

National Biological Assessment  
and Criteria Workshop

Advancing State and Tribal Programs



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# Index 201

## *Selection of Metrics for Index Assembly*

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# Multimetric Index Development

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- Database consisting of reference and stressed populations (sites)
- Classify resource (reference sites, ecoregions)
- Identify and test candidate metrics
- **Select metrics for dimensionless index**
- Select thresholds for assessment (set biocriteria)

# Assembling an Index

1. Identify suites of metrics that meet the following criteria:
  - Ecologically justifiable
  - Responsive
  - Precise
  - Provide unique information
  - Represent a range of metric categories (richness, composition, tolerance, trophic, habit, voltinism)

# Assembling an Index

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2. Set aside a portion of the data for testing / validating the index.
3. Score all potential metrics.
4. Calculate index alternatives by summing or averaging metric scores.

# Assembling an Index

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5. Calculate the DE and precision of each index.
6. Evaluate the alternatives.
7. Test the favored alternatives using the reserved data.

# Identifying Redundancy

- Correlation analysis of potential metrics
- Identify pairs as redundant if the correlation coefficient,  $r$ , is  $> 0.9$
- For  $r > 0.8$ , examine scatterplots
- Avoid suites of metrics containing both metrics in a redundant pair

# Example Correlations

	<i>Total taxa</i>	<i>EPT taxa</i>	<i>Ephemerop. taxa</i>	<i>Plecoptera taxa</i>	<i>Trichoptera taxa</i>	<i>% EPT</i>	<i>% Plecoptera</i>	<i>% Trichoptera</i>	<i>HBI</i>	<i>Intolerant taxa (0-1)</i>
EPT taxa	<b>0.92</b>									
Ephemerop. taxa	<b>0.8</b>	<b>0.87</b>								
Plecoptera taxa	0.77	<b>0.86</b>	0.63							
Trichoptera taxa	<b>0.82</b>	<b>0.86</b>	0.6	0.63						
% EPT	0.26	0.43	0.4	0.4	0.31					
% Plecoptera	0.26	0.38	0.23	0.53	0.24	0.45				
% Trichoptera	0.2	0.23	0.09	0.13	0.37	0.31	0.01			
HBI	-0.38	-0.56	-0.48	-0.54	-0.44	-0.77	-0.53	-0.37		
Intolerant (0-1)	<b>0.84</b>	<b>0.94</b>	<b>0.81</b>	<b>0.83</b>	<b>0.82</b>	0.41	0.37	0.2	-0.59	
Intolerant (0-3)	<b>0.89</b>	<b>0.97</b>	<b>0.82</b>	<b>0.88</b>	<b>0.83</b>	0.41	0.4	0.2	-0.59	<b>0.97</b>

# Attribute groups

INDIVIDUAL CONDITION	TAXONOMIC COMPOSITION	COMMUNITY STRUCTURE	LIFE HISTORY ATTRIBUTES	SYSTEM PROCESSES
DISEASE				TROPHIC DYNAMICS
ANOMALIES	IDENTITY	TAXA RICHNESS	FEEDING GROUPS	PRODUCTIVITY
CONTAMINANT LEVELS	TOLERANCE	RELATIVE ABUNDANCE	HABIT	MATERIAL CYCLES
DEATH	RARE OR ENDANGERED KEY TAXA	DOMINANCE	VOLUNTINISM	PREDATION
METABOLIC RATE				RECRUITMENT

## INTEGRATED BIOASSESSMENT

TOXICITY TESTS

RIVPACS

INVERTEBRATE IBI

FISH IBI





# Select Metric Suites-Examples

Metric	A	B	C	D
Total taxa	X			
EPT taxa		X		
Ephemeroptera taxa			X	X
Plecoptera taxa			X	X
Trichoptera taxa			X	X
% EPT		X	X	
% Plecoptera	X			X
% Trichoptera	X			X
HBI	X	X	X	X
% 3 Dominant			X	
% 5 Dominant	X	X		
Shannon-Weiner				X
Scraper taxa	X	X	X	X
% Scrapers	X	X	X	X
Clinger taxa			X	

# Index Assembly Hands-on Exercise – Selecting metrics

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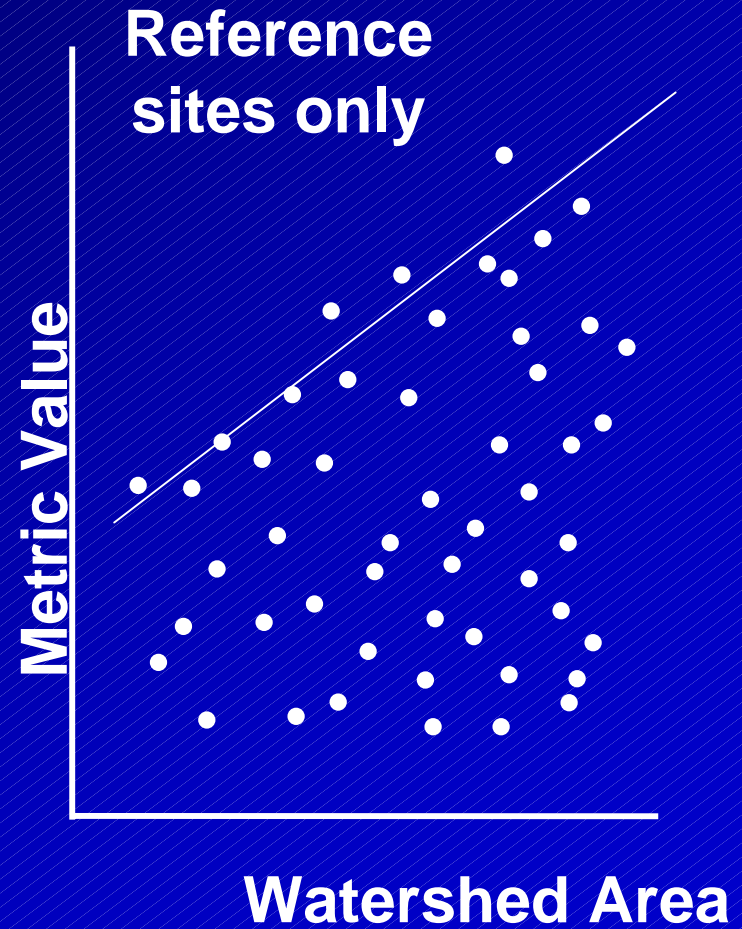
- Discrimination efficiencies (DE) and metric correlations for Idaho data
- **Goal:** Use metric evaluation information to select suites of candidate metrics and test as index alternatives

# Scoring Metrics

- Creates dimensionless values that can be summed or averaged into a single index value.
- Standardizes metric values with respect to some expectation.
- Can be continuous or discrete.
- Can use expectations based on reference distributions or the entire distribution of values in a region.

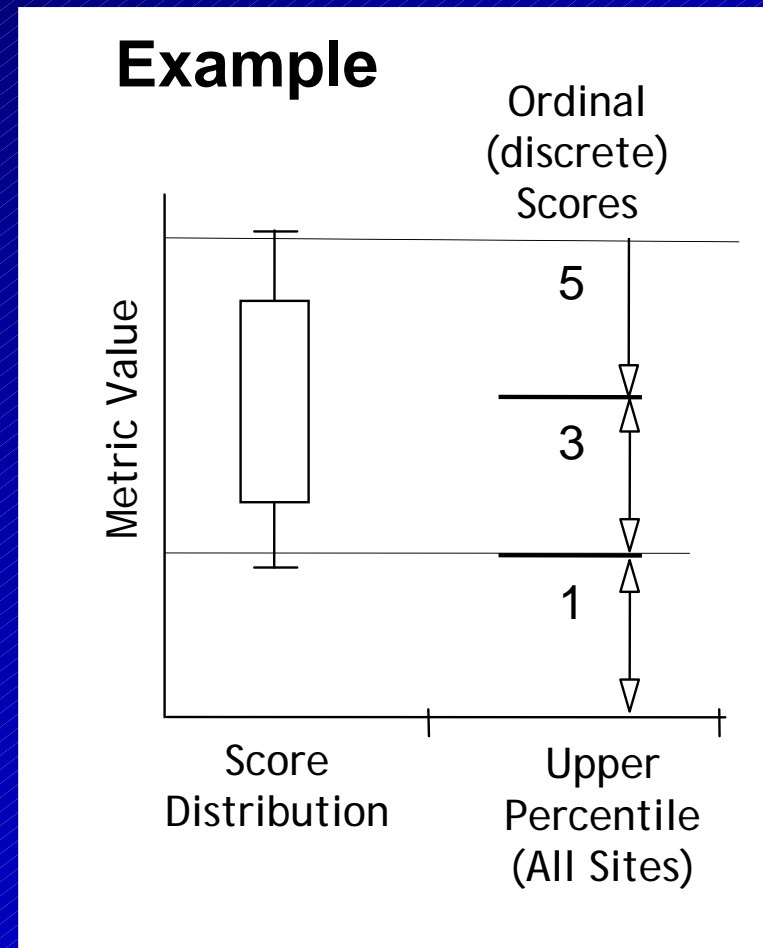
# Scoring metrics – adjustment for natural variation

- Some metrics vary naturally with physical features, such as watershed area, elevation, gradient, and stream order.
- Adjustment for natural factors is necessary to set appropriate expectations for scoring these metrics.
- This type of adjustment can be done by eye or using quantile regression techniques.



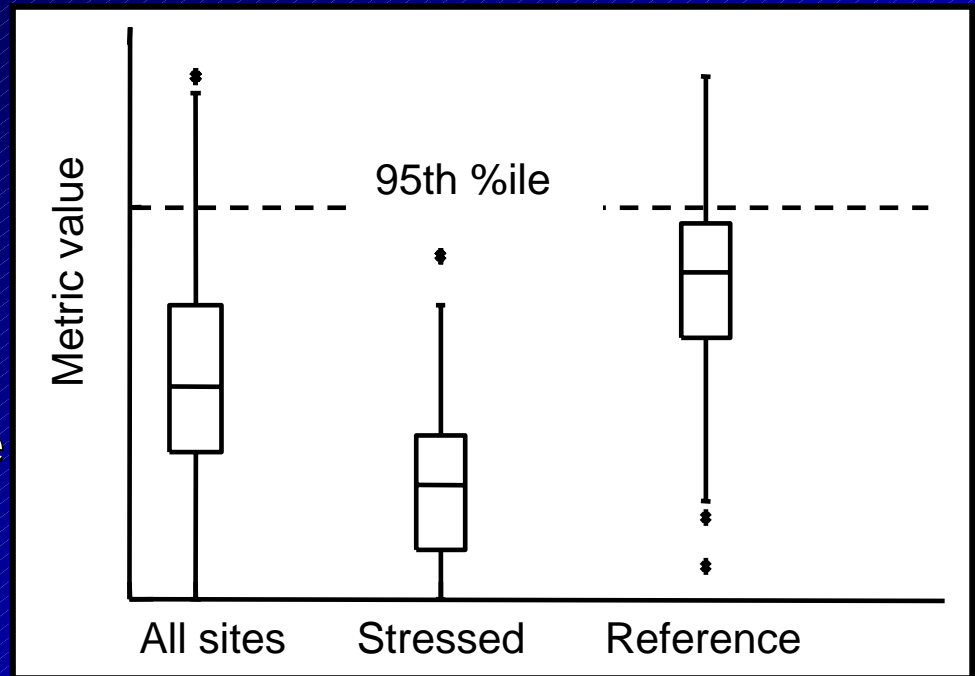
# Scoring Metrics – Discrete

- Metric values receive a discrete score (e.g., 1, 3, or 5) based on comparison to some expectation.
- “Reverse” metrics are scaled so that higher values receive lower scores.
- Metric scores are summed, sometimes rescaled to a sum of 100.



# Scoring Metrics – Continuous

- Scores are scaled to the 95th percentile of all values within each region.
- Scoring is on a continuous scale from 100 to 0.
- “Reverse” metrics are scaled to the 5th percentile.
- Metric scores are averaged (or summed) to obtain index value.



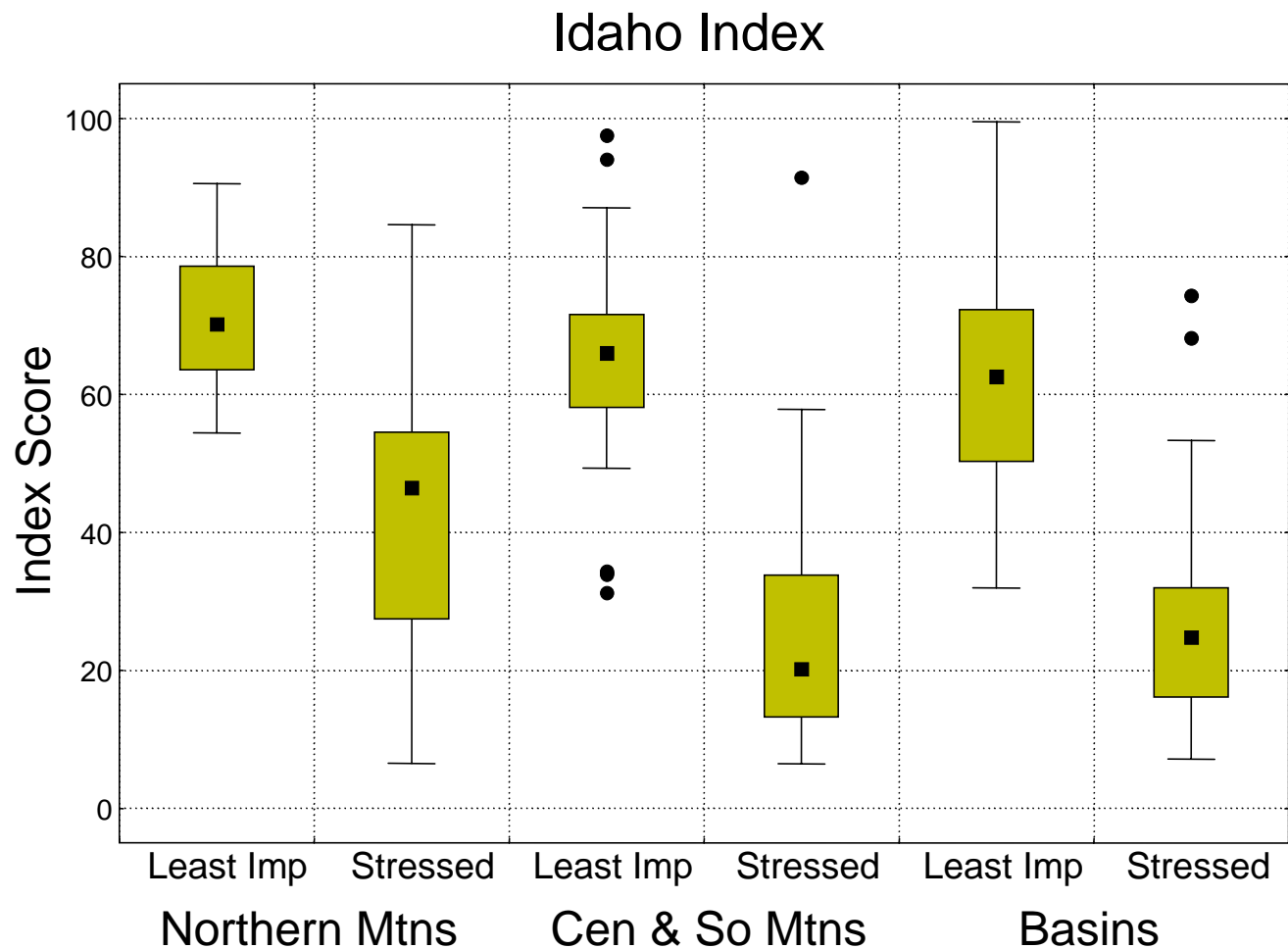
## Formulas:

- $\text{Score} = 100 * (\text{Max} - \text{Value}) / (\text{Max} - 5\text{th}\%)$  (“reverse” metrics)
- $\text{Score} = 100 * \text{Value} / 95\text{th}\%$

# Evaluating Index Alternatives

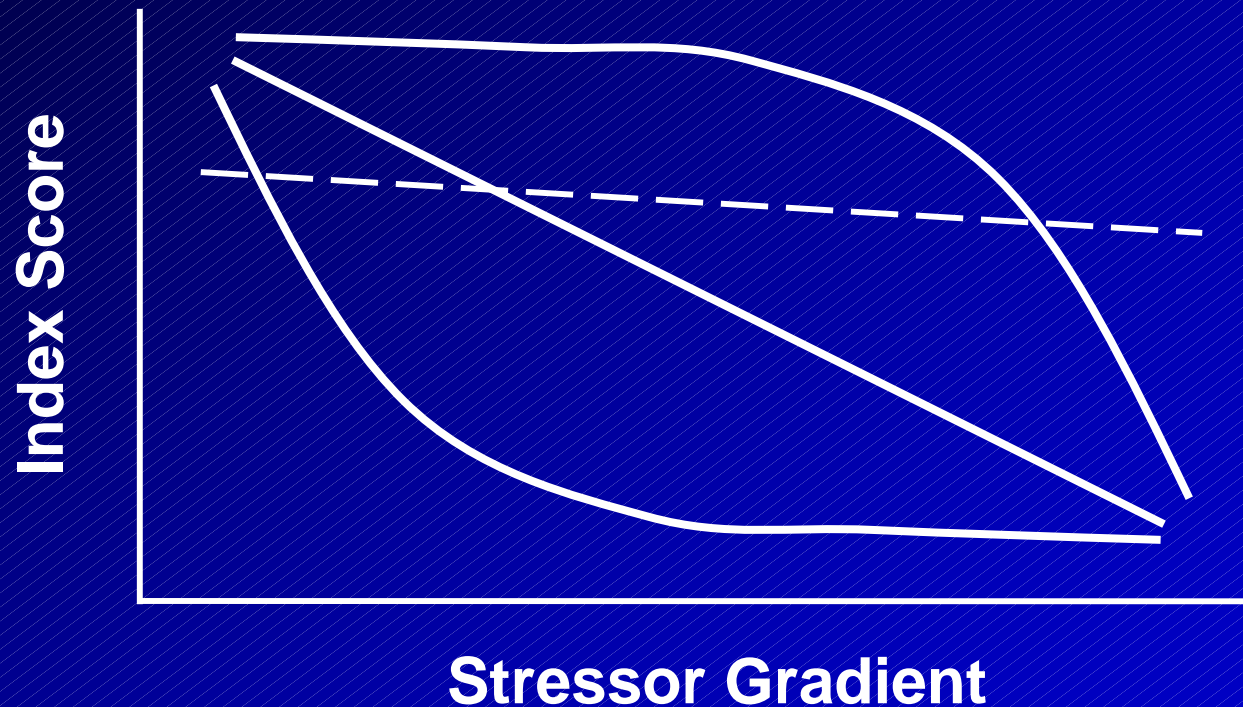
<b>Metric</b>	<b>Model 1</b>	<b>Model 2</b>	<b>Model 3</b>
Total taxa	X	X	X
Ephemeroptera taxa	X	X	X
Plecoptera taxa	X	X	X
Trichoptera taxa	X	X	X
% EPT		X	
% Plecoptera	X		X
% Clingers	X	X	
Clinger taxa			X
Scraper taxa			X
HBI			X
% 5 Dominant taxa			X
Basins DE (25th)	93.1 (47.6)	96.6 (57.1)	96.6 (50.5)
N. Mtns DE (25th)	83.8 (58.1)	89.2 (67.9)	89.2 (65.2)
C&S Mtns DE (25th)	85.7 (55.0)	90.5 (57.3)	90.5 (57.7)
DE (Wtd avg)	88.8	93.1	93.1

# Index Discrimination





# Index Responsiveness to Stressor Gradient



# Index Precision

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1. Find replicated samples.
2. Run ANOVA with Station as the grouping variable.
3. Use the MSE term as an estimate of variance.
4. Take the root of the MSE as an estimate of standard deviation.
5. Calculate CV or CI

# Testing / Evaluating the Index

- Calculate the selected index using the reserved (validation) data.
- Check the “validation” samples against the “calibration” reference 25th percentile.
  - Are approximately 75% of validation reference samples above the threshold?
  - Is the percentage of validation stressed samples below the threshold comparable?

# Summary

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- Select suites of metrics that meet criteria.
- Score metrics and average or sum scores into index value.
- Calculate DE and precision of alternatives.
- Select appropriate index.
- Test the DE of index with validation data.

Reference and Degraded Site Designations (McIntyre)

Environmental data

Taxonomic Data

*A priori* and *a posteriori*  
site classification  
(Gerritsen)

Metric Data  
(Frydenborg)

Metric Exploration  
(Frydenborg)

Select Responsive Metrics  
(Blocksom)

Develop Final Multimetric  
(Blocksom)

Multimetric

*A posteriori*  
site classification

Community  
Cluster Groups

Group Probabilities  
Taxa Frequencies

Expected Taxa

Observed/Expected

RIVPACS

