

Scoring Biological Integrity

the California Stream Condition Index (CSCI)



- **Scoring Tool Enhancements**
 - Update to O/E component
 - Integrating predictive MMI techniques
 - Our recommendations
- **Setting Thresholds**
- **Statewide and Regional Extent Estimates**



Technical Team



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***** *Scientific Review Panel***

How do we convert a list of species
into a condition score?



Qualities of a good scoring tool

Technical Qualities

- precise
- accurate
- responsive

Regulatory Qualities

- universally applicable
- easy to relate to ecological condition
- easy to compare to a standard



Two common approaches for quantifying biotic condition

Species loss indices (e.g., O/E indices)

Ecological structure indices (e.g., multi-metric indices including IBIs)

Scoring Tools Rely on Reference Sites to Establish Expected Conditions

- 485 reference sites used to develop scoring models
- Excellent coverage of CA's natural stream diversity



Species Loss Index (O/E)

Compare number of **observed** (“O”) taxa to
number of **expected** (“E”) taxa

Step 1. Cluster reference sites based on biological similarity

Step 2. Identify natural gradients that best explain clusters (=predictors)

Step 3. Use predictor values at test sites to predict species expected to be observed



O/E Update

- April index performed well and science panel liked it
- **Reference pool adjustments:**
 - added sites to target under-represented areas
 - dropped sites based on stakeholder feedback
- Created new model

Final Model (Random Forests, 10 clusters, 4 predictors):

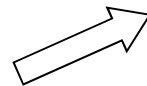
- Average Monthly Temperature (2000-2009)
- Average Monthly Precipitation (2000-2009)
- Log Watershed Area
- Site Elevation

Performance was very similar to our April O/E index

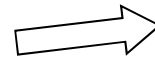
Multi-metric Indices (MMIs)

Species list is converted into metrics representing diversity, ecosystem function, and sensitivity to stress

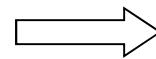
<u>Taxon</u>	<u>Count</u>
Mayfly species 1	43
Mayfly species 2	12
Mayfly species 3	2
Beetle species 1	1
Beetle species 2	1
Midge genus 1	65
Midge species 1	3
Midge species 2	10
Midge genus 2	3
Dragonfly species 1	2
Stonefly species 1	1
Stonefly species 2	14
Worm species 1	9
Worm species 2	2



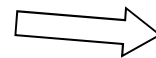
mayfly taxa



predator taxa



% sediment tolerant taxa



% herbivore taxa



% mayfly individuals

Why develop an MMI?

- Science panel recommended exploring MMI
- MMIs have useful qualities
 - Measure ecological attributes other than species loss
 - Very responsive to stress
 - May work well where species-specific predictions are tricky
- **New techniques available** (*see Hawkins and Vander Laan presentation at 2011 CABW*)
 - Adds site-specific adjustments to traditional MMIs



Building a predictive MMI (pMMI)

follows methods of Hawkins and Vander Laan

Step 1. Calculate lots of metrics at reference and stressed sites

***Step 2.** Create models that adjust metric values to account for major natural sources of metric variation

Step 3. Select metrics based on ability to discriminate reference from stressed sites

Step 4. Score metrics (after Cao et al. 2007) and assemble into composite pMMI

Step 1. Calculate metrics at reference sites and stressed sites

- **Sample Information:**
 - 1520 sites had “adequate” samples (i.e., >450 bugs)
 - 514 are reference (same definition as O/E)
 - 175 are highly stressed
- **Calculate Metrics**
 - Used SWAMP’s new bioassessment reporting module
 - Subsample to 500 organisms, calculate at SAFIT Level 1 (midges to family)
- **Use 80% for model development, 20% to validate**

Metrics: the usual suspects

Class	Abundance-based	# Taxa	% Taxa
Taxonomic	% EPT	EPT taxa	% EPT taxa
	[not considered]	Coleoptera taxa	% Coleoptera taxa
	[not considered]	Diptera taxa	% Diptera taxa
	% Chironomidae	[NA]	[NA]
	[not considered]	Non-insect taxa	% Non-insect taxa
	Shannon Diversity	Taxonomic richness	
FFG	% Collectors	Collector taxa	% Collector taxa
	% Predators	Predator taxa	% Predator taxa
	% Scrapers	Scraper taxa	% Scraper taxa
	% Shredders	Shredder taxa	% Shredder taxa
Tolerance	% Intolerant	Intolerant taxa	% Intolerant taxa
	% Tolerant	Tolerant taxa	% Tolerant taxa
	Weighted tolerance value		

Step 2. Adjust metric values to account for influence of natural gradients

- Models allow us to predict site-specific reference expectation for each metric
- Most influential gradients:

- Latitude
- Longitude
- Elevation Range
- Site Elevation
- Precipitation
- Temperature
- log Watershed Area

- Soil Erodability
- Soil Bulk Density
- Soil Permeability
- Hydraulic Conductivity

- MgO_Mean
- Surfur_Mean
- SumAve_Phos
- CaO_Mean
- Mean Phosphorus
- Mean Nitrogen

Step 3. Select most responsive metrics

- Select metrics with the best ability to discriminate reference from stressed (i.e., highest t-value)
- Avoid selecting redundant metrics

Step 4. Score metrics and assemble into composite pMMI (*follows Cao et al. 2007*)

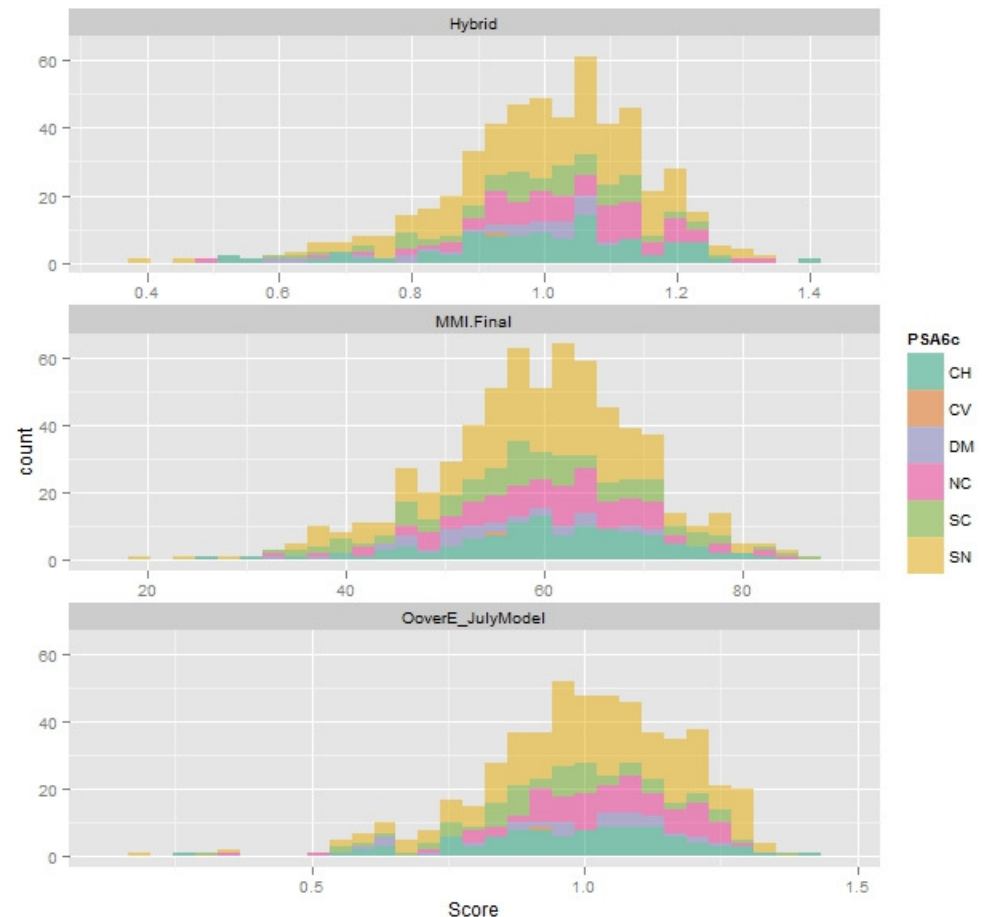
- **Score each metric** by comparing observed value to reference expectation
- **Sum** 10 metrics and **adjust** scale to be equivalent to O/E (divide score by mean of reference)

Final Metrics

Metric	Mod v Null	% explained by RF model	t	Response
Collector taxa	Modeled	11	13.2	Decrease
Coleoptera taxa	Modeled	40	17.6	Decrease
Diptera taxa	Null	7	13.5	Decrease
Intolerant taxa	Modeled	53	32.2	Decrease
Predator taxa	Modeled	11	13.6	Decrease
Scraper taxa	Modeled	38	20.0	Decrease
Shredder taxa	Modeled	42	19.1	Decrease
% Non-Insect Taxa	Modeled	15	18.1	Increase
Shannon diversity	Modeled	16	10.7	Decrease
Tolerance value	Modeled	32	12.4	Increase

Comparing Performance of 3 Scoring Tools

1. Species Loss Index (O/E)
2. Ecological Structure Index (pMMI)
3. Combined Index (“hybrid”)



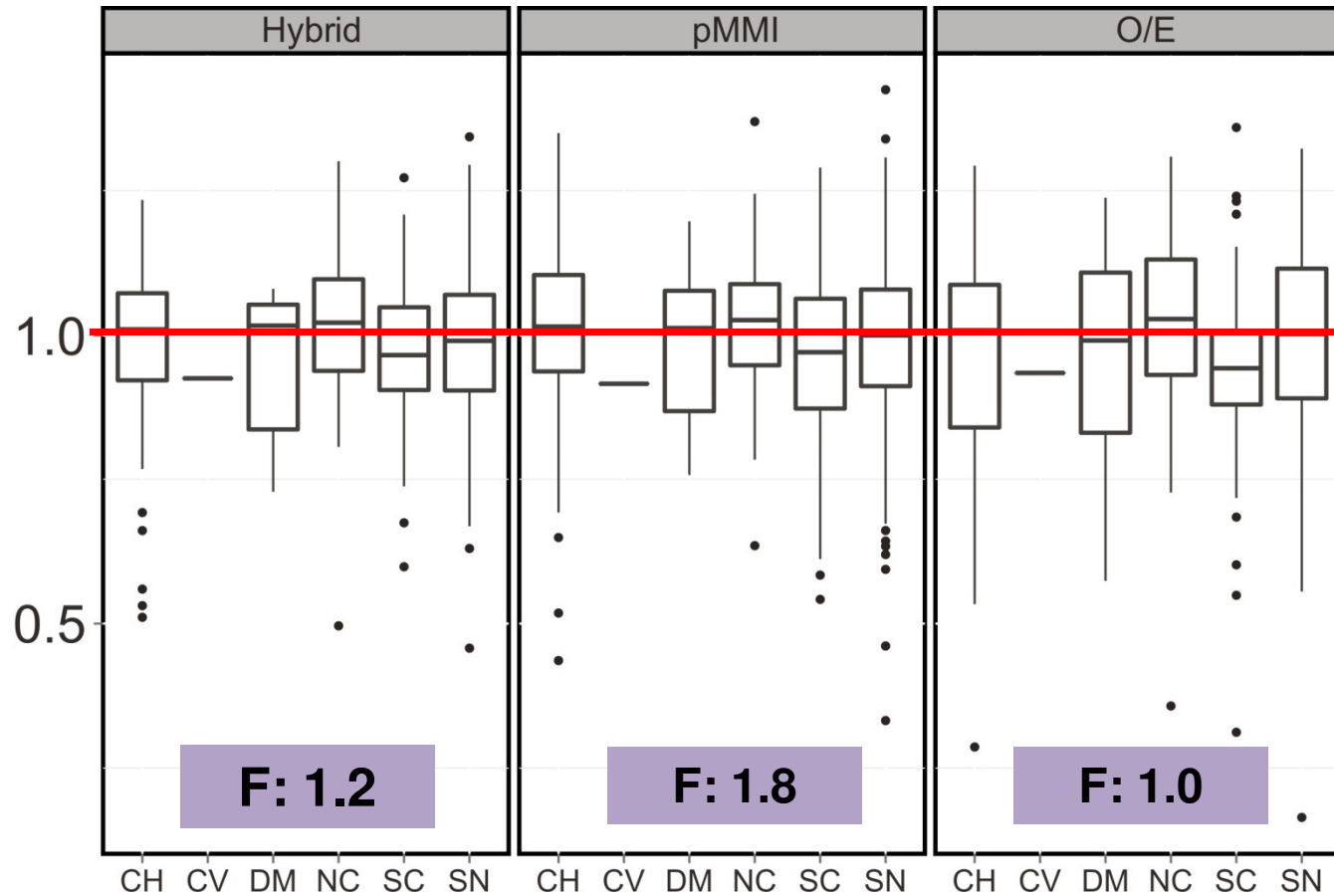
Measuring Performance

All evaluations used a common dataset

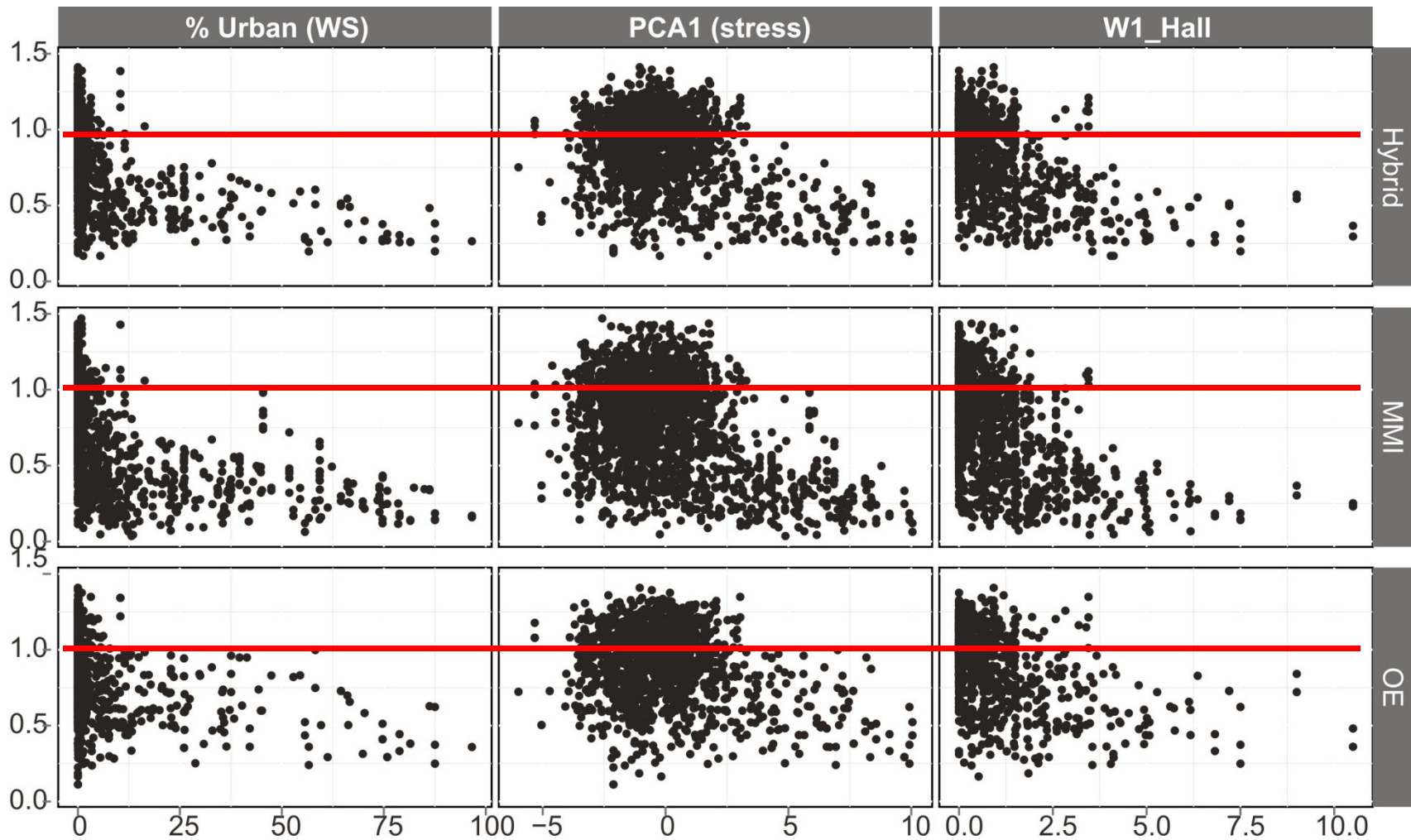
Class	Property	Measure	O/E	pMMI	Hybrid
Precision	Variance of reference sites	SD	0.19	0.15	0.14
Sensitivity/ Responsiveness	Discrimination	t-test	9.5	17.6	15.3
	Variance explained by stress	Random forest model	25%	56%	49%
Accuracy/ Bias	Variance explained by natural gradients (ref sites)	Random forest model	-7%	-9%	-8%
	Difference among PSA regions (ref sites)	ANOVA	1.0	1.8	1.2
Replicability	Within-site variability	Mean within-site SD	0.10	0.10	0.08

Statewide Consistency

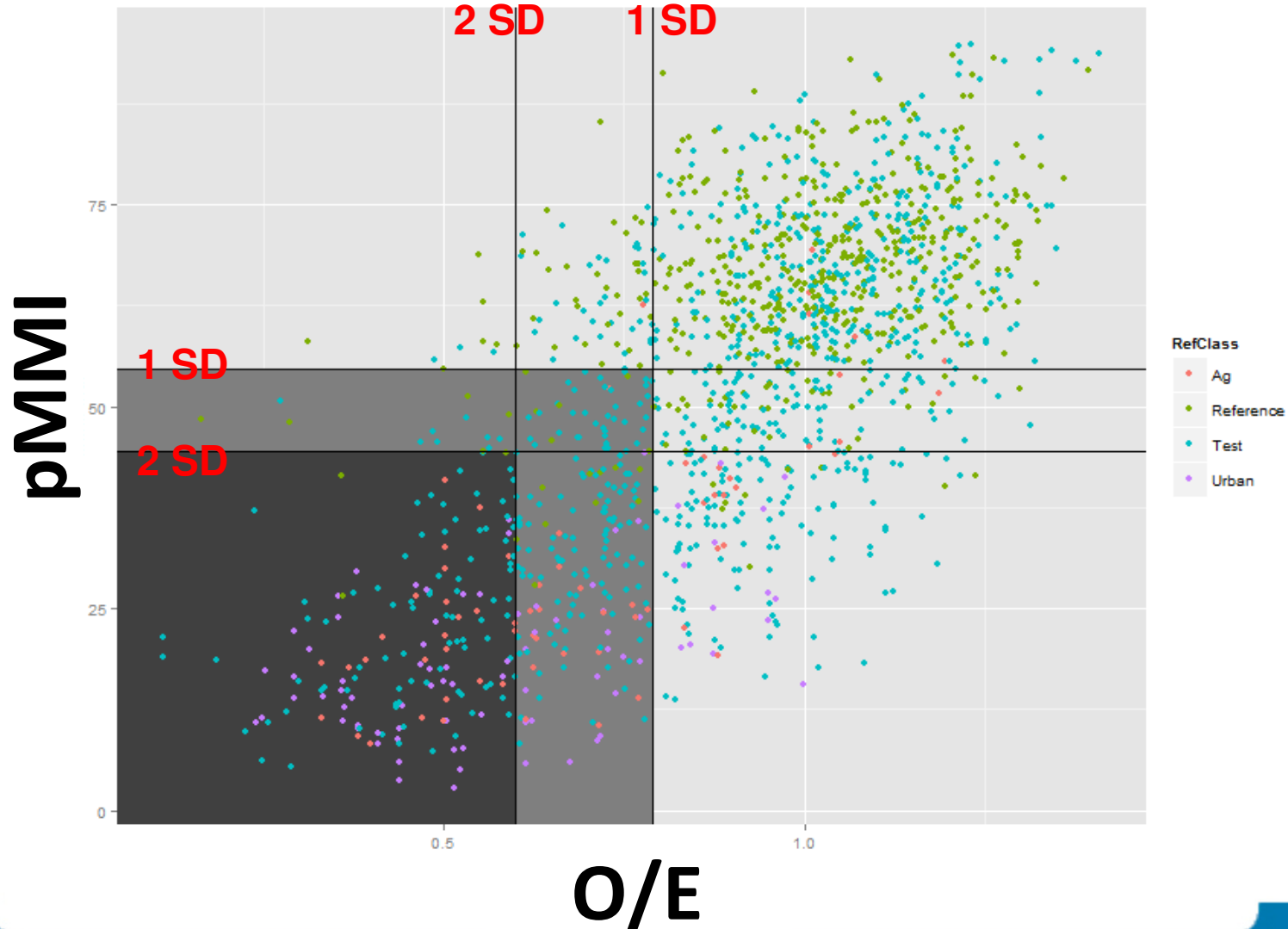
Distribution of reference scores by PSA region



Responsiveness to stress



pMMI and O/E have general agreement, but tell us somewhat different things



Both pMMI and O/E have desirable qualities

- pMMI is precise and very responsive to stress (but it was designed to be)
- % species loss is an intuitive, meaningful measure of condition
- Both are accurate and applicable throughout state
- Potential for complementarity is great -- we explored a few options (see Science Panel)

We recommend a combined index

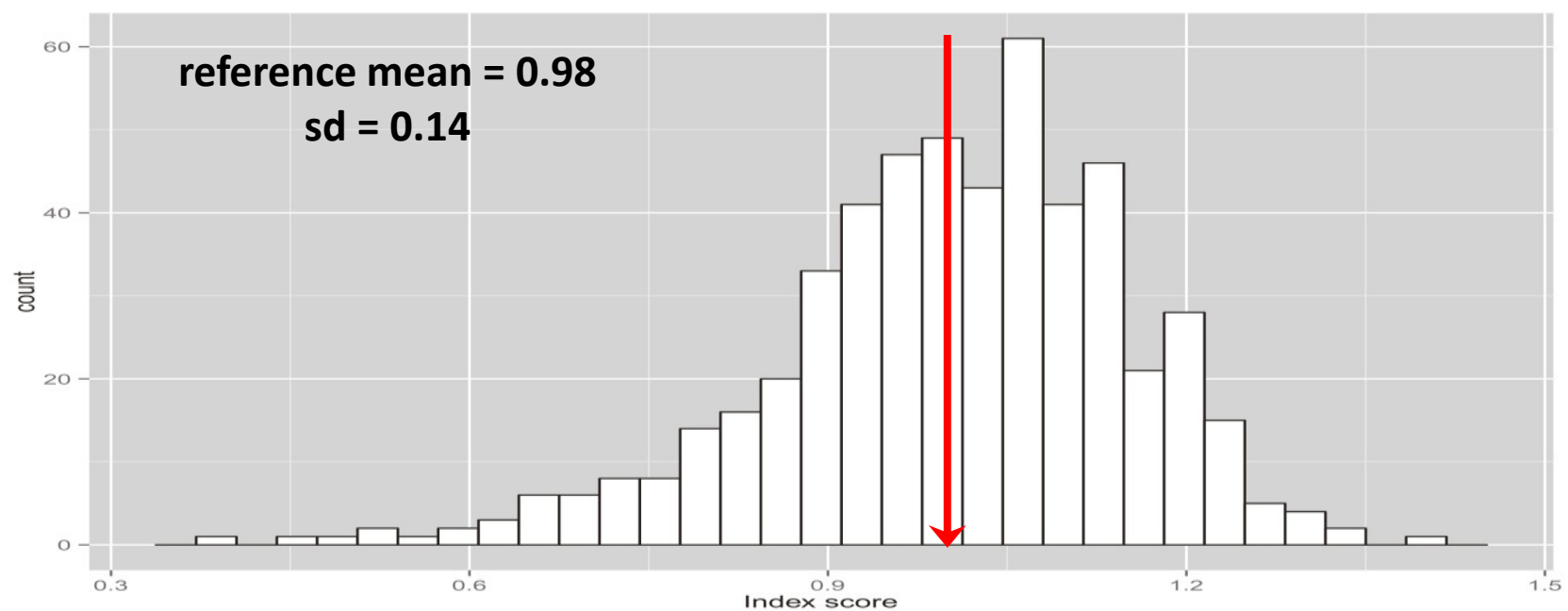
- Retains some of the better qualities of both indices, tempers weaknesses
- Can be disaggregated into component MMI and O/E (i.e., you don't lose information by combining)
- Implementation is easier with a single score

California Stream Condition Index (CSCI)

Part A: Ecological Structure Component (pMMI)

Part B: Taxonomic Loss Component (O/E)

CSCI is a simple average of the two scores



Options for setting thresholds

- Statistical criteria
 - Standard deviation
 - %-ile of reference distribution
- Ecological criteria
 - Acceptable species loss or change in community structure

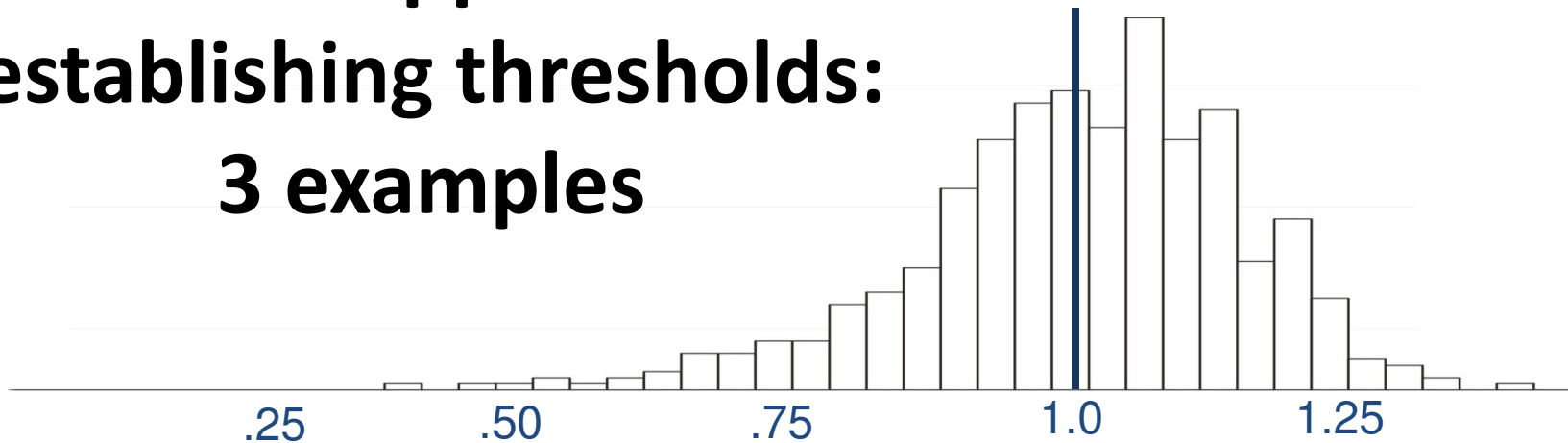


We recommend statistically defined thresholds

- Widely accepted practice
- Setting objective ecological benchmarks is complex
- No technical objections from the science panel



Statistical approaches to establishing thresholds: 3 examples



95% and 85% confidence that site is not equivalent to reference

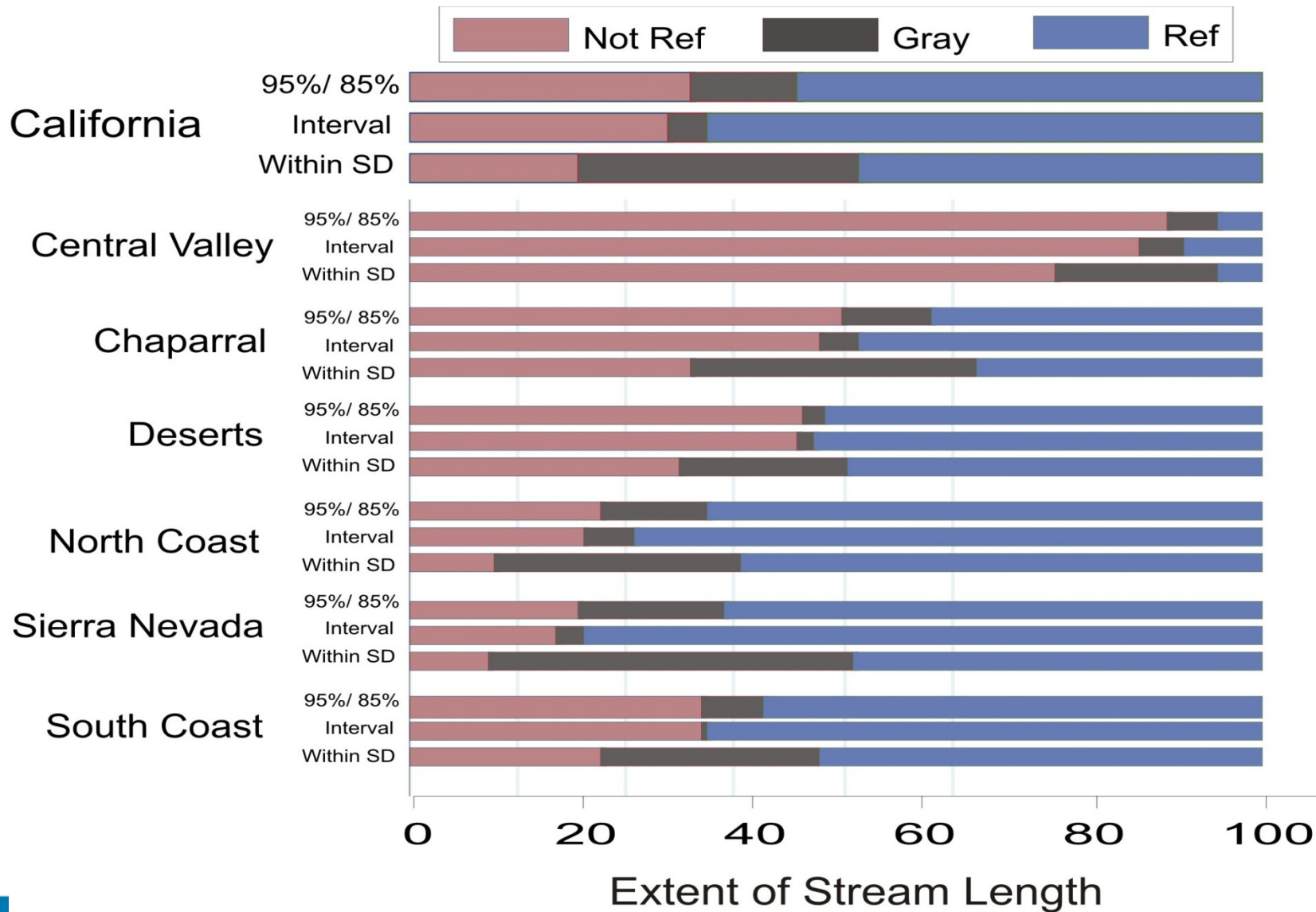


95% confidence that the 95% threshold is where we think it is



Use within-site error rate to establish uncertainty around threshold

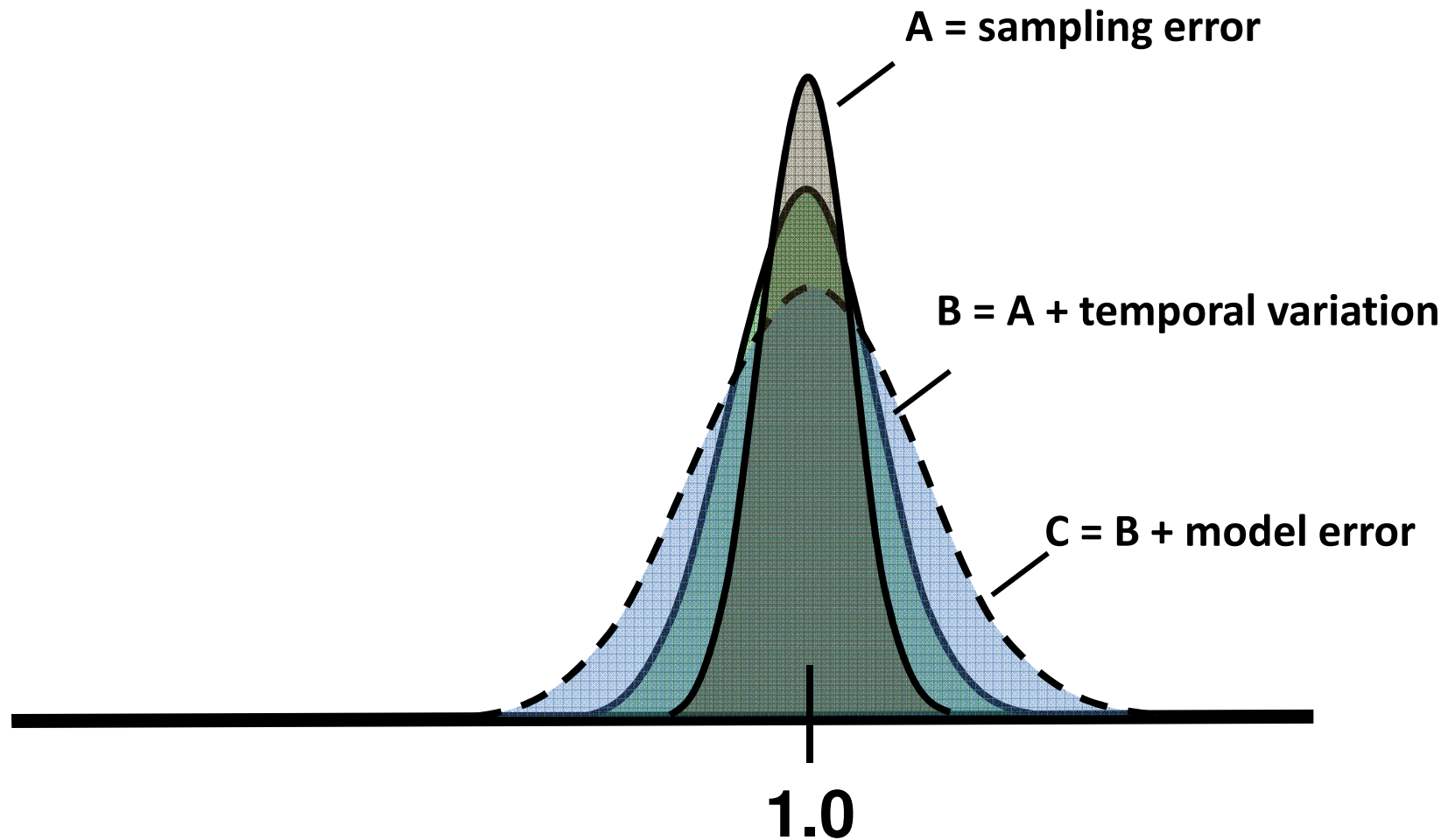
Extent of stream length by region



What's next?

- Explore limits of tool
- Automate calculations
- Document, document, document

Sources of variation in O/E scores



(after Hawkins et al. 2010)