

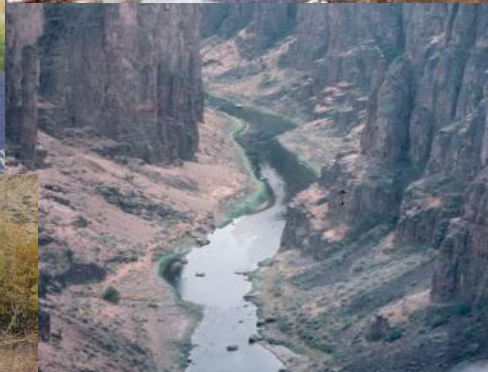
Use of RIVPACS-type Multitaxon Distribution Models to Assess Alterations in Regional Biodiversity

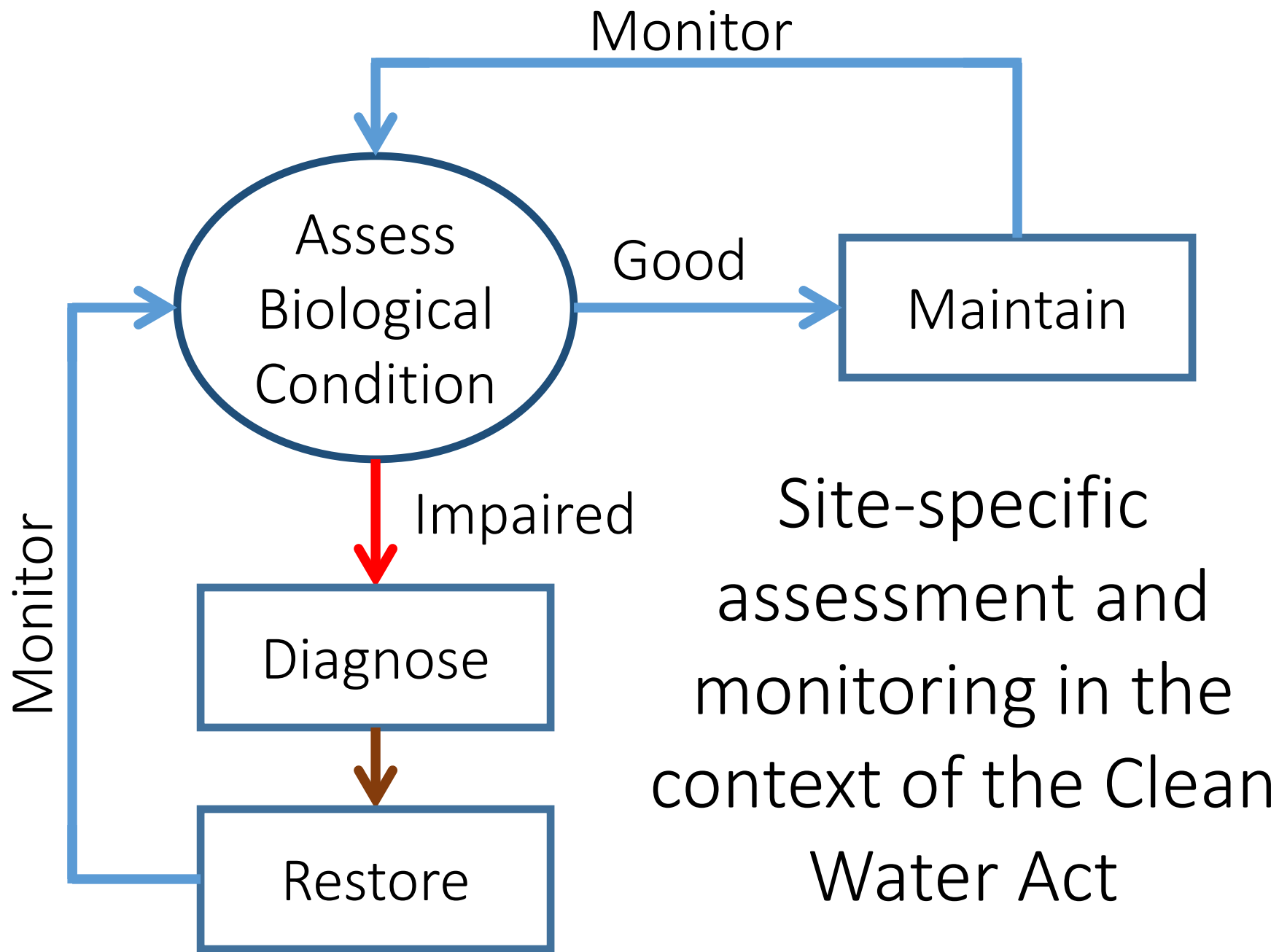
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Traditional application of RIVPACS models

$$p_i = \sum_{j=1}^m p_j c_{j,i} \quad E = \sum p_i$$

O/E

(site-specific measure of
taxonomic completeness)

Traditional RIVPACS-type site-specific assessments

What should be there (E)?



$E = 8$ species

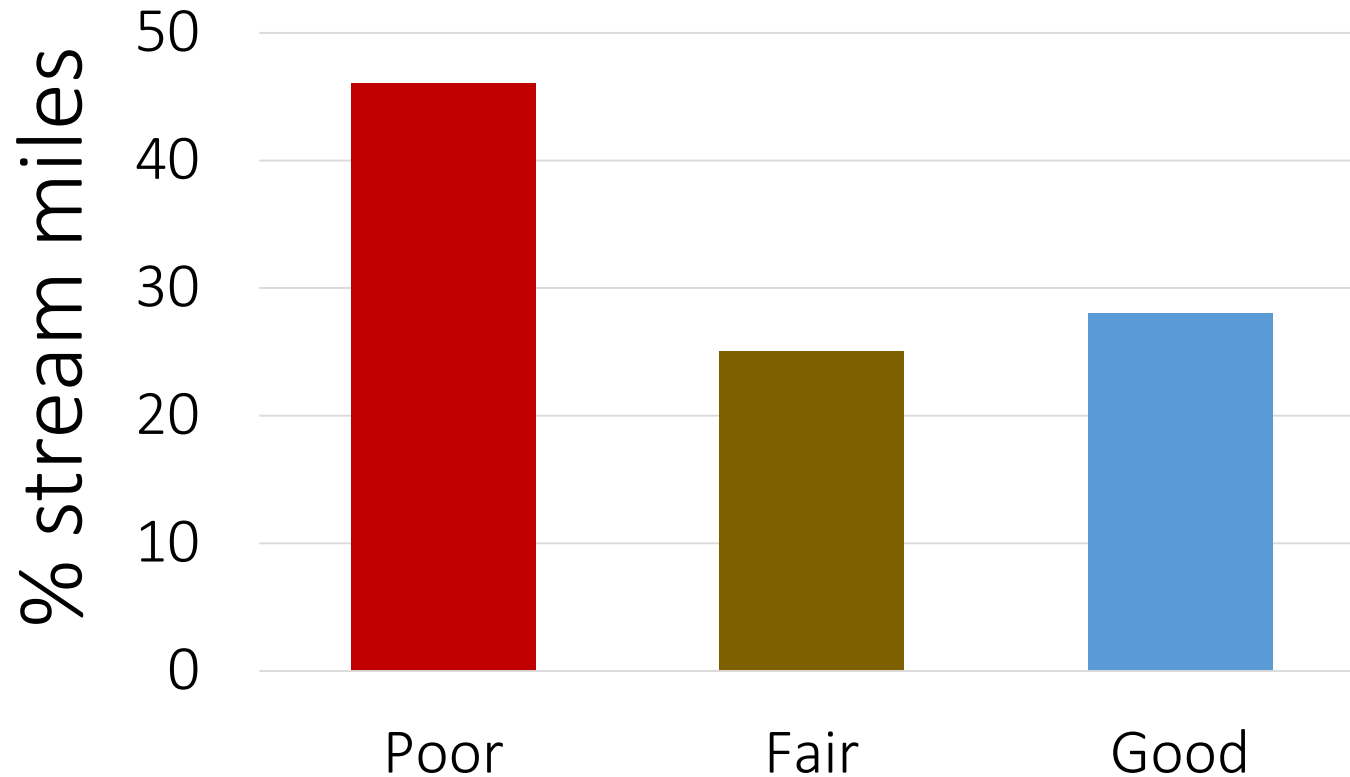
What is actually there (O)?



$O = 3$ species

$$O/E = 0.38$$

NARS scales up site-specific index values
to quantify regional conditions



2008/2009 NRSA

But what about the status of individual species?



With RIVPACS outputs we can also compare observed frequencies of occurrence with expected ones.

$$F_e = \sum p_i / N$$

$$D_i = F_o - F_e$$

Examples from North Carolina and the Mid-Atlantic Highlands

Multitaxon distribution models reveal severe alteration
in the regional biodiversity of freshwater invertebrates

Charles P. Hawkins^{1,3} and Lester L. Yuan^{2,4}

Freshwater Science. 2016/ 35(4) – online
open access

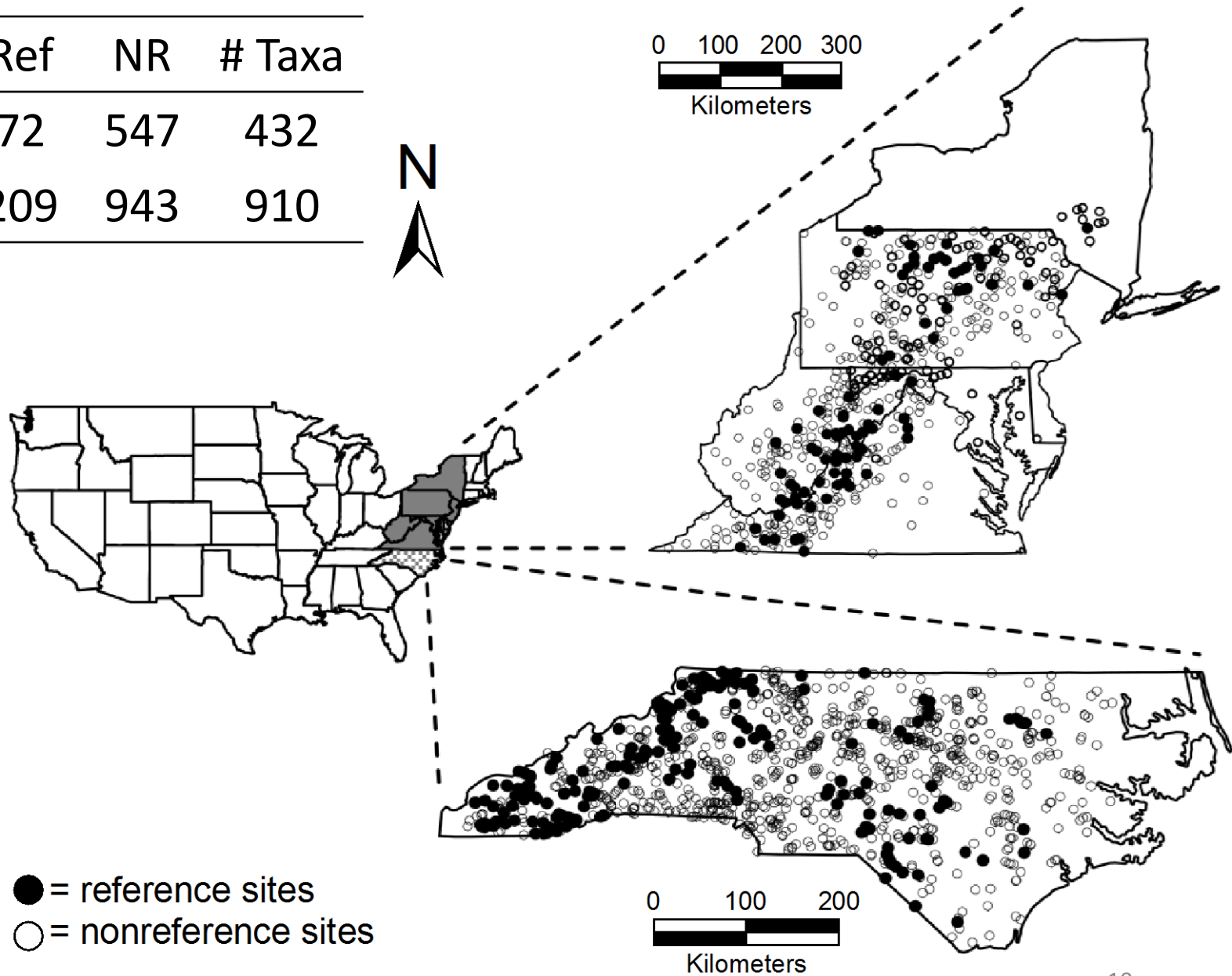
Google - DOI: 10.1086/688848

or

https://www.researchgate.net/profile/Charles_Hawkins

Sites

Region	Ref	NR	# Taxa
MAH	72	547	432
NC	209	943	910



Model Predictors

North Carolina	Mid-Atlantic Highlands
Latitude	Elevation
Longitude	Day of Year
Elevation	Catchment Area
Stream Width	Carbonate Chemistry
Stream Depth	Ecoregion
% Boulder	
% Rubble	
Day of Year	

How well can we predict F_e ?

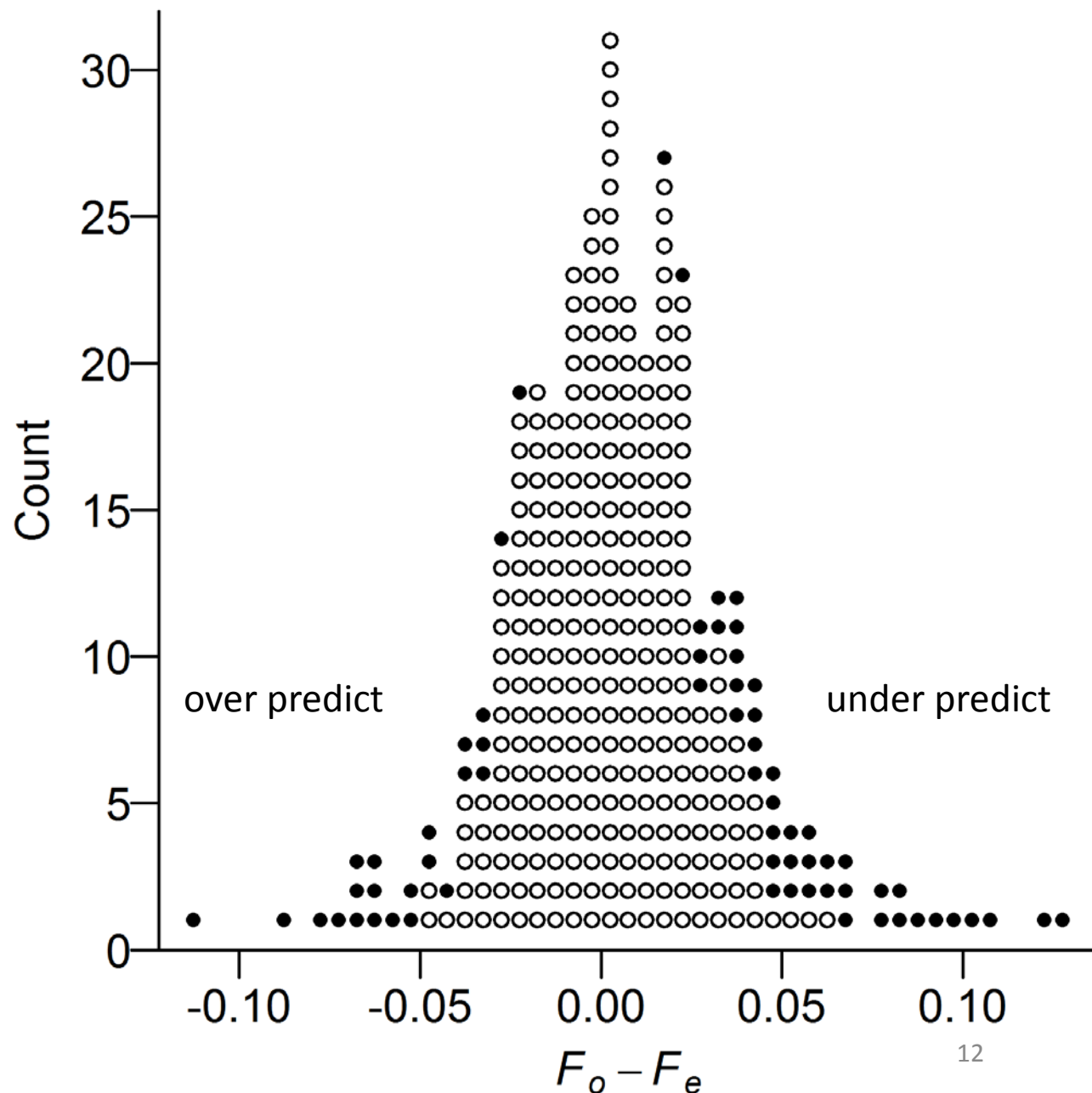
202 NC validation samples.

Used taxa ≥ 5 sites.

D_i values for 350 taxa.

Black dots = 66 taxa (19%) that were either over- or under-predicted.

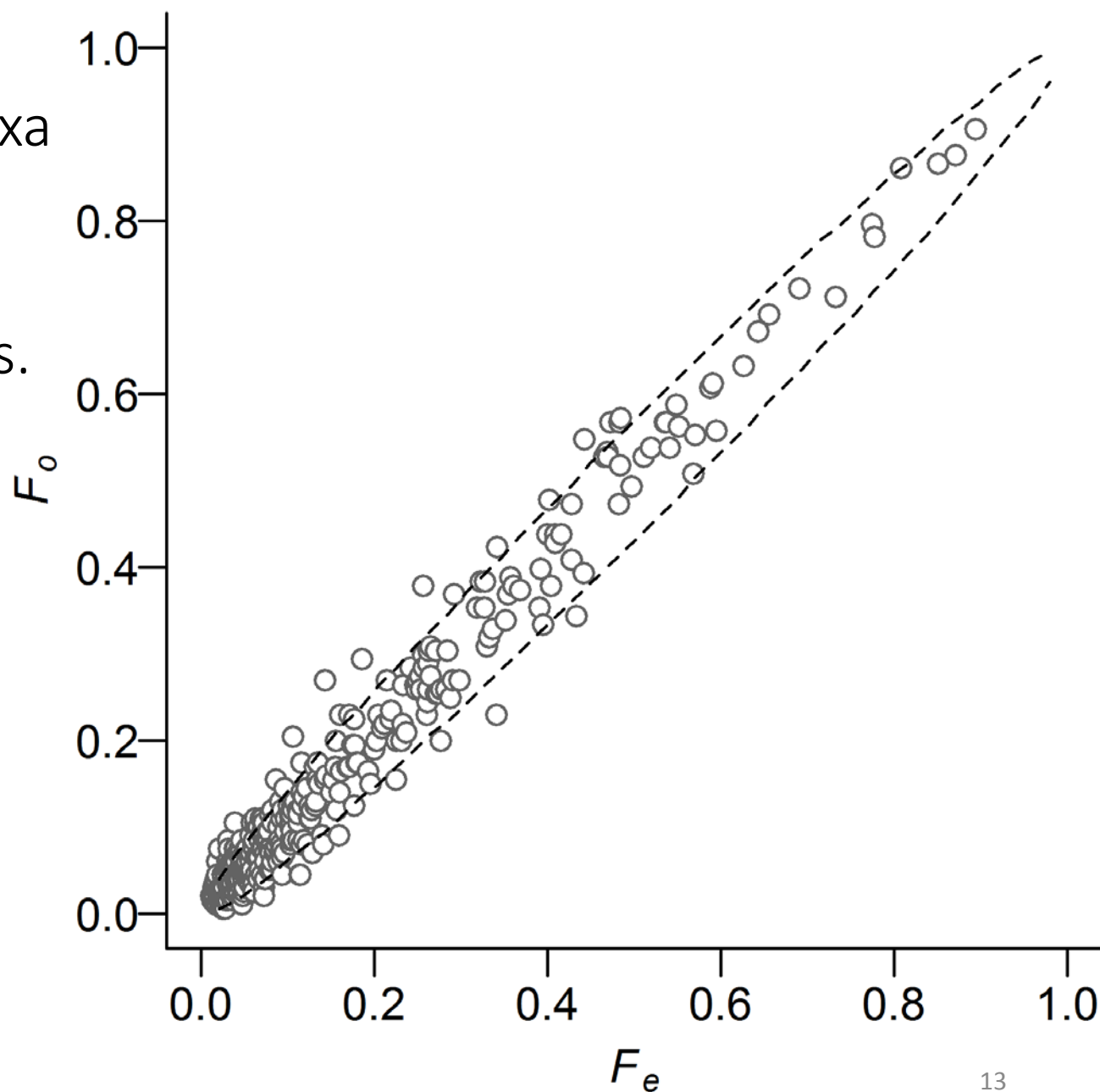
Under-prediction 2X more than over-prediction.

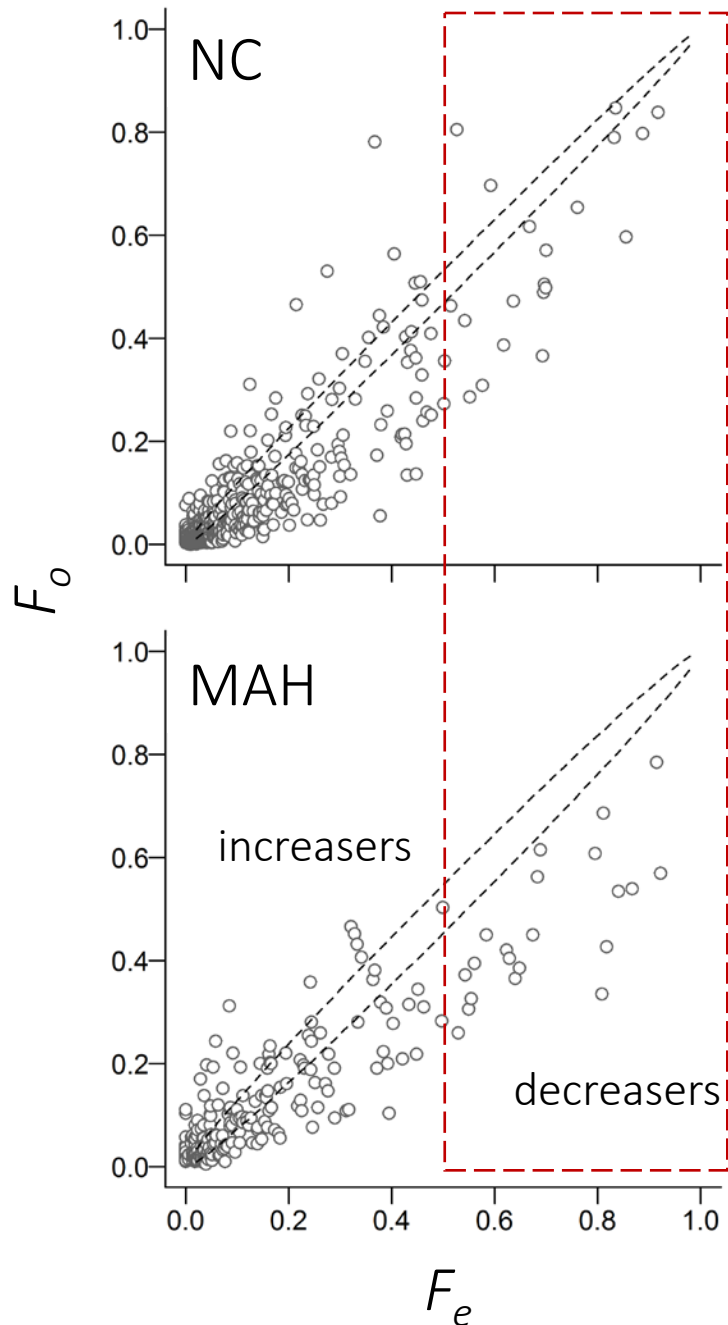


F_o vs F_e for 350 taxa
from 202 NC
reference-quality
validation samples.

Ellipse:
95% CI for F_e

Most under/over
predictions were
for rarer taxa.





Example Taxa

Taxon	D_i
<i>Acroneuria abnormis</i>	-0.26
<i>Isonychia</i>	-0.20
<i>Polycentropus</i>	-0.33
<i>Hexatoma</i>	-0.27
<i>Rheocricotopus robacki</i>	+0.28

Taxon	D_i
<i>Ephemerella</i>	-0.35
<i>Baetis</i>	-0.33
<i>Stempellinella</i>	-0.39
<i>Epeorus</i>	-0.47
<i>Hexatoma</i>	-0.26

Most common taxa were decreaseers!

	North Carolina			Mid-Atlantic Highlands		
Group	# Taxa	%D	%I	# Taxa	%D	%I
Total	547	50	26	251	36	35
Plecoptera	46	74	0	22	68	5
Trichoptera	102	79	7	29	52	21
Ephemeroptera	91	67	10	23	61	13
Diptera	174	36	36	115	30	34
Coleoptera	28	46	36	14	29	57
Odonata	29	2	59	9	11	44
Oligochaeta	25	16	56	18	11	72

So What? / Now What?

- Comparisons of F_e and F_o should be a standard part of NARS summaries of biological condition.
- D_i analyses provide a needed and direct link between CWA and ESA and also informs CBD targets.
- Technical stuff:
 - RIVPACS is a good start but methods that adjust for imperfect detection may be needed.
 - DNA-based IDs could provide needed 'species' level resolution and improve detection estimates.
- California has the data and tools needed to estimate D_i for hundreds of stream invertebrates – who will Pete have do it?

1,318 Probabilistic Sites

Map courtesy of Calvin Yang
SWAMP/SWRCB