common shiner



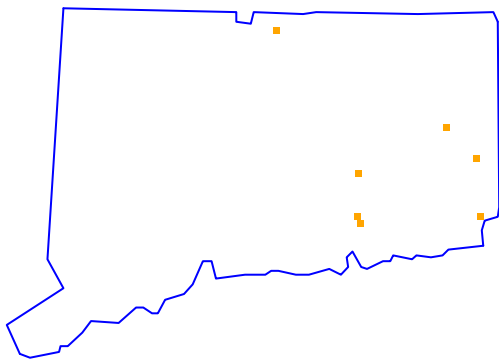
N = 288

Creek chub



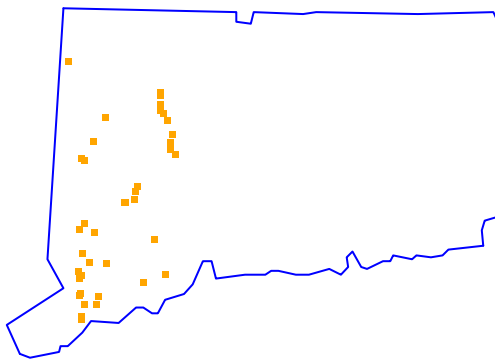
N = 189

Creek chubsucker



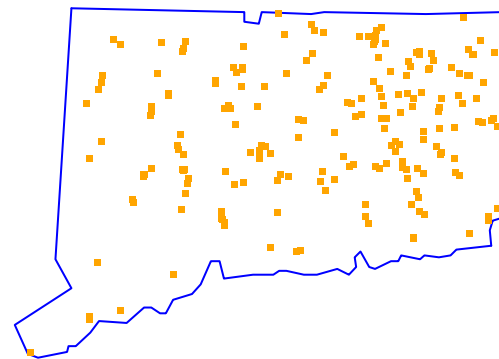
N = 7

Cutlips minnow



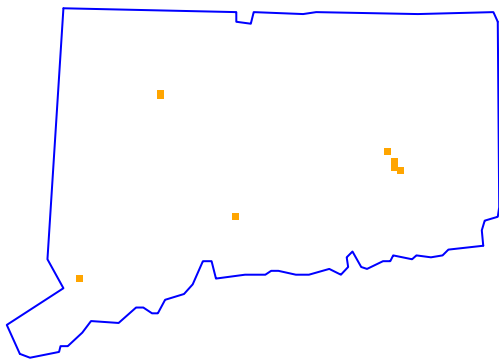
N = 59

Fallfish



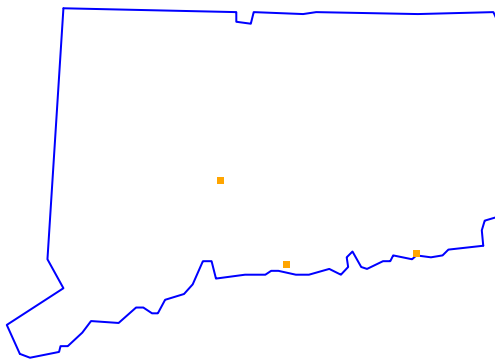
N = 296

Fathead minnow



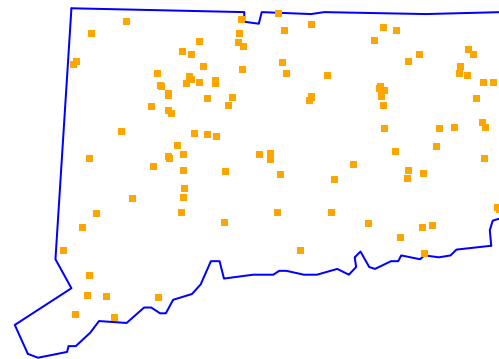
N = 8

Fourspine stickleback



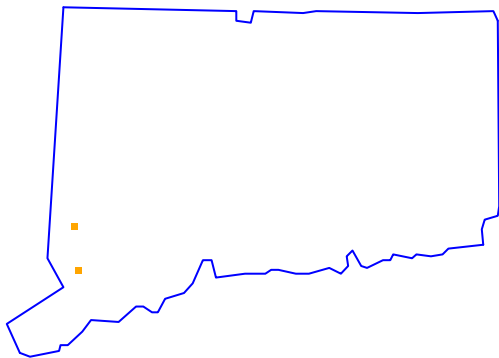
N = 3

Golden shiner



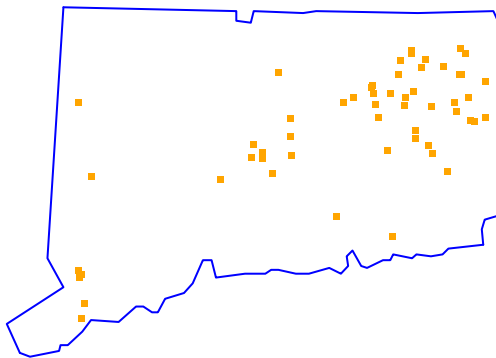
N = 128

Goldfish



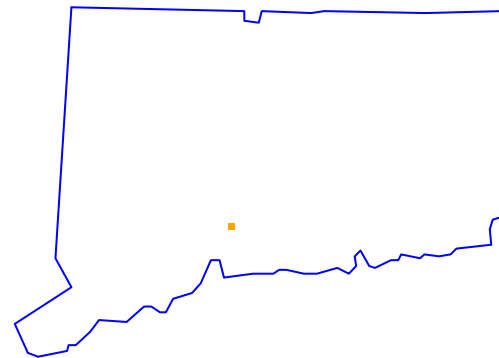
N = 2

Green sunfish



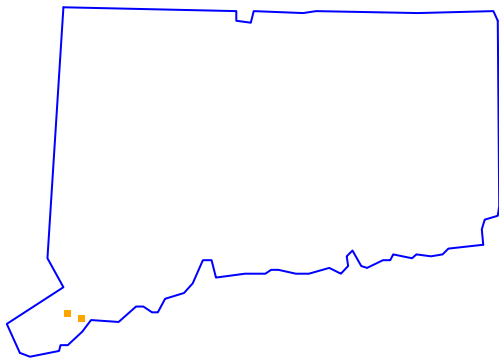
N = 77

Hogchoker



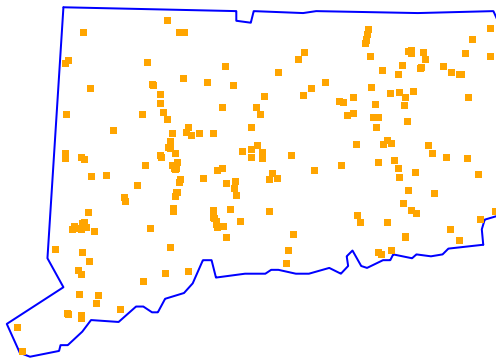
N = 1

Hybrid sunfish



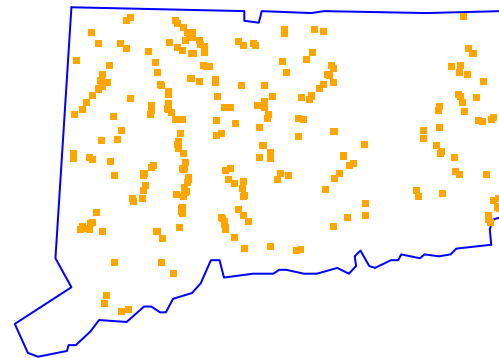
N = 2

Largemouth Bass



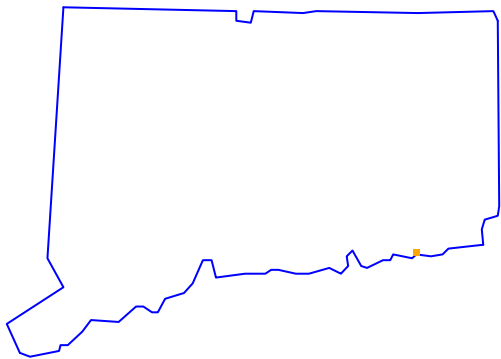
N = 246

Longnose dace



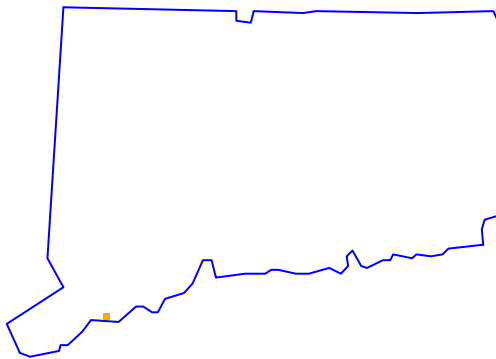
N = 395

Mummichog



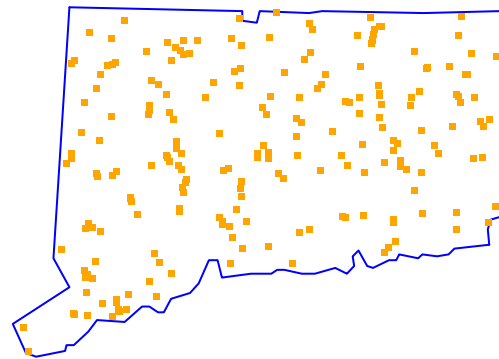
N = 2

Ninespine stickleback



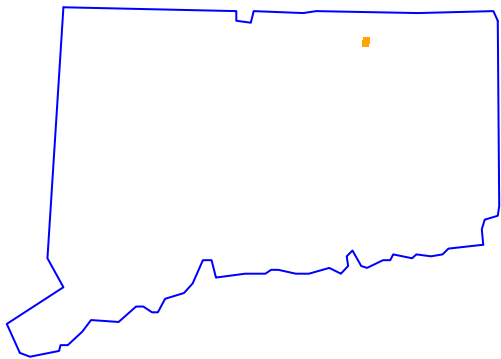
N = 1

Pumpkinseed



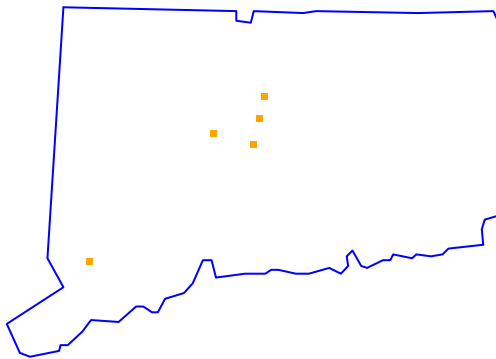
N = 266

Pumpkinseed Hybrid



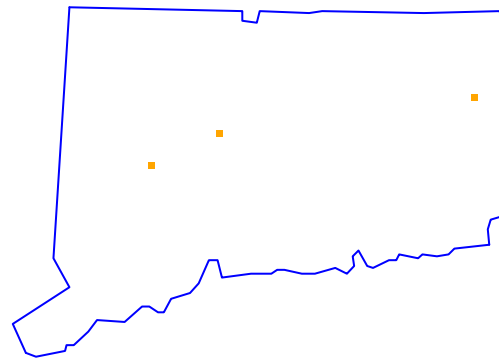
N = 2

Pumpkinseed X Red breast



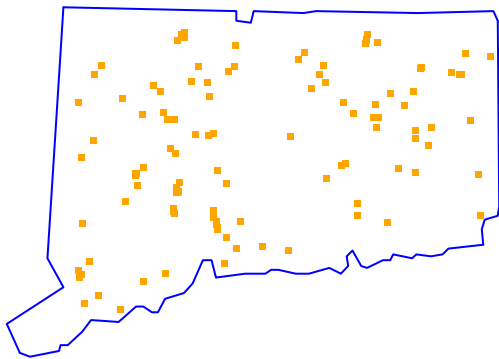
N = 5

Rainbow trout, feral



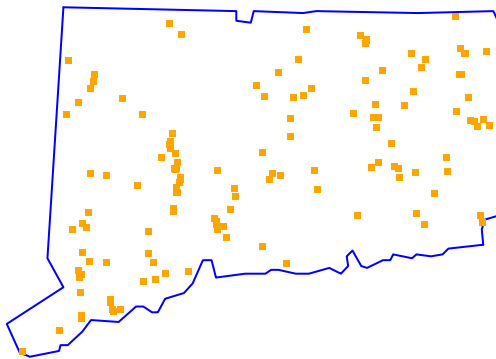
N = 3

Rainbow trout, Stocked



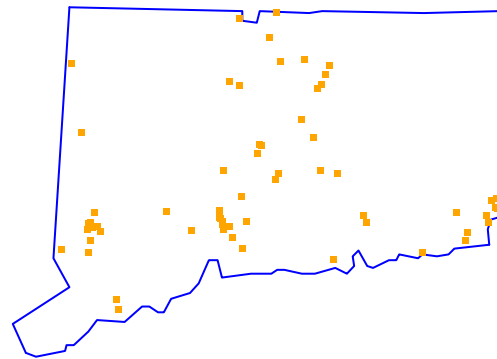
N = 136

Redbreast sunfish



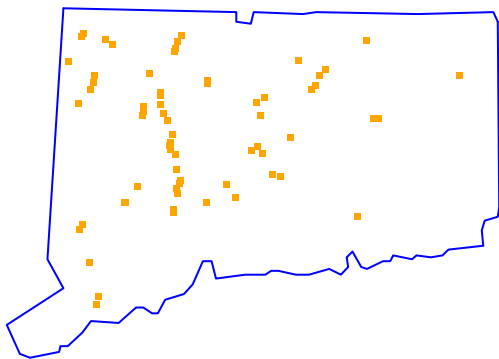
N = 210

Redfin pickerel



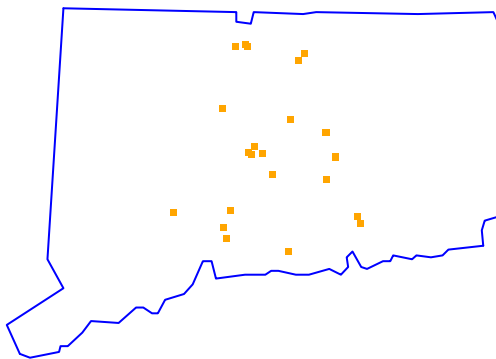
N = 75

Rock Bass



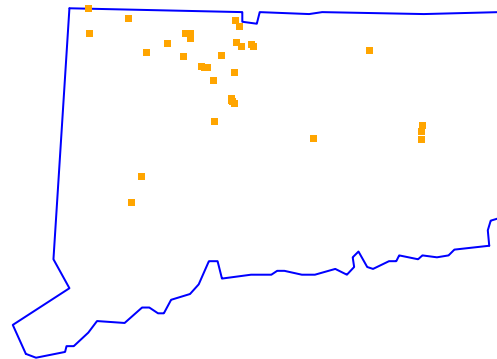
N = 98

Sea lamprey



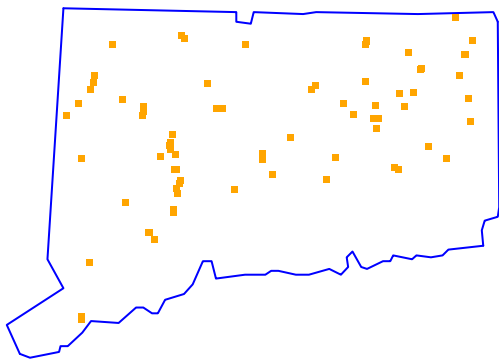
N = 29

Slimy sculpin



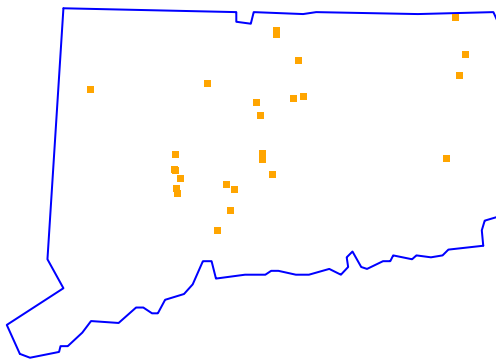
N = 51

Smallmouth bass



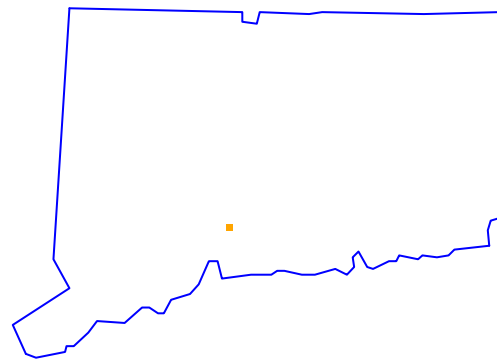
N = 130

Spottail shiner



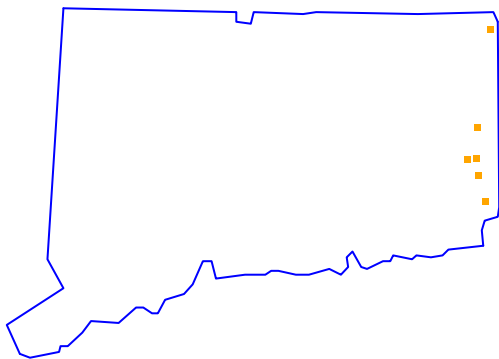
N = 35

Striped bass



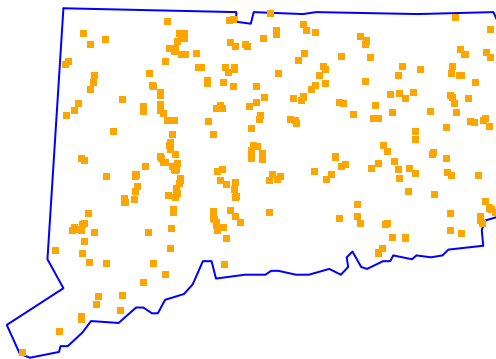
N = 1

Swamp darter



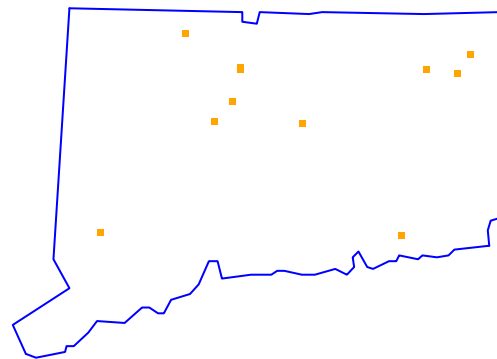
N = 7

Tessellated darter



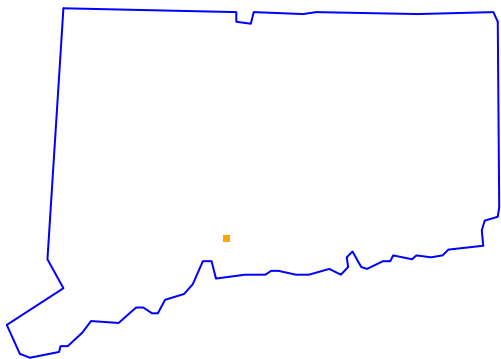
N = 429

Tiger Trout



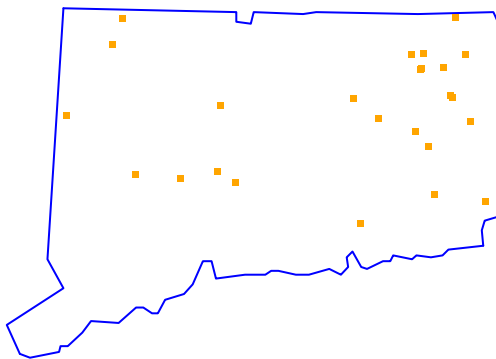
N = 11

Tomcod



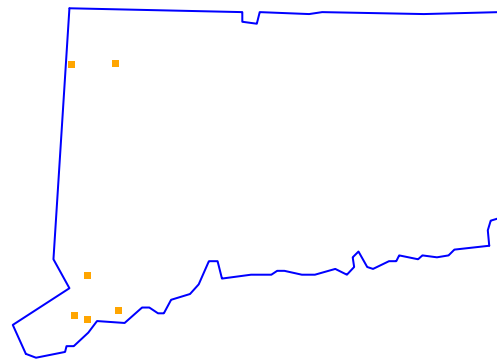
N = 1

Unknown minnow



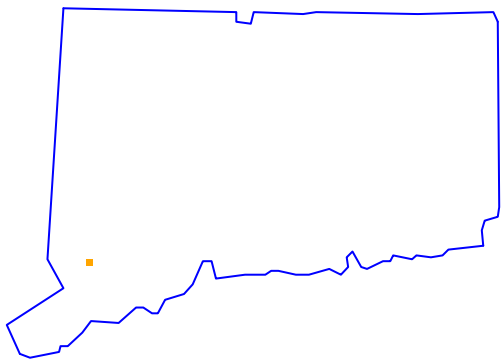
N = 26

Unknown sunfish



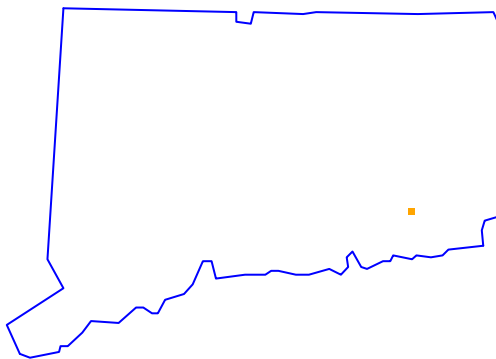
N = 6

Walleye



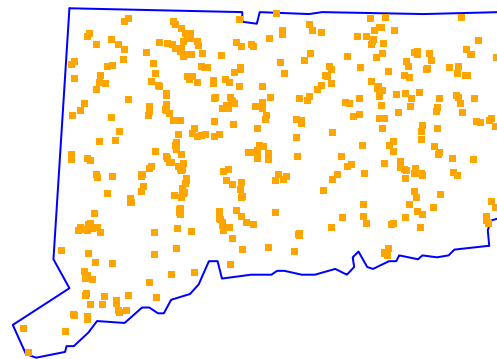
N = 2

White catfish



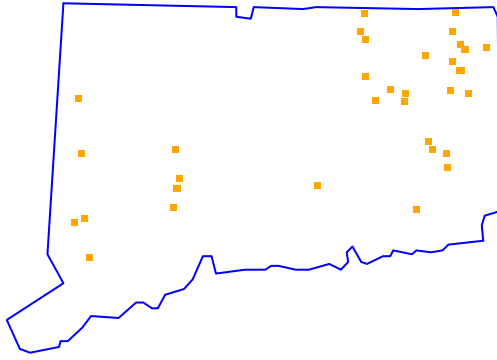
N = 1

White sucker



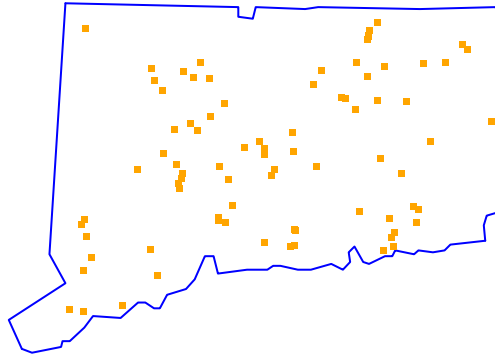
N = 604

Yellow bullhead



N = 51

Yellow perch



N = 100

APPENDIX B

Additional BCG Background Information

Table B1. Narrative descriptions of the 10 attributes that distinguish the six tiers of the Biological Condition Gradient (Davies and Jackson 2006).

| Biological Condition Gradient Tiers | | | | | | |
|---|---|--|---|--|---|---|
| | 1 | 2 | 3 | 4 | 5 | 6 |
| | Natural or native condition | Minimal changes in structure of the biotic community and minimal changes in ecosystem function | Evident changes in structure of the biotic community and minimal changes in ecosystem function | Moderate changes in structure of the biotic community and minimal changes in ecosystem function | Major changes in structure of the biotic community and moderate changes in ecosystem function | Severe changes in structure of the biotic community and major loss of ecosystem function |
| General Description of Biological Condition | | | | | | |
| Attributes | Native structural, functional and taxonomic integrity is preserved; ecosystem function is preserved within the range of natural variability | Virtually all native taxa are maintained with some changes in biomass and/or abundance; ecosystem functions are fully maintained within the range of natural variability | Some changes in structure due to loss of some rare native taxa; shifts in relative abundance of taxa but sensitive-ubiquitous taxa are common and abundant; ecosystem functions are fully maintained through redundant attributes of the system | Moderate changes in structure due to replacement of some sensitive-ubiquitous taxa by more tolerant taxa, but reproducing populations of some sensitive taxa are maintained; overall balanced distribution of all expected major groups; ecosystem functions largely maintained through redundant attributes | Sensitive taxa are markedly diminished; conspicuously unbalanced distribution of major groups from that expected; organism condition shows signs of physiological stress; system function shows reduced complexity and redundancy; increased build-up or export of unused materials | Extreme changes in structure; wholesale changes in taxonomic composition; extreme alterations from normal densities and distributions; organism condition is often poor; ecosystem functions are severely altered |

Table B1. Continued...

| | 1 | 2 | 3 | 4 | 5 | 6 |
|--|--|--|--|---|---|--|
| | Natural or native condition | Minimal changes in structure of the biotic community and minimal changes in ecosystem function | Evident changes in structure of the biotic community and minimal changes in ecosystem function | Moderate changes in structure of the biotic community and minimal changes in ecosystem function | Major changes in structure of the biotic community and moderate changes in ecosystem function | Severe changes in structure of the biotic community and major loss of ecosystem function |
| I Historically documented, sensitive, long-lived or regionally endemic taxa | As predicted for natural occurrence except for global extinctions | As predicted for natural occurrence except for global extinctions | Some may be absent due to global extinction or local extirpation | Some may be absent due to global, regional or local extirpation | Usually absent | Absent |
| II Sensitive-rare taxa | As predicted for natural occurrence, with at most minor changes from natural densities | Virtually all are maintained with some changes in densities | Some loss, with replacement by functionally equivalent sensitive-ubiquitous taxa | May be markedly diminished | Absent | Absent |
| III Sensitive-ubiquitous taxa | As predicted for natural occurrence, with at most minor changes from natural densities | Present and may be increasingly abundant | Common and abundant; relative abundance greater than sensitive-rare taxa | Present with reproducing populations maintained; some replacement by functionally equivalent taxa of intermediate tolerance | Frequently absent or markedly diminished | Absent |

Table B1. Continued...

| | 1 | 2 | 3 | 4 | 5 | 6 |
|---|---|--|---|---|---|---|
| | Natural or native condition | Minimal changes in structure of the biotic community and minimal changes in ecosystem function | Evident changes in structure of the biotic community and minimal changes in ecosystem function | Moderate changes in structure of the biotic community and minimal changes in ecosystem function | Major changes in structure of the biotic community and moderate changes in ecosystem function | Severe changes in structure of the biotic community and major loss of ecosystem function |
| IV Taxa of intermediate tolerance | As predicted for natural occurrence, with at most minor changes from natural densities | As naturally present with slight increases in abundance | Often evident increases in abundance | Common and often abundant; relative abundance may be greater than sensitive-ubiquitous taxa | Often exhibit excessive dominance | May occur in extremely high or extremely low densities; richness of all taxa is low |
| V Tolerant taxa | As predicted for natural occurrence, at most minor changes from natural densities | As naturally present with slight increases in abundance | May be increases in abundance of functionally diverse tolerant taxa | May be common but do not exhibit significant dominance | Often occur in high densities and may be dominant | Usually comprise the majority of the assemblage; often extreme departures from normal densities (high or low) |
| VI Non-native or intentionally introduced taxa | Non-native taxa, if present, do not displace native taxa or alter native structural or functional integrity | Non-native taxa may be present, but occurrence has a non-detrimental effect on native taxa | Sensitive or intentionally introduced non-native taxa may dominate some assemblages (e.g., fish or macrophytes) | Some replacement of sensitive non-native taxa with functionally diverse assemblage of non-native taxa of intermediate tolerance | Some assemblages (e.g., fish or macrophytes) are dominated by tolerant non-native taxa | Often dominant; may be the only representative of some assemblages (e.g., plants, fish, bivalves) |

Table B1. Continued...

| | 1 | 2 | 3 | 4 | 5 | 6 |
|--|---|--|--|---|---|---|
| | Natural or native condition | Minimal changes in structure of the biotic community and minimal changes in ecosystem function | Evident changes in structure of the biotic community and minimal changes in ecosystem function | Moderate changes in structure of the biotic community and minimal changes in ecosystem function | Major changes in structure of the biotic community and moderate changes in ecosystem function | Severe changes in structure of the biotic community and major loss of ecosystem function |
| VII Organism condition (especially of long-lived organisms) | Any anomalies are consistent with naturally occurring incidence and characteristics | Any anomalies are consistent with naturally occurring incidence and characteristics | Anomalies are infrequent | Incidence of anomalies may be slightly higher than expected | Biomass may be reduced; anomalies increasingly common | Long-lived taxa may be absent; biomass reduced; anomalies common and serious; minimal reproduction except for extremely tolerant groups |
| VIII Ecosystem functions | All are maintained within the range of natural variability | All are maintained within the range of natural variability | Virtually all are maintained through functionally redundant system attributes; minimal increase in export except at high storm flows | Virtually all are maintained through functionally redundant system attributes though there is evidence of loss of efficiency (e.g., increased export or decreased import) | Apparent loss of some ecosystem functions manifested as increased export or decreased import of some resources, and changes in energy exchange rates (e.g., P/R, decomposition) | Most functions show extensive and persistent disruption |
| IX Spatial and temporal extent of detrimental effects | N/A A natural disturbance regime is maintained | Limited to small pockets and short duration | Limited to the reach scale and/or limited to within a season | Mild detrimental effects may be detectable beyond the reach scale and may include more than one season | Detrimental effects extend far beyond the reach scale leaving only a few islands of adequate conditions; effect extends across multiple seasons | Detrimental effects may eliminate all refugia and colonization sources within the catchment and affect multiple seasons |

Table B1. Continued...

| | 1 | 2 | 3 | 4 | 5 | 6 |
|----------------------------|---|--|--|---|---|---|
| | Natural or native condition | Minimal changes in structure of the biotic community and minimal changes in ecosystem function | Evident changes in structure of the biotic community and minimal changes in ecosystem function | Moderate changes in structure of the biotic community and minimal changes in ecosystem function | Major changes in structure of the biotic community and moderate changes in ecosystem function | Severe changes in structure of the biotic community and major loss of ecosystem function |
| X Ecosystem connectance | System is highly connected in space and time, at least annually | Ecosystem connectance is unimpaired | Slight loss of connectance but there are adequate local recolonization sources | Some loss of connectance but colonization sources and refugia exist within the catchment | Significant loss of ecosystem connectance is evident; recolonization sources do not exist for some taxa | Complete loss of ecosystem connectance in at least one dimension (i.e., longitudinal, lateral, vertical, or temporal) lowers reproductive success of most groups; frequent failures in reproduction and recruitment |

Table B2. Ecological attributes used to develop the BCG.

| Attribute | Description |
|-----------|--|
| I | <p>Historically documented, sensitive, long-lived or regionally endemic taxa</p> <p><i>“Historically documented” refers to taxa known to have been supported in a waterbody or region prior to enactment of the 1972 Clean Water Act, according to historical records compiled by State or federal agencies or published scientific literature.</i></p> |
| II | <p>Highly Sensitive Taxa (note: this was identified as “Sensitive-Rare taxa” in Davies and Jackson 2006)</p> <p><i>These are taxa that naturally occur in low numbers relative to total population density but may make up a large relative proportion of richness. In high quality sites, they may be ubiquitous in occurrence or may be restricted to certain micro-habitats. Many of these species commonly occur at low densities, thus their occurrence is dependent on sample effort. Often stenothermic (having a narrow range of thermal tolerance) or cold-water obligates; commonly k-strategists (populations maintained at a fairly constant level; slower development; longer life-span). They may have specialized food resource needs or feeding strategies, and are generally intolerant to significant alteration of the physical or chemical environment. They are often the first taxa observed to be lost from a community following moderate disturbance or pollution (Figure 3-1).</i></p> |
| III | <p>Intermediate Sensitive Taxa (or Sensitive and Common Taxa)</p> <p><i>These taxa are ordinarily common and abundant in natural communities when conventional sampling methods are used (Figure 3-1). They often have a broader range of tolerances than highly sensitive taxa, and usually occur in reduced abundance and reduced frequencies at disturbed or polluted sites. These are taxa that comprise a substantial portion of natural communities, and that often exhibit negative response (loss of population, richness) at mild pollution loads or habitat alteration.</i></p> |
| IV | <p>Taxa of Intermediate Tolerance</p> <p><i>Taxa that comprise a substantial portion of natural communities; may be r-strategists (early colonizers with rapid turn-over times; e.g., “boom/bust population characteristics or they may be eurythermal (having a broad thermal tolerance range). Many have generalist or facultative feeding strategies enabling utilization of diverse food types. They are readily collected with conventional sample methods. These species have little or no detectable response to a stress gradient (Figure 3-1), and are often equally abundant in both reference and stressed sites. Some intermediate taxa may show an “intermediate disturbance” response, where densities and frequency of occurrence are highest at intermediate levels of stress but are intolerant of excessive pollution loads or habitat alteration. These taxa are readily collected with conventional sample methods.</i></p> |

| | |
|----|---|
| V | <p>Tolerant Taxa</p> <p><i>Taxa that comprise a low proportion of natural communities. Taxa often are tolerant of a greater degree of disturbance and stress than other organisms and are thus resistant to a variety of pollution or habitat induced stress. They may increase in number (sometimes greatly) under severely altered or stressed conditions, and may possess adaptations for highly enriched conditions, hypoxia, or toxic substances (Figure 3-1). Commonly r-strategists (early colonizers with rapid turn-over times: e.g., “boom/bust” population characteristics), these are the last survivors in severely disturbed systems.</i></p> |
| VI | <p>Non-native taxa</p> <p><i>With respect to a particular ecosystem, any species not native to that ecosystem. Species introduced or spread from one region of the U.S. to another outside their normal range are non-native or non-indigenous, as are species introduced from other continents.</i></p> <p>Vla – Highly tolerant non-native taxa</p> |
| X | <p>Catadromous fish, indicating ecosystem connectivity</p> <p><i>Access or linkage (in space/time) to materials, locations, and conditions required for maintenance of interacting populations of aquatic life; the opposite of fragmentation; necessary for metapopulation maintenance and natural flows of energy and nutrients across ecosystem boundaries.</i></p> |

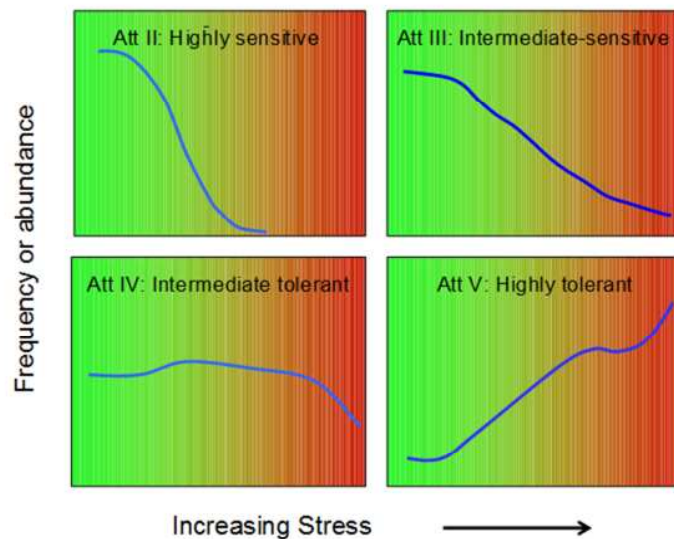


Figure B1. Diagram showing responses of attributes to a stress gradient.

APPENDIX C

Fish BCG Attribute Assignments

Appendix C. Table C -1. BCG attribute assignments for fish. Assignments are the same for both small-cold and medium-large subclasses. This list is sorted first by family, then by common name.

| BCG Attribute | Order | Family | Scientific Name | Common Name | Total # individuals |
|---------------|-------------------|---------------|------------------------|--------------------------|---------------------|
| x | Pleuronectiformes | Achiridae | Trinectes maculatus | Hogchoker | 4 |
| 10 | Anguilliformes | Anguillidae | Anguilla rostrata | American eel | 10397 |
| 2 | Cypriniformes | Catostomidae | Erimyzon oblongus | Creek chubsucker | 15 |
| 4 | Cypriniformes | Catostomidae | Catostomus commersoni | White sucker | 23426 |
| x | Perciformes | Centrarchidae | Centrarchidae | Unknown sunfish | 11 |
| x | Perciformes | Centrarchidae | Enneacanthus obesus | Banded Sunfish | 5 |
| 6 | Perciformes | Centrarchidae | Pomoxis nigromaculatus | Black Crappie | 19 |
| 6a | Perciformes | Centrarchidae | Lepomis macrochirus | Bluegill sunfish | 2784 |
| x | Perciformes | Centrarchidae | | Bluegill X Pumpkinseed | 2 |
| 6a | Perciformes | Centrarchidae | Lepomis cyanellus | Green sunfish | 364 |
| x | Perciformes | Centrarchidae | | Hybrid sunfish | 4 |
| 6a | Perciformes | Centrarchidae | Micropterus salmoides | Largemouth Bass | 1648 |
| 4 | Perciformes | Centrarchidae | Lepomis gibbosus | Pumpkinseed | 1764 |
| x | Perciformes | Centrarchidae | | Pumpkinseed Hybrid | 7 |
| x | Perciformes | Centrarchidae | | Pumpkinseed X Red breast | 20 |
| 4 | Perciformes | Centrarchidae | Lepomis auritus | Redbreast sunfish | 4485 |
| 6 | Perciformes | Centrarchidae | Ambloplites rupestris | Rock Bass | 801 |
| 6 | Perciformes | Centrarchidae | Micropterus dolomieu | Smallmouth bass | 2285 |
| 2 | Scorpaeniformes | Cottidae | Cottus cognatus | Slimy sculpin | 2194 |
| 4 | Cypriniformes | Cyprinidae | Rhinichthys atratulus | Blacknose dace | 55137 |
| 6a | Cypriniformes | Cyprinidae | Pimephales notatus | Bluntnose minnow | 11 |

Table C -1 continued...

| BCG Attribute | Order | Family | Scientific Name | Common Name | Total # individuals |
|----------------------|--------------------|----------------|-------------------------|-----------------------|----------------------------|
| x | Cypriniformes | Cyprinidae | Notropis bifrenatus | Bridled shiner | 3 |
| 6a | Cypriniformes | Cyprinidae | Cyprinus carpio | Carp | 13 |
| 3 | Cypriniformes | Cyprinidae | Luxilus cornutus | Common shiner | 9046 |
| 4 | Cypriniformes | Cyprinidae | Semotilus atromaculatus | Creek chub | 4974 |
| 4 | Cypriniformes | Cyprinidae | Exoglossum maxillingua | Cutlips minnow | 2552 |
| 3 | Cypriniformes | Cyprinidae | Semotilus corporalis | Fallfish | 10020 |
| 6a | Cypriniformes | Cyprinidae | Pimephales promelas | Fathead minnow | 45 |
| 5 | Cypriniformes | Cyprinidae | Notemigonus crysoleucas | Golden shiner | 595 |
| 6a | Cypriniformes | Cyprinidae | Carassius auratus | Goldfish | 3 |
| 3 | Cypriniformes | Cyprinidae | Rhinichthys cataractae | Longnose dace | 15499 |
| 4 | Cypriniformes | Cyprinidae | Notropis hudsonius | Spottail shiner | 1344 |
| x | Cypriniformes | Cyprinidae | Cyprinidae | Unknown minnow | 383 |
| 4 | Esociformes | Esocidae | Esox niger | Chain pickerel | 521 |
| 4 | Esociformes | Esocidae | Esox americanus | Redfin pickerel | 612 |
| 5 | Cyprinodontiformes | Fundulidae | Fundulus diaphanus | Banded killifish | 259 |
| x | Cyprinodontiformes | Fundulidae | Fundulus heteroclitus | Mummichog | 18 |
| x | Gadiformes | Gadidae | Lota lota | Burbot | 29 |
| x | Gadiformes | Gadidae | Microgadus tomcod | Tomcod | 2 |
| 4 | Gasterosteiformes | Gasterosteidae | Apeltes quadracus | Fourspine stickleback | 118 |
| x | Gasterosteiformes | Gasterosteidae | Pungitius pungitius | Ninespine stickleback | 1 |
| 5 | Siluriformes | Ictaluridae | Ameiurus nebulosus | Brown bullhead | 479 |
| x | Siluriformes | Ictaluridae | Ameiurus catus | White catfish | 1 |
| 6a | Siluriformes | Ictaluridae | Ameiurus natalis | Yellow bullhead | 187 |
| 10 | Perciformes | Moronidae | Morone saxatilis | Striped bass | 1 |

Table C -1 continued...

| BCG Attribute | Order | Family | Scientific Name | Common Name | Total # individuals |
|----------------------|--------------------|-----------------|---------------------------------|------------------------|----------------------------|
| 4 | Perciformes | Moronidae | Morone americana | White perch | 2 |
| x | Perciformes | Percidae | Etheostoma fusiforme | Swamp darter | 21 |
| 4 | Perciformes | Percidae | Etheostoma olmstedi | Tesselated darter | 8832 |
| x | Perciformes | Percidae | Stizostedion vitreum | Walleye | 5 |
| x | Perciformes | Percidae | Perca flavescens | Yellow perch | 744 |
| 2 | Petromyzontiformes | Petromyzontidae | Lampetra appendix | American Brook lamprey | 19 |
| 10 | Petromyzontiformes | Petromyzontidae | Petromyzon marinus | Sea lamprey | 174 |
| 6b | Salmoniformes | Salmonidae | Salmo salar (stocked) | Atlantic salmon | 3237 |
| 6b | Salmoniformes | Salmonidae | Salvelinus fontinalis (stocked) | Brook trout (stocked) | 246 |
| 2 | Salmoniformes | Salmonidae | Salvelinus fontinalis (wild) | Brook trout (wild) | 8221 |
| 6b | Salmoniformes | Salmonidae | Salmo trutta | Brown trout | 5321 |
| 6b | Salmoniformes | Salmonidae | Oncorhynchus mykiss | Rainbow trout | 475 |
| x | Salmoniformes | Salmonidae | Sibericus sabertoothii | Tiger Trout | 13 |
| 6a | Esociformes | Umbridae | Umbra limi | Central mudminnow | 32 |

APPENDIX D

Capture probability modeled vs. disturbance
gradient

Generalized additive models (GAM) were used to characterize the relationships between species presence/absence and the percent of the watershed comprised of developed land¹. Capture probability plots were generated for species that occurred at 20 or more sites. Capture probability refers to the likelihood of occurrence along the gradient of interest.

Curve shapes generally fall into three categories: increasing, decreasing, or unimodal. In the example in Figure D-1, the American eel has an increasing curve shape; this means that the probability of capturing the American eel increases as % developed area increases.

The black solid line in the plots is the modeled capture probability (based on the GAM model) and the circles are the mean observed probability in equal distance bins. For example, if you divide the stressor gradient into 50 equal distance bins, the mean of any data points (1 or 0) within that bin are taken as the probability of capture and plotted against the mean stressor value in the bin. The black dashed lines represent the 90% confidence interval, and the vertical red dashed line and number associated with it represent the optima, which in these plots equals the 50% area under the modeled curve. In the example in Figure D-1, the American eel has a 50% chance of occurring at sites with 18% developed area.

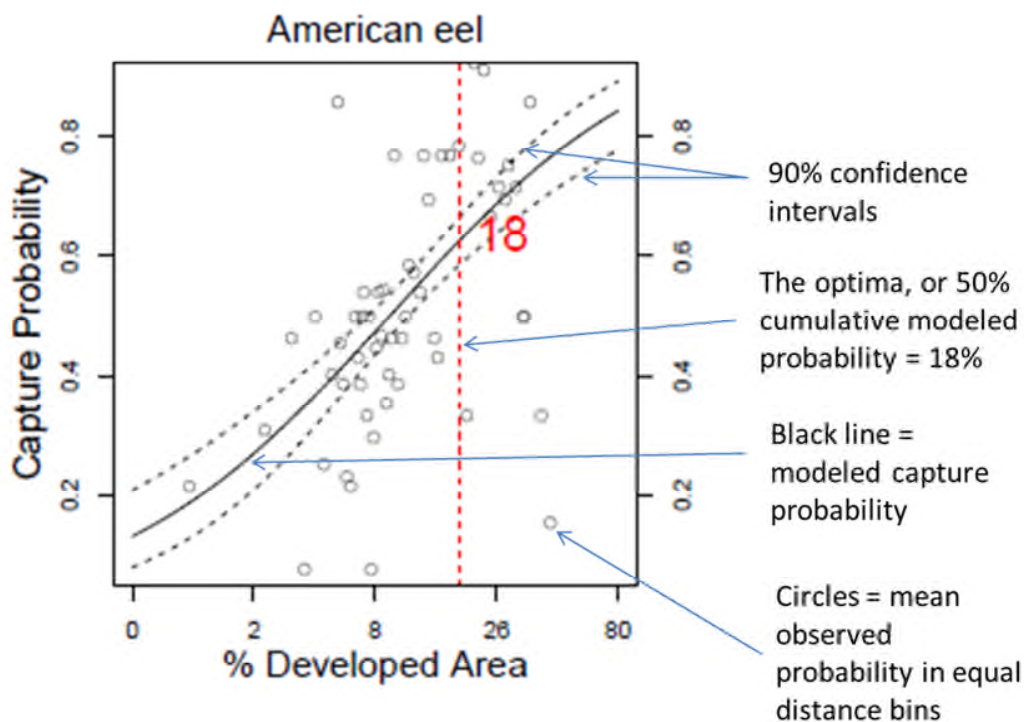
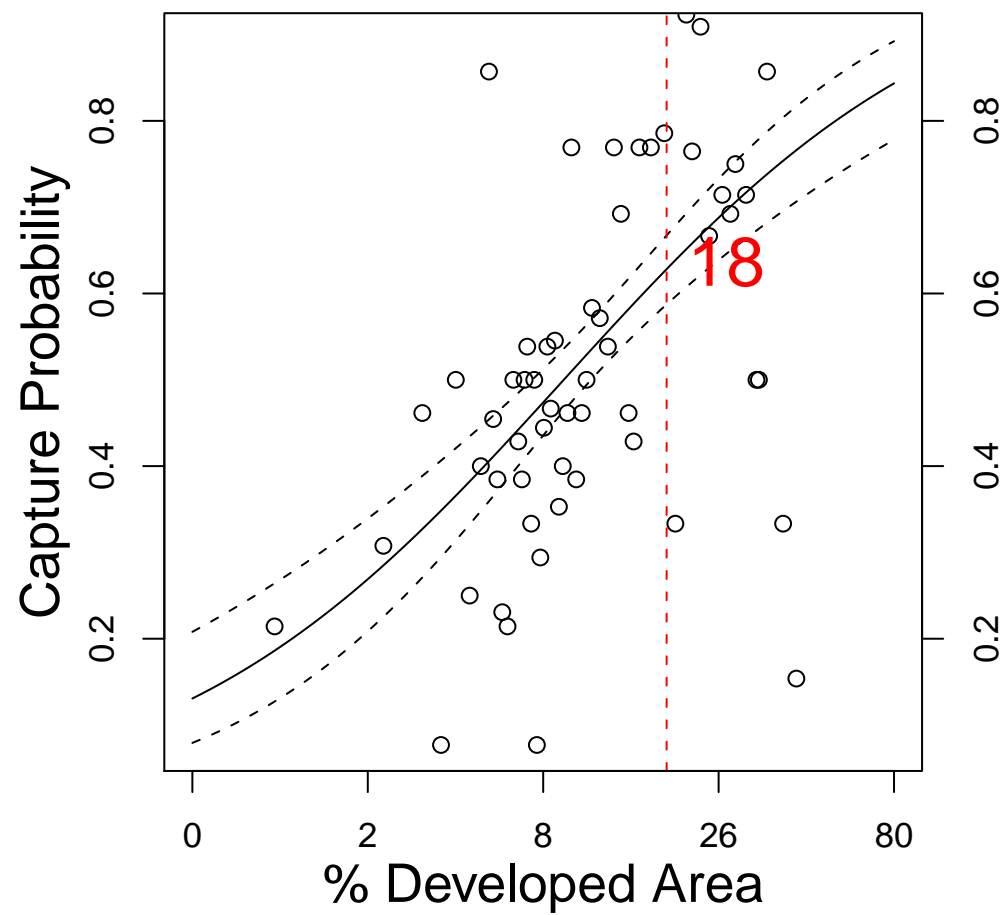


Figure D-1. Percent developed tolerance plot for the American eel. The probability of capturing this species at a site increases as % developed area increases.

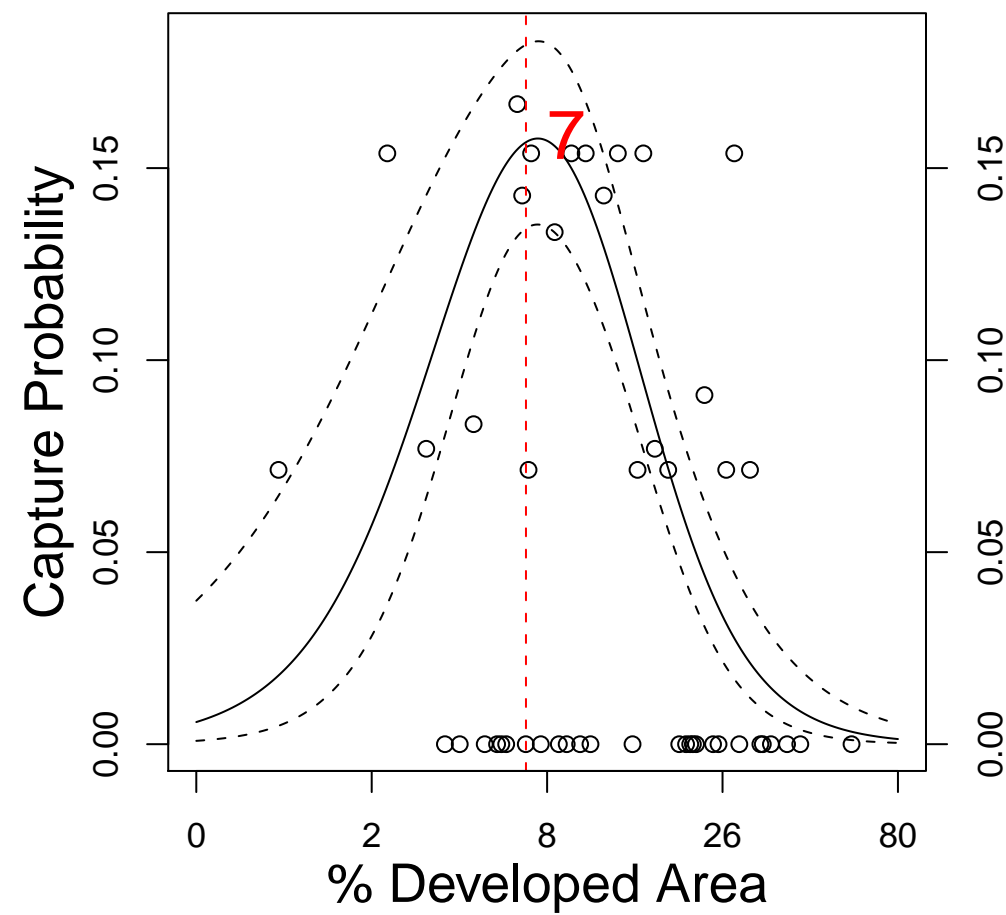
¹ The developed land data were provided by CT DEEP.

Taxon responses to Disturbance Gradient

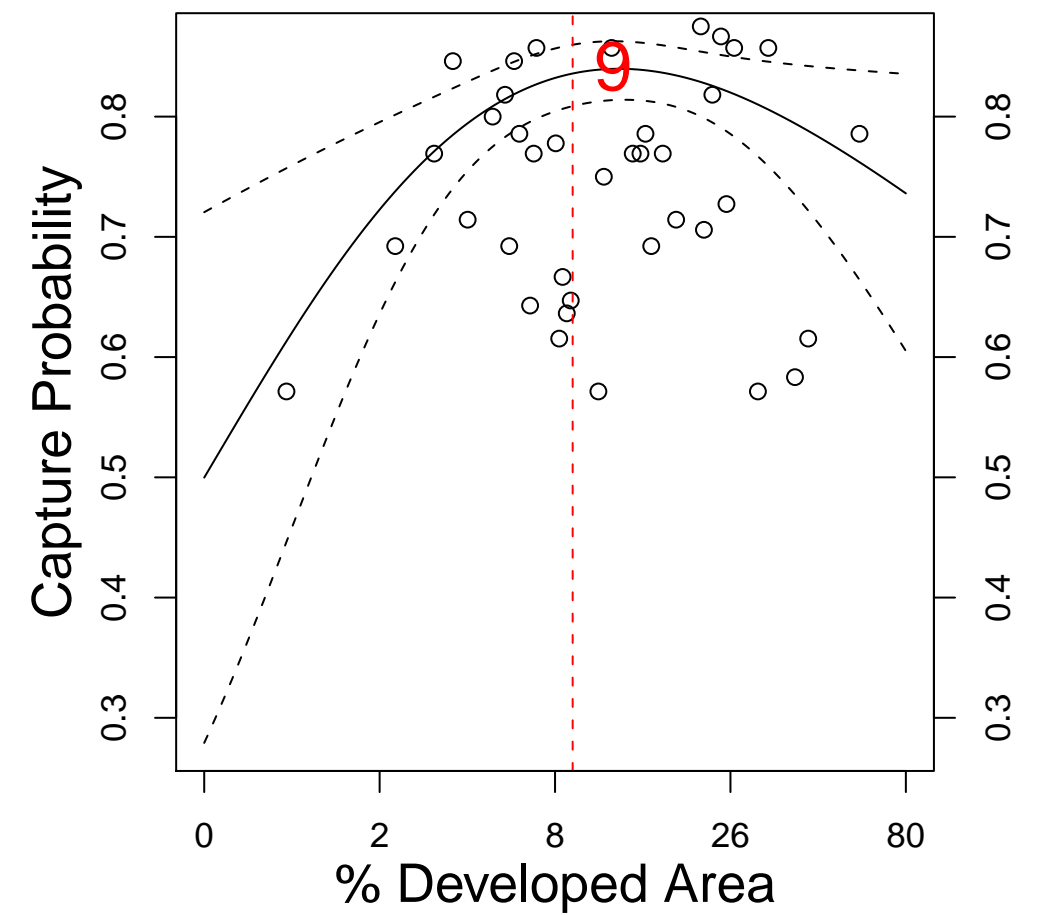
American eel



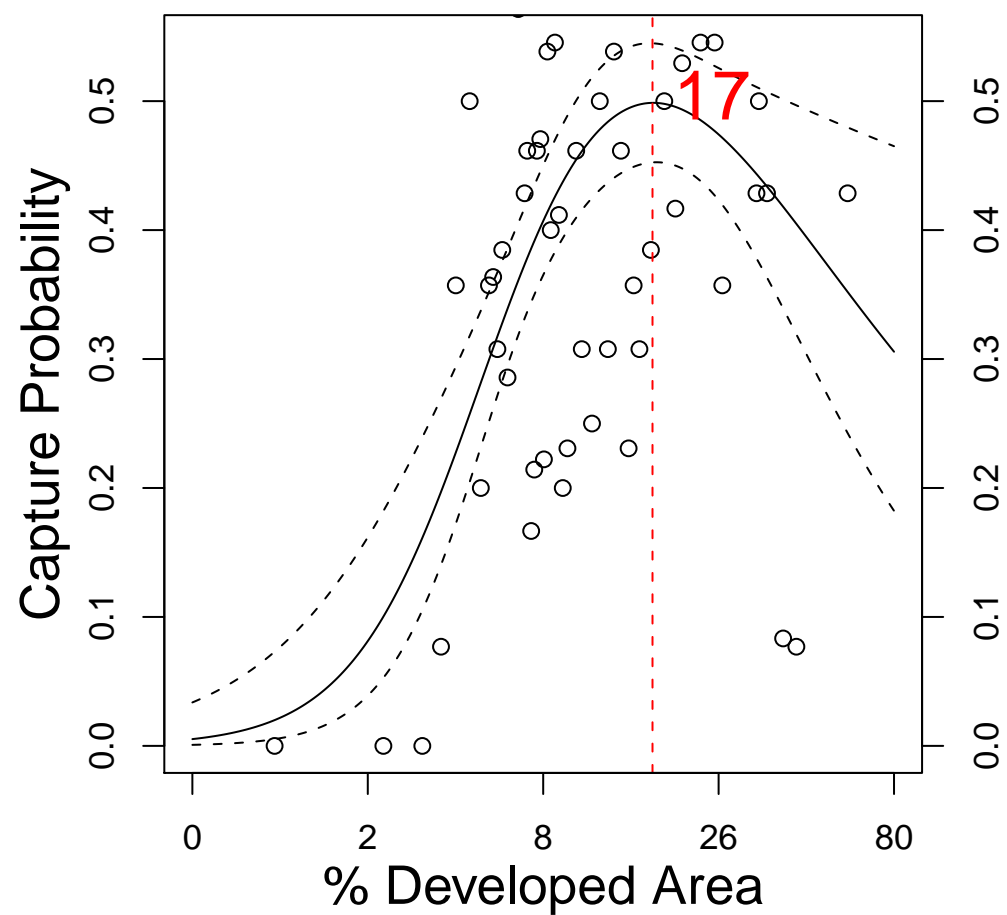
Atlantic salmon, fry stocked



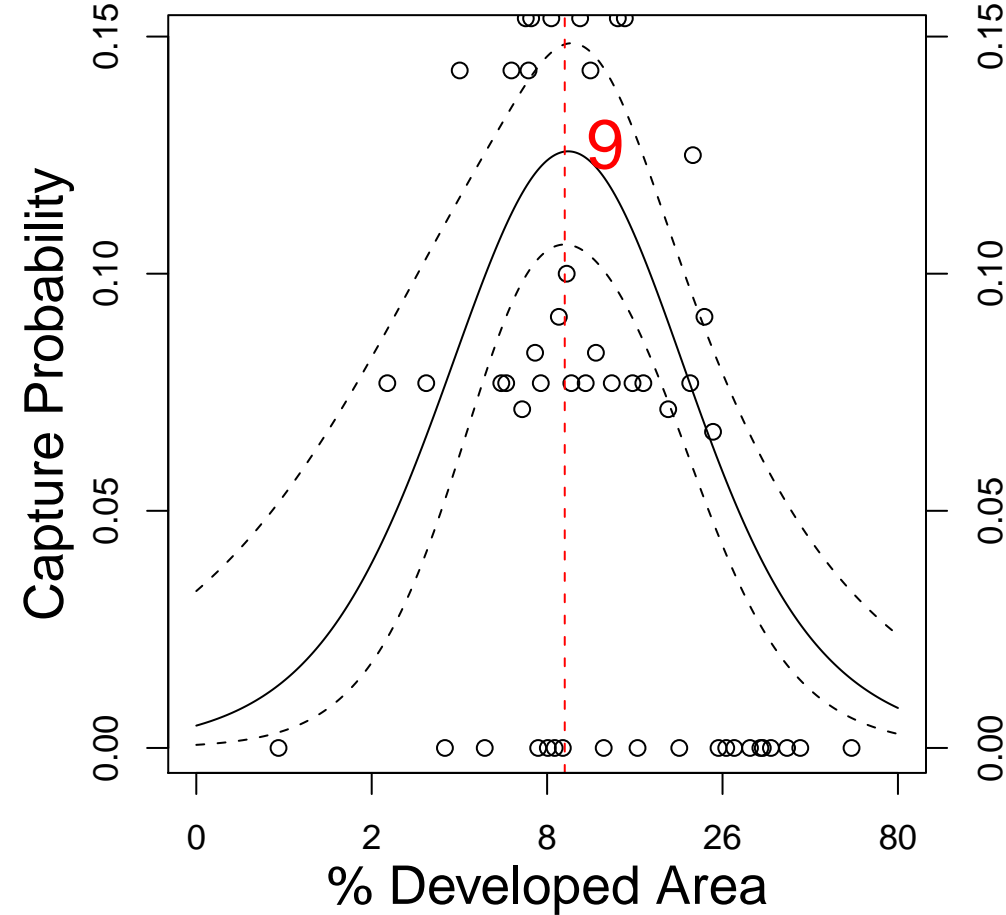
Blacknose dace



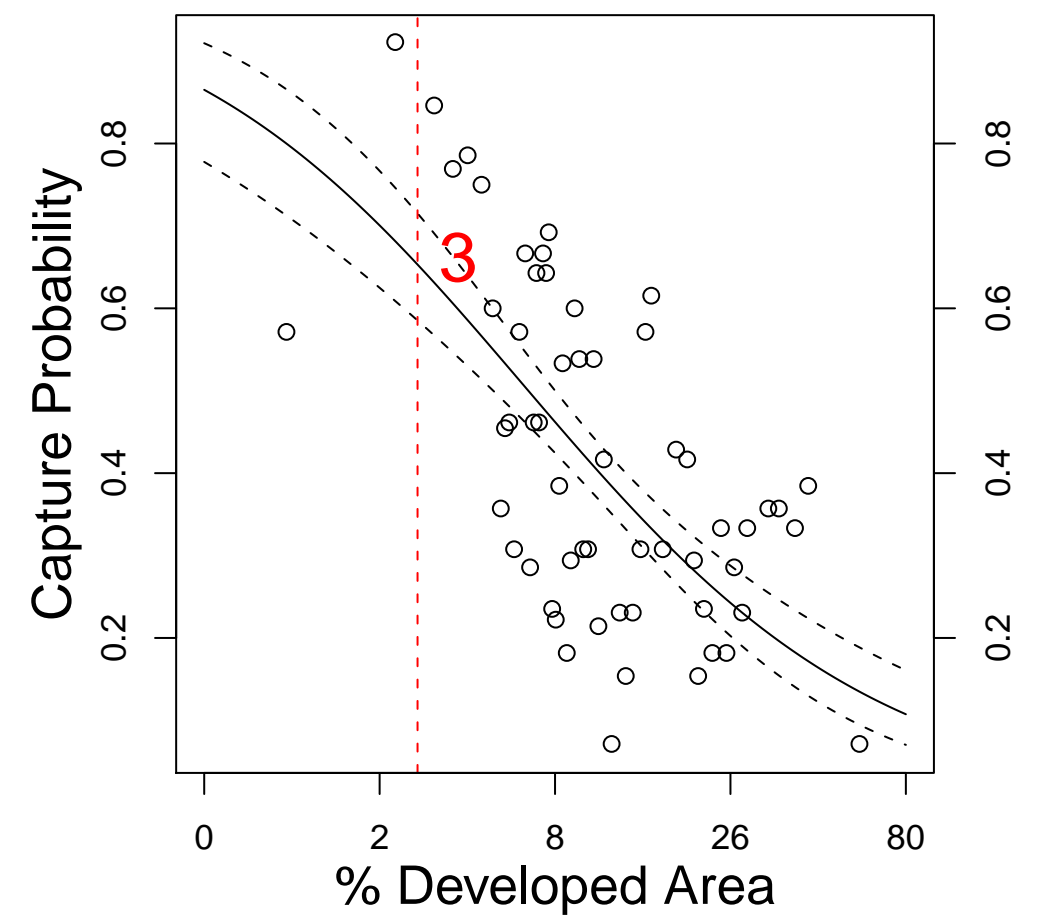
Bluegill sunfish



Brook trout, stocked

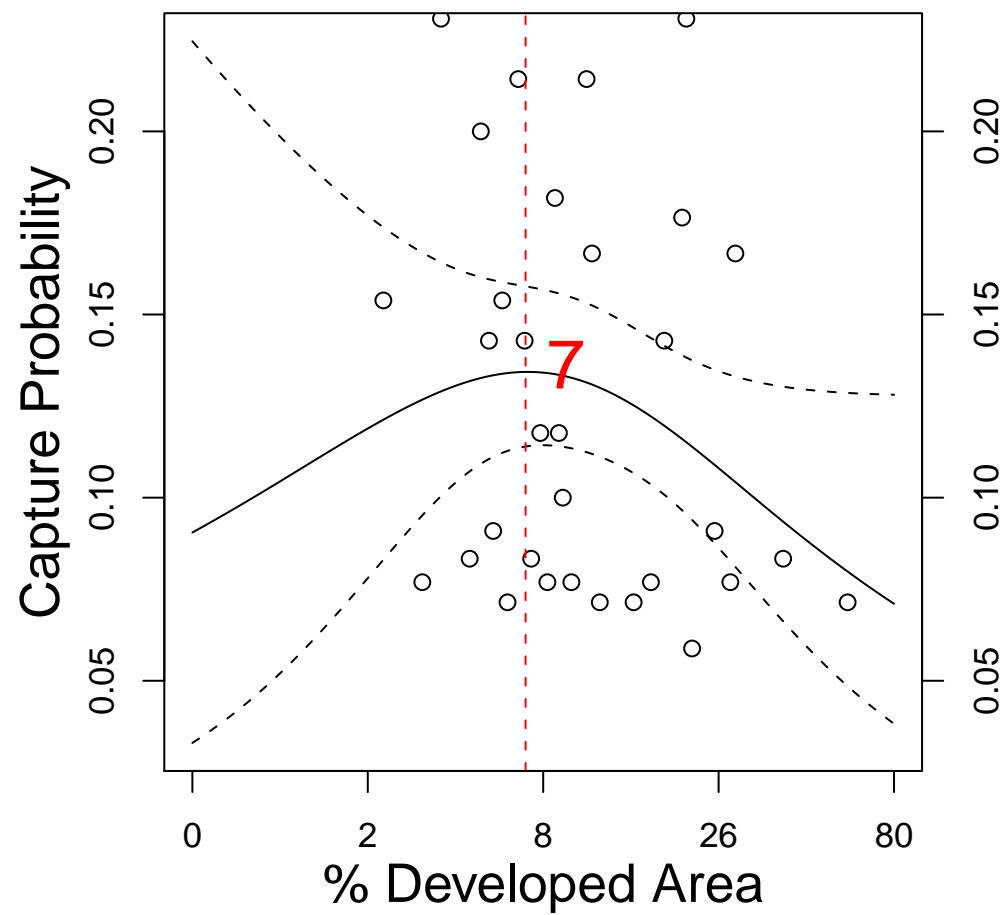


Brook trout, wild

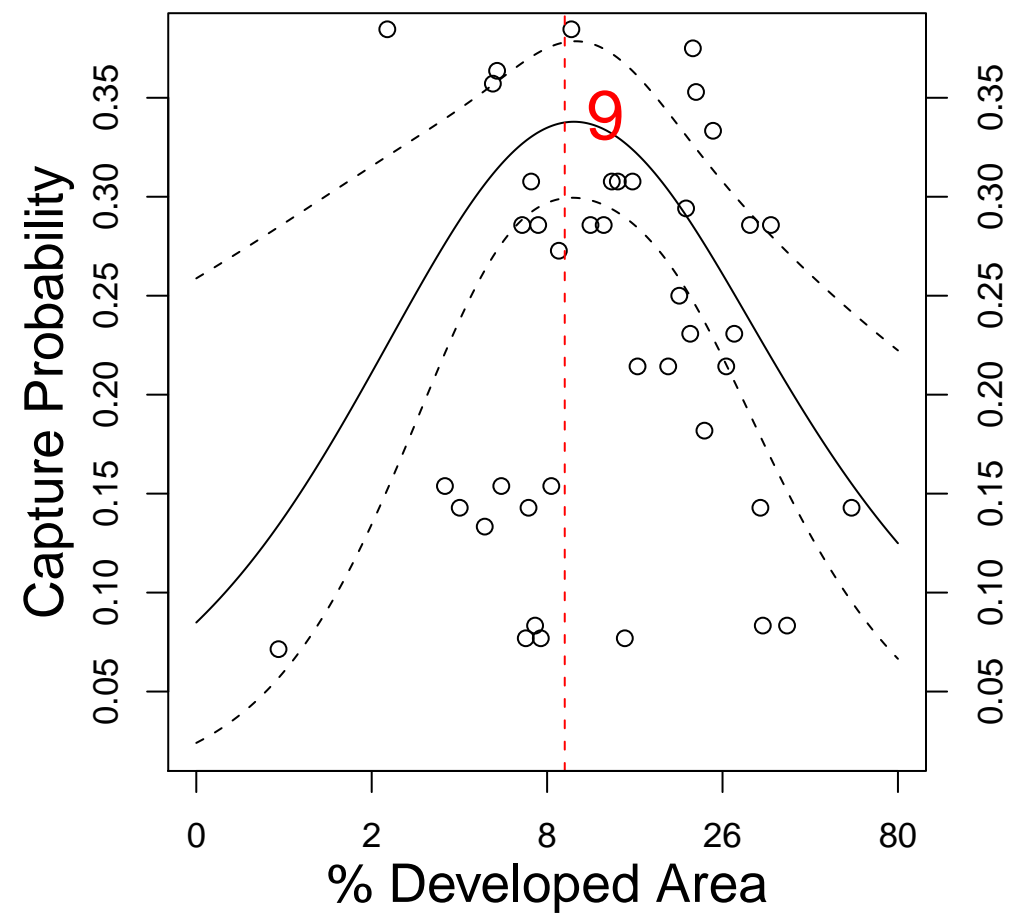


Taxon responses to Disturbance Gradient

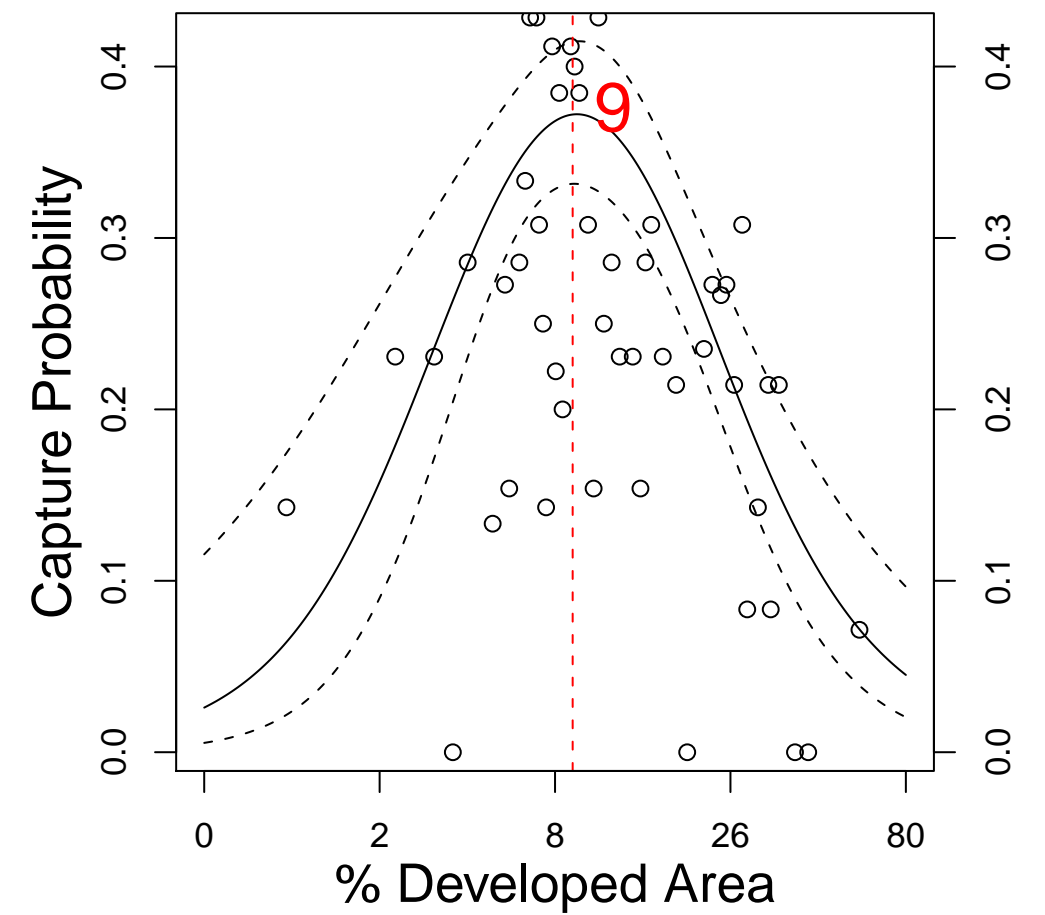
Brown bullhead



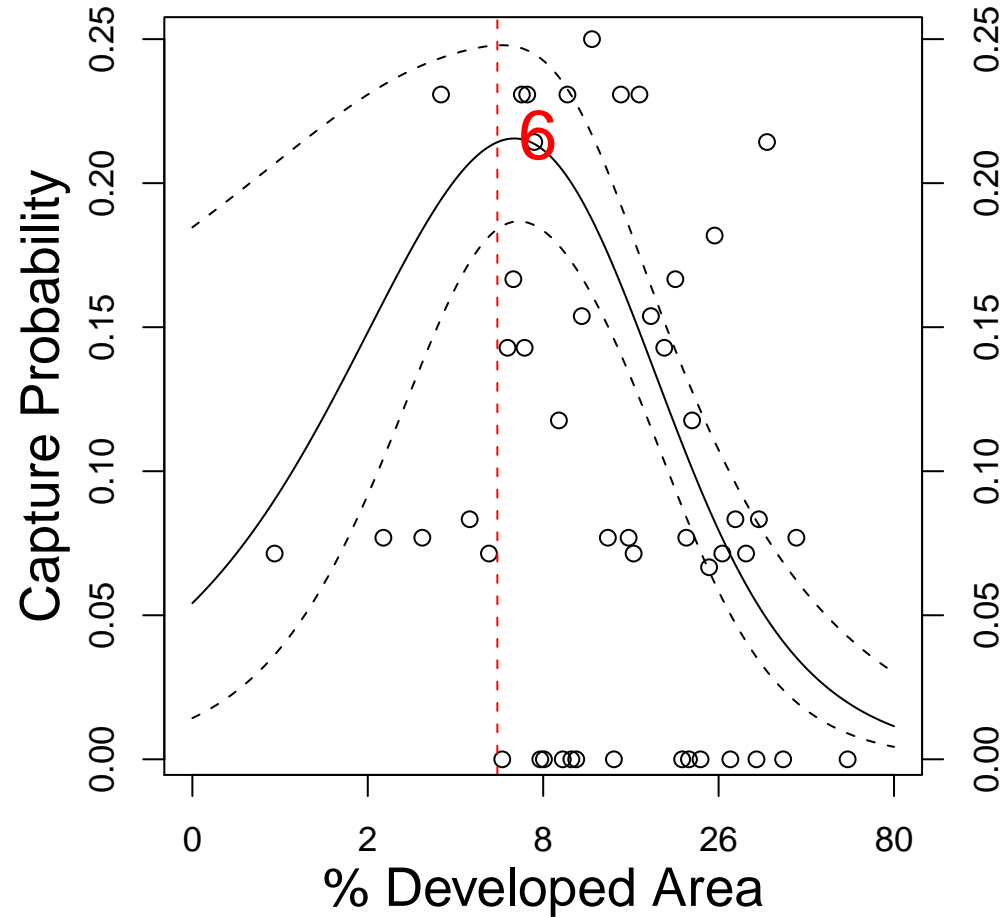
Brown trout, naturalized



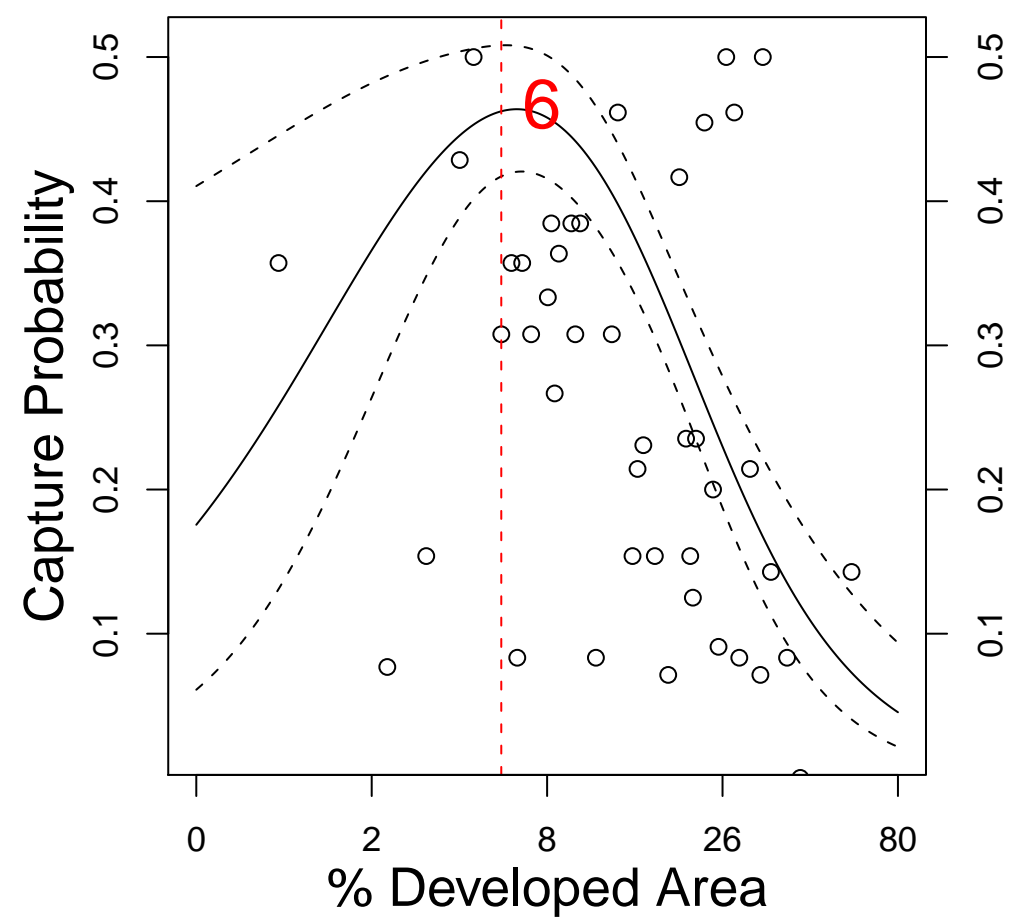
Brown trout, stocked



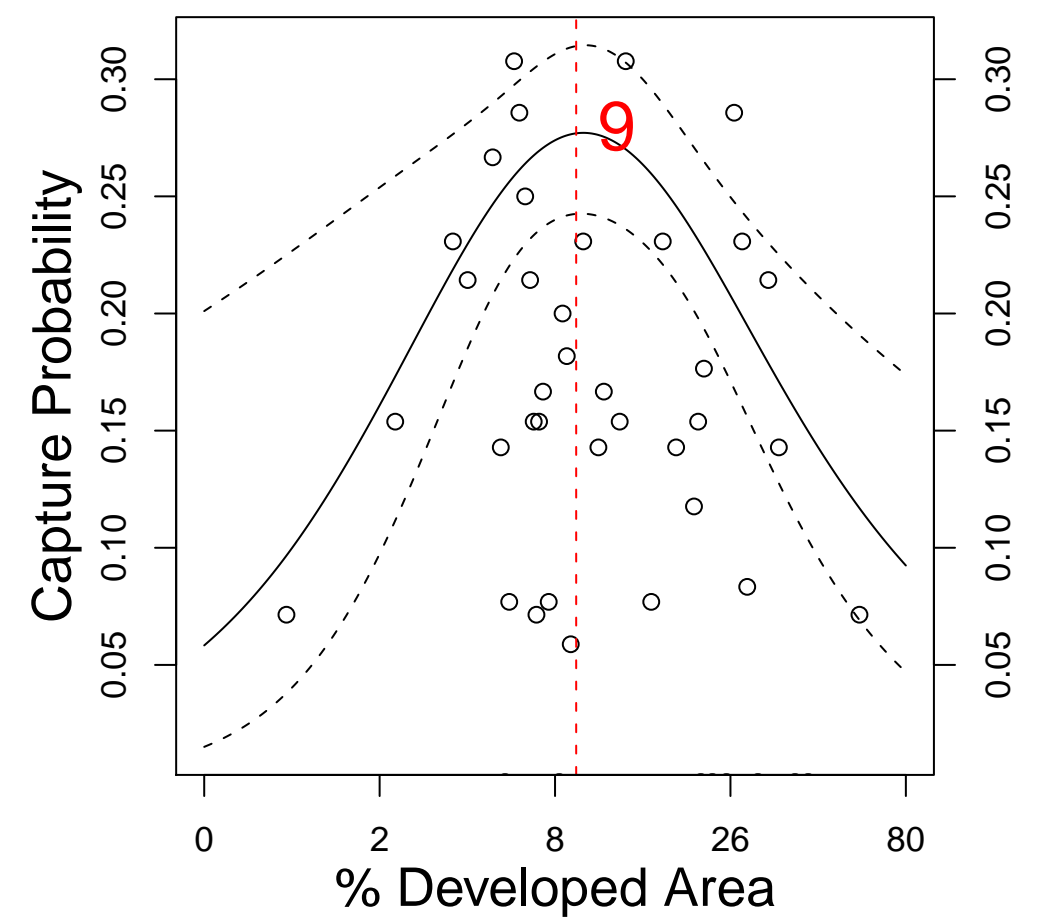
Chain pickerel



Common shiner

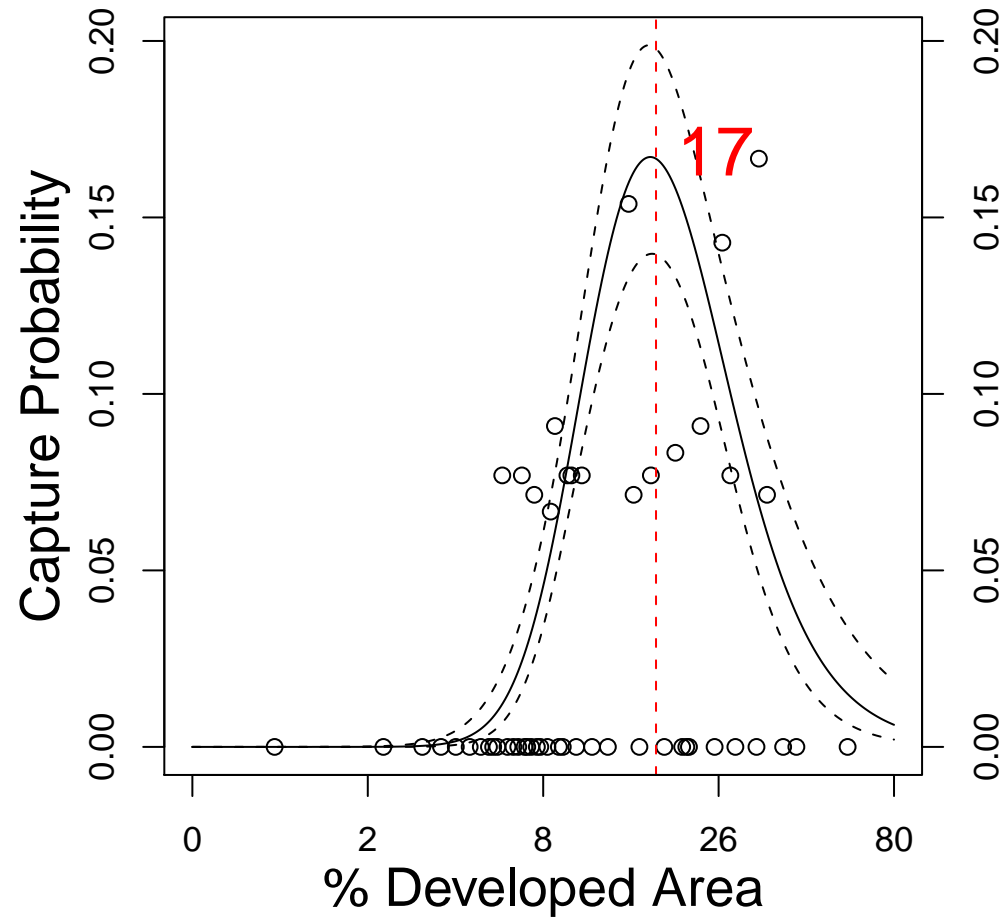


Creek chub

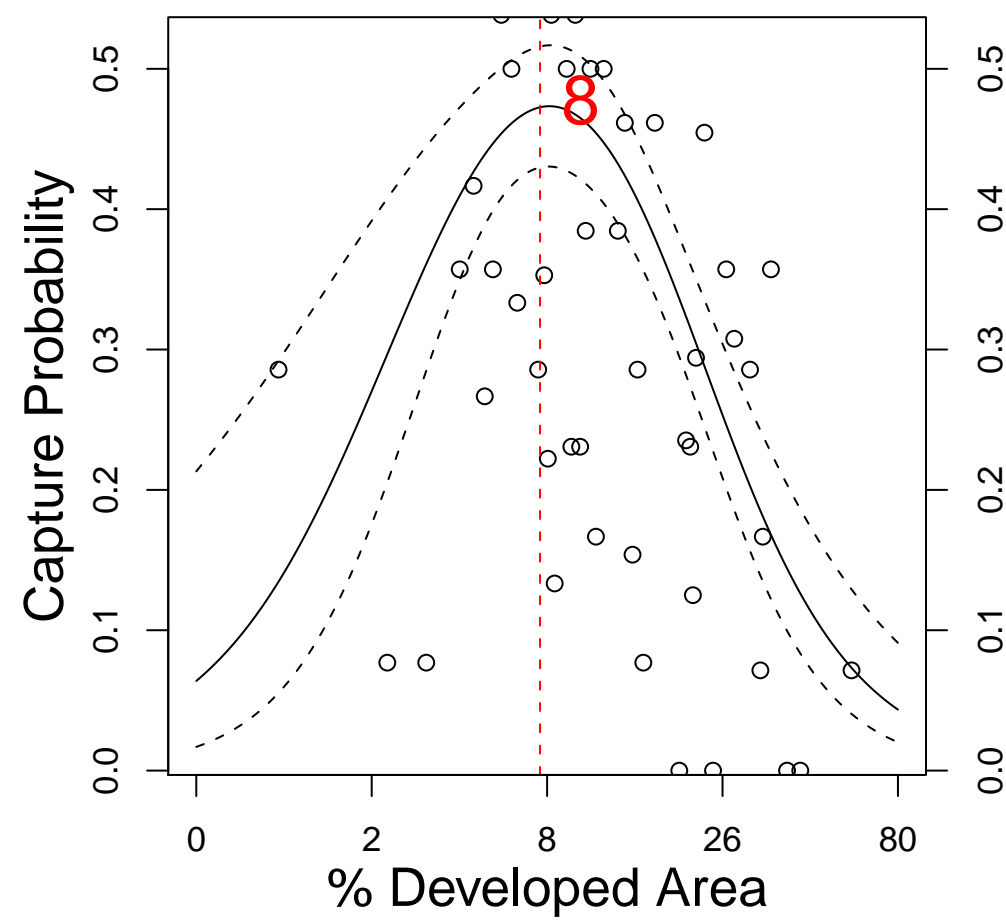


Taxon responses to Disturbance Gradient

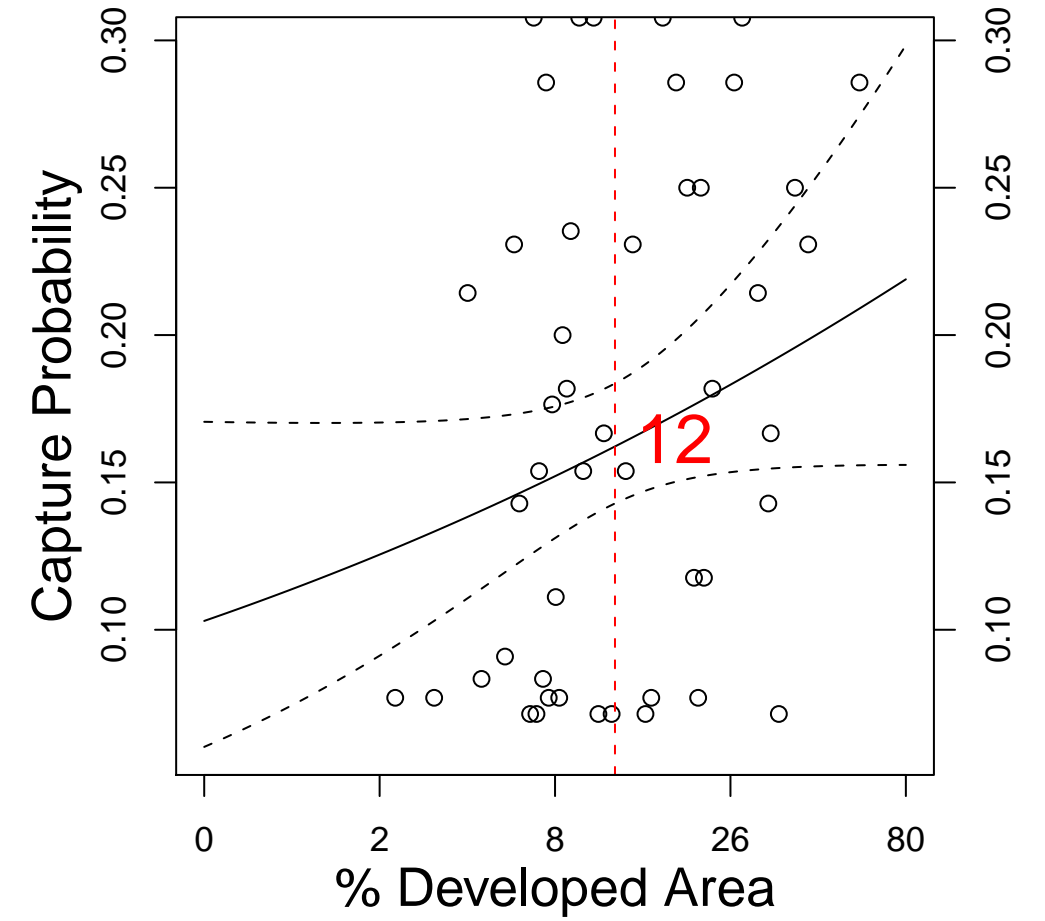
Cutlips minnow



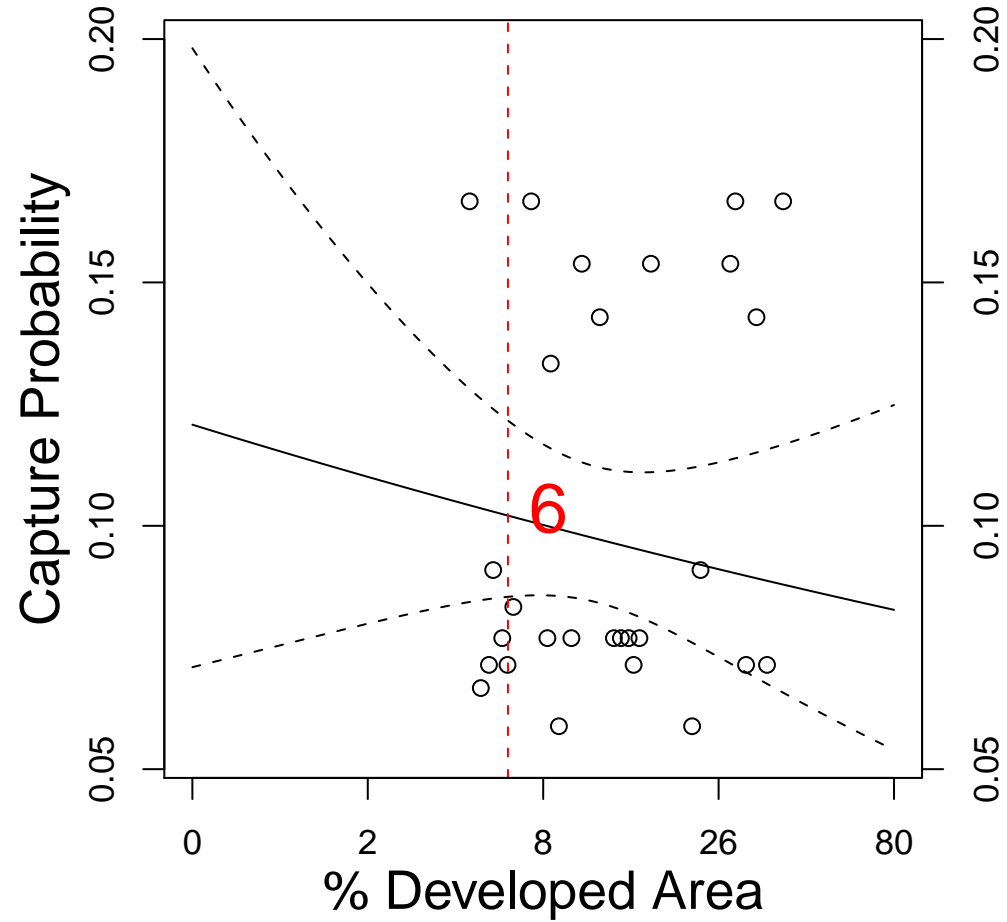
Fallfish



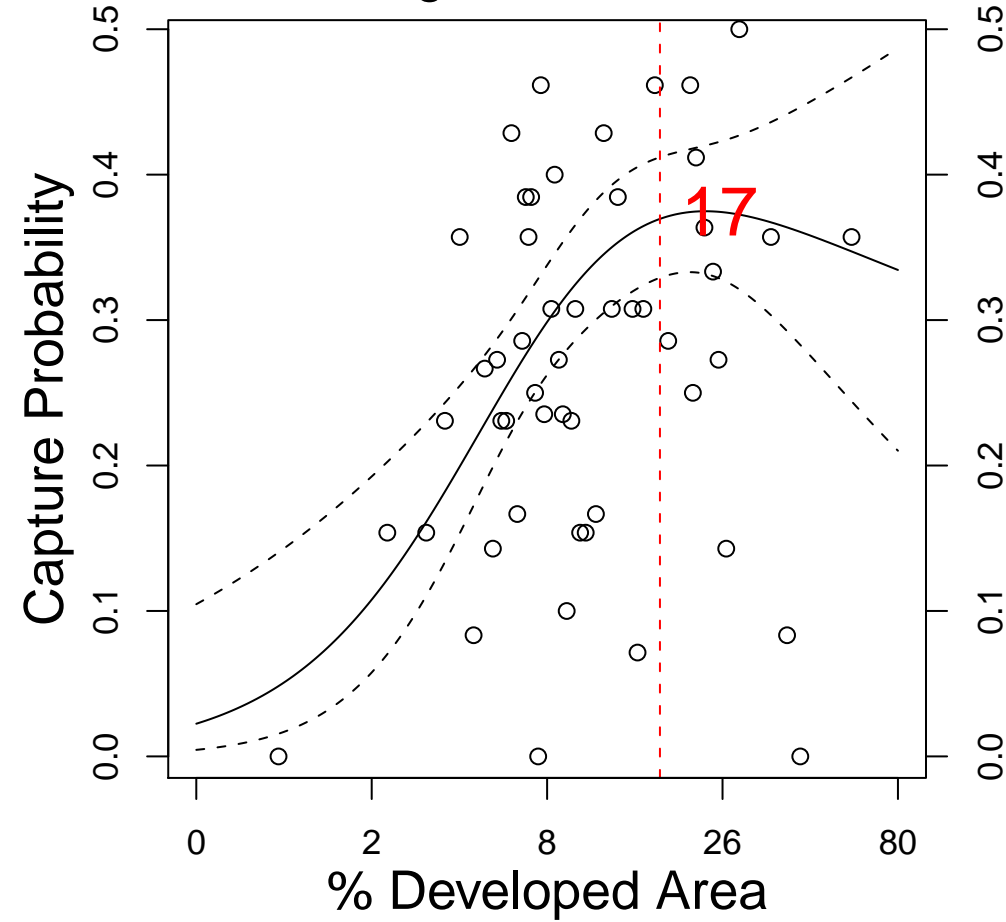
Golden shiner



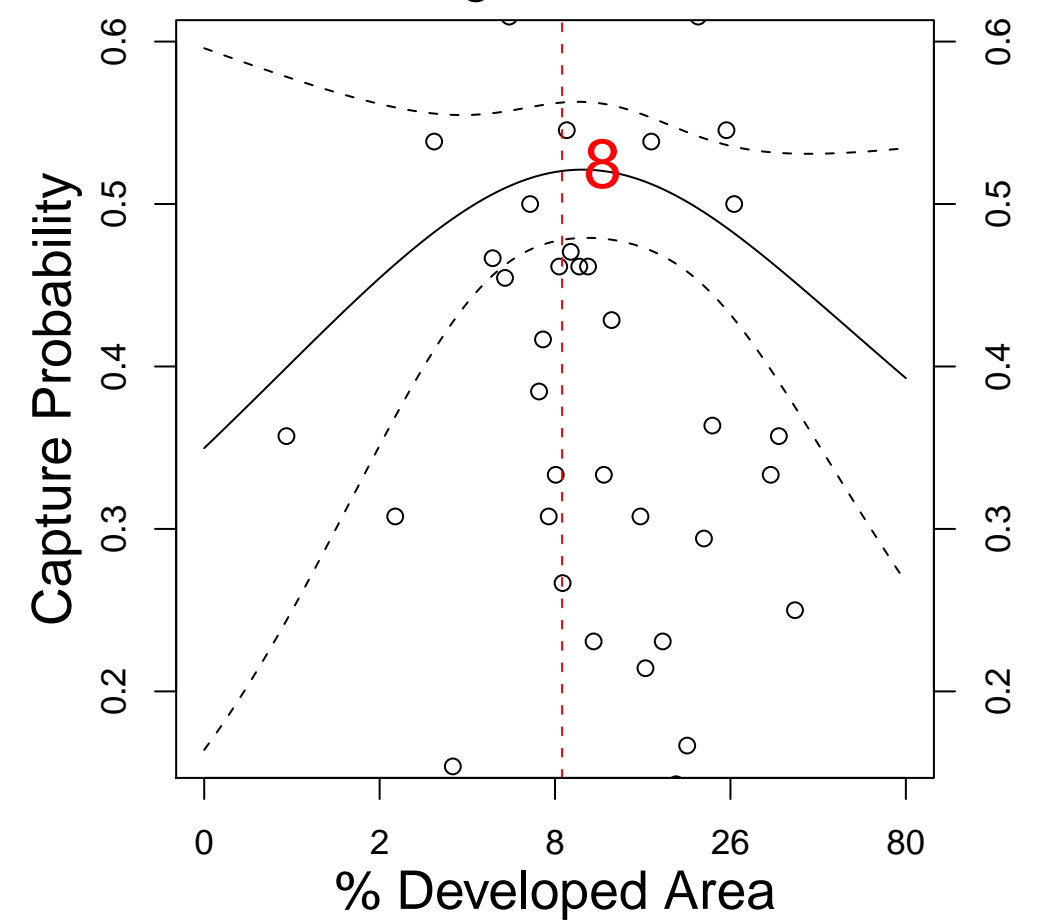
Green sunfish



Largemouth Bass

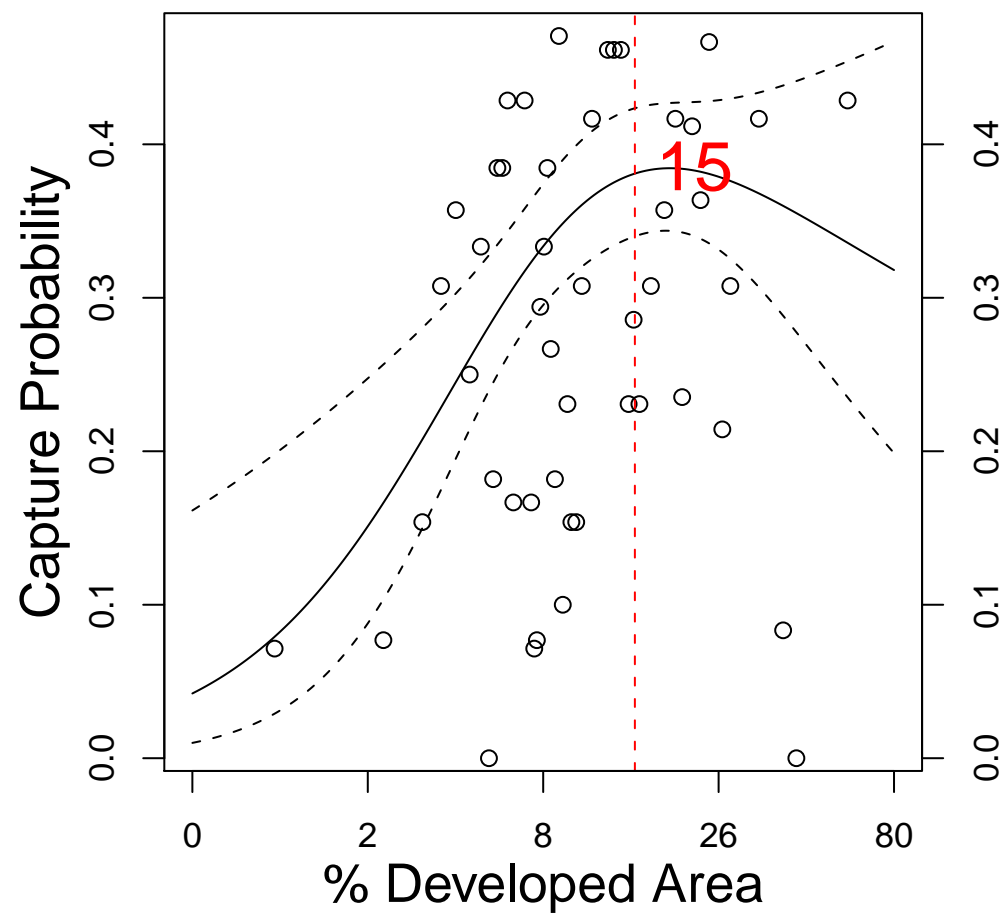


Longnose dace

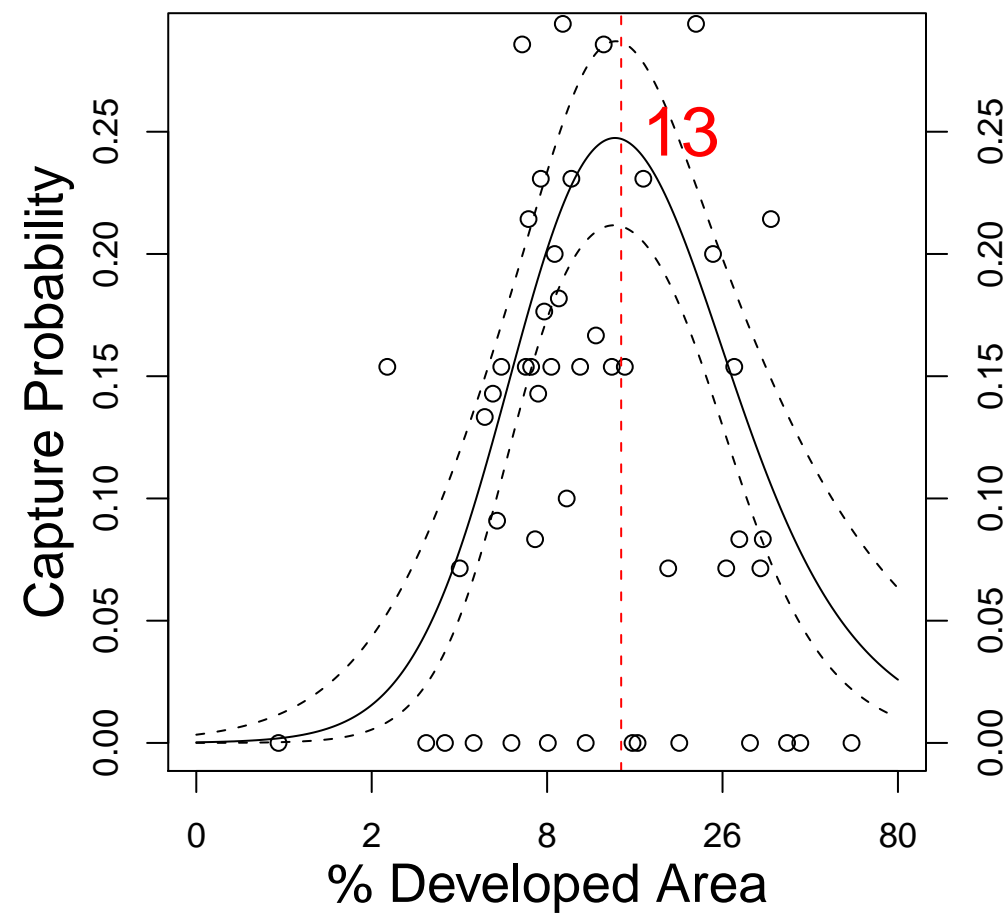


Taxon responses to Disturbance Gradient

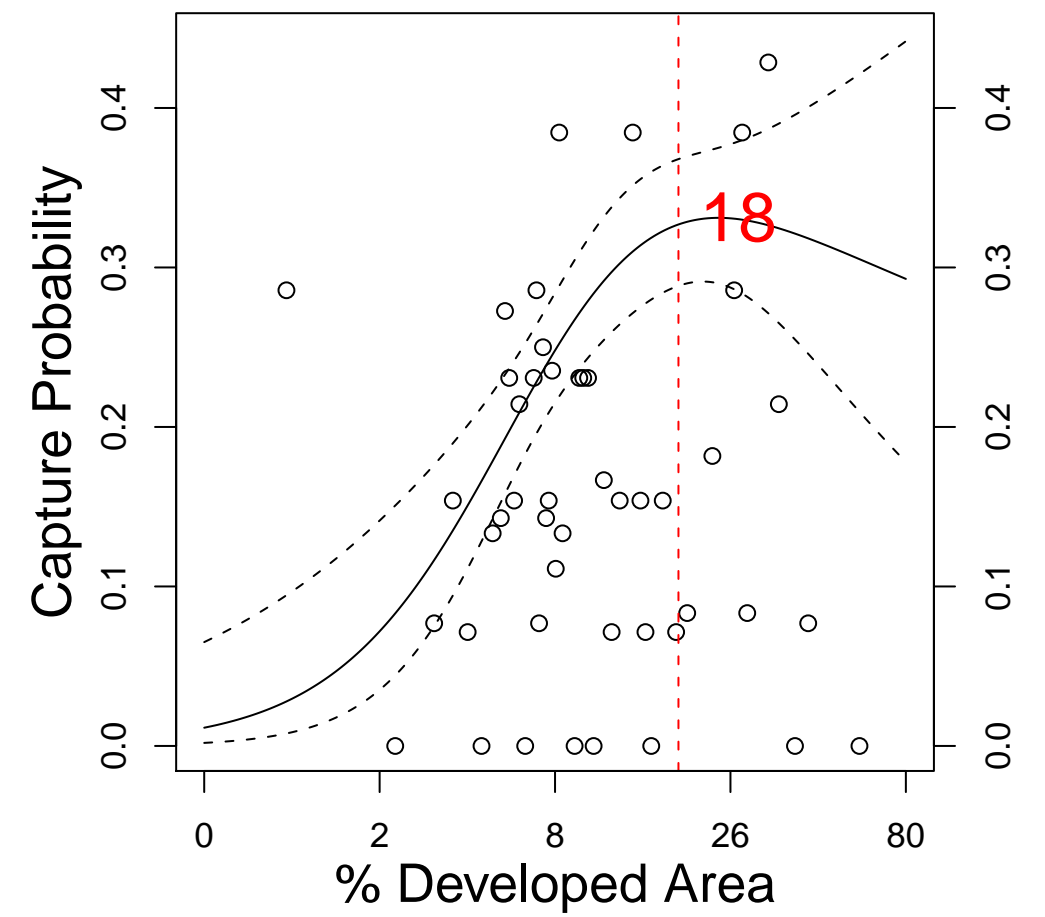
Pumpkinseed



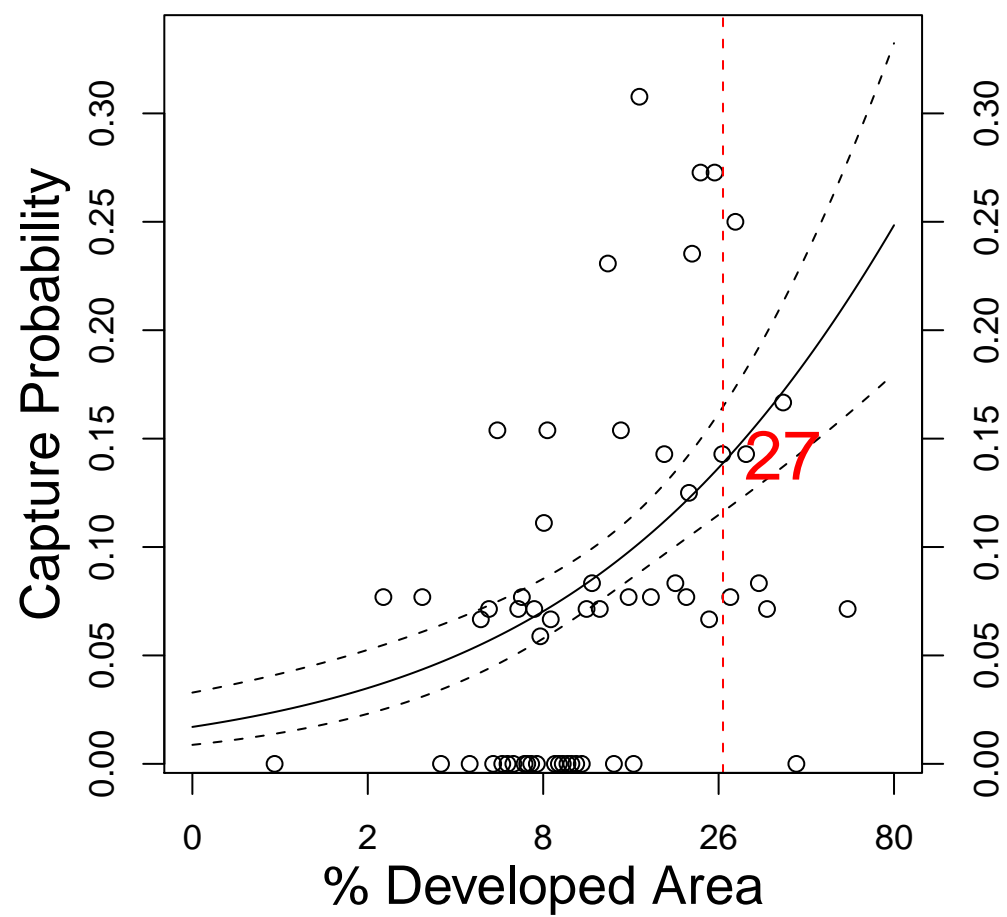
Rainbow trout, Stocked



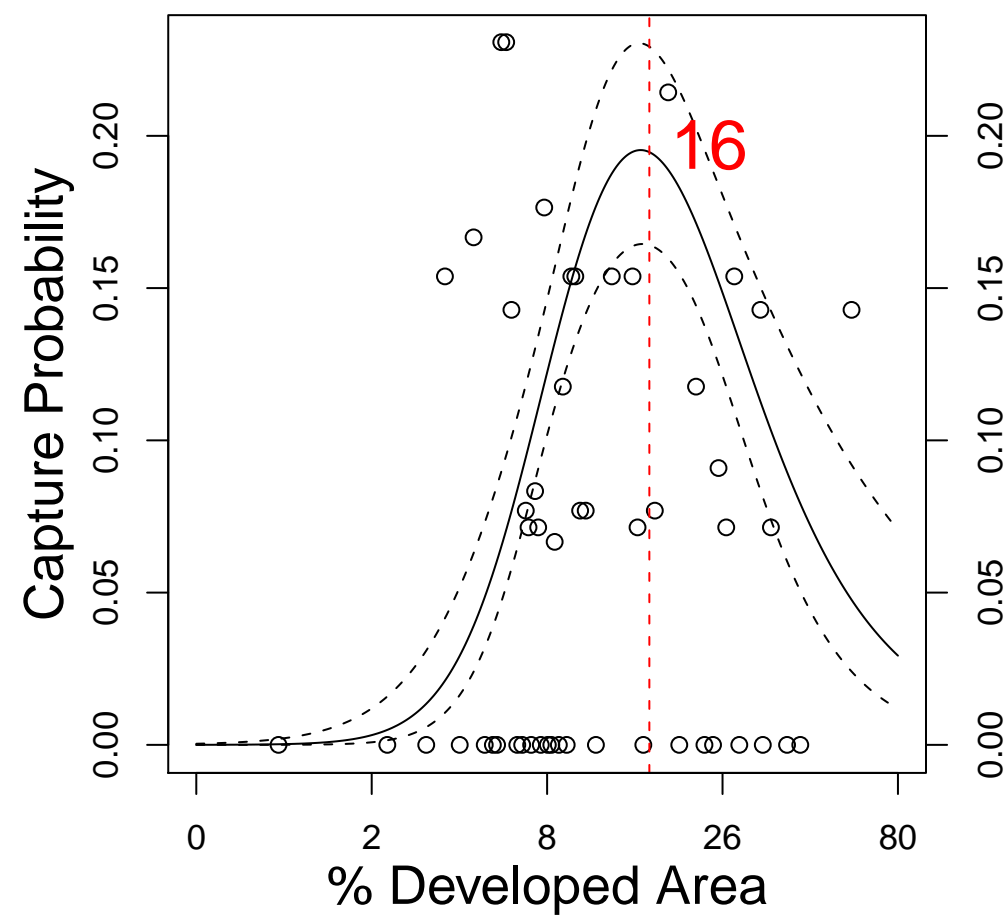
Redbreast sunfish



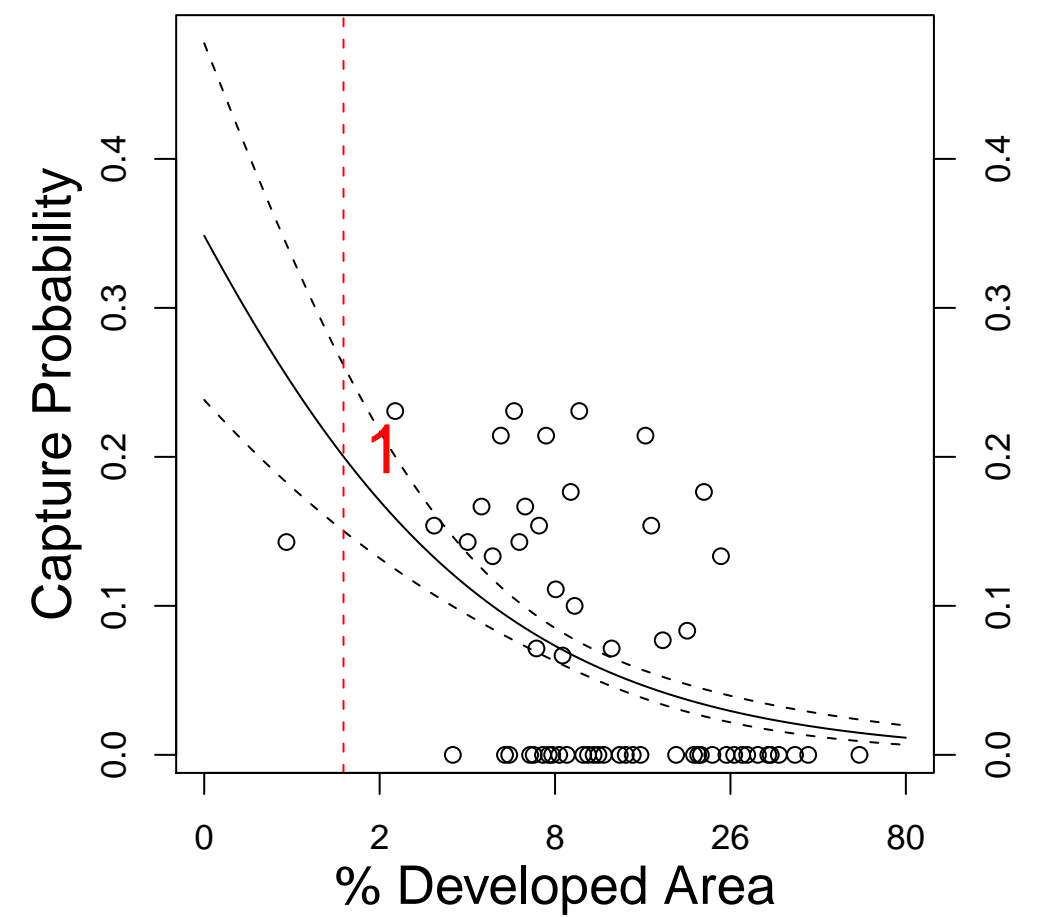
Redfin pickerel



Rock Bass

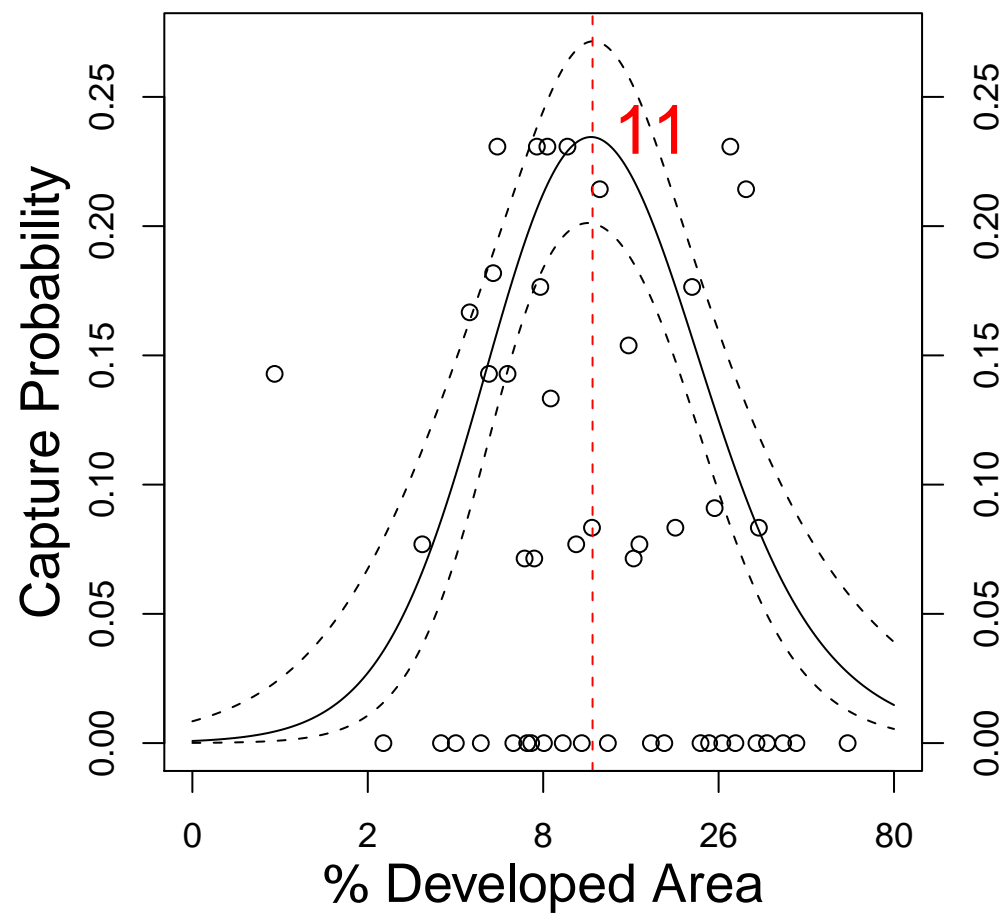


Slimy sculpin

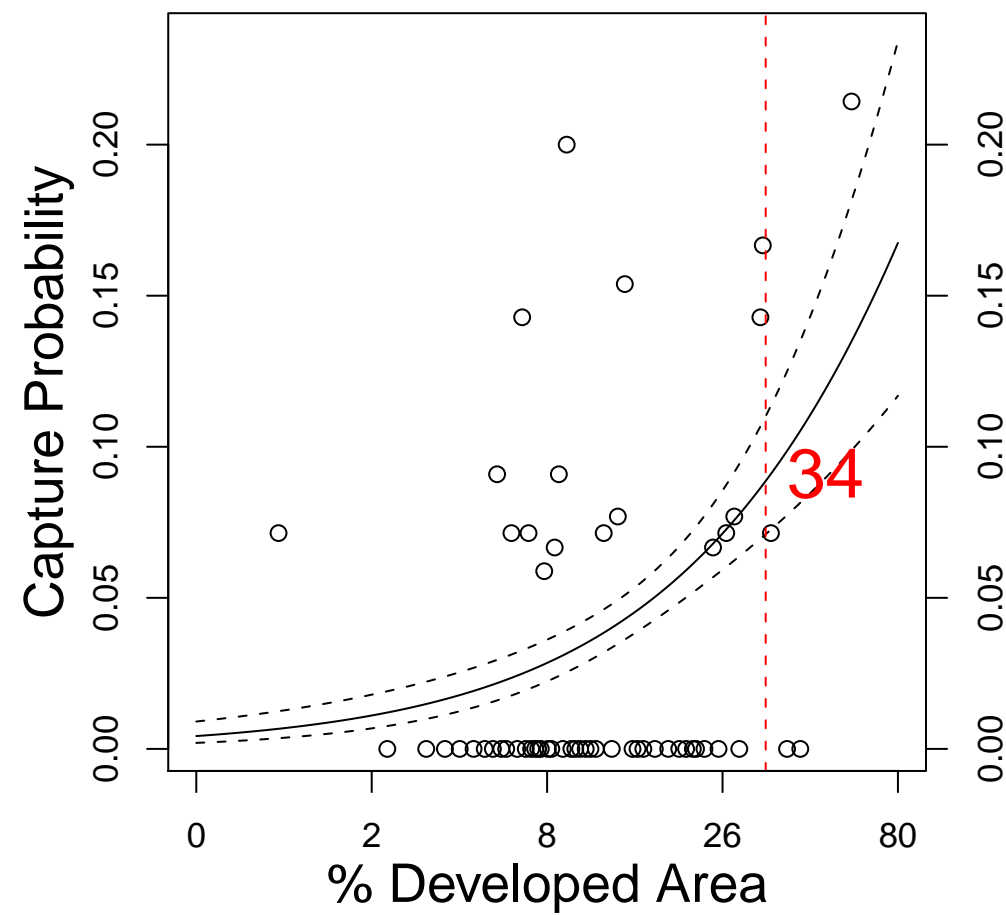


Taxon responses to Disturbance Gradient

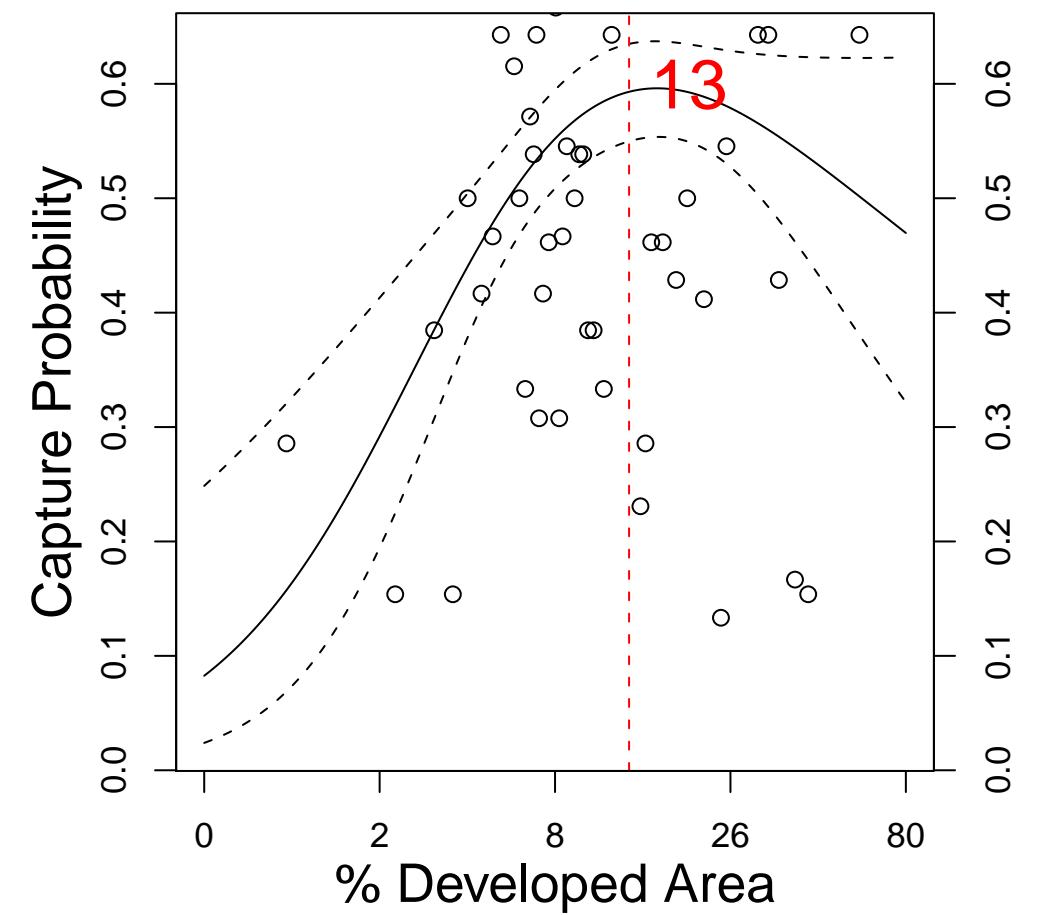
Smallmouth bass



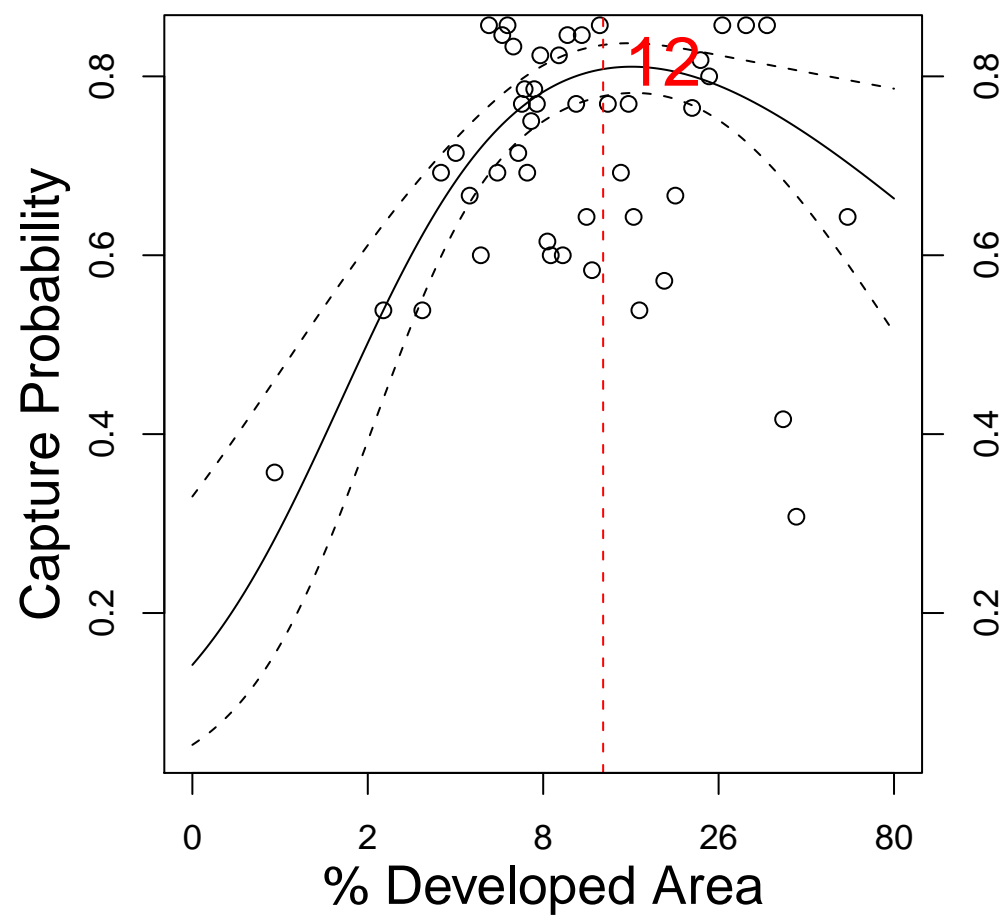
Spottail shiner



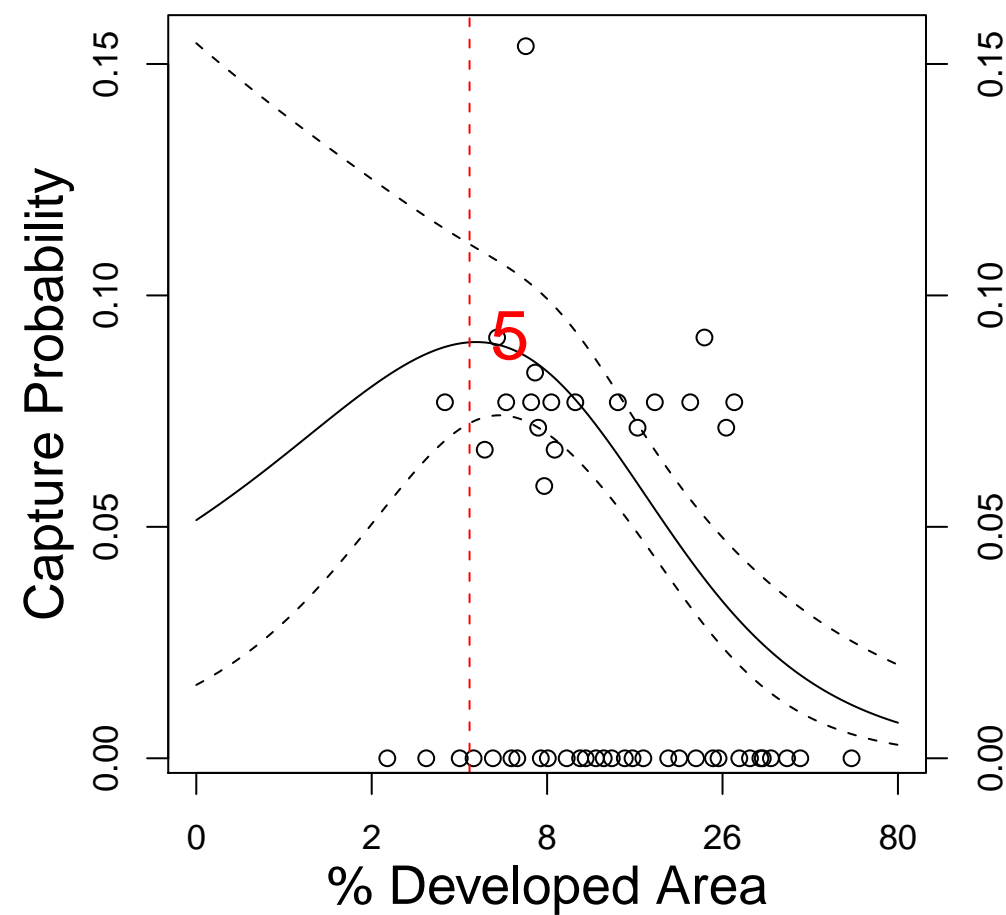
Tesselated darter



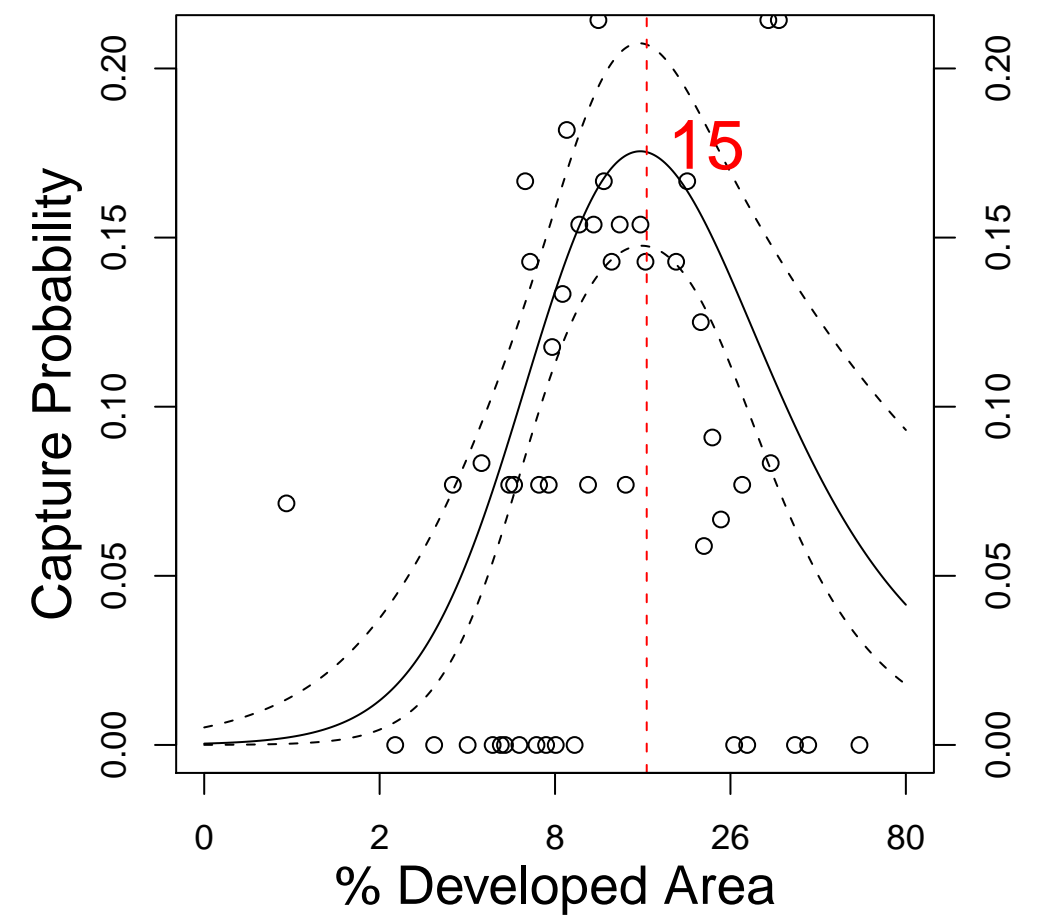
White sucker



Yellow bullhead



Yellow perch



APPENDIX E

Sample worksheet

Figure E1. Example of a worksheet that was used when making BCG level assignments.

| | | | |
|-------------------|---------------------|----------------------------|--|
| ExerciseID | Samp309 | Consensus BCG Level | Range of BCG calls & reasoning |
| CollMethod | single pass wadable | 3 | 3-, 2- to 4; even split 3/4; like cyprinid diversity, looks warm |

| BCG Att | # Taxa | # Ind | Pct Taxa | Pct Ind |
|--------------|--------|-------|----------|---------|
| 1 | 0 | 0.00 | 0.00 | 0.00 |
| 2 | 0 | 0.00 | 0.00 | 0.00 |
| 3 | 1 | 0.02 | 0.13 | 0.11 |
| 4 | 6 | 0.19 | 0.75 | 0.88 |
| 5 | 0 | 0.00 | 0.00 | 0.00 |
| 5a | 0 | 0.00 | 0.00 | 0.00 |
| 6 | 1 | 0.00 | 0.13 | 0.00 |
| 6a | 0 | 0.00 | 0.00 | 0.00 |
| x | 0 | 0.00 | 0.00 | 0.00 |
| Total | 8 | 0.22 | | |

| BCG Att | Common Name | Scientific Name | Individs_m2 | Individs |
|---------|----------------------|--------------------------------|-------------|------------|
| 10 | American eel | <i>Anguilla rostrata</i> | 0.02 | 28 |
| 4 | Blacknose dace | <i>Rhinichthys atratulus</i> | 0.09 | 141 |
| 6 | Brown trout, stocked | <i>Salmo trutta hatcheryis</i> | 0.00 | 1 |
| 4 | Creek chub | <i>Semotilus atromaculatus</i> | 0.05 | 68 |
| 4 | Cutlips minnow | <i>Exoglossum maxillingua</i> | 0.03 | 44 |
| 3 | Longnose dace | <i>Rhinichthys cataractae</i> | 0.02 | 37 |
| 4 | Redbreast sunfish | <i>Lepomis auritus</i> | 0.01 | 17 |
| 4 | Tessellated darter | <i>Etheostoma olmstedi</i> | 0.01 | 12 |
| 4 | White sucker | <i>Catostomus commersoni</i> | 0.00 | 7 |
| | | Total | 0.24 | 355 |

| | |
|--|------------------------|
| StationID | 2685 |
| StreamName | Aspetuck River |
| Parameter | Value |
| Watershed Area (mi²) | 7.8278 |
| SampleArea_m2 | 1500 |
| SizeTempClass | medium_cool |
| CT_JulyMeanTemp | NA |
| TNC_Temp | Transitional Cool |
| TNC_Gradient | Moderate-High Gradient |
| TNC_Geology | Low Buffered, Acidic |
| Stress_Cat | minimal |
| pctNatural | 81.6 |
| Comments | |

| Expert | BCG Level | Reasons |
|----------|-----------|---|
| Chris B | 3+ | would like to see att 3 or 2 for higher |
| Mike B | 2- | balanced list |
| Neal | 4 | |
| Brian | 3- | like diversity cyprinids |
| Mike H | 4 | looks warm |
| Yoichiro | 4 | |
| Dave H | 3 | Moderate stream spp. Diversity = 5 |
| Rich | 3 | Like diversity and that there are 3 Bis, but needs more BKT |

APPENDIX F

Box plots of metrics for small-cold samples that were assessed

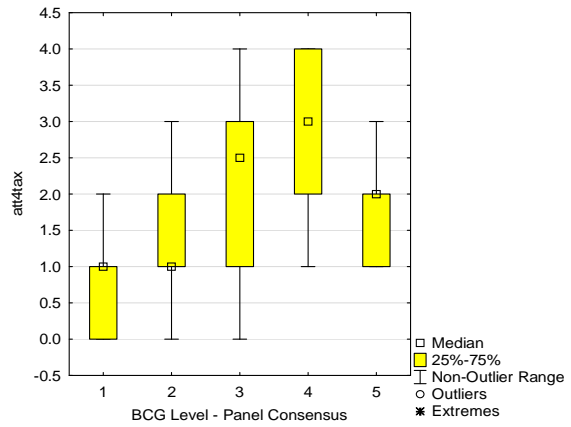
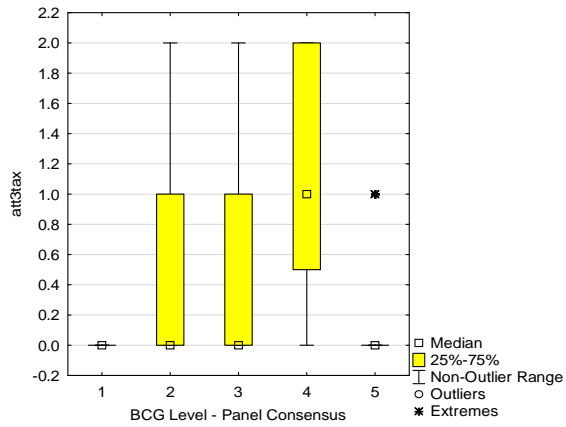
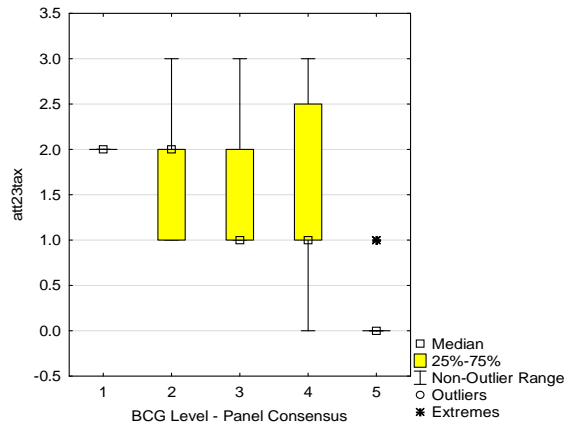
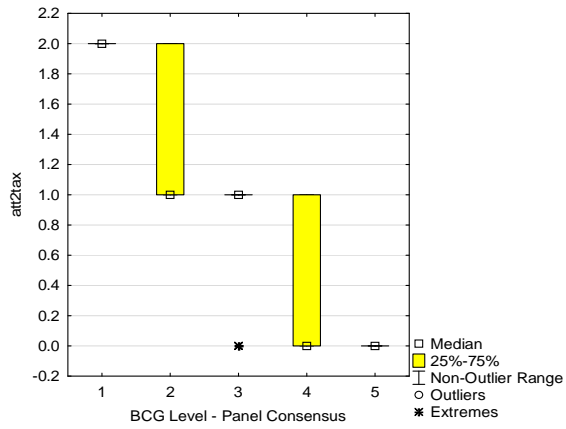
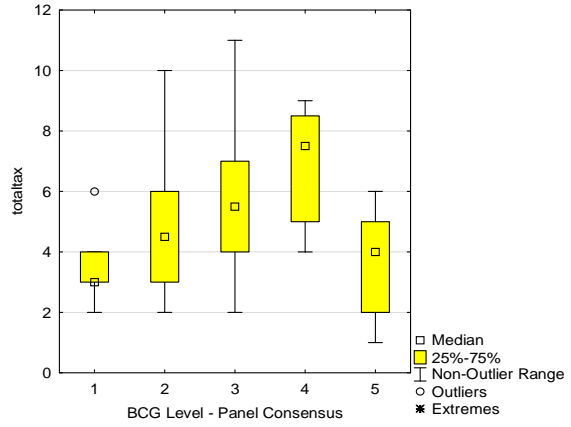
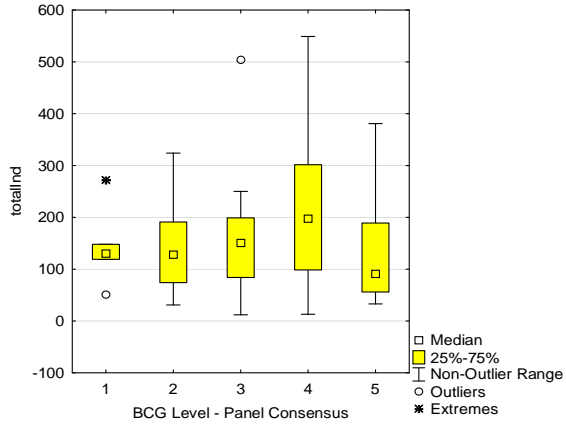
Box plots were generated to examine the distributions of metric values across BCG levels. Table F1 contains descriptions of the metric codes that are on the y-axes of the plots.

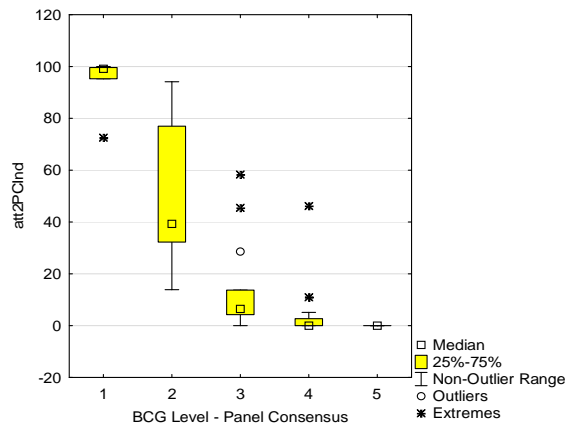
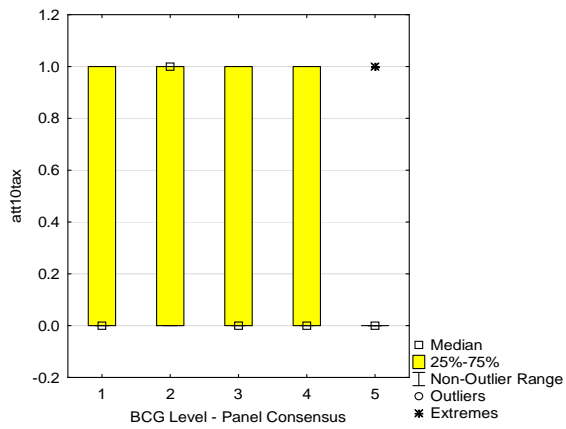
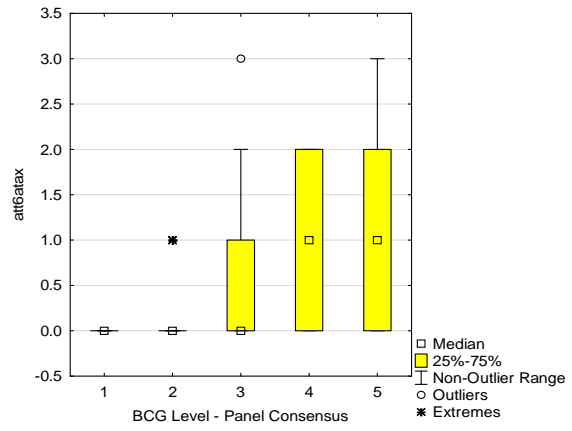
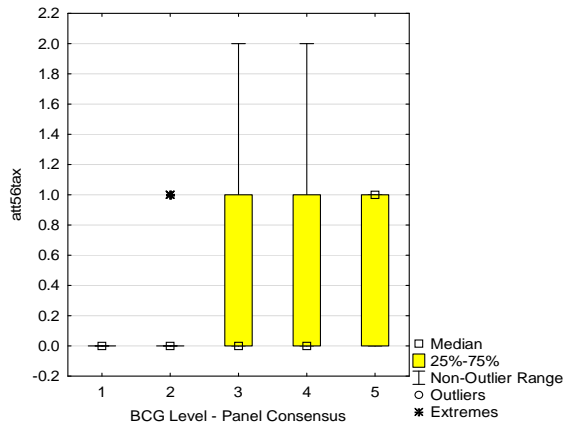
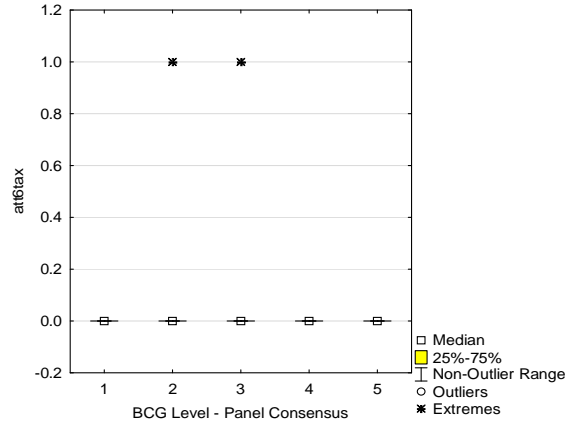
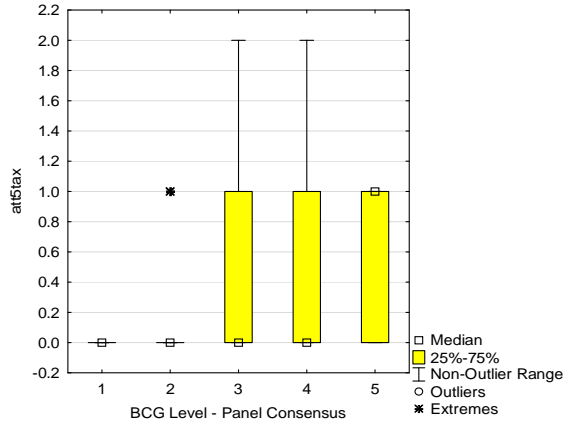
Table F1. Descriptions of the metric codes that are on the y-axes of the box plots.

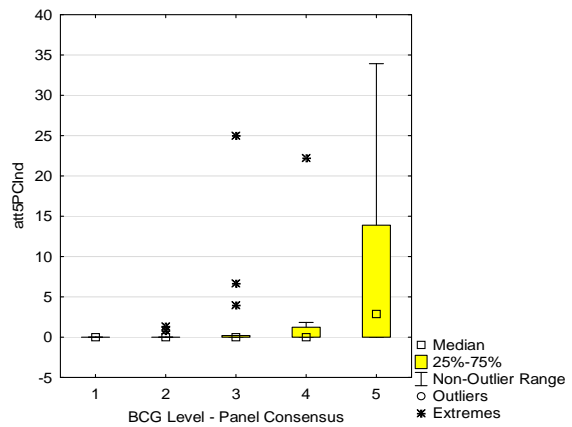
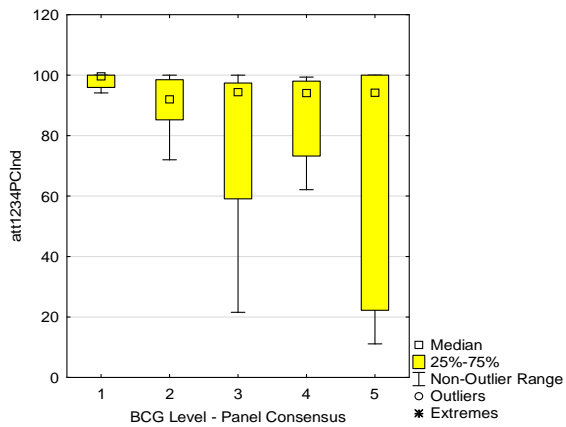
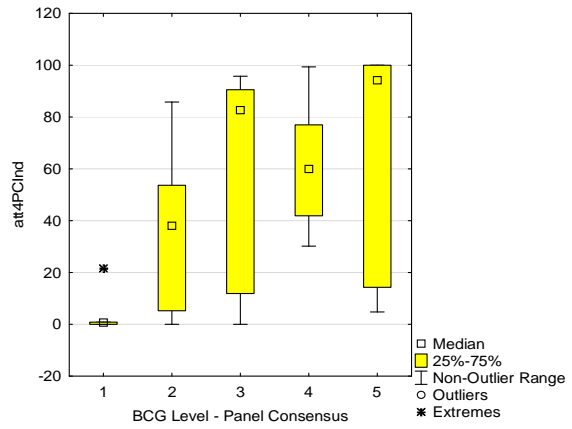
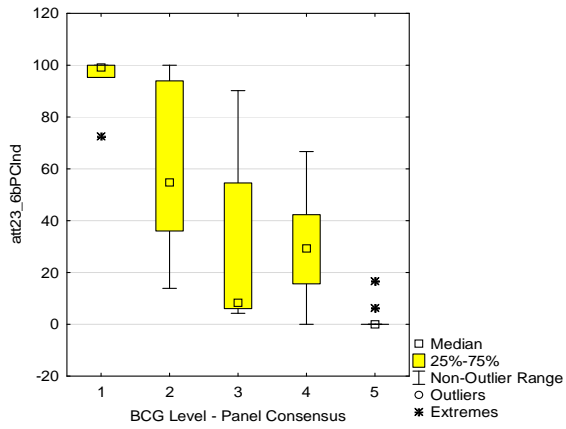
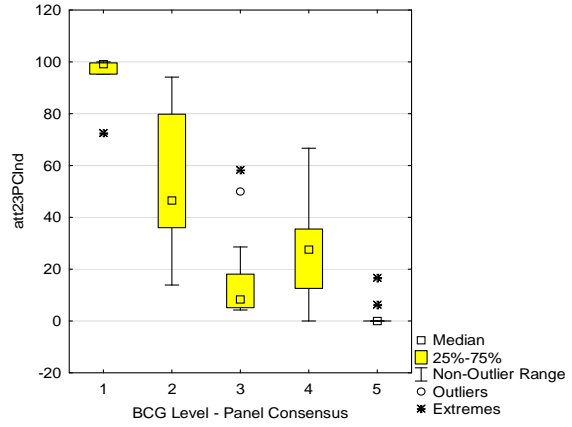
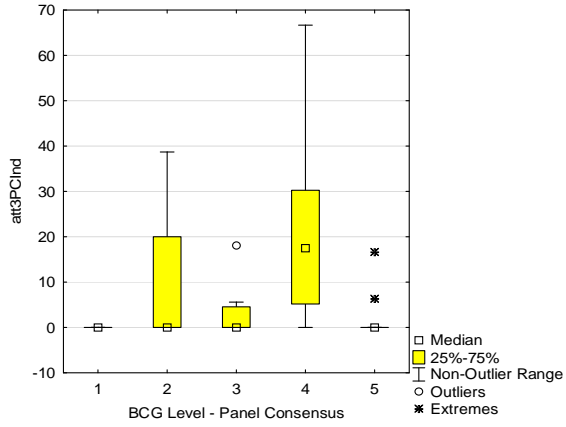
| Metric Code | Description |
|--------------------|---|
| totalInd | Number of total individuals |
| totaltax | Number of total taxa |
| att2tax | Number of Attribute II taxa |
| att23tax | Number of Attribute II + III taxa |
| att3tax | Number of Attribute III taxa |
| att4tax | Number of Attribute IV taxa |
| att5tax | Number of Attribute V taxa |
| att6tax | Number of Attribute VI taxa |
| att56tax | Number of Attribute V + VI taxa |
| att6atax | Number of Attribute VIa taxa |
| att10tax | Number of Attribute X taxa |
| att2PCInd | % Attribute II individuals |
| att23PCInd | % Attribute II + III individuals |
| att3PCInd | % Attribute III individuals |
| att23_6bPCInd | % Attribute II + III + VIb individuals |
| att4PCInd | % Attribute IV individuals |
| att1234PCInd | % Attribute I + II + III + IV individuals |
| att5PCInd | % Attribute V individuals |
| att6PCInd | % Attribute VI individuals |
| att5_6PCInd | % Attribute V + VI individuals |
| att5_6aPCInd | % Attribute V + VIa individuals |
| att6_6aPCInd | % Attribute VI + VIa individuals |
| att6b_PCInd | % Attribute VIb individuals |
| att6ab_PCInd | % Attribute VIa + VIb individuals |
| att10PCInd | % Attribute X individuals |
| att2Pctax | % Attribute II taxa |
| att3Pctax | % Attribute III taxa |
| att23Pctax | % Attribute II + III taxa |
| att23_6bPctax | % Attribute II + III + VIb taxa |
| att4Pctax | % Attribute IV taxa |
| att234_6bPctax | % Attribute I + II + III + IV + VIb taxa |
| att5Pctax | % Attribute V taxa |
| att6Pctax | % Attribute VI taxa |
| att6aPctax | % Attribute VIa taxa |
| att6bPctax | % Attribute VIb taxa |
| att10Pctax | % Attribute X taxa |
| att4Dom | % Most dominant Attribute IV taxon |

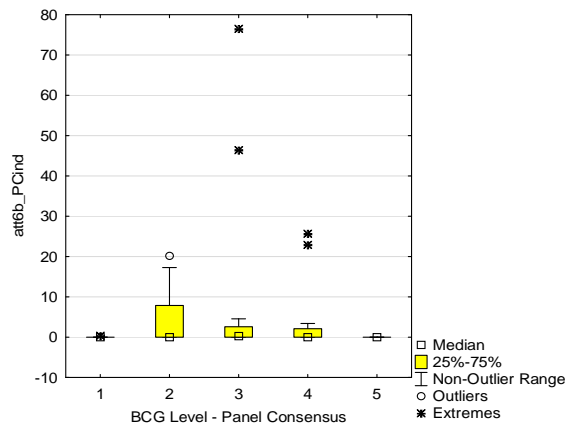
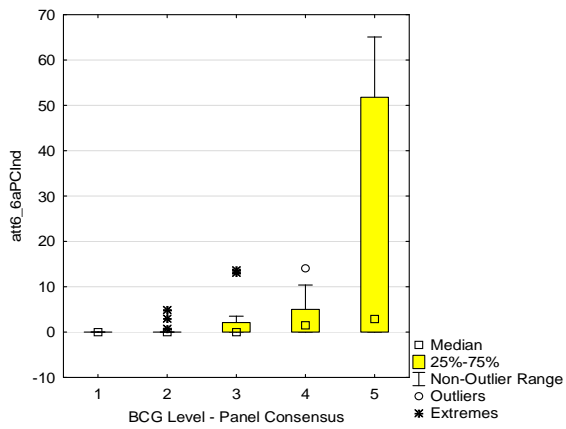
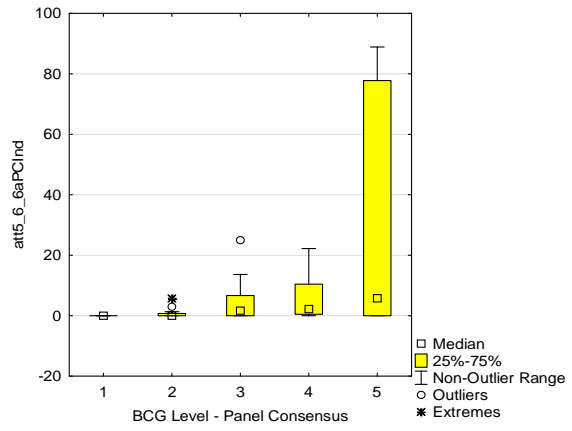
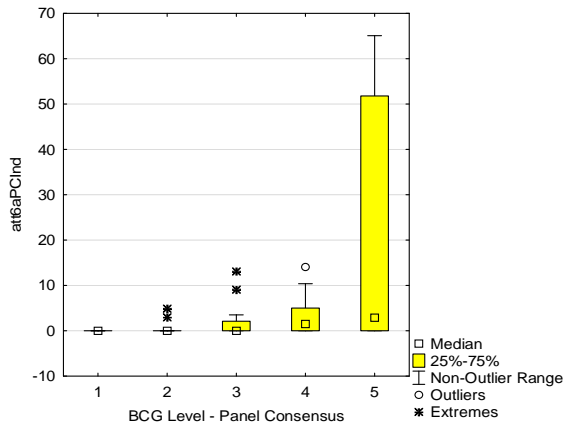
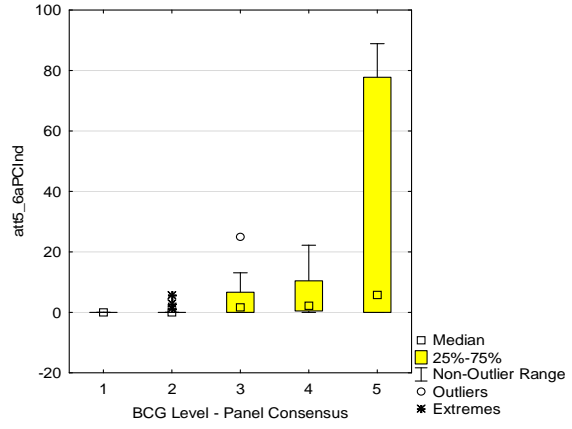
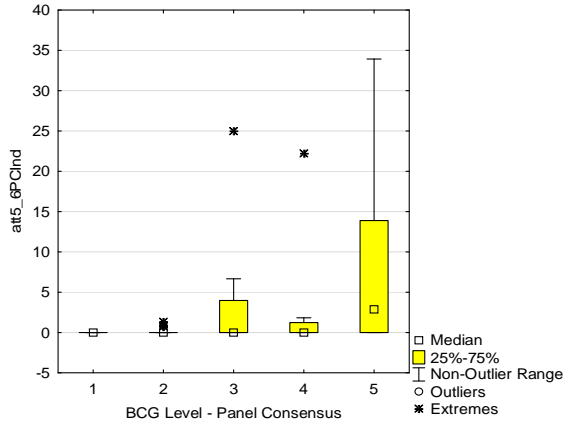
Table F1 continued...

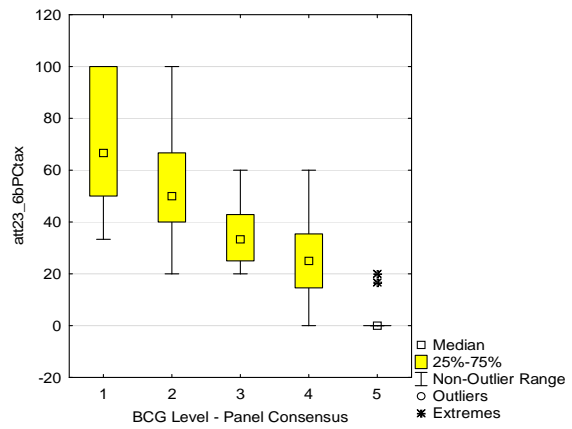
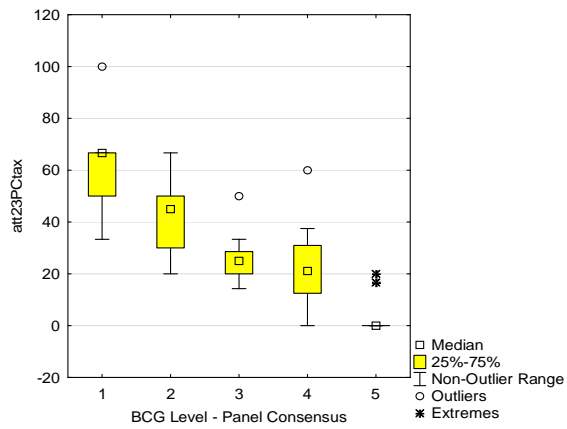
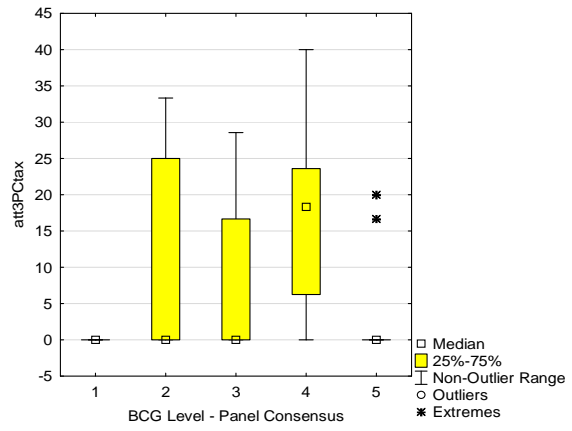
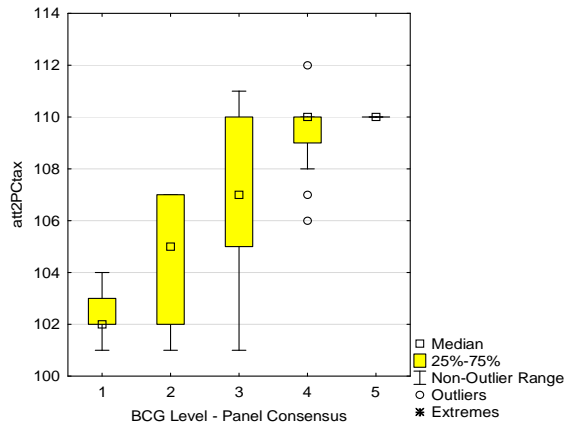
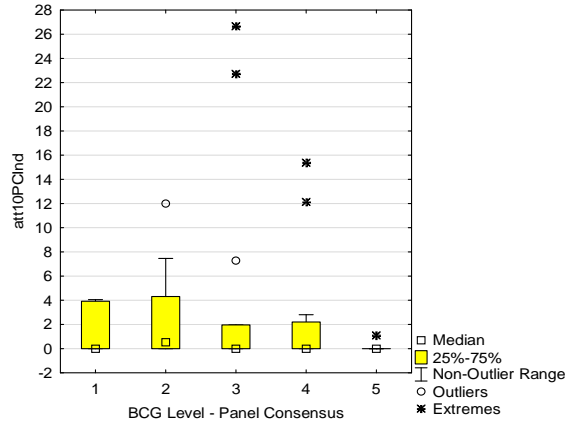
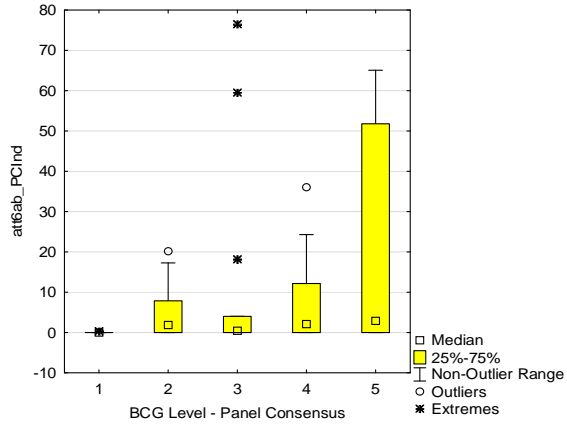
| Metric Code | Description |
|---------------------------|--|
| att5Dom | % Most dominant Attribute V taxon |
| att6Dom | % Most dominant Attribute VI taxon |
| att6aDom | % Most dominant Attribute VIa taxon |
| att566aDom | % Most dominant Attribute V + VI + VIa taxon |
| BrookTroutPCInd | % Wild brook trout individuals |
| BrownTroutPCInd | % Brown trout individuals |
| BKT_TotalSalm_PCInd | (# Wild brook trout individuals/# total salmonid individuals)*100 |
| att6bSalm_TotalSalm_PCInd | (# Attribute 6b salmonid individuals/# total salmonid individuals)*100 |
| Centrarchid_tax | Number of Centrarchidae taxa |
| Centrarchid_PCtax | Percent Centrarchidae taxa |
| Centrarchid_PCInd | Percent Centrarchidae individuals |
| Salmonid_tax | Number of Salmonidae taxa |
| Salmonid_PCtax | Percent Salmonidae taxa |
| Salmonid_PCInd | Percent Salmonidae individuals |
| Cyprin_tax | Number of Cyprinidae taxa |
| Cyprin_PCtax | Percent Cyprinidae taxa |
| Cyprin_PCInd | Percent Cyprinidae individuals |
| BNDCCCMWS_PCInd | % Black nose dace + % creek chub + % cutlips minnow + % white sucker individuals |
| totaltax5Plus | Number of total taxa, counting only taxa with 5 or more individuals |
| Shan_base_2 | Shannon-wiener diversity index (base 2) |
| Evenness | Evenness |
| totalDens_m2 | Total density/meter ² |
| totalDens_100m2 | Total density/100 meter ² |
| totaltaxNoStocked | Total number of taxa, not counting stocked taxa |

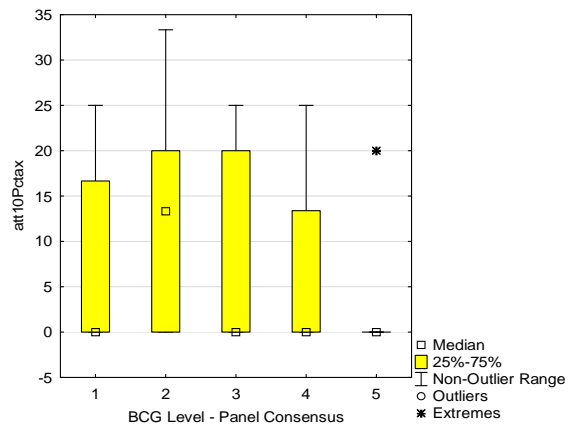
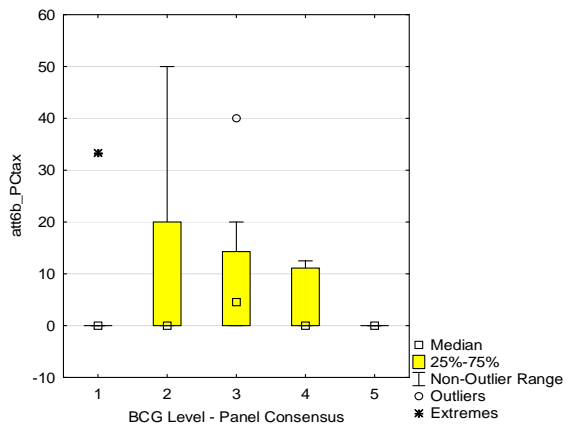
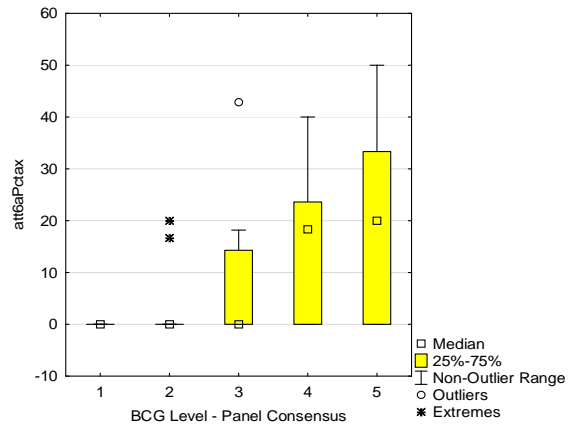
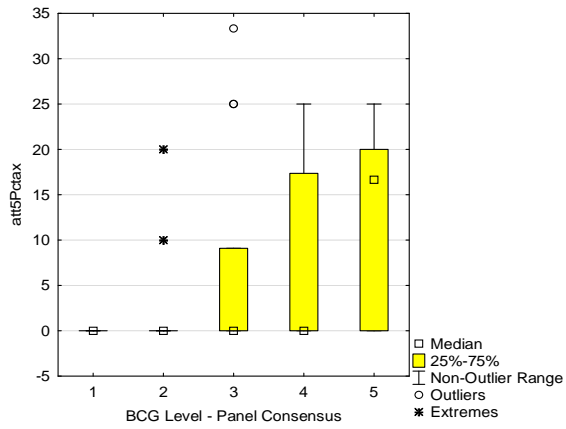
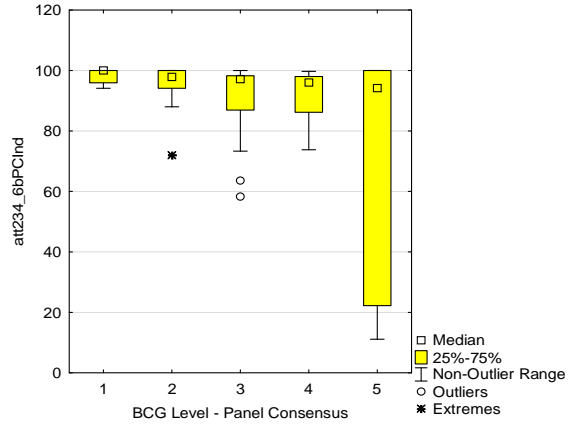
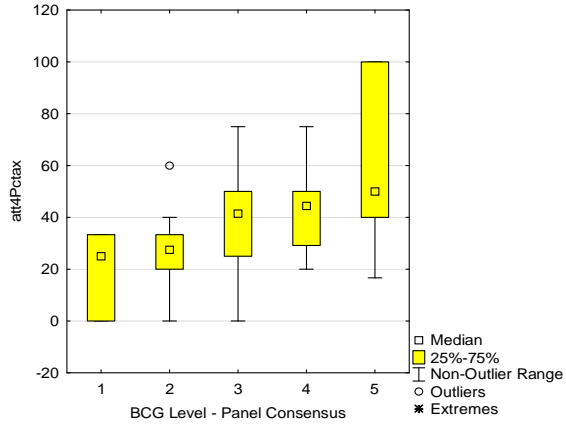


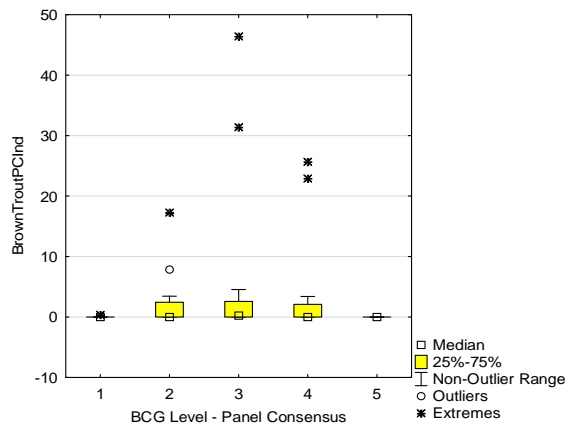
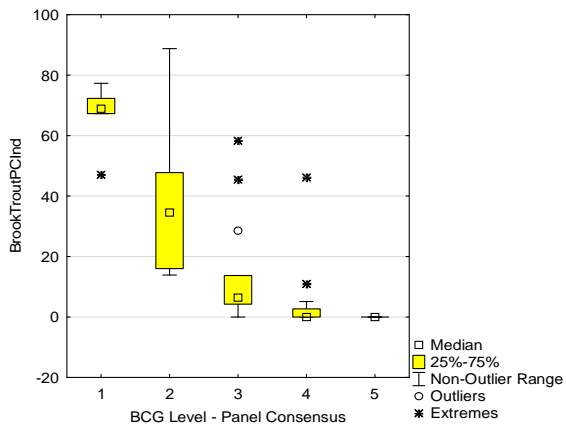
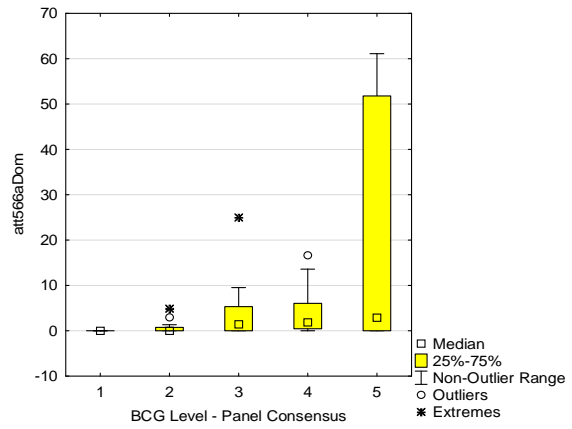
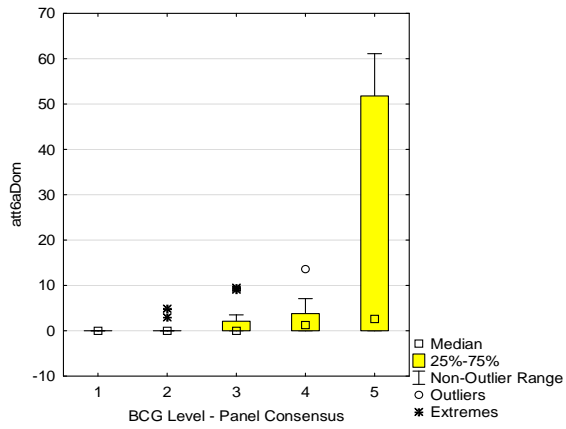
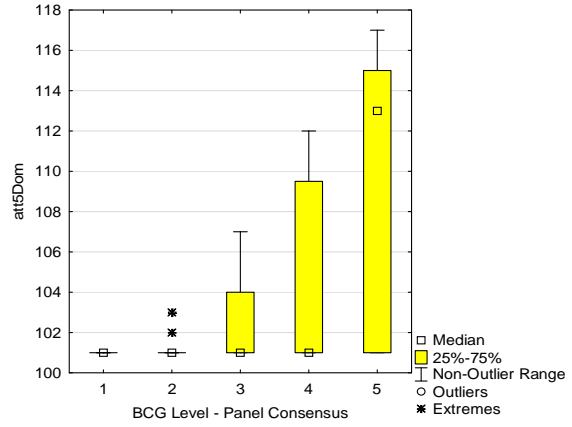
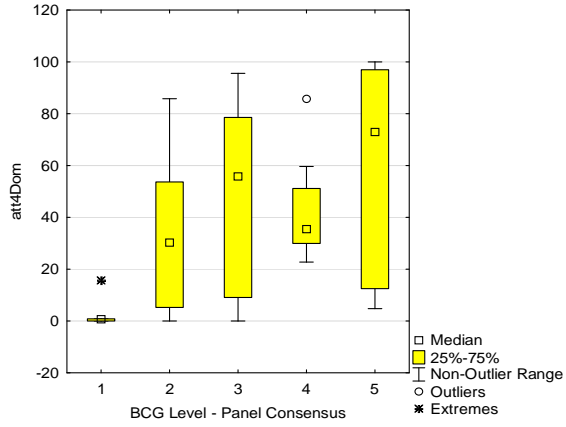


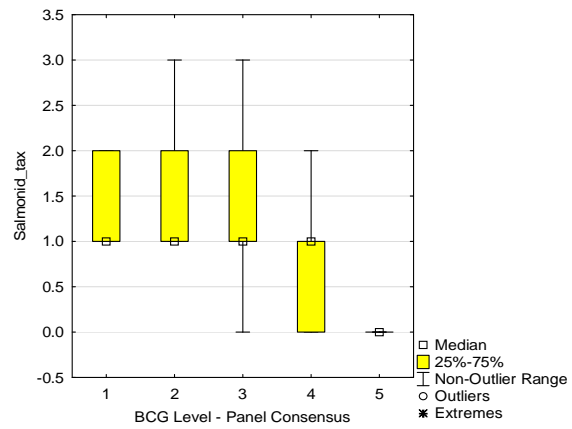
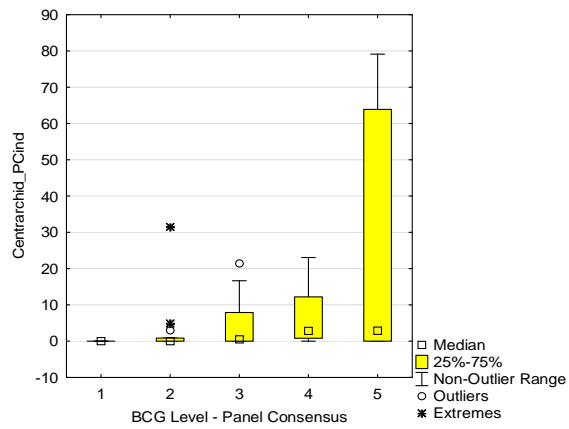
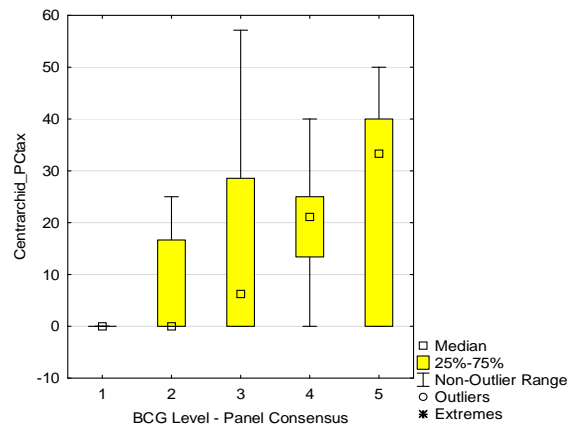
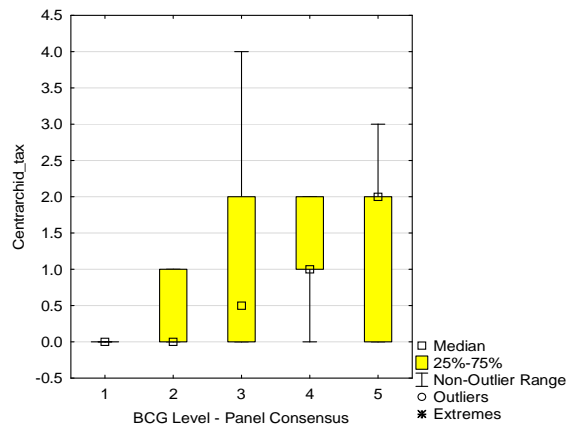
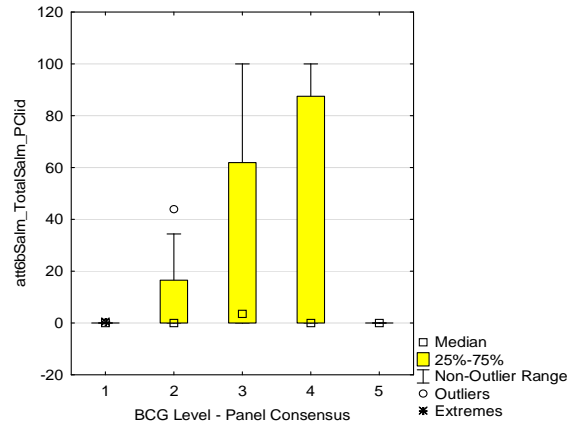
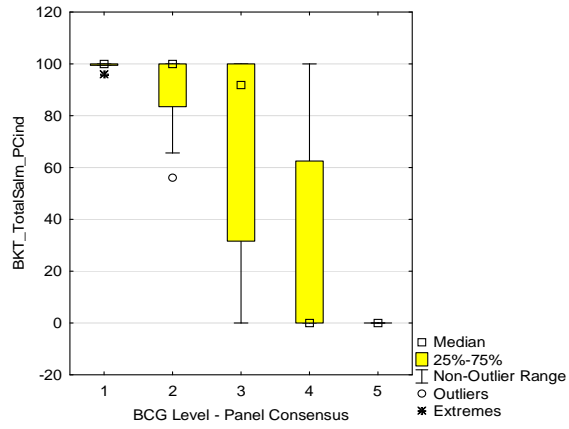


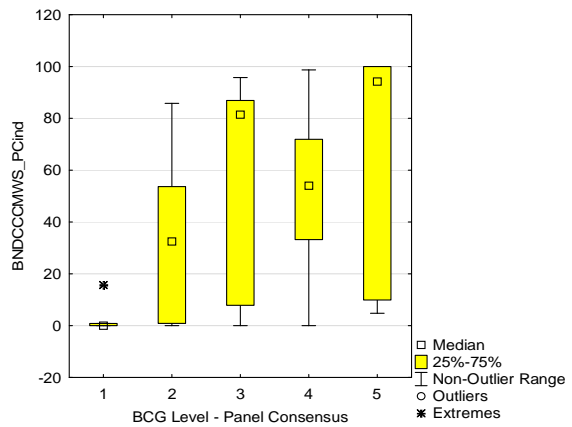
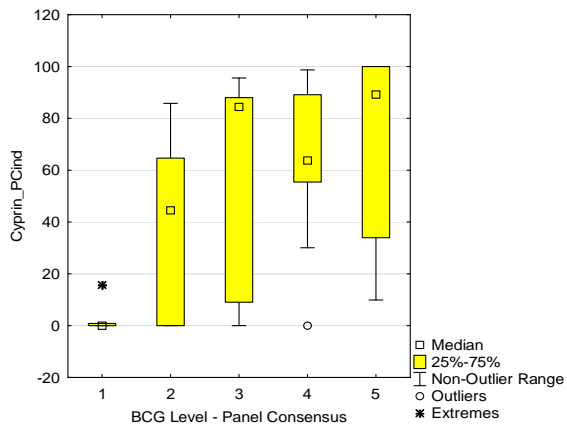
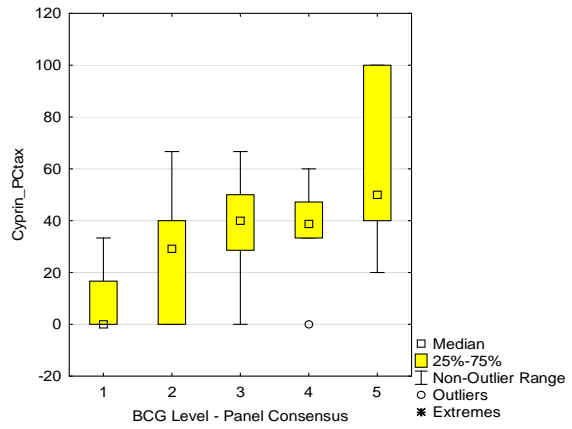
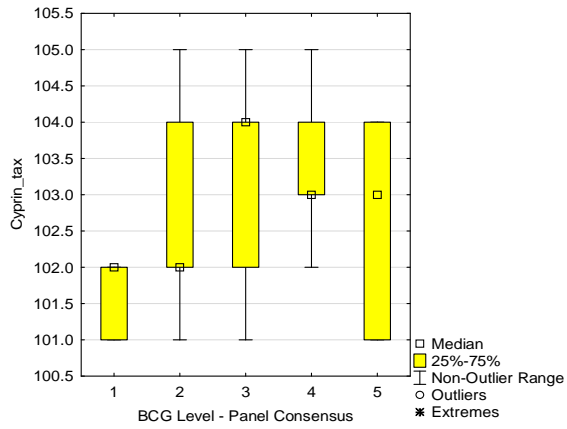
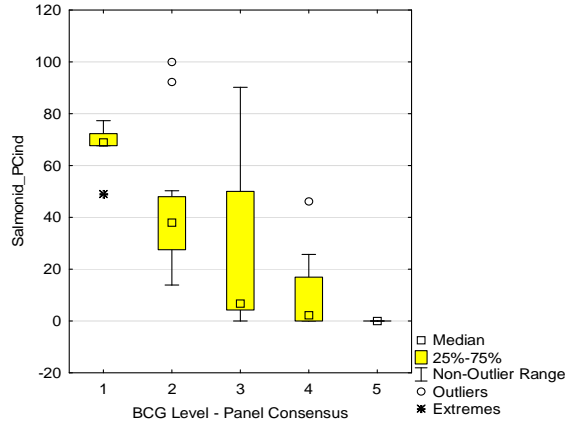
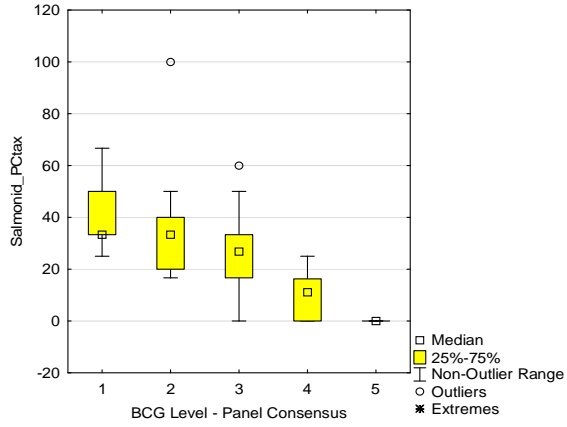


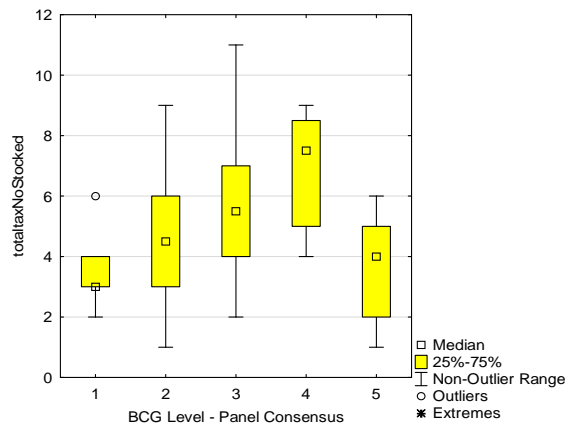
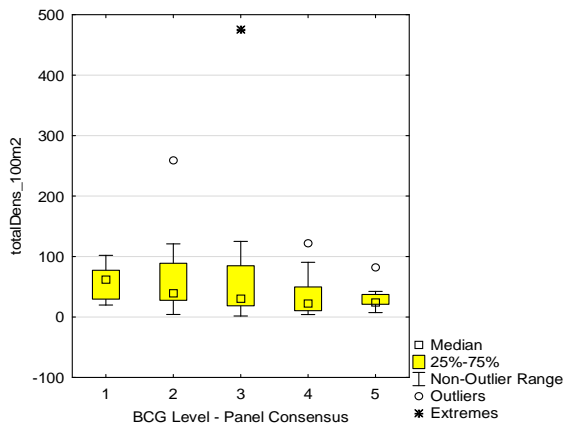
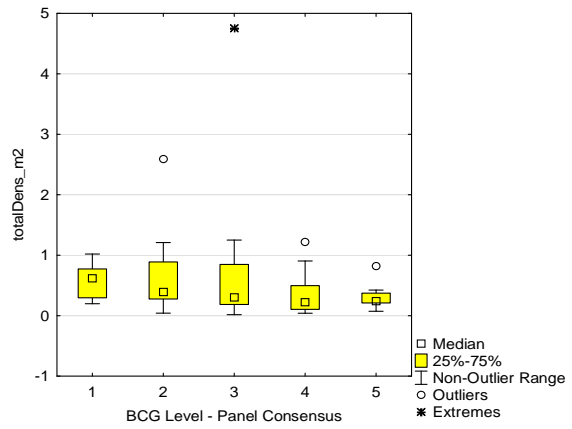
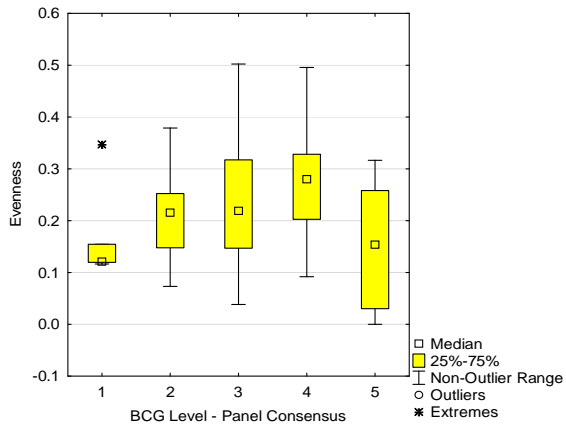
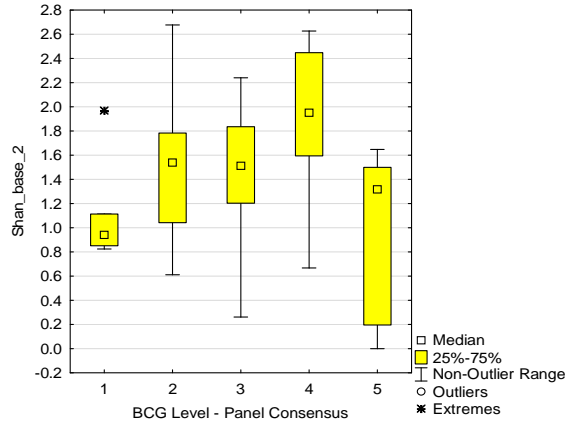
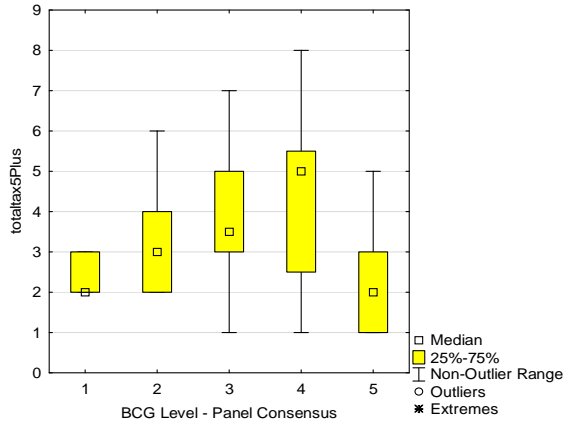












APPENDIX G

Small-cold BCG Level Assignments

Appendix G. Participants made BCG level assignments on 40 small-cold samples for the calibration exercise and 14 samples for the validation exercise. Samples were assessed using the scoring scale shown in Table G1.

Table G1. Scoring scale that was used for making BCG level assignments.

| | |
|-------|----|
| best | 1 |
| | 1- |
| | 2+ |
| | 2 |
| | 2- |
| | 3+ |
| | 3 |
| | 3- |
| | 4+ |
| | 4 |
| | 4- |
| | 5+ |
| | 5 |
| | 5- |
| | 6+ |
| | 6 |
| worst | 6- |

Table G-2. BCG level assignments and sample information for *small-cold* samples that were assessed during the *calibration* exercise. BCG level assignments are as follows: Final=consensus BCG level (=the assignment made by the majority of participants), without the + or - (2+ and 2- were assigned to level 2, etc.); Best= the best BCG level assignment assigned by a participant (based on the scoring scale in Table G1); Worst=the worst BCG level assignment given by a participant; Samples are highlighted in yellow if the consensus call from the panelists is different from the primary call from the model.

| Fish SampID | Collection Date | Station ID | Waterbody Name | Phase 1 Class | Panelist consensus | | | Primary Model | Notes |
|-------------|-----------------|------------|----------------------|---------------|--------------------|------|-------|---------------|-------------------------------------|
| | | | | | Final | Best | Worst | | |
| 13245 | 6/10/2008 | 23 | Bunnell Brook | small_cold | 5 | 4 | 5 | 5 | |
| 3045 | 7/9/2001 | 131 | Hubbard Brook | small_cold | 1 | 1 | 3+ | 2/3 (tie) | repeat sample (round1=2; round 2=1) |
| 3044 | 7/9/2001 | 132 | Hubbard Brook | small_cold | 2 | 2+ | 3 | 2 | |
| 13313 | 8/1/2008 | 428 | Coginchaug River | small_cool | 3 | 2 | 4+ | 4 | |
| 21557 | 7/29/2010 | 670 | Brides Brook | small_cool | 2 | 1 | 2- | 2 | |
| 22345 | 9/10/2010 | 697 | Steele Brook | small_cool | 4 | 3 | 5 | 5 | |
| 14439 | 6/4/2009 | 763 | Rocky Brook | small_cold | 3 | 1 | 3- | 3 | |
| 4519 | 7/22/2003 | 911 | Beach Brook | small_cold | 2 | 1 | 3 | 1 | |
| 4468 | 7/2/2003 | 924 | Clark Creek | small_cool | 3 | 2 | 3- | 3 | |
| 8647 | 7/19/2006 | 927 | Fivemile Brook | small_cold | 3 | 3 | 4 | 3 | |
| 4430 | 6/24/2003 | 933 | Wood Creek | small_cold | 2 | 2 | 3 | 2 | |
| 4514 | 7/17/2003 | 971 | Jefferson Hill Brook | small_cold | 2 | 2+ | 3- | 2 | |
| 21547 | 7/28/2010 | 1035 | Meetinghouse Brook | small_cool | 4 | 3- | 5 | 4 | repeat sample (both rounds=4) |
| 13159 | 7/17/2008 | 1257 | Webster Brook | small_cool | 3 | 2 | 3 | 4 | |
| 5249 | 6/28/2004 | 1440 | Sages Ravine Brook | small_cold | 1 | 1 | 2+ | 1 | |
| 21322 | 7/21/2010 | 1456 | Bone Mill Brook | small_cold | 1 | 1 | 2 | 1 | |
| 6557 | 7/12/2005 | 1659 | Cedar Swamp Brook | small_cold | 5 | 5 | 6 | 5 | |
| 6558 | 7/12/2005 | 1660 | Cedar Swamp Brook | small_cold | 5 | 5+ | 6+ | 5 | |
| 20495 | 6/11/2010 | 1916 | Thompson Brook | small_cool | 2 | 1 | 2 | 2 | repeat sample (both rounds=2) |
| 8561 | 7/11/2006 | 1939 | Sanford Brook | small_cold | 2 | 2+ | 3 | 2 | |

Table G-2. continued...

| Fish SampID | Collection Date | Station ID | Waterbody Name | Phase 1 Class | Panelist consensus | | | Primary Model | Notes |
|-------------|-----------------|------------|----------------------------------|---------------|--------------------|------|-------|---------------|---|
| | | | | | Final | Best | Worst | | |
| 8682 | 7/20/2006 | 1951 | Town Farm Brook (Clatter Valley) | small_cold | 3 | 2 | 4 | 3 | repeat sample (both rounds=3) |
| 8815 | 7/31/2006 | 1966 | Ekonk Brook | small_cool | 4 | 3 | 5+ | 4 | |
| 8919 | 8/4/2006 | 1976 | Menunketesuck River | small_cool | 4 | 3 | 4 | 2 | |
| 11067 | 7/19/2007 | 2295 | Mott Hill Brook | small_cold | 1 | 1 | 2 | 1 | |
| 20315 | 6/2/2010 | 2295 | Mott Hill Brook | small_cold | 2 | 1 | 2 | 2 | |
| 20630 | 6/17/2010 | 2342 | Brown Brook | small_cold | 2 | 2+ | 3+ | 2/3 (tie) | |
| 12733 | 6/18/2008 | 2343 | Bruce Brook | small_cool | 5 | 4- | 6 | 5 | repeat sample (both rounds=5) |
| 20256 | 6/1/2010 | 2532 | Branch Brook | small_cold | 4 | 2- | 4- | 3 | |
| 13038 | 7/7/2008 | 2533 | Straddle Brook | small_cold | 4 | 3- | 5+ | 4 | |
| 21127 | 7/9/2010 | 2634 | Green Fall River | small_cool | 2 | 2+ | 3- | 2 | repeat sample (round1=2; round 2=2/3 tie) |
| 14944 | 7/17/2009 | 2672 | Sumner Brook | small_cold | 3 | 2 | 4 | 4 | repeat sample (round1=4; round 2=3) |
| 12676 | 6/11/2008 | 2680 | Jacks Brook | small_cool | 4 | 4 | 5 | 5 | |
| 12595 | 6/9/2008 | 2693 | Lydall Brook | small_cold | 5 | 5 | 6 | 5 | |
| 12597 | 6/9/2008 | 2709 | Bigelow Brook | small_cool | 4 | 2+ | 4- | 4 | repeat sample (round1=2; round 2=4) |
| 13244 | 6/10/2008 | 2710 | Punch Brook | small_cold | 3 | 2 | 4 | 3 | |
| 13105 | 7/11/2008 | 2714 | East Mountain Brook | small_cold | 2 | 2+ | 3- | 4 | |
| 21307 | 7/16/2010 | 5845 | Stony Brook | small_cool | 2 | 1- | 4+ | 3 | repeat sample (both rounds=2) |
| 15551 | 9/9/2009 | 5923 | Gulf Brook | small_cold | 5 | 3 | 5- | 5 | |
| 20632 | 6/17/2010 | 6125 | Beebe Brook | small_cool | 3 | 2- | 4 | 4 | repeat sample (both rounds=3) |
| 21440 | 7/22/2010 | 6163 | Pease Brook | small_cool | 3 | 3 | 4 | 4 | |

Table G-3. Site information for *small-cold* fish samples that were analyzed during the BCG *calibration* exercise. Area refers to the upstream watershed area. Land use (%Devl=% developed, % Imperv= % impervious, % Natl= % natural) is for the upstream catchment area. TNC fields are derived from The Nature Conservancy’s Northeast Aquatic Habitat Classification (Olivero and Anderson 2008). TITAN thermal classes were based on Beauchene et al. 2012. Additional information (i.e. nutrient and habitat data) may available for some of the sites.

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|------------------|----------|---------|------------|---|--------|----------|--------|---|------------------------------|-------------------|---------------------|
| 23 | Bunnell Brook | -72.9657 | 41.7846 | 4.2 | Southern New England Coastal Plains and Hills | 11.7 | 4.4 | 67.3 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Cold | |
| 131 | Hubbard Brook | -72.5803 | 41.7082 | 2.2 | Connecticut Valley | 27.5 | 10.5 | 50.3 | High Gradient: $\geq 2 < 5\%$ | Low Buffered, Acidic | Cold | |
| 132 | Hubbard Brook | -72.5843 | 41.7089 | 2.2 | Connecticut Valley | 27.5 | 10.5 | 50.3 | High Gradient: $\geq 2 < 5\%$ | Low Buffered, Acidic | Cold | |
| 428 | Coginchaug River | -72.6882 | 41.4435 | 3.6 | Connecticut Valley | 5.1 | 2.7 | 78.8 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | |
| 670 | Brides Brook | -72.2419 | 41.3360 | 0.3 | Long Island Sound Coastal Lowland | 15.0 | 4.8 | 79.3 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Transitional Cool | |
| 697 | Steele Brook | -73.1153 | 41.6105 | 5.9 | Southern New England Coastal Plains and Hills | 19.3 | 8.1 | 40.7 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Transitional Cool | 1 |
| 763 | Rocky Brook | -71.8020 | 42.0134 | 0.5 | Southern New England Coastal Plains and Hills | 2.2 | 4.2 | 14.8 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Cold | |
| 911 | Beach Brook | -72.8575 | 41.9460 | 2.1 | Berkshire Transition | 1.3 | 1.6 | 95.5 | Very High Gradient: $> 5\%$ | Moderately Buffered, Neutral | Cold | |
| 924 | Clark Creek | -72.4735 | 41.4426 | 2.4 | Long Island Sound Coastal Lowland | 3.6 | 2.0 | 94.2 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Transitional Cool | |
| 927 | Fivemile Brook | -73.1597 | 41.3846 | 1.9 | Southern New England Coastal Plains and Hills | 15.9 | 4.9 | 74.3 | Very High Gradient: $> 5\%$ | Moderately Buffered, Neutral | Cold | |

Table G-3. continued...

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|----------------------|----------|---------|------------|--|--------|----------|--------|---|------------------------------|-------------------|---------------------|
| 933 | Wood Creek | -73.2362 | 41.6387 | 3.4 | Southern New England Coastal Plains and Hills | 7.9 | 3.3 | 75.9 | High Gradient: $\geq 2 < 5\%$ | Moderately Buffered, Neutral | Cold | |
| 971 | Jefferson Hill Brook | -73.1195 | 41.7477 | 2.5 | Berkshire Transition | 4.5 | 2.5 | 79.9 | Very High Gradient: $> 5\%$ | Moderately Buffered, Neutral | Cold | 1 |
| 1035 | Meetinghouse Brook | -72.8160 | 41.4913 | 4.3 | Connecticut Valley | 45.2 | 21.1 | 16.6 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Transitional Cool | |
| 1257 | Webster Brook | -72.7421 | 41.6405 | 5.4 | Connecticut Valley | 57.1 | 27.8 | 15.1 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | |
| 1440 | Sages Ravine Brook | -73.4245 | 42.0497 | 3.5 | | 0.7 | 1.8 | 22.1 | Very High Gradient: $> 5\%$ | Moderately Buffered, Neutral | Cold | 1 |
| 1456 | Bone Mill Brook | -72.3162 | 41.9250 | 2.5 | Lower Worcester Plateau/Eastern Connecticut Upland | 5.3 | 2.6 | 88.6 | High Gradient: $\geq 2 < 5\%$ | Low Buffered, Acidic | Cold | 1 |
| 1659 | Cedar Swamp Brook | -72.2790 | 41.8164 | 2.5 | Southern New England Coastal Plains and Hills | 16.4 | 5.2 | 70.0 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | 3 |
| 1660 | Cedar Swamp Brook | -72.2841 | 41.8110 | 2.5 | Southern New England Coastal Plains and Hills | 16.4 | 5.2 | 70.0 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | |
| 1916 | Thompson Brook | -72.8497 | 41.7681 | 3.9 | Connecticut Valley | 22.3 | 9.7 | 53.6 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | 1 |
| 1939 | Sanford Brook | -72.9364 | 41.4723 | 1.0 | Connecticut Valley | 14.8 | 6.6 | 76.9 | High Gradient: $\geq 2 < 5\%$ | Moderately Buffered, Neutral | Cold | |

Table G-3. continued...

| Station ID | Waterbody Name | Long | Lat | Area (mi ²) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|----------------------------------|----------|---------|-------------------------|--|--------|----------|--------|--------------------------------------|------------------------------|-------------------|---------------------|
| 1951 | Town Farm Brook (Clatter Valley) | -73.3889 | 41.5477 | 3.8 | Southern New England Coastal Plains and Hills | 14.1 | 4.9 | 60.7 | High Gradient: ≥2 < 5% | Moderately Buffered, Neutral | Cold | |
| 1966 | Ekonk Brook | -71.8652 | 41.6952 | 3.7 | Southern New England Coastal Plains and Hills | 3.8 | 2.5 | 76.5 | Moderate-High Gradient: ≥0.5 < 2% | Low Buffered, Acidic | Transitional Cool | |
| 1976 | Menunketesuck River | -72.5519 | 41.3898 | 1.9 | Southern New England Coastal Plains and Hills | 10.8 | 4.0 | 78.6 | Moderate-High Gradient: ≥0.5 < 2% | Low Buffered, Acidic | Transitional Cool | |
| 2295 | Mott Hill Brook | -72.5365 | 41.6615 | 2.8 | Southern New England Coastal Plains and Hills | 3.1 | 2.2 | 90.1 | High Gradient: ≥2 < 5% | Low Buffered, Acidic | Cold | 1 |
| 2342 | Brown Brook | -73.2799 | 41.9267 | 5.6 | Berkshire Transition | 0.5 | 1.2 | 98.6 | High Gradient: ≥2 < 5% | Moderately Buffered, Neutral | Cold | 1.5 |
| 2343 | Bruce Brook | -73.1551 | 41.1899 | 2.7 | Long Island Sound Coastal Lowland | 79.9 | 35.9 | 5.8 | Moderate-High Gradient: ≥0.5 < 2% | Moderately Buffered, Neutral | Transitional Cool | |
| 2532 | Branch Brook | -72.1256 | 41.9199 | 4.9 | Lower Worcester Plateau/Eastern Connecticut Upland | 3.8 | 2.0 | 90.4 | Moderate-High Gradient: ≥0.5 < 2% | Moderately Buffered, Neutral | Cold | 1 |
| 2533 | Straddle Brook | -72.3782 | 41.7270 | 4.4 | Southern New England Coastal Plains and Hills | 8.1 | 3.3 | 78.1 | Moderate-High Gradient: ≥0.5 < 2% | Moderately Buffered, Neutral | Cold | |

Table G-3. continued...

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|---------------------|----------|---------|------------|---|--------|----------|--------|---------------------------------------|------------------------------|-------------------|---------------------|
| 2634 | Green Fall River | -71.8159 | 41.4816 | 4.0 | Southern New England Coastal Plains and Hills | 2.8 | 2.0 | 93.5 | Moderate-High Gradient: >=0.5 < 2% | Low Buffered, Acidic | Transitional Cool | |
| 2672 | Sumner Brook | -72.6375 | 41.4835 | 1.5 | Southern New England Coastal Plains and Hills | 7.6 | 3.1 | 86.3 | High Gradient: >=2 < 5% | Moderately Buffered, Neutral | Cold | |
| 2680 | Jacks Brook | -73.1170 | 41.4420 | 1.8 | Southern New England Coastal Plains and Hills | 7.9 | 3.3 | 78.5 | Moderate-High Gradient: >=0.5 < 2% | Moderately Buffered, Neutral | Transitional Cool | |
| 2693 | Lydall Brook | -72.5215 | 41.7952 | 2.8 | Connecticut Valley | 39.0 | 29.8 | 45.8 | Moderate-High Gradient: >=0.5 < 2% | Low Buffered, Acidic | Cold | 2 |
| 2709 | Bigelow Brook | -72.5530 | 41.7846 | 3.2 | Connecticut Valley | 73.8 | 32.5 | 14.1 | Moderate-High Gradient: >=0.5 < 2% | Low Buffered, Acidic | Transitional Cool | 1 |
| 2710 | Punch Brook | -72.9259 | 41.7815 | 1.7 | Southern New England Coastal Plains and Hills | 8.2 | 3.4 | 84.7 | High Gradient: >=2 < 5% | Moderately Buffered, Neutral | Cold | |
| 2714 | East Mountain Brook | -72.9778 | 41.8772 | 3.1 | Berkshire Transition | 9.6 | 3.5 | 80.2 | High Gradient: >=2 < 5% | Moderately Buffered, Neutral | Cold | |
| 5845 | Stony Brook | -72.1730 | 41.3691 | 2.8 | Long Island Sound Coastal Lowland | 19.7 | 7.3 | 65.5 | Moderate-High Gradient: >=0.5 < 2% | Low Buffered, Acidic | Transitional Cool | |
| 5923 | Gulf Brook | -72.7807 | 41.3805 | 1.1 | Connecticut Valley | 0.0 | 1.3 | 100.0 | High Gradient: >=2 < 5% | Moderately Buffered, Neutral | Cold | |

Table G-3. continued...

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|-------------------|-----------------------|-------------|------------|-------------------|---|---------------|-----------------|---------------|---------------------------------------|---------------------------------|--------------------------|----------------------------|
| 6125 | Beebe Brook | -73.4807 | 41.8406 | 2.7 | Western New England Marble Valleys | 2.1 | 1.5 | 96.2 | Very Low Gradient: <0.02% | Moderately Buffered, Neutral | Transitional Cool | |
| 6163 | Pease Brook | -72.2349 | 41.6418 | 2.7 | Southern New England Coastal Plains and Hills | 5.6 | 2.9 | 72.3 | Moderate-High Gradient: >=0.5 < 2% | Moderately Buffered, Neutral | Transitional Cool | |

Table G-4. BCG level assignments and sample information for *small-cold* samples that were assessed during the *validation* exercise. BCG level assignments are as follows: Final=consensus BCG level (=the assignment made by the majority of participants), without the + or - (2+ and 2- were assigned to level 2, etc.); Best= the best BCG level assignment assigned by a participant (based on the scoring scale in Table G1); Worst=the worst BCG level assignment given by a participant; Samples are highlighted in yellow if the consensus call from the panelists is different from the primary call from the model (model assignment 1=primary; 2=secondary; tie=tie between primary and secondary).

| Fish SampID | Collection Date | Station ID | Waterbody Name | Phase 1 Class | Panelist consensus | | | Primary Model | Notes |
|-------------|-----------------|------------|-----------------------------|---------------|--------------------|------|-------|---------------|----------------------------------|
| | | | | | Final | Best | Worst | | |
| 12725 | 6/16/2008 | 260 | Pequabuck River | small_cold | 5 | 4+ | 5 | 5 | |
| 12686 | 6/13/2008 | 1004 | Wash Brook | small_cool | 4 | 3+ | 4- | 4 | |
| 13155 | 7/17/2008 | 1338 | Belcher Brook | small_cool | 4 | 3- | 4- | 4 | |
| 5255 | 6/29/2004 | 1445 | Cobble Brook | small_cold | 4 | 3- | 4- | 4 | |
| 5324 | 7/7/2004 | 1454 | Shepaug River, tributary to | small_cold | 2 | 1- | 2- | 2 | |
| 5373 | 7/9/2004 | 1456 | Bone Mill Brook | small_cold | 1 | 1 | 1 | 1 | |
| 8678 | 7/20/2006 | 1518 | Lee Brook | small_cold | 3 | 2- | 5+ | 3 | |
| 13279 | 7/25/2008 | 1625 | Beaver Brook | small_cold | 3/4 (tie) | 3 | 4 | 3/4 (tie) | |
| 6559 | 7/12/2005 | 1661 | Nelson Brook | small_cool | 5 | 4- | 6 | 5 | |
| 14337 | 8/24/2007 | 2291 | Branch Brook | small_cold | 4 | 1- | 4+ | 3/4 (tie) | |
| 15533 | 9/10/2009 | 2407 | Steele Brook | small_cold | 5 | 4 | 6+ | 5 | |
| 13150 | 7/14/2008 | 2719 | Cherry Brook, tributary to | small_cold | 3 | 3 | 4+ | 3 | model assignment is close to a 4 |
| 13203 | 7/18/2008 | 2724 | North Brook | small_cold | 3 | 3+ | 4+ | 3 | |
| 21473 | 7/27/2010 | 5346 | Jordan Brook | small_cool | 2 | 2+ | 3 | 2/3 (tie) | |

Table G-5. Site information for *small-cold* fish samples that were analyzed during the BCG *validation* exercise. Area refers to the upstream watershed area. Land use (%Devl=% developed, % Imperv= % impervious, % Natl= % natural) is for the upstream catchment area. TNC fields are derived from The Nature Conservancy's Northeast Aquatic Habitat Classification (Olivero and Anderson 2008). Additional information (i.e. nutrient and habitat data) may available for some of the sites.

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class (median) |
|------------|-----------------------------|-----------|----------|------------|--|--------|----------|--------|---|------------------------------|-------------------|------------------------------|
| 260 | Pequabuck River | -73.01503 | 41.67887 | 2.3 | Southern New England Coastal Plains and Hills | 21.8 | 8.8 | 48.2 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Cold | |
| 1004 | Wash Brook | -72.73766 | 41.81698 | 3.8 | Connecticut Valley | 26.1 | 9.2 | 34.4 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | |
| 1338 | Belcher Brook | -72.75766 | 41.60498 | 3.9 | Connecticut Valley | 28.8 | 9.1 | 52.9 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Transitional Cool | 3 |
| 1445 | Cobble Brook | -73.45424 | 41.74538 | 4.5 | Berkshire Transition | 5.9 | 2.9 | 78.0 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | 1 |
| 1454 | Shepaug River, tributary to | -73.34472 | 41.59025 | 2.1 | Southern New England Coastal Plains and Hills | 9.2 | 3.9 | 55.1 | High Gradient: $\geq 2 < 5\%$ | Moderately Buffered, Neutral | Cold | |
| 1456 | Bone Mill Brook | -72.31624 | 41.92499 | 2.5 | Lower Worcester Plateau/Eastern Connecticut Upland | 5.3 | 2.6 | 88.6 | High Gradient: $\geq 2 < 5\%$ | Low Buffered, Acidic | Cold | 1 |

Table G-5. continued...

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Dev 1 | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class (median) |
|------------|----------------------------|-----------|----------|------------|--|---------|----------|--------|---|------------------------------|-------------------|------------------------------|
| 1518 | Lee Brook | -73.22893 | 41.43336 | 1.2 | Southern New England Coastal Plains and Hills | 17.9 | 5.3 | 74.9 | High Gradient: $\geq 2 < 5\%$ | Moderately Buffered, Neutral | Cold | |
| 1625 | Beaver Brook | -72.99575 | 41.95125 | 2.0 | Lower Berkshire Hills | 3.9 | 2.0 | 89.8 | High Gradient: $\geq 2 < 5\%$ | Moderately Buffered, Neutral | Cold | |
| 1661 | Nelson Brook | -72.28434 | 41.81076 | 1.8 | Southern New England Coastal Plains and Hills | 14.9 | 4.8 | 74.4 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | |
| 2291 | Branch Brook | -72.12450 | 41.91081 | 4.9 | Lower Worcester Plateau/Eastern Connecticut Upland | 3.8 | 2.0 | 90.4 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | 1 |
| 2407 | Steele Brook | -73.13100 | 41.61750 | 1.6 | Southern New England Coastal Plains and Hills | 14.5 | 7.1 | 30.1 | High Gradient: $\geq 2 < 5\%$ | Low Buffered, Acidic | Cold | |
| 2719 | Cherry Brook, tributary to | -72.89249 | 41.89822 | 1.8 | Berkshire Transition | 4.8 | 2.4 | 91.6 | High Gradient: $\geq 2 < 5\%$ | Moderately Buffered, Neutral | Cold | |
| 2724 | North Brook | -73.01374 | 41.84648 | 1.9 | Berkshire Transition | 10.3 | 3.6 | 80.1 | High Gradient: $\geq 2 < 5\%$ | Moderately Buffered, Neutral | Cold | |
| 5346 | Jordan Brook | -72.15078 | 41.36723 | 3.5 | Long Island Sound Coastal Lowland | 18.7 | 6.1 | 64.9 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Transitional Cool | |

APPENDIX H

Box plots of metrics for medium-large samples
that were assessed

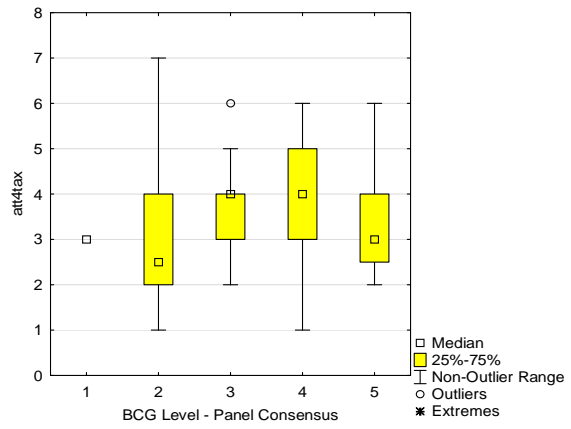
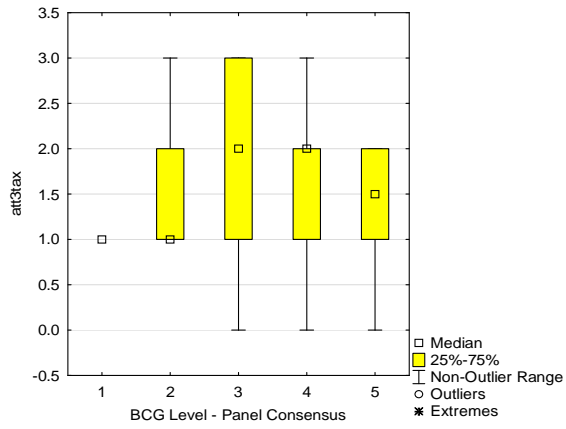
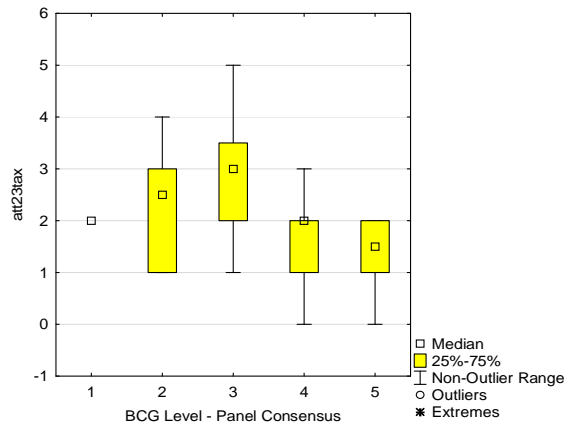
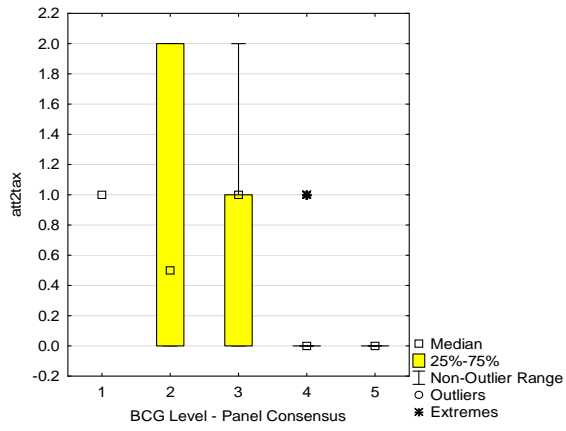
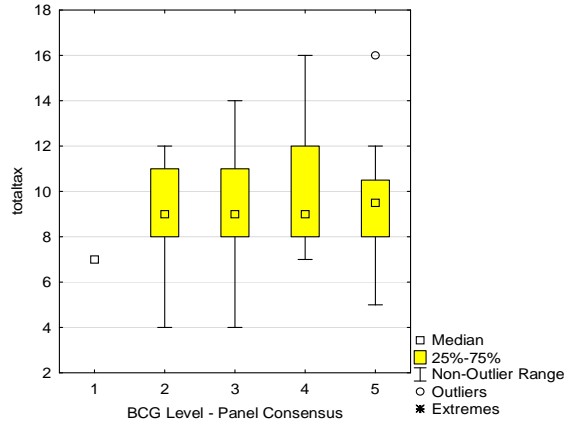
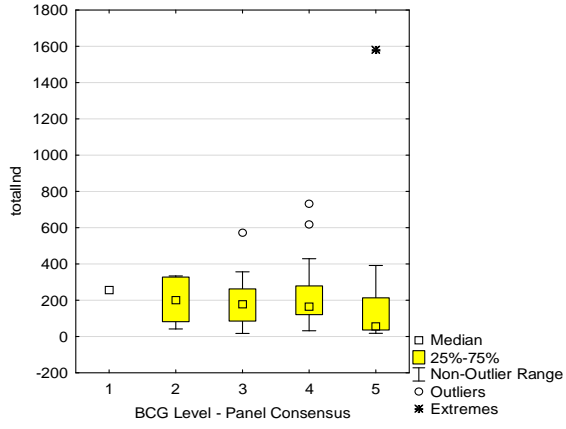
Box plots were generated to examine the distributions of metric values across BCG levels. Table F1 contains descriptions of the metric codes that are on the y-axes of the plots.

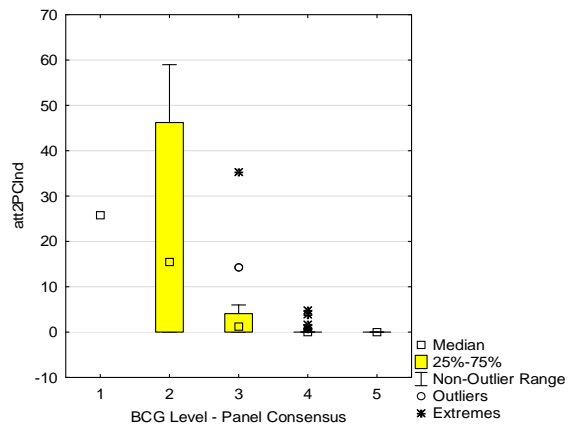
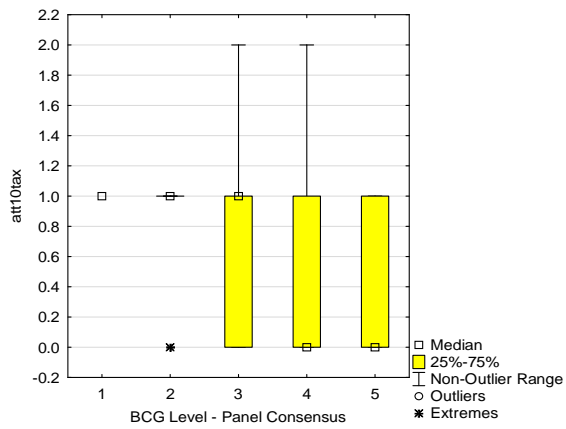
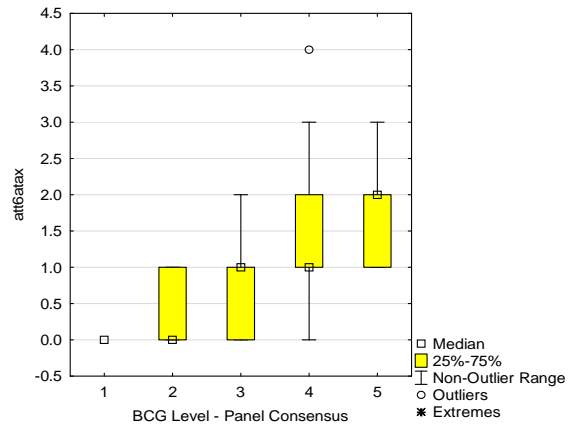
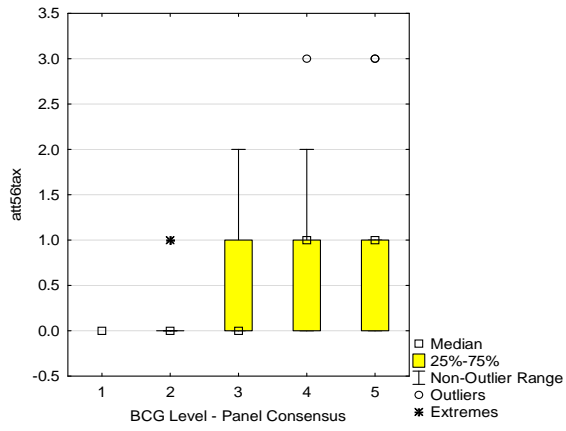
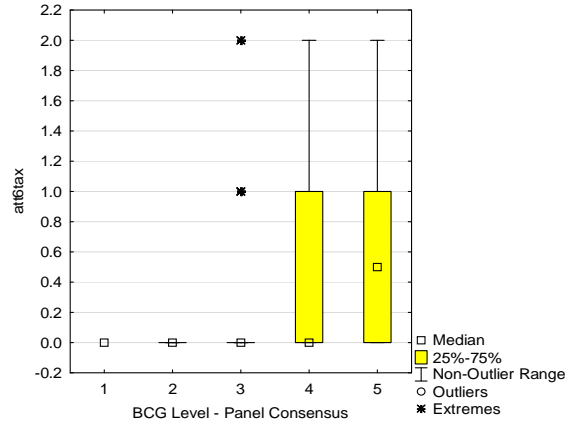
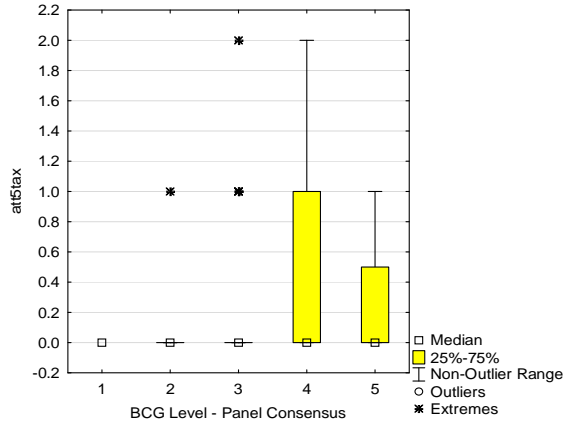
Table H1. Descriptions of the metric codes that are on the y-axes of the box plots.

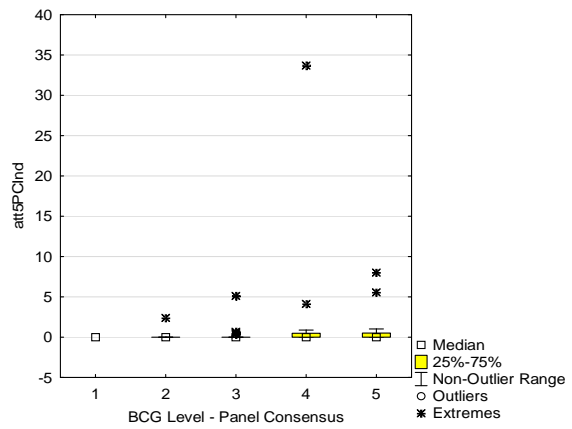
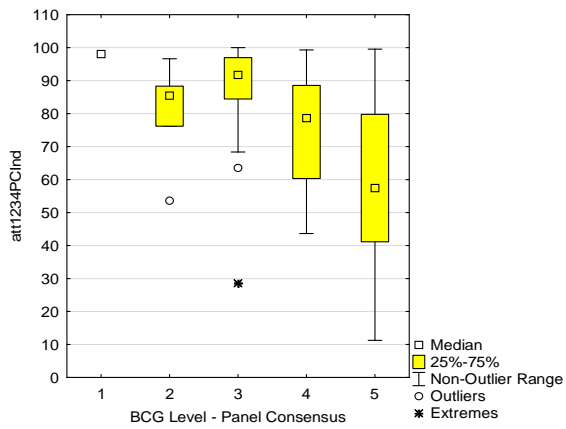
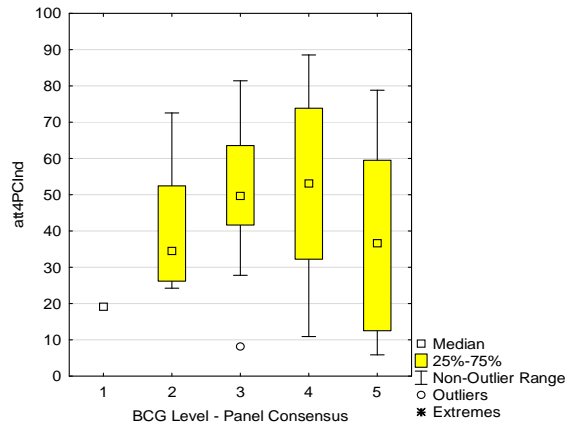
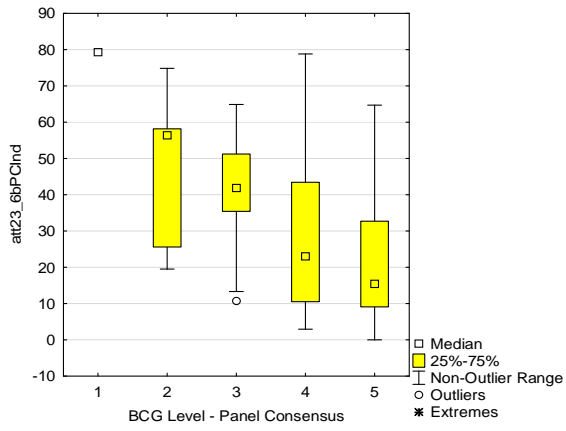
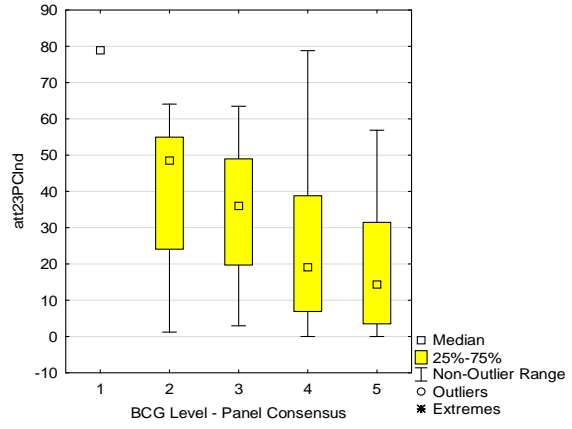
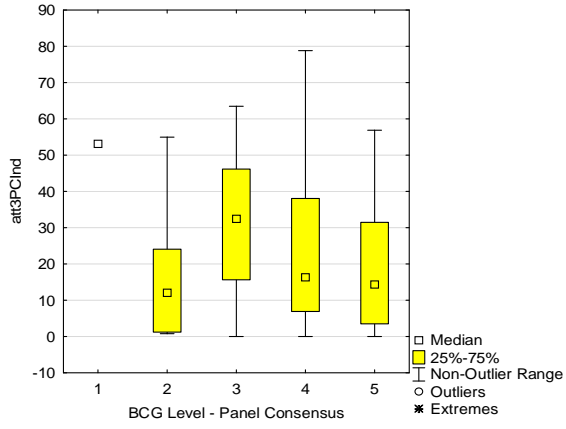
| Metric Code | Description |
|--------------------|---|
| totalInd | Number of total individuals |
| totaltax | Number of total taxa |
| att2tax | Number of Attribute II taxa |
| att23tax | Number of Attribute II + III taxa |
| att3tax | Number of Attribute III taxa |
| att4tax | Number of Attribute IV taxa |
| att5tax | Number of Attribute V taxa |
| att6tax | Number of Attribute VI taxa |
| att56tax | Number of Attribute V + VI taxa |
| att6atax | Number of Attribute VIa taxa |
| att10tax | Number of Attribute X taxa |
| att2PCInd | % Attribute II individuals |
| att23PCInd | % Attribute II + III individuals |
| att3PCInd | % Attribute III individuals |
| att23_6bPCInd | % Attribute II + III + VIb individuals |
| att4PCInd | % Attribute IV individuals |
| att1234PCInd | % Attribute I + II + III + IV individuals |
| att5PCInd | % Attribute V individuals |
| att6PCInd | % Attribute VI individuals |
| att5_6PCInd | % Attribute V + VI individuals |
| att5_6aPCInd | % Attribute V + VIa individuals |
| att6_6aPCInd | % Attribute VI + VIa individuals |
| att6b_PCInd | % Attribute VIb individuals |
| att6ab_PCInd | % Attribute VIa + VIb individuals |
| att10PCInd | % Attribute X individuals |
| att2Pctax | % Attribute II taxa |
| att3Pctax | % Attribute III taxa |
| att23Pctax | % Attribute II + III taxa |
| att23_6bPctax | % Attribute II + III + VIb taxa |
| att4Pctax | % Attribute IV taxa |
| att234_6bPctax | % Attribute I + II + III + IV + VIb taxa |
| att5Pctax | % Attribute V taxa |
| att6Pctax | % Attribute VI taxa |
| att6aPctax | % Attribute VIa taxa |
| att6bPctax | % Attribute VIb taxa |
| att10Pctax | % Attribute X taxa |
| att4Dom | % Most dominant Attribute IV taxon |

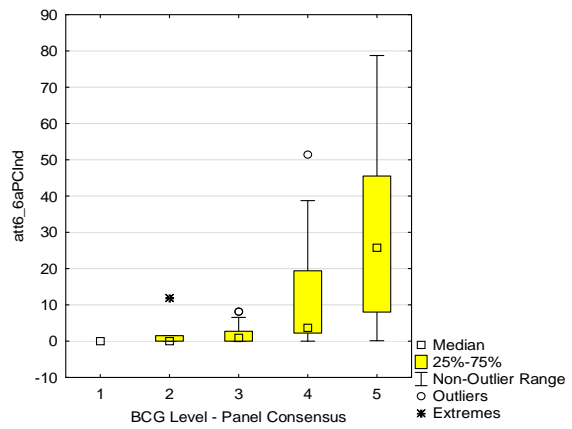
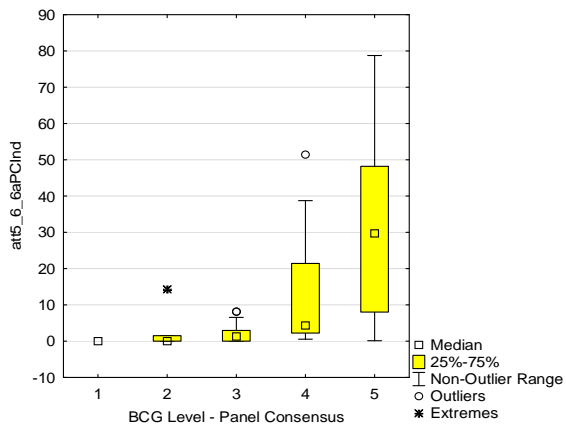
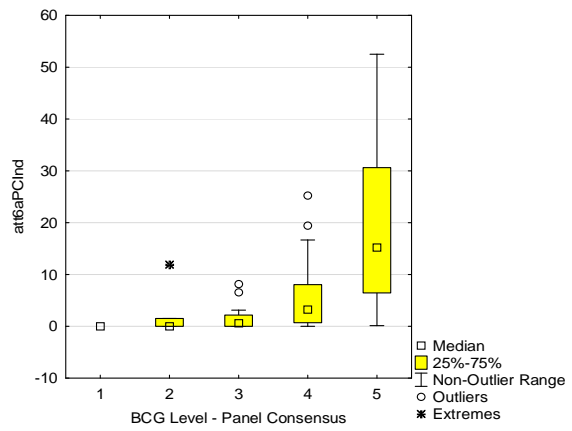
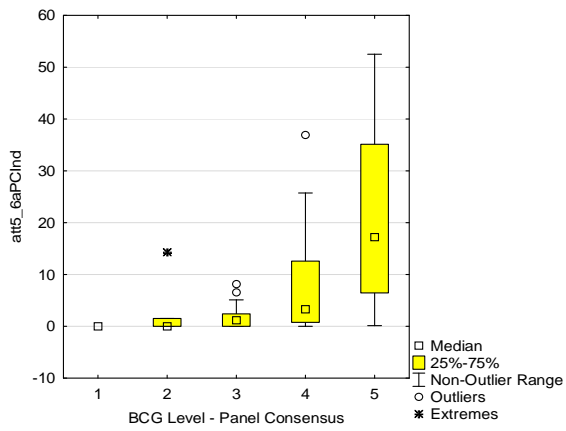
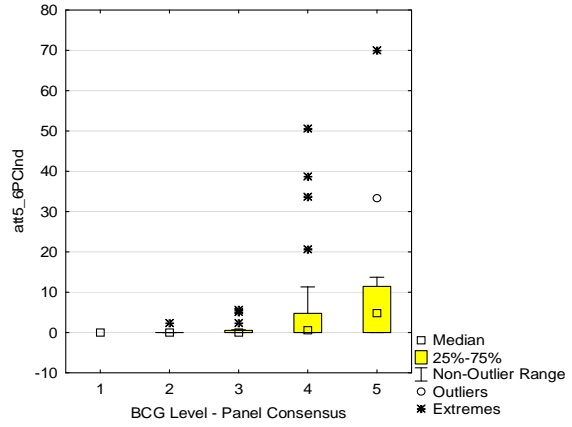
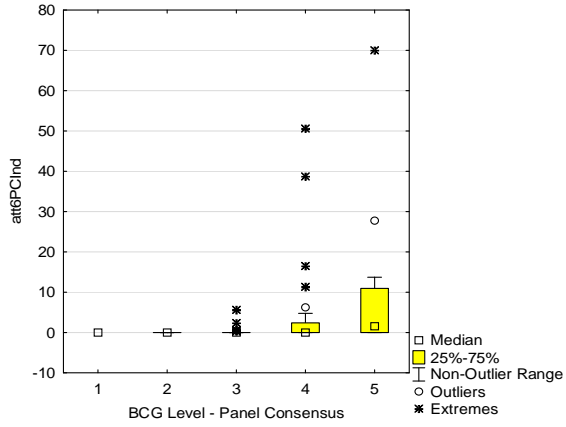
Table H1 continued...

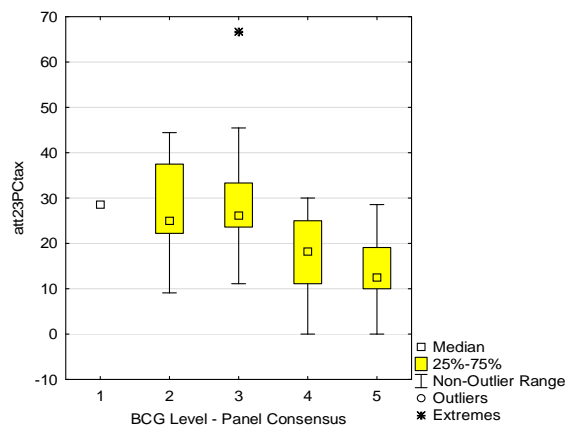
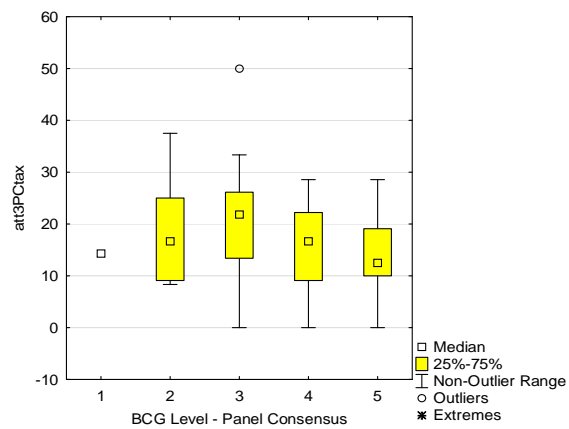
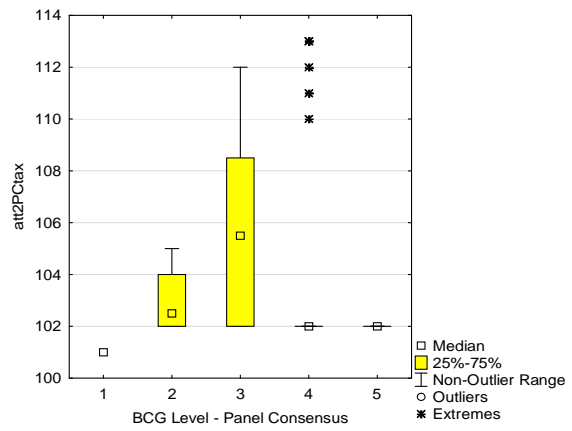
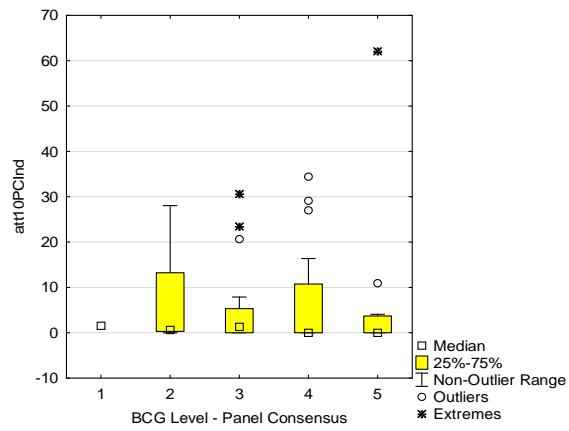
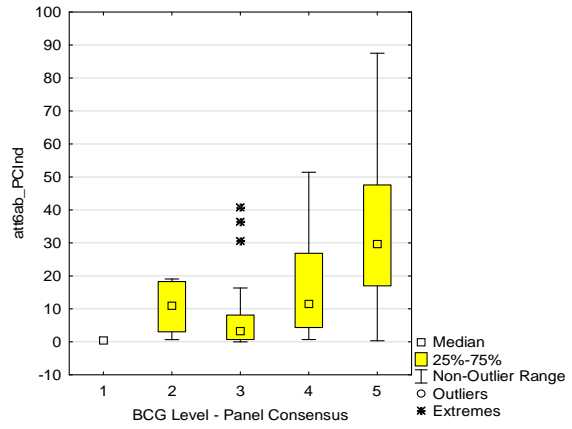
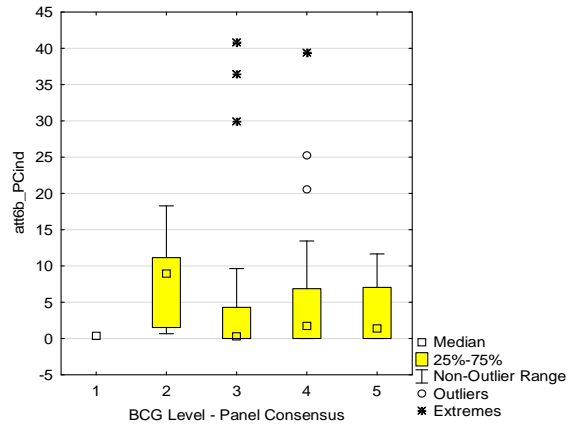
| Metric Code | Description |
|---------------------------|--|
| att5Dom | % Most dominant Attribute V taxon |
| att6Dom | % Most dominant Attribute VI taxon |
| att6aDom | % Most dominant Attribute VIa taxon |
| att566aDom | % Most dominant Attribute V + VI + VIa taxon |
| BrookTroutPCInd | % Wild brook trout individuals |
| BrownTroutPCInd | % Brown trout individuals |
| BKT_TotalSalm_PCInd | (# Wild brook trout individuals/# total salmonid individuals)*100 |
| att6bSalm_TotalSalm_PCInd | (# Attribute 6b salmonid individuals/# total salmonid individuals)*100 |
| Centrarchid_tax | Number of Centrarchidae taxa |
| Centrarchid_PCtax | Percent Centrarchidae taxa |
| Centrarchid_PCInd | Percent Centrarchidae individuals |
| Salmonid_tax | Number of Salmonidae taxa |
| Salmonid_PCtax | Percent Salmonidae taxa |
| Salmonid_PCInd | Percent Salmonidae individuals |
| Cyprin_tax | Number of Cyprinidae taxa |
| Cyprin_PCtax | Percent Cyprinidae taxa |
| Cyprin_PCInd | Percent Cyprinidae individuals |
| BNDCCCMWS_PCInd | % Black nose dace + % creek chub + % cutlips minnow + % white sucker individuals |
| totaltax5Plus | Number of total taxa, counting only taxa with 5 or more individuals |
| Shan_base_2 | Shannon-wiener diversity index (base 2) |
| Evenness | Evenness |
| totalDens_m2 | Total density/meter ² |
| totalDens_100m2 | Total density/100 meter ² |
| totaltaxNoStocked | Total number of taxa, excluding stocked taxa |

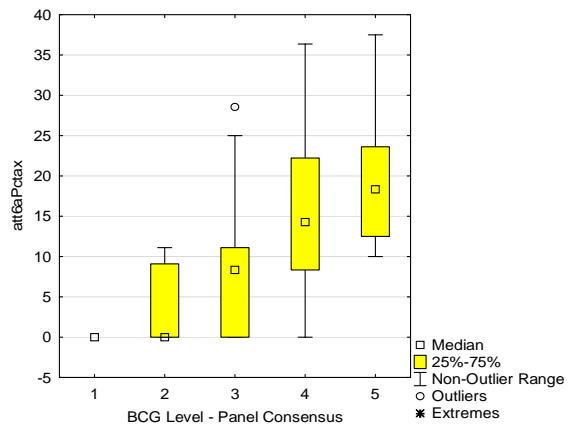
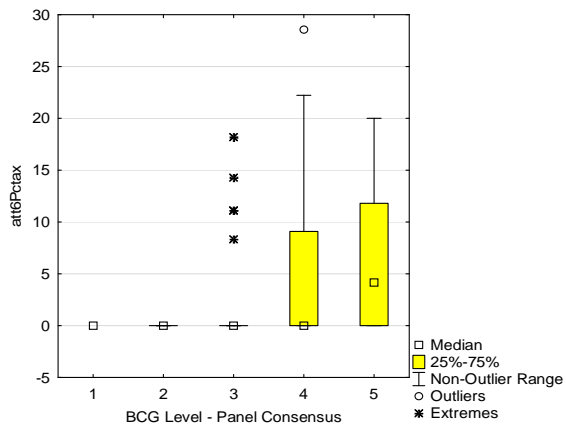
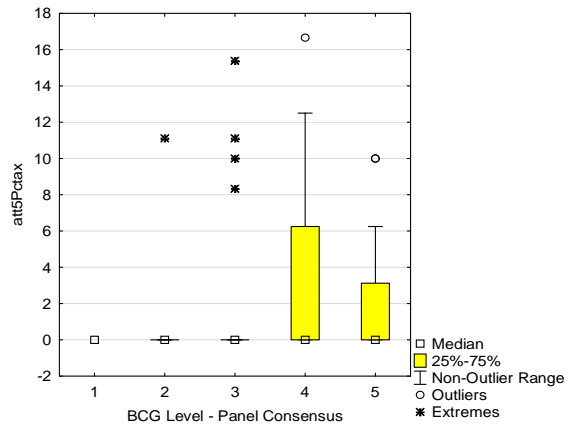
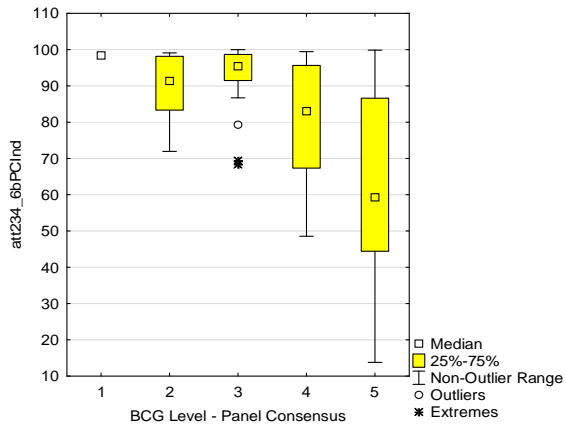
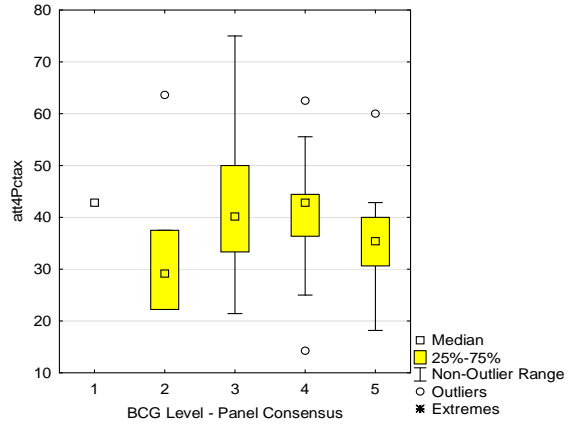
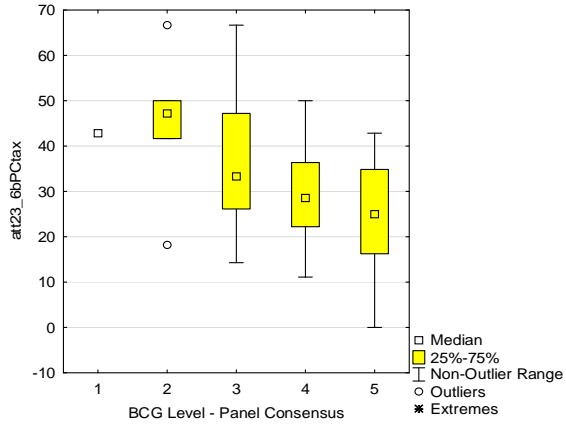


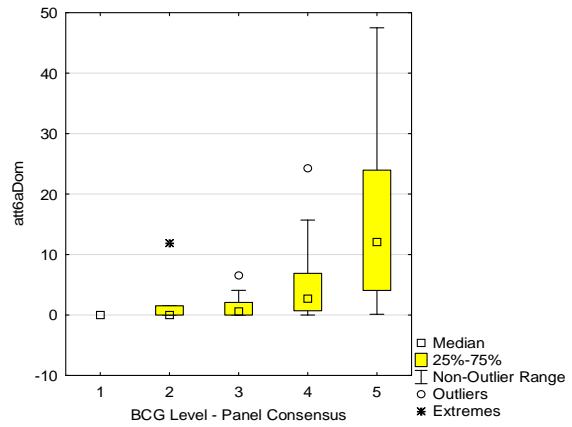
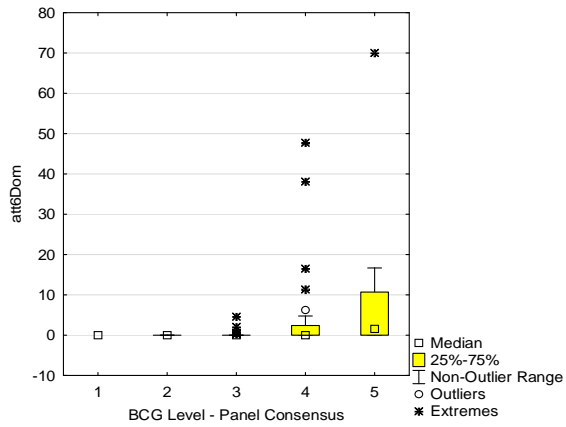
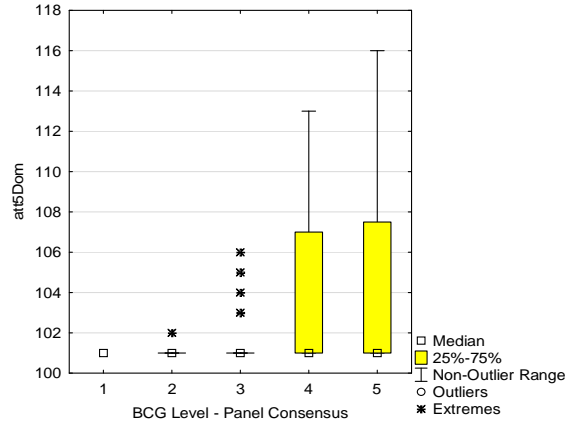
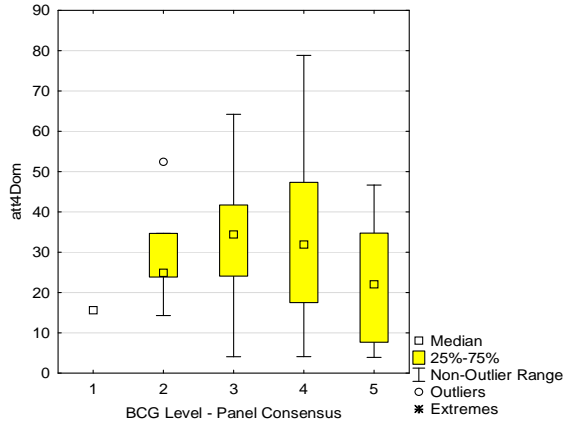
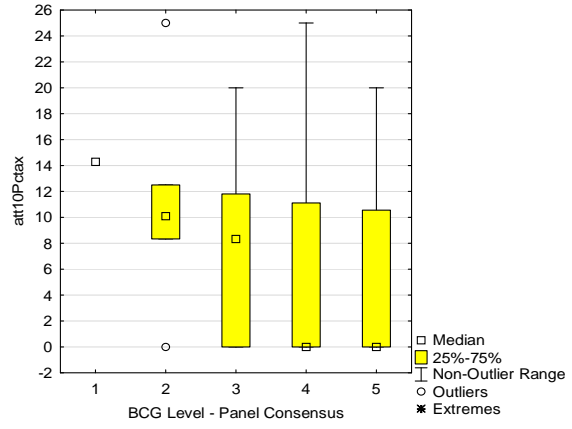
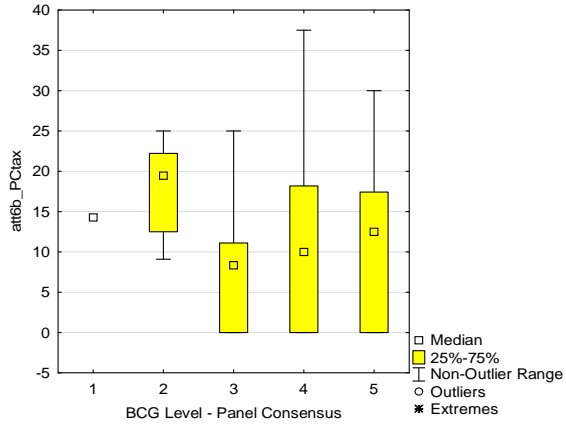


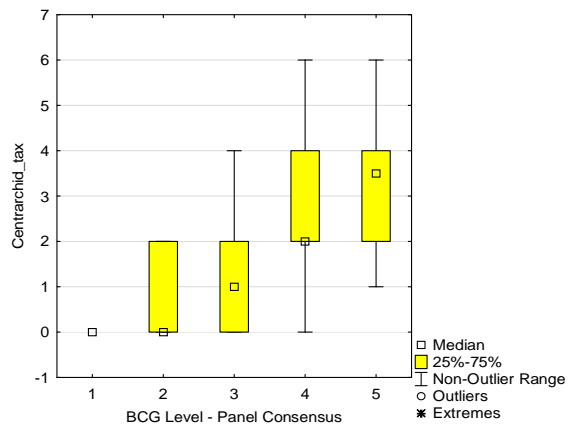
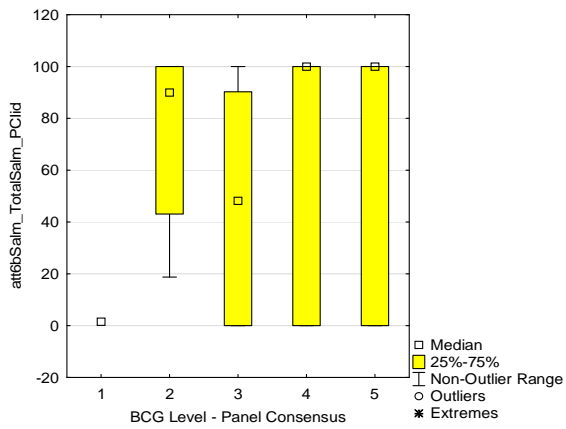
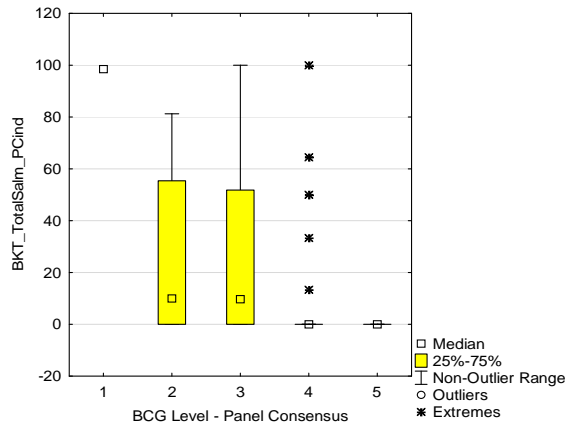
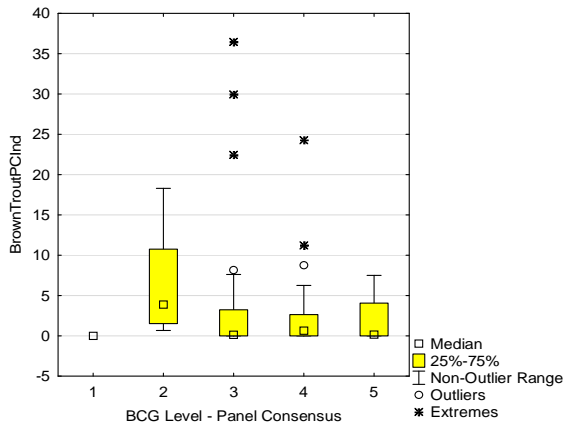
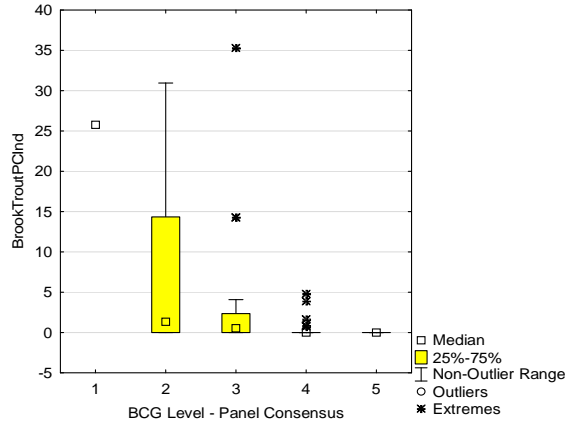
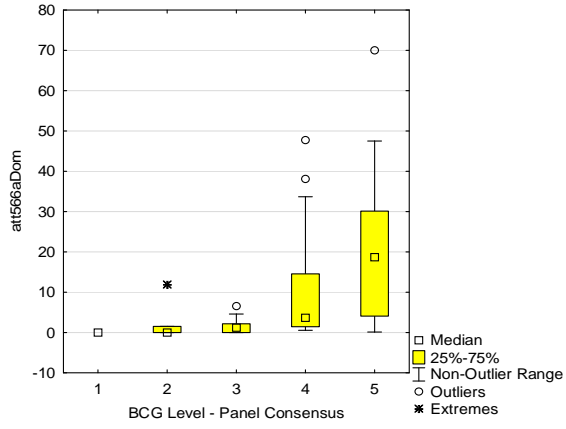


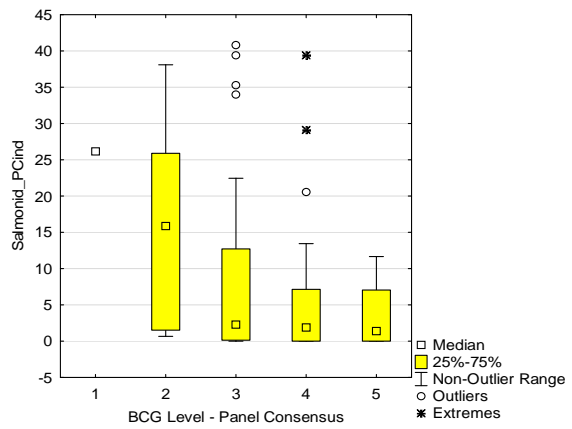
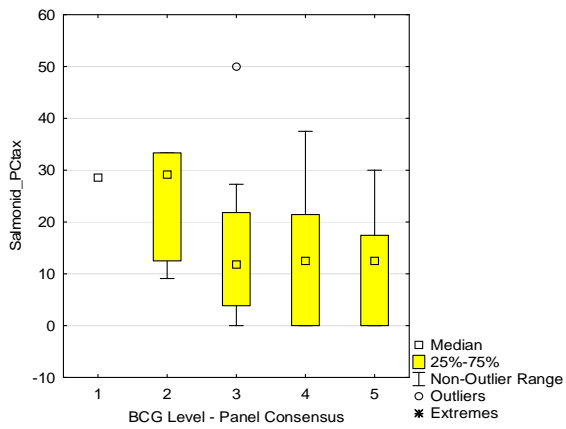
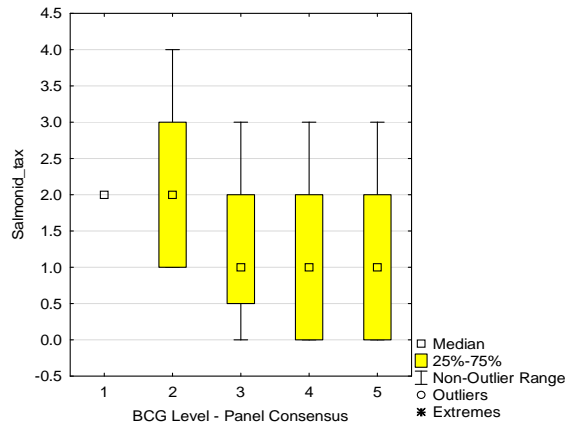
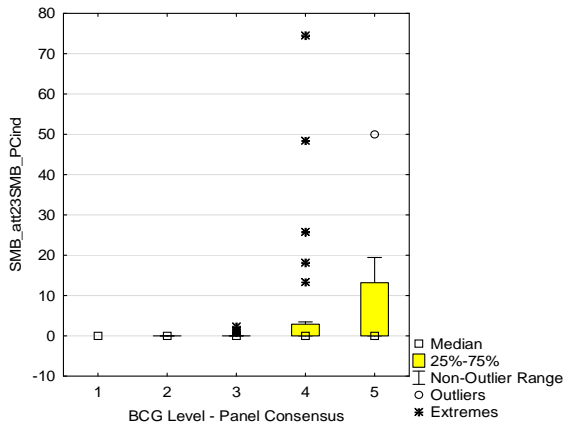
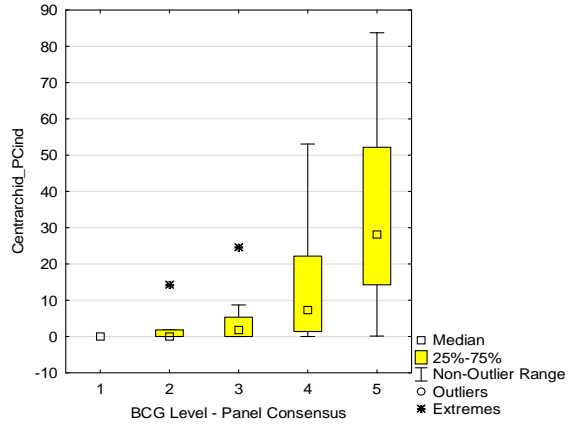
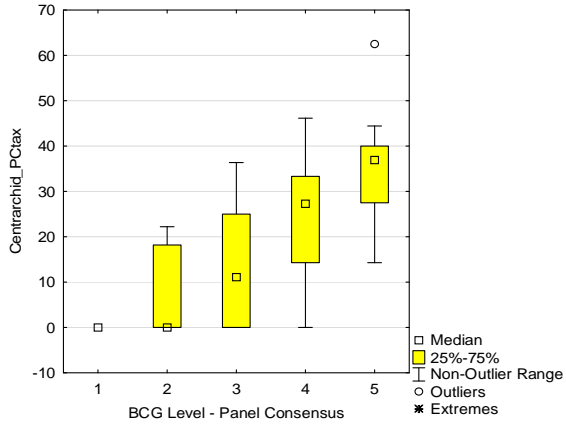


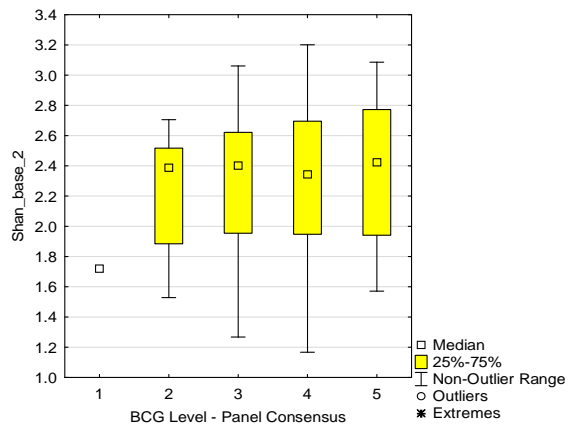
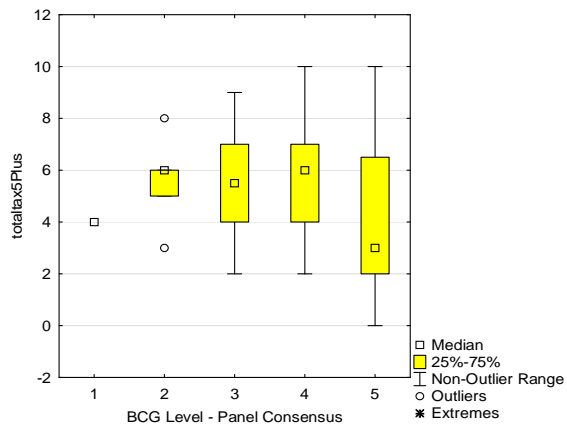
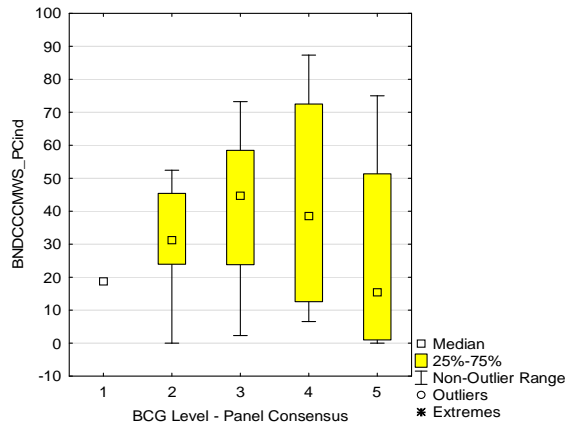
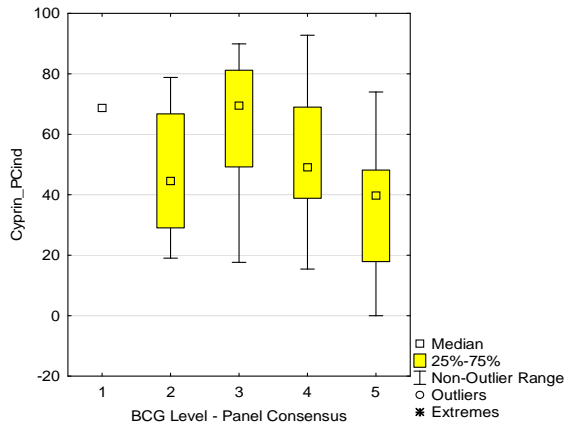
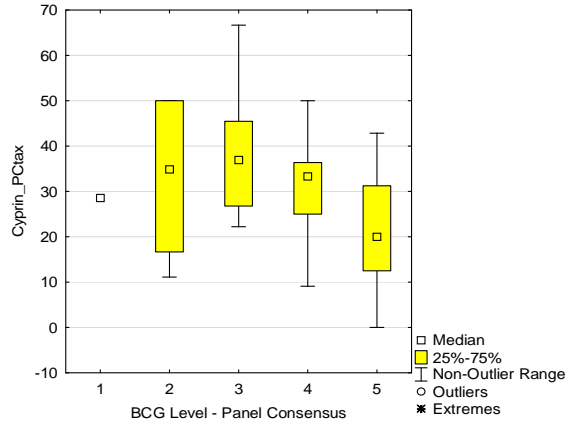
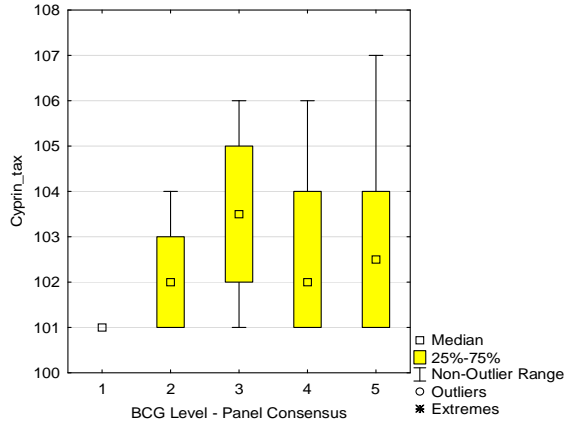


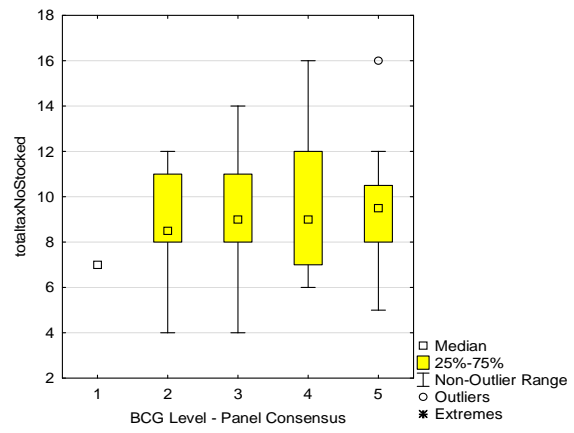
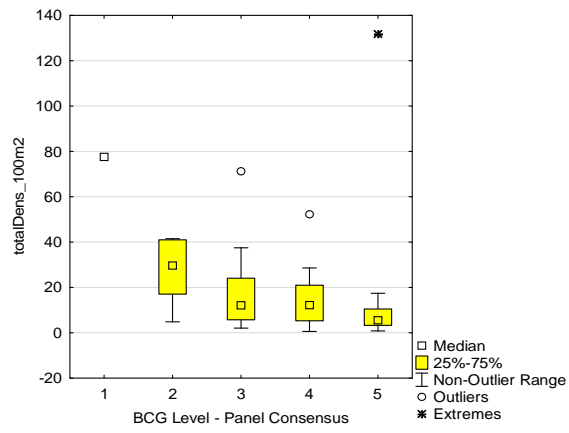
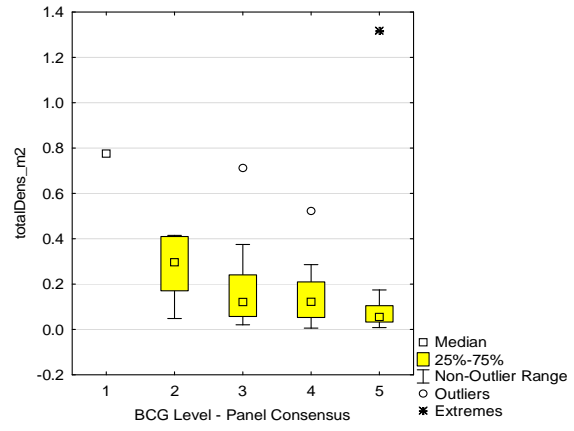
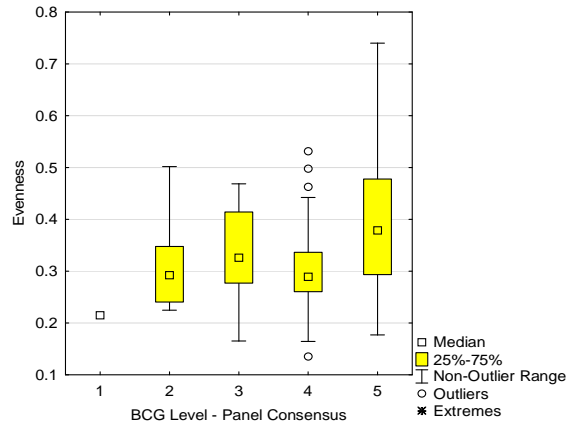












APPENDIX I

Medium-large BCG Level Assignments

Appendix I. Participants made BCG level assignments on 54 medium-large samples for the calibration exercise and 16 samples for the validation exercise. Samples were assessed using the scoring scale shown in Table II.

Table II. Scoring scale that was used for making BCG level assignments.

| | |
|-------|----|
| best | 1 |
| | 1- |
| | 2+ |
| | 2 |
| | 2- |
| | 3+ |
| | 3 |
| | 3- |
| | 4+ |
| | 4 |
| | 4- |
| | 5+ |
| | 5 |
| | 5- |
| | 6+ |
| | 6 |
| worst | 6- |

Table I-2. BCG level assignments and sample information for *medium-large* samples that were assessed during the *calibration* exercise. BCG level assignments are as follows: Final=consensus BCG level (=the assignment made by the majority of participants), without the + or - (2+ and 2- were assigned to level 2, etc.); Best= the best BCG level assignment assigned by a participant (based on the scoring scale in Table G1); Worst=the worst BCG level assignment given by a participant; Samples are highlighted in yellow if the consensus call from the panelists is different from the primary call from the model.

| Fish SampID | Collection Date | Station ID | Waterbody Name | Phase 1 Class | Panelist consensus | | | Primary Model | Notes |
|-------------|-----------------|------------|-------------------|---------------|--------------------|------|-------|---------------|---------------------------------------|
| | | | | | Final | Best | Worst | | |
| 1922 | 6/19/2000 | 272 | Piper Brook | medium_cool | 5 | 4 | 5 | 4 | |
| 3692 | 7/3/2002 | 761 | Latimer Brook | medium_cool | 3 | 2- | 4- | 3 | |
| 4473 | 7/8/2003 | 910 | Hollenbeck River | medium_cool | 4 | 3 | 4 | 4 | |
| 4516 | 7/18/2003 | 918 | Moosup River | large_cool | 3 | 3 | 5 | 4 | |
| 5256 | 6/29/2004 | 1446 | Bantam River | medium_cool | 3 | 2 | 3 | 3 | |
| 5308 | 7/1/2004 | 1449 | Yantic River | large_cool | 4 | 3 | 5 | 4 | |
| 5311 | 7/1/2004 | 1450 | Blackledge River | medium_cold | 4 | 4+ | 4- | 3/4 (tie) | |
| 5433 | 7/16/2004 | 1088 | Natchaug River | large_cool | 4 | 3+ | 5 | 4 | |
| 6520 | 8/26/2004 | 1498 | Roaring Brook | medium_cold | 3 | 3 | 4 | 4 | |
| 6539 | 7/15/2005 | 472 | Moosup River | large_cool | 3 | 2- | 3- | 3 | |
| 8174 | 6/12/2006 | 20 | Branch Brook | medium_cool | 5 | 4- | 6 | 5 | |
| 8510 | 7/10/2006 | 319 | Saugatuck River | medium_cool | 5 | 5 | 5- | 5 | |
| 8788 | 7/28/2006 | 216 | Naugatuck River | large_cool | 3 | 3+ | 4- | 3 | repeat sample (round1=3; round 2=3/4) |
| 9628 | 9/8/2006 | 331 | Steele Brook | medium_cool | 4 | 3+ | 5 | 4 | repeat sample (round1=3; round 2=4) |
| 10501 | 6/14/2007 | 101 | Harbor Brook | medium_cool | 3 | 3+ | 4 | 4 | |
| 10590 | 6/21/2007 | 1281 | Sasco Brook | medium_cool | 2 | 2+ | 3- | 4 | |
| 10714 | 6/27/2007 | 241 | Norwalk River | medium_cool | 4 | 3- | 4- | 4 | repeat sample (both rounds=4) |
| 10717 | 6/28/2007 | 235 | Norwalk River | medium_cool | 4 | 4+ | 4- | 4 | |
| 10802 | 7/5/2007 | 2311 | Hall Meadow Brook | medium_cold | 3 | 3 | 4- | 3 | |
| 10979 | 7/11/2007 | 1806 | Muddy River | medium_cool | 3 | 1 | 3- | 3 | |

Table I-2. continued...

| Fish SampID | Collection Date | Station ID | Waterbody Name | Phase 1 Class | Panelist consensus | | | Primary Model | Notes |
|-------------|-----------------|------------|-----------------------------|---------------|--------------------|------|-------|----------------|---|
| | | | | | Final | Best | Worst | | |
| 12540 | 6/4/2008 | 1088 | Natchaug River | large_cool | 5 | 4 | 5- | 5 | |
| 12726 | 6/16/2008 | 264 | Pequabuck River | medium_cold | 4 | 4+ | 5 | 4 | |
| 12826 | 6/19/2008 | 2642 | Pequabuck River | medium_cold | 4 | 4 | 5 | 4 | |
| 12833 | 6/25/2008 | 2662 | Bass Brook | medium_cool | 3 | 3 | 4 | 4 | |
| 13202 | 7/18/2008 | 1644 | Nepaug River | medium_cold | 3 | 2 | 4 | 3 (close to 2) | repeat sample (both rounds=3); model assignment is close to a 2 |
| 13763 | 9/2/2008 | 127 | Housatonic River | large_cool | 4 | 2- | 4 | 4 (close to 5) | model assignment is close to a 5 |
| 13764 | 9/2/2008 | 914 | Housatonic River | large_cool | 4 | 3 | 4- | 5 | |
| 14242 | 7/21/2008 | 2658 | Coppermine Brook | medium_cold | 2 | 1 | 3 | 2 | |
| 14243 | 8/23/2008 | 1081 | Roaring Brook | medium_cool | 3 | 2 | 4 | 3 | repeat sample (round1=3; round 2=3/4) |
| 14246 | 8/30/2008 | 2641 | Hop River | medium_cold | 4 | 4+ | 4- | 3 (close to 2) | model assignment is close to a 2 |
| 14248 | 8/31/2008 | 354 | Trout Brook | medium_cool | 4 | 3 | 4- | 4 | |
| 14401 | 6/1/2009 | 90 | Furnace Brook | medium_cool | 5 | 4+ | 6 | 5 | repeat sample (round1=5; round 2=4/5) |
| 14618 | 6/25/2009 | 2659 | Pequabuck River | medium_cold | 5 | 5+ | 5- | 4 | |
| 14688 | 6/26/2009 | 1701 | Patagansett River | medium_cool | 5 | 5 | 6+ | 5 | |
| 14717 | 6/29/2009 | 2679 | West Branch Naugatuck River | medium_cold | 4 | 3- | 5+ | 4 | |
| 14719 | 6/29/2009 | 2673 | East Aspetuck River | medium_cold | 4 | 4+ | 5+ | 3/4 (tie) | |
| 14970 | 7/22/2009 | 470 | Fenton River | medium_cold | 4 | 3+ | 5 | 4 | |
| 14986 | 7/23/2009 | 464 | Hop River | large_cool | 5 | 5+ | 5- | 4 | |
| 15044 | 7/27/2009 | 1656 | Little River | medium_cool | 3 | 1- | 3- | 2 | |
| 15057 | 7/28/2009 | 1125 | Beaver Brook | medium_cold | 3 | 2- | 3- | 2/3 (tie) | |
| 15461 | 9/1/2009 | 2685 | Aspetuck River | medium_cool | 3 | 2- | 4 | 3 | |
| 15462 | 9/1/2009 | 2681 | Saugatuck River | medium_cool | 2 | 2 | 3- | 3 | |

Table I-2. continued...

| Fish SampID | Collection Date | Station ID | Waterbody Name | Phase 1 Class | Panelist consensus | | | Primary Model | Notes |
|-------------|-----------------|------------|-----------------------------|---------------|--------------------|------|-------|---------------|--|
| | | | | | Final | Best | Worst | | |
| 15464 | 8/5/2009 | 477 | Mashamoquet Brook | medium_cool | 3 | 2 | 3- | 3 | |
| 15740 | 8/13/2009 | 367 | Willimantic River | large_cool | 4 | 3 | 5 | 4 | |
| 15834 | 8/10/2009 | 189 | Natchaug River | large_cool | 4 | 4+ | 4- | 5 | |
| 18772 | 9/2/2009 | 152 | Little River | medium_cold | 5 | 4 | 5 | 4 | |
| 20928 | 6/30/2010 | 606 | Green Fall River | medium_cool | 1 | 1 | 2- | 2 | repeat sample (round1=2; round 2=1) |
| 21232 | 7/15/2010 | 49 | East Branch Eightmile River | medium_cool | 4 | 3 | 4- | 3 | |
| 21233 | 7/15/2010 | 1236 | Beaver Brook | medium_cool | 3 | 2 | 4 | 2 | |
| 21300 | 7/16/2010 | 475 | Myron Kinney Brook | medium_cool | 2 | 2 | 3 | 2 | |
| 21302 | 7/16/2010 | 1841 | Broad Brook | medium_cool | 2 | 2 | 3 | 2 | |
| 21303 | 7/16/2010 | 650 | Ashaway River | medium_cool | 3 | 3 | 4+ | 2 | |
| 21438 | 7/22/2010 | 6161 | Bartlett Brook | medium_cold | 5 | 4+ | 5 | 4 | |
| 22114 | 8/30/2010 | 289 | Quinnipiac River | large_cool | 4 | 3 | 4- | 4 | repeat sample (both rounds=4) |

Table I-3. Site information for *medium-large* fish samples that were analyzed during the BCG *calibration* exercise. Area refers to the upstream watershed area. Land use (%Devl=% developed, % Imperv= % impervious, % Natl= % natural) is for the upstream catchment area. TNC fields are derived from The Nature Conservancy’s Northeast Aquatic Habitat Classification (Olivero and Anderson 2008). TITAN thermal classes were based on Beauchene et al. 2012. Additional information (i.e. nutrient and habitat data) may available for some of the sites.

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|-----------------------------|----------|---------|------------|--|--------|----------|--------|---|---|-------------------|---------------------|
| 20 | Branch Brook | -73.0810 | 41.6434 | 21.6 | Southern New England Coastal Plains and Hills | 9.8 | 4.2 | 64.6 | Low Gradient: >= 0.02 < 0.1% | Moderately Buffered, Neutral | Transitional Cool | |
| 49 | East Branch Eightmile River | -72.3375 | 41.4309 | 16.4 | Southern New England Coastal Plains and Hills | 8.2 | 3.4 | 77.6 | Low Gradient: >= 0.02 < 0.1% | Moderately Buffered, Neutral | Transitional Cool | 3 |
| 90 | Furnace Brook | -72.2979 | 41.9679 | 15.3 | Lower Worcester Plateau/Eastern Connecticut Upland | 6.6 | 2.9 | 70.5 | Very Low Gradient: <0.02% | Moderately Buffered, Neutral | Transitional Cool | 2 |
| 101 | Harbor Brook | -72.8218 | 41.5314 | 12.1 | Connecticut Valley | 48.2 | 19.1 | 27.5 | Low-Moderate Gradient: >= 0.1 < 0.5% | Moderately Buffered, Neutral | Transitional Cool | 2 |
| 127 | Housatonic River | -73.3815 | 41.8280 | 90.5 | Berkshire Transition | 6.1 | 3.0 | 72.3 | Low-Moderate Gradient: >= 0.1 < 0.5% | Assume Moderately Buffered (Size 3+ rivers) | Transitional Cool | |

Table I-3. continued...

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|-----------------|----------|---------|------------|---|--------|----------|--------|---|------------------------------|-------------------|---------------------|
| 152 | Little River | -72.0582 | 41.6393 | 34.5 | Southern New England Coastal Plains and Hills | 6.8 | 3.1 | 78.7 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | |
| 189 | Natchaug River | -72.1182 | 41.8008 | 73.2 | Southern New England Coastal Plains and Hills | 6.8 | 2.8 | 83.0 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | 3 |
| 216 | Naugatuck River | -73.1145 | 41.7891 | 52.9 | Berkshire Transition | 13.9 | 6.4 | 75.1 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | 3 |
| 235 | Norwalk River | -73.4414 | 41.2675 | 6.1 | Southern New England Coastal Plains and Hills | 22.1 | 6.9 | 69.7 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | 3 |
| 241 | Norwalk River | -73.4341 | 41.2460 | 9.4 | Southern New England Coastal Plains and Hills | 24.0 | 7.8 | 66.3 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | 3 |
| 264 | Pequabuck River | -72.9936 | 41.6693 | 14.0 | Southern New England Coastal Plains and Hills | 15.6 | 7.2 | 69.2 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Cold | |
| 272 | Piper Brook | -72.7274 | 41.7186 | 17.2 | Connecticut Valley | 52.9 | 28.2 | 28.1 | Low Gradient: $\geq 0.02 < 0.1\%$ | Moderately Buffered, Neutral | Transitional Cool | |

Table I-3. continued...

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|-------------------|----------|---------|------------|---|--------|----------|--------|---|------------------------------|-------------------|---------------------|
| 289 | Quinnipiac River | -72.8407 | 41.4501 | 111.0 | Connecticut Valley | 33.6 | 14.7 | 43.2 | Very Low Gradient: <0.02% | Moderately Buffered, Neutral | Transitional Cool | 2 |
| 319 | Saugatuck River | -73.3948 | 41.2945 | 20.4 | Southern New England Coastal Plains and Hills | 10.7 | 4.3 | 79.3 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Transitional Cool | 3 |
| 331 | Steele Brook | -73.0703 | 41.5805 | 17.0 | Southern New England Coastal Plains and Hills | 33.8 | 13.8 | 38.0 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Transitional Cool | 2 |
| 354 | Trout Brook | -72.7231 | 41.7314 | 17.7 | Connecticut Valley | 46.5 | 22.8 | 32.4 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | 3 |
| 367 | Willimantic River | -72.3079 | 41.8326 | 99.6 | Southern New England Coastal Plains and Hills | 9.4 | 3.7 | 75.0 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | |
| 464 | Hop River | -72.2548 | 41.7212 | 79.8 | Southern New England Coastal Plains and Hills | 11.0 | 4.0 | 72.9 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | |
| 470 | Fenton River | -72.2100 | 41.7925 | 28.0 | Southern New England Coastal Plains and Hills | 9.7 | 4.0 | 79.1 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | |

Table I-3. continued...

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|--------------------|----------|---------|------------|---|--------|----------|--------|---|------------------------------|-------------------|---------------------|
| 472 | Moosup River | -71.8931 | 41.7148 | 75.6 | Southern New England Coastal Plains and Hills | 5.4 | 3.7 | 42.8 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Transitional Cool | 3 |
| 475 | Myron Kinney Brook | -71.8619 | 41.5533 | 6.1 | Southern New England Coastal Plains and Hills | 3.9 | 2.4 | 78.7 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Transitional Cool | |
| 477 | Mashamoquet Brook | -71.9359 | 41.8499 | 28.9 | Southern New England Coastal Plains and Hills | 7.0 | 3.2 | 71.9 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | |
| 606 | Green Fall River | -71.8169 | 41.4568 | 10.4 | Long Island Sound Coastal Lowland | 3.4 | 2.3 | 80.8 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Transitional Cool | 1 |
| 650 | Ashaway River | -71.7963 | 41.4433 | 23.2 | | 4.4 | 2.5 | 78.8 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Low Buffered, Acidic | Transitional Cool | |
| 761 | Latimer Brook | -72.2209 | 41.4209 | 10.2 | Southern New England Coastal Plains and Hills | 8.3 | 3.8 | 77.1 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Low Buffered, Acidic | Transitional Cool | |
| 910 | Hollenbeck River | -73.3316 | 41.9581 | 28.1 | Western New England Marble Valleys | 3.8 | 2.2 | 85.8 | Very Low Gradient: $< 0.02\%$ | Moderately Buffered, Neutral | Transitional Cool | |

Table I-3. continued...

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|------------------|----------|---------|------------|---|--------|----------|--------|---|---|-------------------|---------------------|
| 914 | Housatonic River | -73.3906 | 41.8072 | 186.4 | Berkshire Transition | 6.1 | 2.9 | 75.0 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Assume Moderately Buffered (Size 3+ rivers) | Transitional Cool | |
| 918 | Moosup River | -71.8422 | 41.7172 | 67.4 | Southern New England Coastal Plains and Hills | 4.4 | 3.4 | 40.2 | Low Gradient: $\geq 0.02 < 0.1\%$ | Low Buffered, Acidic | Transitional Cool | |
| 1081 | Roaring Brook | -72.8808 | 41.7594 | 7.6 | Connecticut Valley | 23.9 | 8.3 | 63.1 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | 3 |
| 1088 | Natchaug River | -72.1523 | 41.7569 | 88.7 | Southern New England Coastal Plains and Hills | 6.9 | 2.9 | 82.7 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | 2 |
| 1125 | Beaver Brook | -72.1092 | 41.6841 | 7.8 | Southern New England Coastal Plains and Hills | 5.9 | 2.9 | 78.7 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | 1 |
| 1236 | Beaver Brook | -72.3289 | 41.4100 | 8.3 | Southern New England Coastal Plains and Hills | 4.5 | 2.4 | 86.5 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Transitional Cool | 3 |
| 1281 | Sasco Brook | -73.3012 | 41.1457 | 8.4 | Long Island Sound Coastal Lowland | 25.2 | 8.8 | 43.6 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Transitional Cool | 2.5 |

Table I-3. continued...

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|-------------------|----------|---------|------------|--|--------|----------|--------|---|------------------------------|-------------------|---------------------|
| 1446 | Bantam River | -73.1823 | 41.7417 | 20.7 | Southern New England Coastal Plains and Hills | 7.8 | 3.5 | 69.3 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Low Buffered, Acidic | Transitional Cool | |
| 1449 | Yantic River | -72.1759 | 41.5702 | 53.8 | Southern New England Coastal Plains and Hills | 8.4 | 3.4 | 70.0 | Low Gradient: $\geq 0.02 < 0.1\%$ | Moderately Buffered, Neutral | Transitional Cool | 2 |
| 1450 | Blackledge River | -72.4261 | 41.6069 | 21.9 | Southern New England Coastal Plains and Hills | 13.7 | 4.7 | 70.8 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | 2 |
| 1498 | Roaring Brook | -72.2656 | 41.9152 | 18.4 | Lower Worcester Plateau/Eastern Connecticut Upland | 7.4 | 2.9 | 85.0 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | |
| 1644 | Nepaug River | -73.0271 | 41.8382 | 9.9 | Berkshire Transition | 9.2 | 3.5 | 77.4 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | |
| 1656 | Little River | -72.0473 | 41.7561 | 16.2 | Southern New England Coastal Plains and Hills | 7.4 | 3.2 | 76.3 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | 1 |
| 1701 | Patagansett River | -72.2124 | 41.3557 | 6.2 | Long Island Sound Coastal Lowland | 18.3 | 1.4 | 69.6 | Low Gradient: $\geq 0.02 < 0.1\%$ | Low Buffered, Acidic | Transitional Cool | |

Table I-3. continued...

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|-------------------|----------|---------|------------|---|--------|----------|--------|---|------------------------------|-------------------|---------------------|
| 1806 | Muddy River | -72.8012 | 41.4151 | 12.3 | Connecticut Valley | 15.8 | 5.9 | 48.8 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Transitional Cool | 3 |
| 1841 | Broad Brook | -71.9703 | 41.5538 | 11.8 | Southern New England Coastal Plains and Hills | 4.8 | 2.7 | 80.8 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | |
| 2311 | Hall Meadow Brook | -73.1689 | 41.8861 | 12.0 | Lower Berkshire Hills | 4.3 | 2.2 | 89.0 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | 2 |
| 2641 | Hop River | -72.4089 | 41.7712 | 8.2 | Southern New England Coastal Plains and Hills | 14.8 | 4.8 | 68.2 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | |
| 2642 | Pequabuck River | -72.9272 | 41.6698 | 24.9 | Southern New England Coastal Plains and Hills | 27.8 | 12.1 | 53.1 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Cold | |
| 2658 | Coppermine Brook | -72.9261 | 41.7137 | 8.5 | Southern New England Coastal Plains and Hills | 15.0 | 6.5 | 75.6 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | |
| 2659 | Pequabuck River | -72.9082 | 41.6740 | 26.1 | Connecticut Valley | 29.2 | 12.8 | 52.0 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Cold | |

Table I-3. continued...

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|-----------------------------|----------|---------|------------|---|--------|----------|--------|---|------------------------------|-------------------|---------------------|
| 2662 | Bass Brook | -72.7587 | 41.6931 | 8.5 | Connecticut Valley | 40.6 | 26.4 | 41.0 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Transitional Cool | |
| 2673 | East Aspetuck River | -73.3785 | 41.6546 | 19.0 | Southern New England Coastal Plains and Hills | 8.6 | 3.4 | 74.4 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | |
| 2679 | West Branch Naugatuck River | -73.1602 | 41.8561 | 19.1 | Berkshire Transition | 3.7 | 2.2 | 90.8 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | |
| 2681 | Saugatuck River | -73.4229 | 41.3220 | 13.1 | Southern New England Coastal Plains and Hills | 10.8 | 4.4 | 81.8 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Low Buffered, Acidic | Transitional Cool | |
| 2685 | Aspetuck River | -73.3303 | 41.2933 | 7.8 | Southern New England Coastal Plains and Hills | 8.1 | 3.4 | 81.6 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Transitional Cool | |
| 6161 | Bartlett Brook | -72.2563 | 41.5883 | 13.4 | Southern New England Coastal Plains and Hills | 6.7 | 3.0 | 73.3 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | |

Table I-4. BCG level assignments and sample information for *medium-large* samples that were assessed during the *validation* exercise. BCG level assignments are as follows: Final=consensus BCG level (=the assignment made by the majority of participants), without the + or - (2+ and 2- were assigned to level 2, etc.); Best= the best BCG level assignment assigned by a participant (based on the scoring scale in Table G1); Worst=the worst BCG level assignment given by a participant; Samples are highlighted in yellow if the consensus call from the panelists is different from the primary call from the model.

| Fish SampID | Collection Date | Station ID | Waterbody Name | Phase 1 Class | Panelist consensus | | | Primary Model | Notes |
|-------------|-----------------|------------|--------------------------|---------------|--------------------|------|-------|---------------|-----------------------------------|
| | | | | | Final | Best | Worst | | |
| 327 | 7/19/1999 | 163 | Mattabesset River | medium_cool | 4 | 3- | 4- | 4 | |
| 4228 | 7/26/1999 | 246 | Norwalk River | large_cool | 4 | 4 | 5+ | 4 | |
| 4537 | 7/29/2003 | 916 | Hockanum River | medium_cool | 5 | 4 | 5- | 5 | |
| 5257 | 6/30/2004 | 359 | West Branch Salmon Brook | medium_cold | 2 | 1 | 2- | 2 | |
| 5434 | 7/16/2004 | 189 | Natchaug River | large_cool | 4 | 3 | 5 | 4 | |
| 5548 | 7/27/2004 | 310 | Salmon Brook | large_cool | 3 | 2+ | 3- | 3 | |
| 6606 | 7/25/2005 | 478 | Blackwell Brook | medium_cool | 4 | 3- | 4- | 4 | |
| 7263 | 7/29/2005 | 1671 | Mount Hope River | medium_cool | 5 | 3+ | 5- | 3 | has direct agricultural influence |
| 8782 | 7/24/2006 | 122 | Hollenbeck River | medium_cold | 3 | 2+ | 3- | 3 | |
| 12991 | 7/2/2008 | 325 | Shepaug River | large_cool | 4 | 3 | 5- | 4 | |
| 14244 | 8/23/2008 | 1513 | Cherry Brook | medium_cold | 3 | 1- | 3- | 4 | |
| 14723 | 7/1/2009 | 458 | Willimantic River | large_cool | 3 | 3+ | 5+ | 3 | |
| 15058 | 7/28/2009 | 480 | Merrick Brook | medium_cold | 3 | 2 | 3- | 2/3 (tie) | |
| 15734 | 8/4/2009 | 1482 | Pease Brook | medium_cool | 4 | 3+ | 4+ | 3 | panel call was very close to a 3 |
| 20873 | 6/29/2010 | 278 | Pomperaug River | large_cool | 3 | 2- | 3 | 2 | |
| 21183 | 7/1/2010 | 621 | Yantic River | medium_cool | 4 | 3 | 4 | 4 | |

Table I-5. Site information for *medium-large* fish samples that were analyzed during the BCG *validation* exercise. Area refers to the upstream watershed area. Land use (%Devl=% developed, % Imperv= % impervious, % Natl= % natural) is for the upstream catchment area. TNC fields are derived from The Nature Conservancy's Northeast Aquatic Habitat Classification (Olivero and Anderson 2008). TITAN thermal classes were based on Beauchene et al. 2012. Additional information (i.e. nutrient and habitat data) may available for some of the sites.

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|------------------|----------|---------|------------|---|--------|----------|--------|---|------------------------------|-------------------|---------------------|
| 122 | Hollenbeck River | -73.3058 | 41.9431 | 17.6 | Berkshire Transition | 4.8 | 2.5 | 81.6 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | 1 |
| 163 | Mattabeset River | -72.7127 | 41.6189 | 45.8 | Connecticut Valley | 31.4 | 13.9 | 44.3 | Very Low Gradient: $< 0.02\%$ | Moderately Buffered, Neutral | Transitional Cool | 3 |
| 189 | Natchaug River | -72.1182 | 41.8008 | 73.2 | Southern New England Coastal Plains and Hills | 6.8 | 2.8 | 83.0 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | 3 |
| 246 | Norwalk River | -73.4295 | 41.1267 | 56.5 | Long Island Sound Coastal Lowland | 27.6 | 12.0 | 52.7 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Transitional Cool | |
| 278 | Pomperaug River | -73.2165 | 41.5491 | 56.8 | Connecticut Valley | 9.1 | 3.8 | 64.9 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | |
| 310 | Salmon Brook | -72.7749 | 41.9366 | 65.2 | Connecticut Valley | 8.1 | 3.7 | 69.7 | Very Low Gradient: $< 0.02\%$ | Moderately Buffered, Neutral | Transitional Cool | 1 |

Table I-5. continued...

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|--------------------------|----------|---------|------------|--|--------|----------|--------|---|------------------------------|-------------------|---------------------|
| 325 | Shepaug River | -73.3308 | 41.5489 | 131.4 | Southern New England Coastal Plains and Hills | 7.9 | 3.4 | 72.6 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | 3 |
| 359 | West Branch Salmon Brook | -72.8215 | 41.9372 | 23.8 | Connecticut Valley | 7.1 | 3.0 | 84.6 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | 1 |
| 458 | Willimantic River | -72.3058 | 41.9423 | 53.8 | Lower Worcester Plateau/Eastern Connecticut Upland | 9.0 | 3.7 | 71.5 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Low Buffered, Acidic | Transitional Cool | 3 |
| 478 | Blackwell Brook | -71.9488 | 41.7407 | 22.7 | Southern New England Coastal Plains and Hills | 7.8 | 3.4 | 73.7 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | |
| 480 | Merrick Brook | -72.1101 | 41.6610 | 13.0 | Southern New England Coastal Plains and Hills | 6.5 | 3.1 | 72.4 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | 1 |
| 621 | Yantic River | -72.1918 | 41.5766 | 39.2 | Southern New England Coastal Plains and Hills | 8.6 | 3.4 | 74.4 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Transitional Cool | |
| 916 | Hockanum River | -72.5204 | 41.8078 | 49.1 | Connecticut Valley | 25.0 | 10.4 | 49.9 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Low Buffered, Acidic | Transitional Cool | 2.5 |

Table I-5. continued...

| Station ID | Waterbody Name | Long | Lat | Area (mi2) | Level 4 Ecoregion | % Devl | % Imperv | % Natl | TNC Gradient | TNC Geology | TNC Thermal Class | TITAN thermal class |
|------------|------------------|----------|---------|------------|--|--------|----------|--------|---|------------------------------|-------------------|---------------------|
| 1482 | Pease Brook | -72.1923 | 41.5947 | 11.7 | Southern New England Coastal Plains and Hills | 7.0 | 3.5 | 55.0 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | 1 |
| 1513 | Cherry Brook | -72.9295 | 41.8365 | 13.8 | Berkshire Transition | 7.5 | 3.1 | 82.9 | Moderate-High Gradient: $\geq 0.5 < 2\%$ | Moderately Buffered, Neutral | Cold | 1 |
| 1671 | Mount Hope River | -72.1603 | 41.8772 | 12.4 | Lower Worcester Plateau/Eastern Connecticut Upland | 8.0 | 3.1 | 83.2 | Low-Moderate Gradient: $\geq 0.1 < 0.5\%$ | Moderately Buffered, Neutral | Transitional Cool | 2 |