

# California Statewide Algae Plan: Past, Present, and Future

Susie Theroux  
SCCWRP



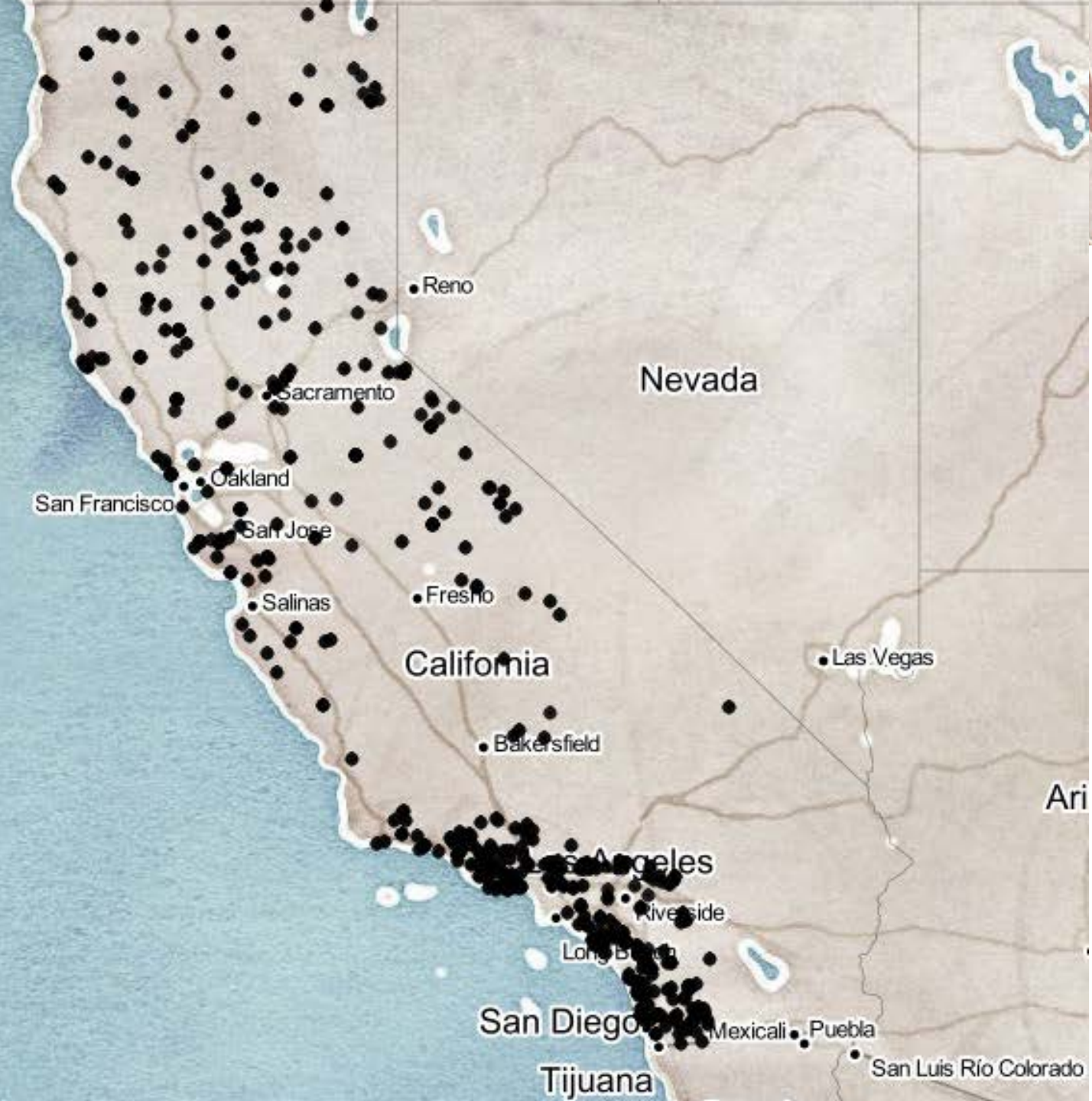


2008



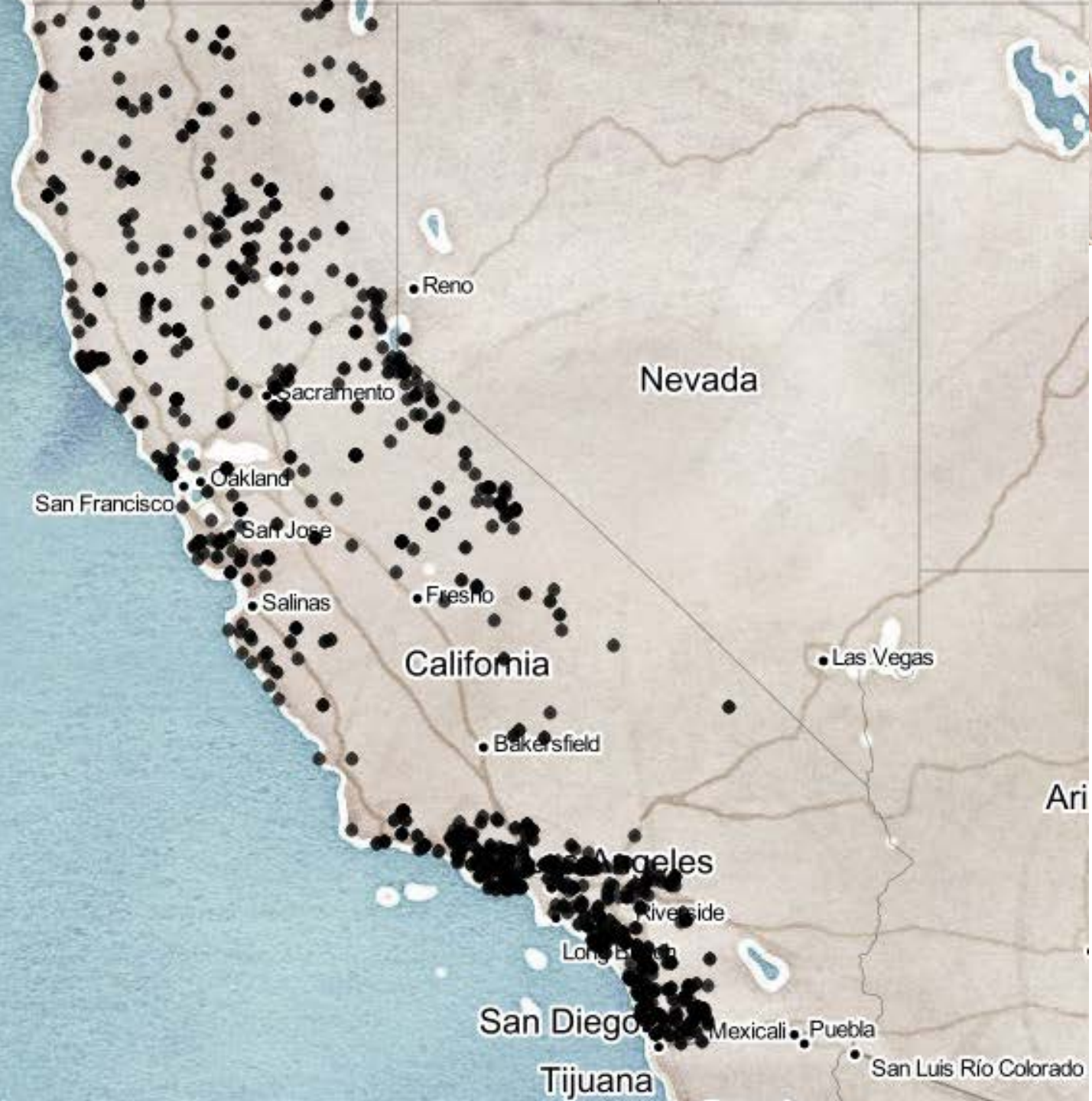


**2008-2009**



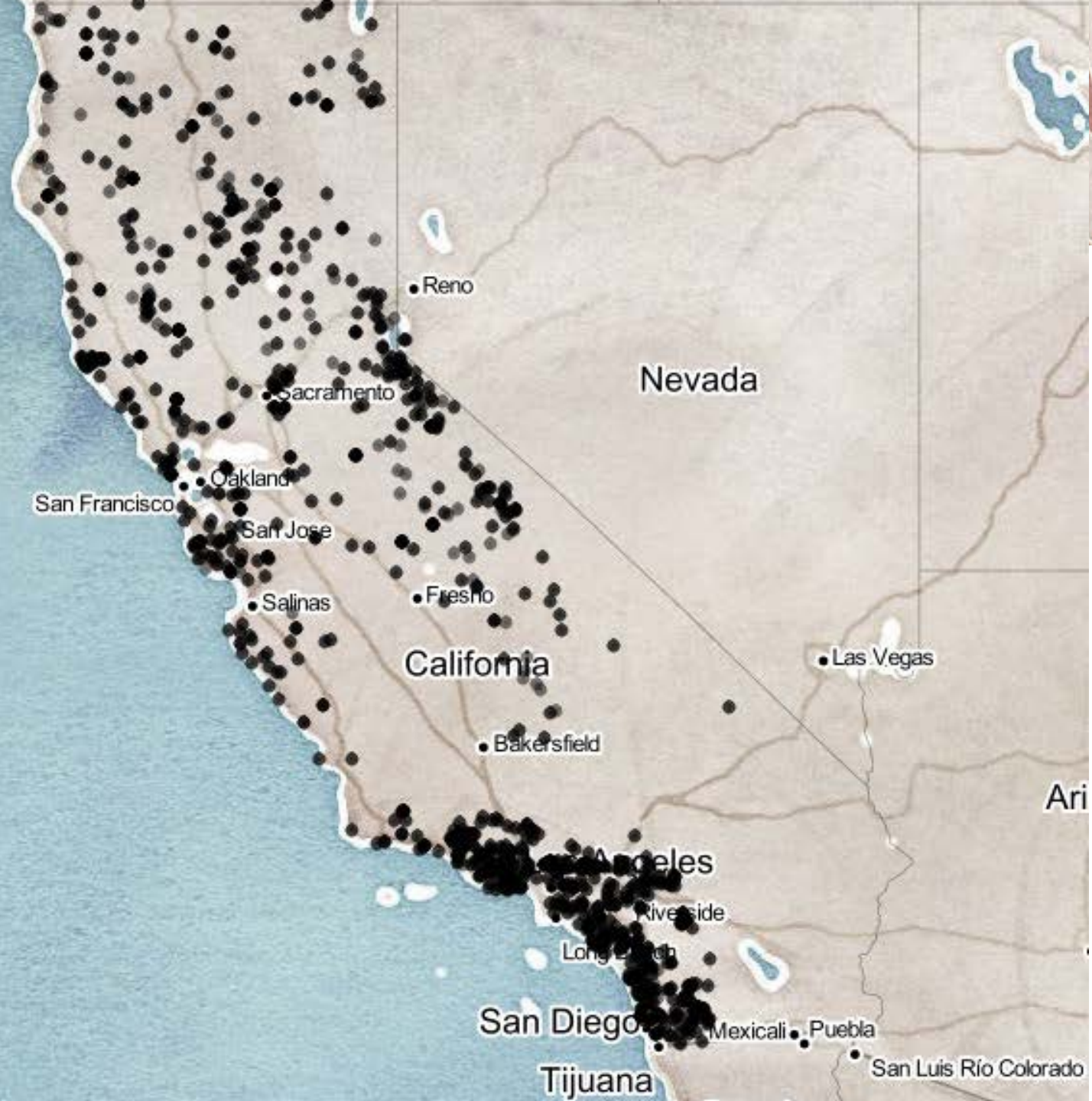


**2008-2010**



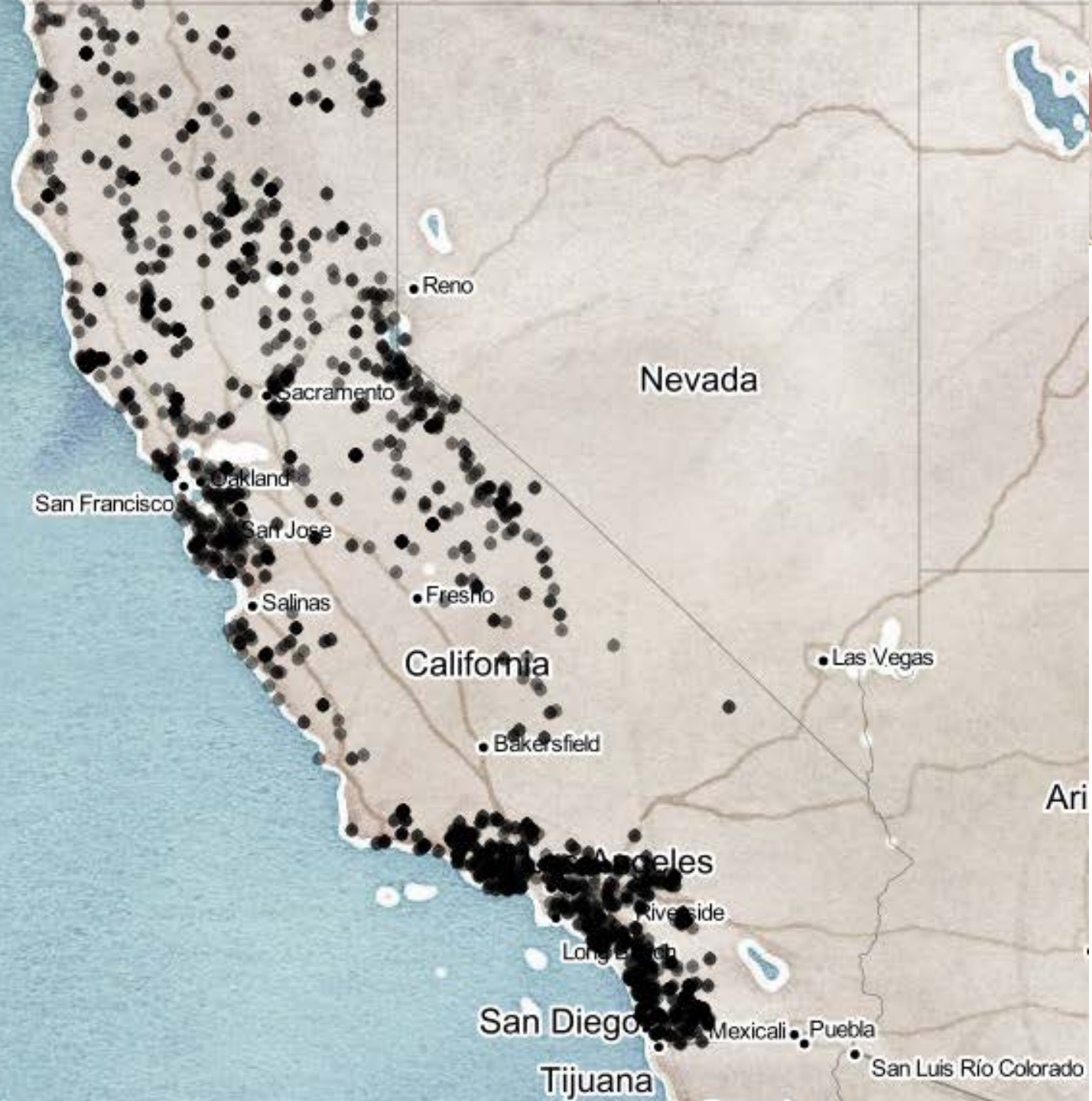


**2008-2011**



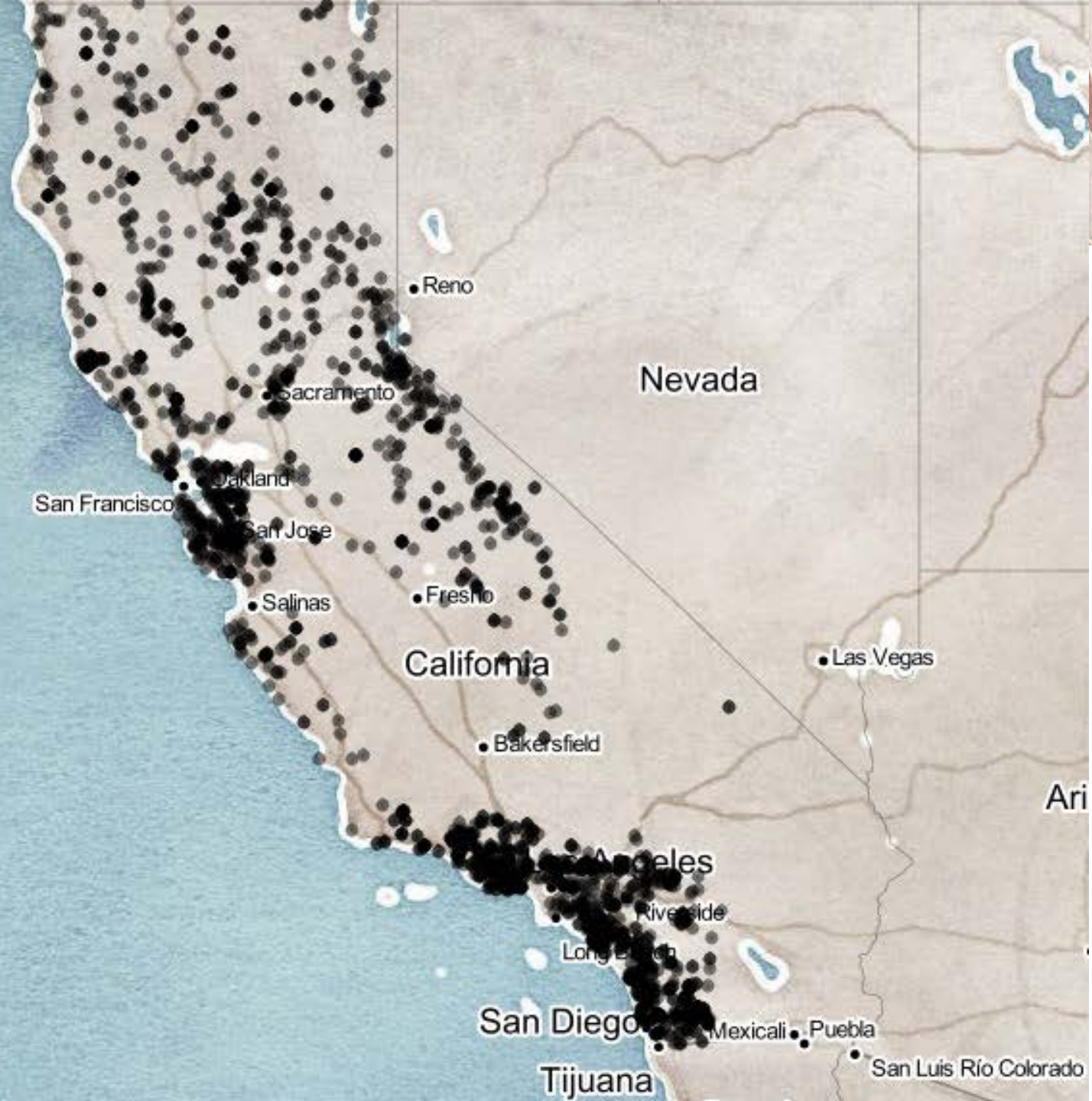


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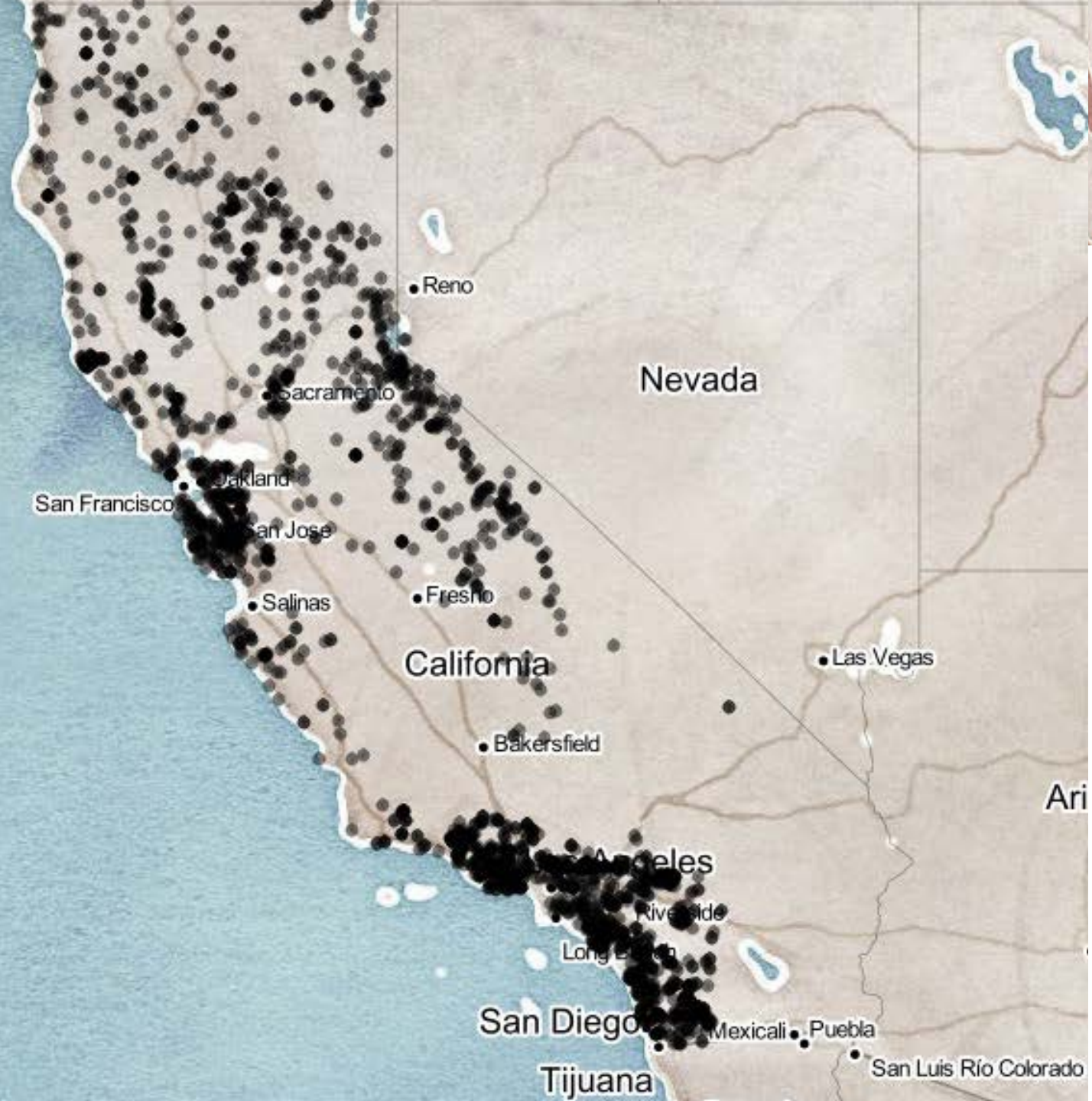


**2008-2013**



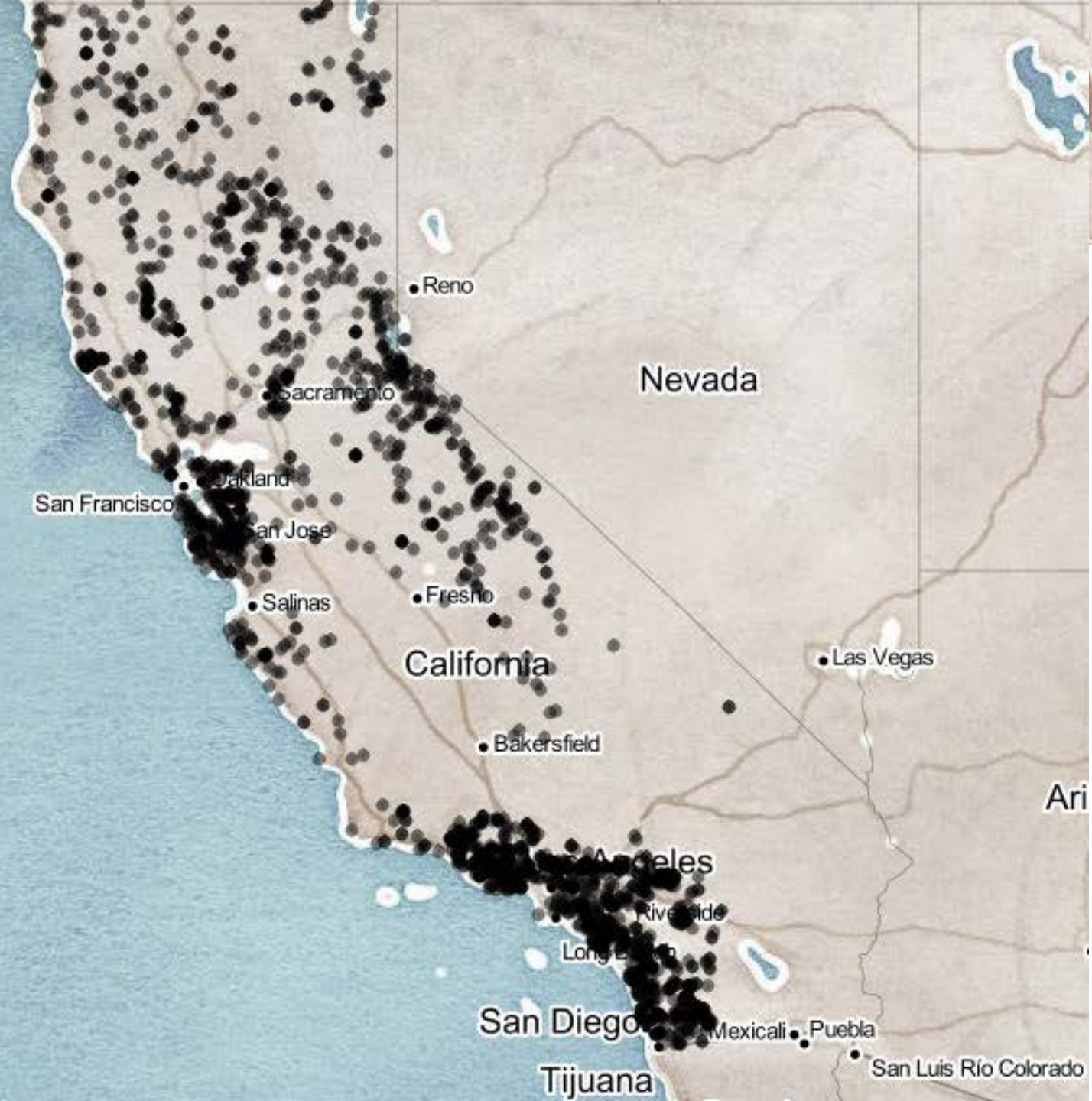


**2008-2014**





**2008-2015**





# Statewide Algae Plan

## Past – Standard methods/tools

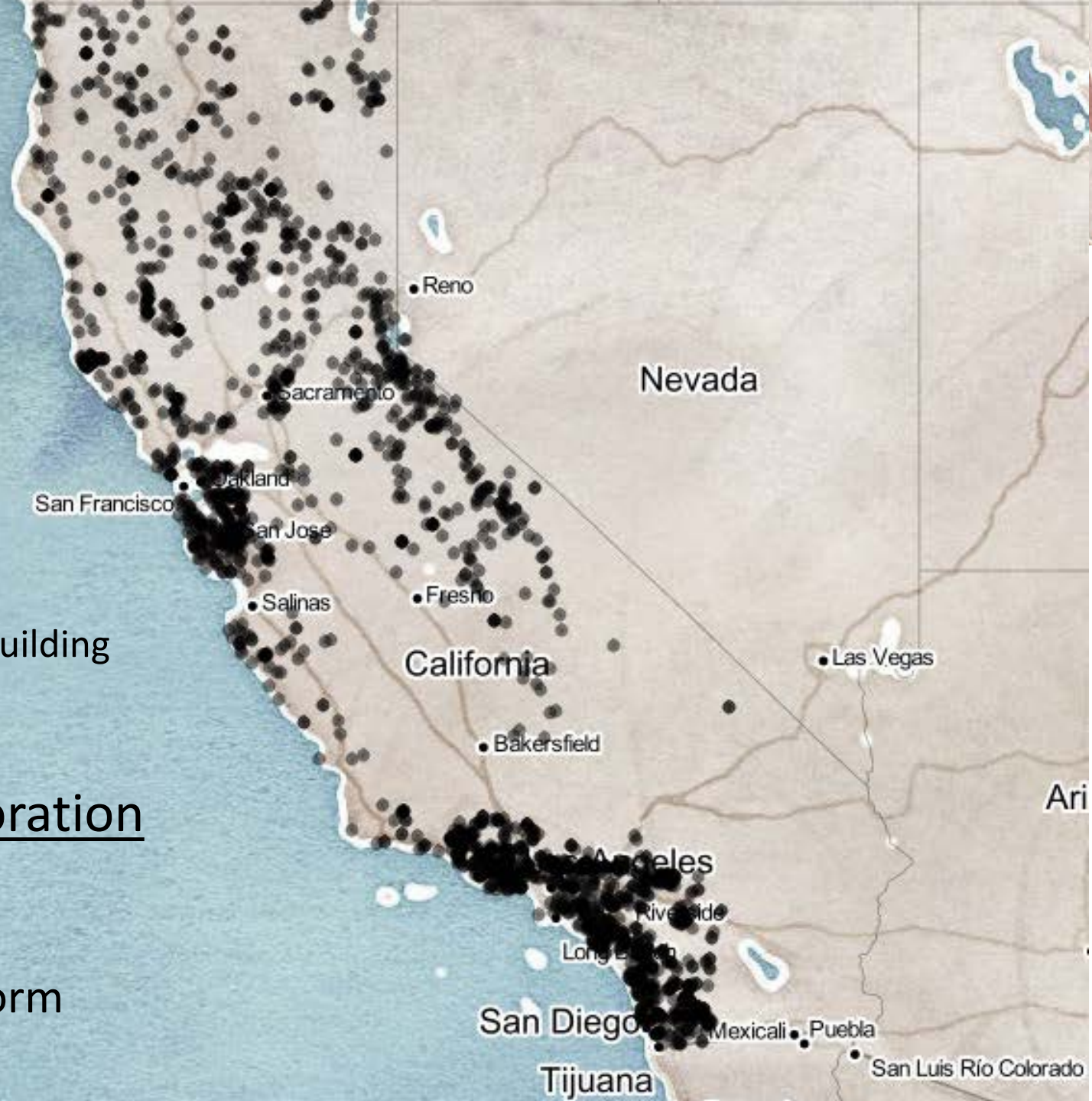
- Sampling and lab protocols
- S. California Algal IBI

## Present - Multiuse datasets

- Statewide Algal Index
  - STE
  - Molecular methods
- } Capacity building

## Future – Communication/collaboration

- Integrating tools into State infrastructure
- Using algal bioassessment to inform regulatory policy





# Why algae?

- Provide a direct link to nutrient concentrations and imbalances
- Sensitive to changes in water chemistry
- Short life span, rapid growth rate and rapid response to stress
- High dispersal rates and high species numbers
- Toxic species





# Algal bioassessment protocols



SWAMP Bioassessment Procedures May 2016

## STANDARD OPERATING PROCEDURES (SOP) FOR THE COLLECTION OF FIELD DATA FOR BIOASSESSMENTS OF CALIFORNIA WADEABLE STREAMS: BENTHIC MACROINVERTEBRATES, ALGAE, AND PHYSICAL HABITAT

Prepared by:

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**A. Elizabeth Fetscher**, Senior Environmental Scientist, San Diego Regional Water Quality Control Board

**Lilian B. Busse**, Environmental Scientist, San Diego Regional Water Quality Control Board

SWAMP-SOP-SB-2016-0001



SWAMP Bioassessment Procedures 2015

## Standard Operating Procedures for Laboratory Processing, Identification, and Enumeration of Stream Algae

September 2015

**Rosalina Stancheva<sup>1</sup>, Lilian Busse<sup>2</sup>, Patrick Kociolek<sup>3</sup> and Robert Sheath<sup>1</sup>**

<sup>1</sup> California Primary Algae Laboratory  
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California State University San Marcos  
333 S. Twin Oaks Valley Road  
San Marcos, CA 92096

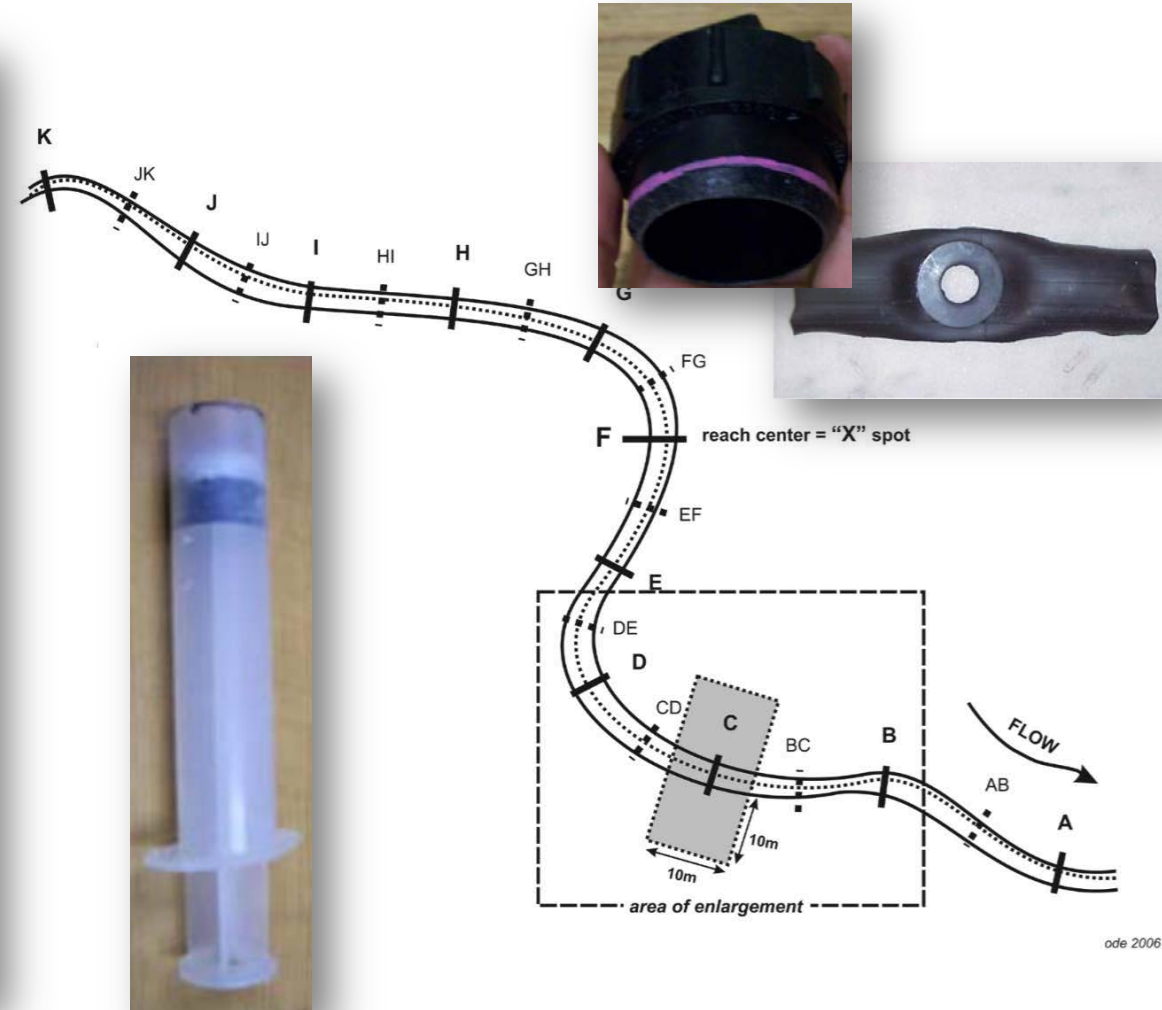
<sup>2</sup> San Diego Regional Water Quality Control Board  
State Water Resources Control Board  
9174 Sky Park Court  
San Diego, CA 92123

<sup>3</sup> Museum of Natural History and Department of Ecology and Evolutionary Biology, University of Colorado  
UCB 218, Boulder, CO 80309

SWAMP-SOP-2015-0003



[www.waterboards.ca.gov/swamp](http://www.waterboards.ca.gov/swamp)



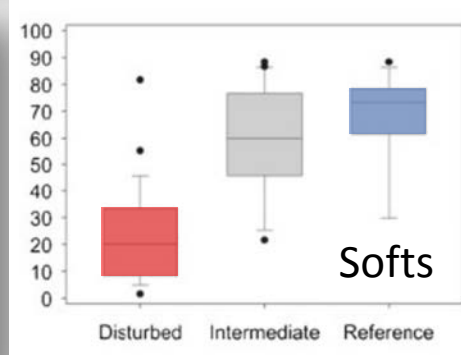
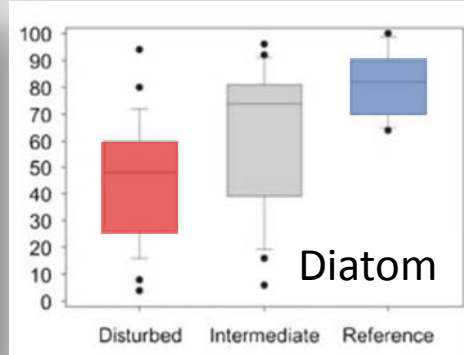
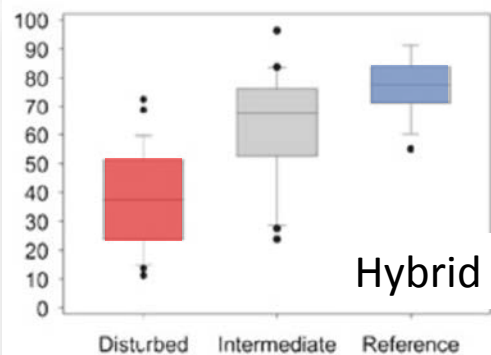


# Algal Index of Biotic Integrity (IBI)

J Appl Phycol (2014) 26:433–450  
DOI 10.1007/s10811-013-0088-2

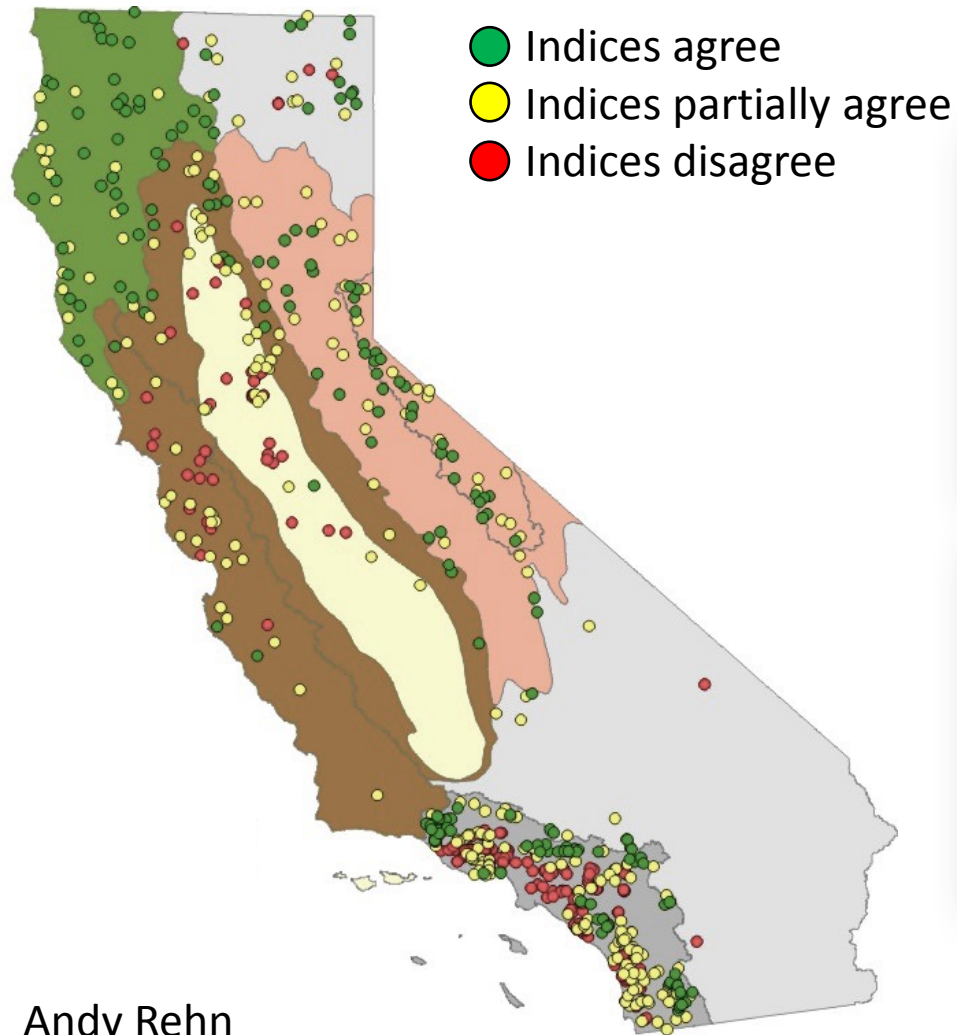
## Development and comparison of stream indices of biotic integrity using diatoms vs. non-diatom algae vs. a combination

A. Elizabeth Fetscher • Rosalina Stancheva •  
J. Patrick Kociolek • Robert G. Sheath • Eric D. Stein •  
Raphael D. Mazor • Peter R. Ode • Lilian B. Busse





# Two indices are better than one



- California Stream Condition Index (CSCI): benthic macroinvertebrate (bugs)
- Algal IBI (H2O)
- CRAM<sub>bio-phys</sub> measures biotic structure and physical structure
- **CSCI was more sensitive to physical habitat conditions, whereas IBI was more sensitive to chemistry**



# Current efforts: Statewide Algal Index

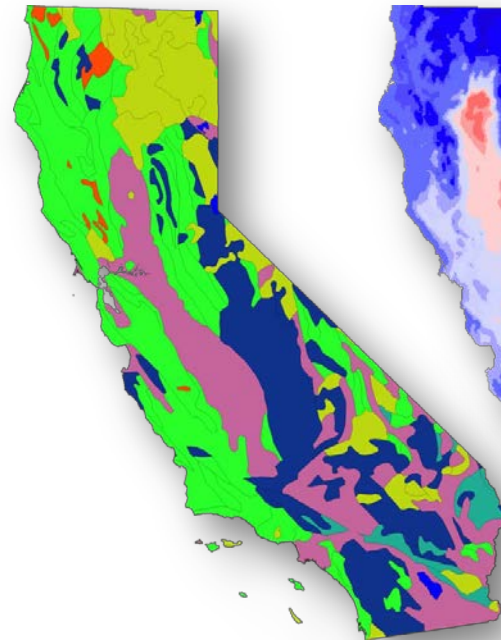
***Problem:*** Algal IBI tuned for Southern California

***Solution:*** Develop predictive statewide Algal Index

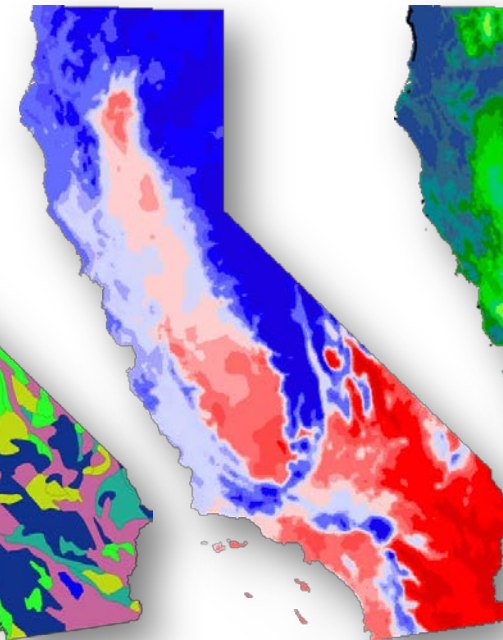
## Dataset

- Stormwater Monitoring Coalition (SMC)
- Perennial Stream Assessment (PSA)
- Reference Condition Management Program (RCMP)
- Regional Monitoring Coalition(RMC)
- 2000 stations, 3800 taxa

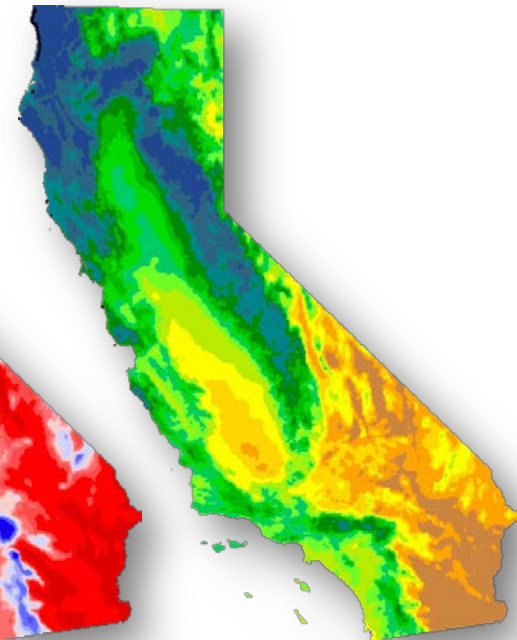
## Geology



## Temperature



## Precipitation





# Statewide Algal Index

## Reference-based approach

- Site-specific expectations for each site based on taxa found at groups of similar reference sites

## Observed/expected (O/E)

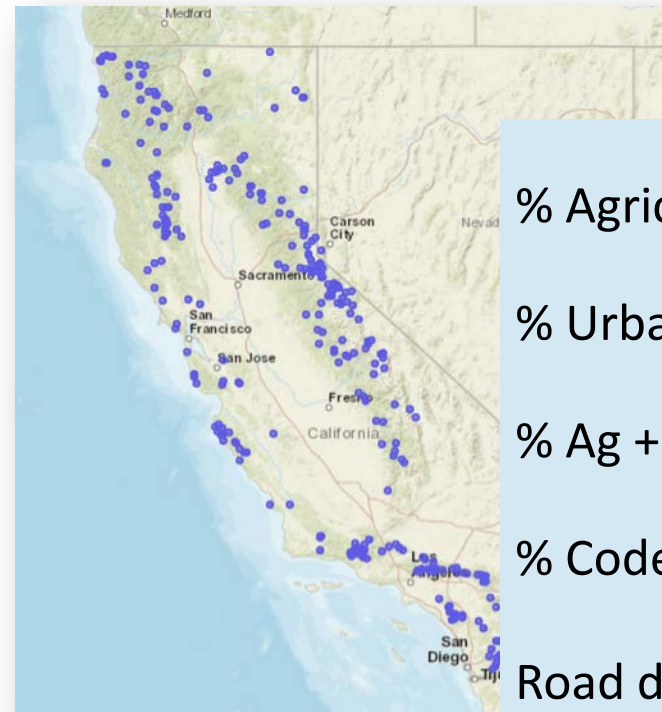
- Taxonomic completeness

## Multi-Metric Index (MMI)

- Ecological structure
- Comprised of several metrics that represent community structure

- ETA Fall 2017

## Reference



% Agriculture	Dam distance
% Urban	% Canals and pipelines
% Ag + % Urban	In-stream gravel mines
% Code 21	Producer mines
Road density	Specific conductance
Road crossings	W1_HALL

Mazor et al., 2016; Ode et al., 2016

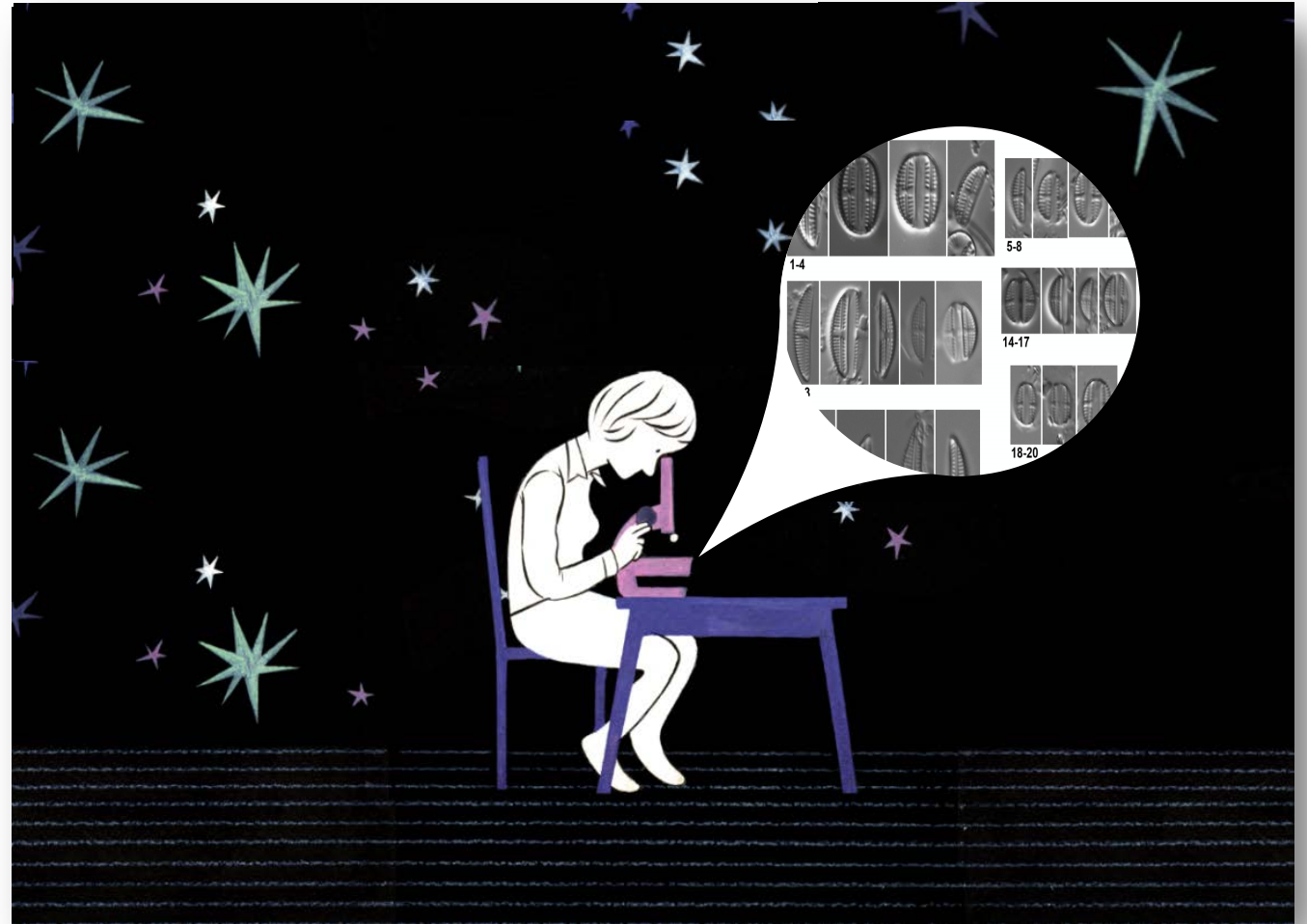


# Capacity building - STE

**Problem:** Taxonomy data not always comparable, variable levels of taxonomic resolution

**Solution:** Standard Taxonomic Effort (STE)

- Standard taxa list and level of taxonomic resolution
- Minimize time spent harmonizing algae taxonomy data from different labs

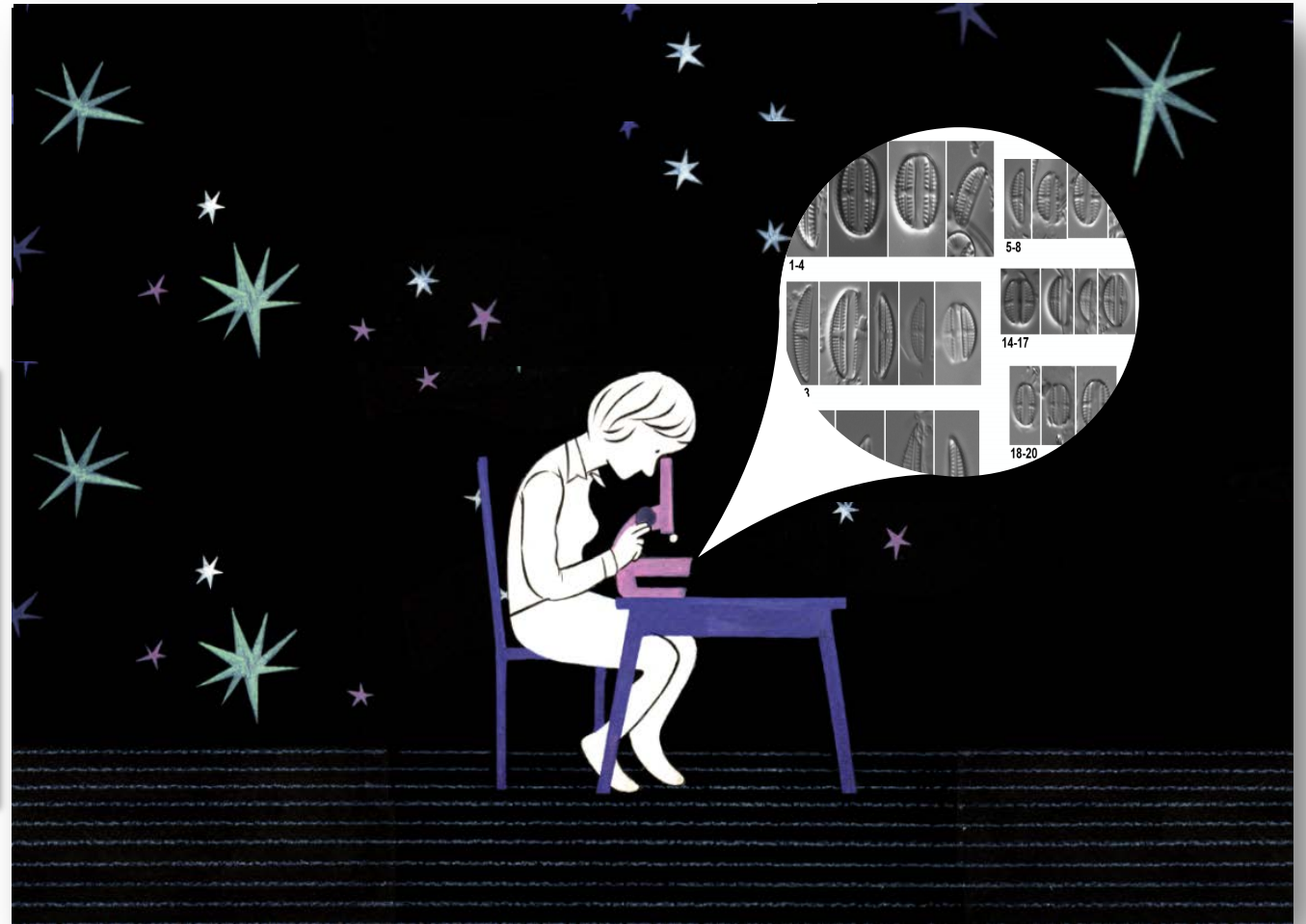




# Capacity building - STE

- Build upon the SAFIT model for bugs
- ETA: June 2017

Class	STE 1	STE 2	Low Abd	Trait	Hard to ID
Prasinophyceae	Genus	Species	x		
Cyanophyceae	Species	Species		x	
Zygnematophyceae	Genus	Genus			x





# Capacity building - Molecular methods

***Problem:*** Limited number of labs capable of performing algae taxonomic analyses; results in backlog of taxonomy data and long wait times





# Capacity building - Molecular methods

***Solution:*** Explore DNA-based approach to algae taxonomy

- Illuminate previously overlooked species
- Dozens of commercial and academic labs can perform analyses
- Inexpensive

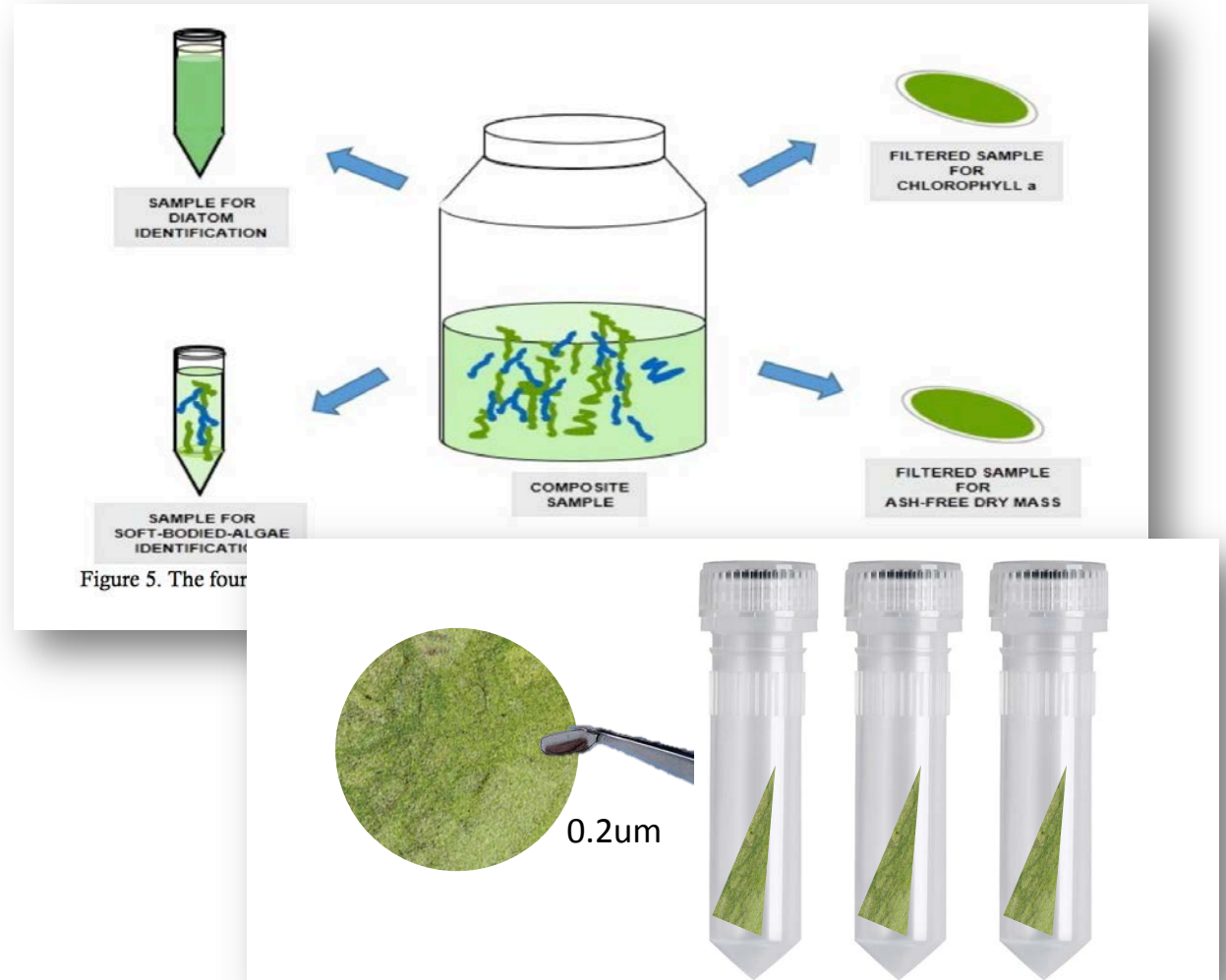




# Capacity building - Molecular methods

## DNA barcode approach

- Easily integrated into current field sampling protocols
- Sample stable (frozen) for months to years
- Pilot studies: ~200 Paired DNA/morphology samples collected during 2016





# Capacity building - Molecular methods



## Key questions for pilot studies

1. How do morphology-based and DNA-based algae taxonomy data compare?
2. What algal species are missing from DNA reference databases?
3. How well do algal indices perform with DNA data?

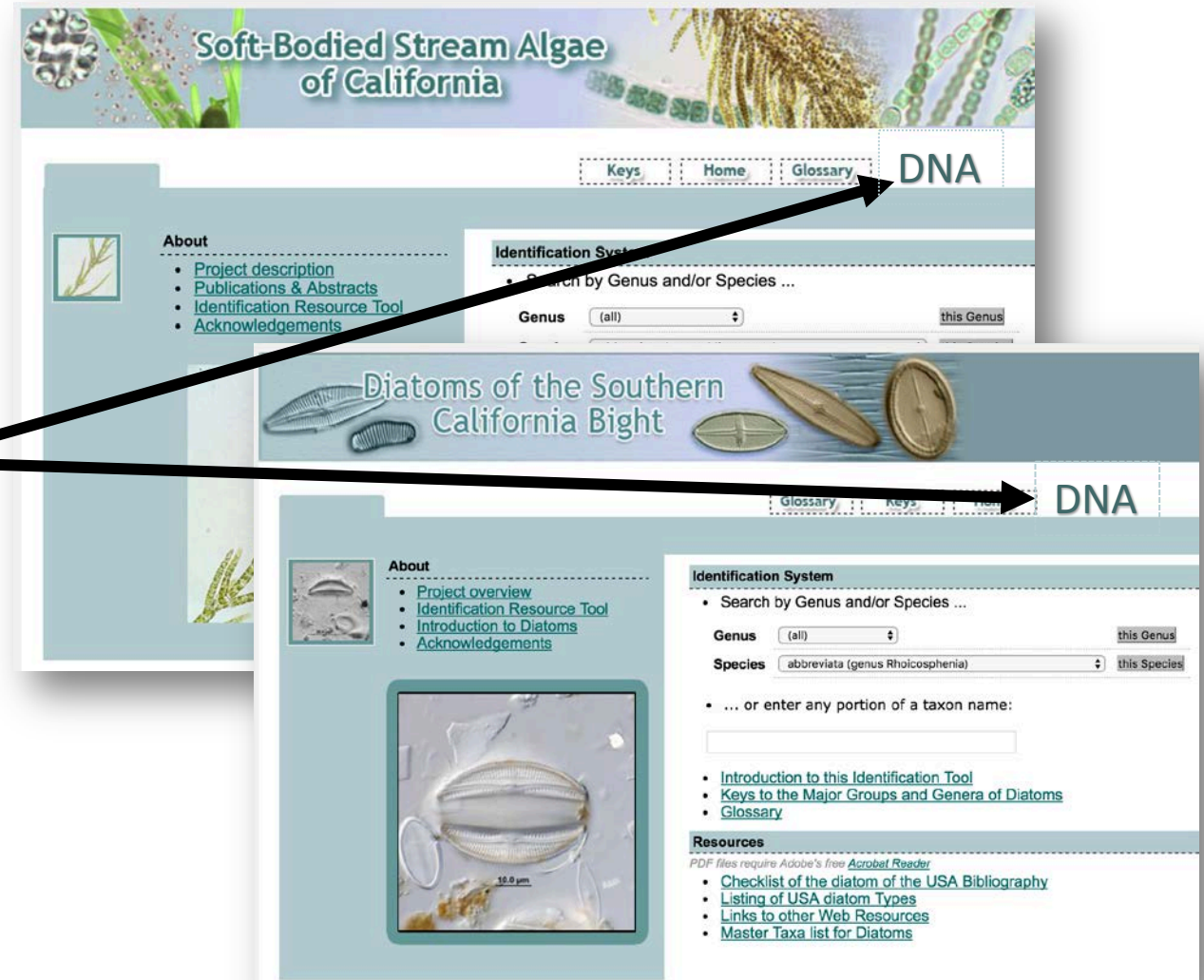
ETA: Early 2018



# Future plans

## Making algae tools accessible

- Incorporating Statewide Algal Index into online resources (SWAMP)
- Building California Algae DNA Reference Database
  - Isolating and sequencing priority taxa absent from existing DNA databases
- Guidance documents





# Future plans

## Integration of algal toolkit

- Support State Water Board's combined nutrient and biointegrity policy
  - Set nutrient thresholds?
  - How to use as one of multiple lines of evidence? In concert with other indices?
  - How to use in water programs, e.g. ambient assessment and 303(d) listing, NPDES, etc. ?
  - Required technical elements for its implementation?





# Thanks!

Contact: [susannat@sccwrp.org](mailto:susannat@sccwrp.org)

- Interested in algae DNA sampling? Have DNA samples to send to SCCWRP?
- Do you have algal bioassessment data not in SWAMP that could be useful in the statewide index development?

## Acknowledgements

- |                   |                            |
|-------------------|----------------------------|
| - SWAMP/SWRCB     | -Pete Ode, Andy Rehn       |
| - Regional Boards | -Rafi Mazor                |
| - CSUSM           | -Eric Stein, Martha Sutula |
| - SMC             | -Betty Fetscher            |
| - DFW             |                            |



# BONUS SLIDES





# Capacity building - Molecular methods

## How will Algal Indices perform with DNA barcode data?



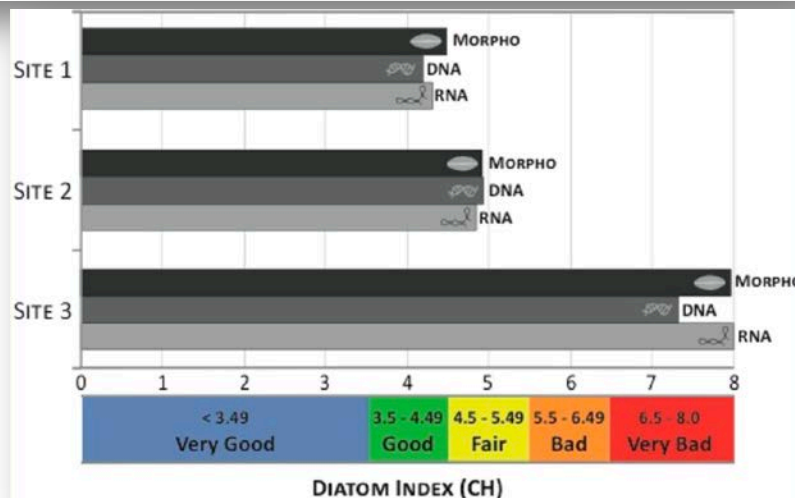
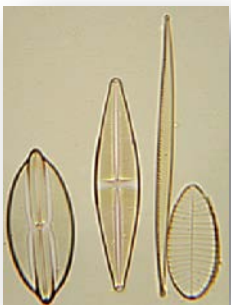
DOI: 10.1021/es506158m  
*Environ. Sci. Technol.* 2015, 49, 7597–7605

pubs.acs.org/est

### Environmental Monitoring: Inferring the Diatom Index from Next-Generation Sequencing Data

Joana Amorim Visco,<sup>†</sup> Laure Apothéloz-Perret-Gentil,<sup>†</sup> Arielle Cordonier,<sup>‡</sup> Philippe Esling,<sup>†,§</sup> Loïc Pillet,<sup>†,||</sup> and Jan Pawlowski\*,<sup>†</sup>

Swiss Diatom Index  
(DI-CH)



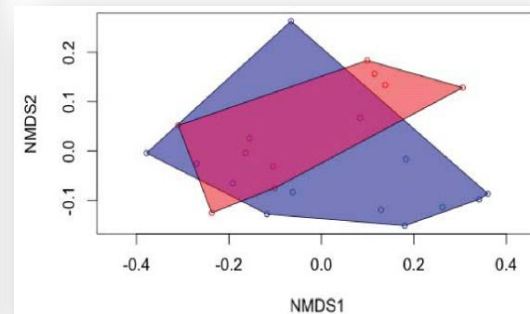
PLOS ONE | DOI:10.1371/journal.pone.0138432 October 21, 2015

### RESEARCH ARTICLE

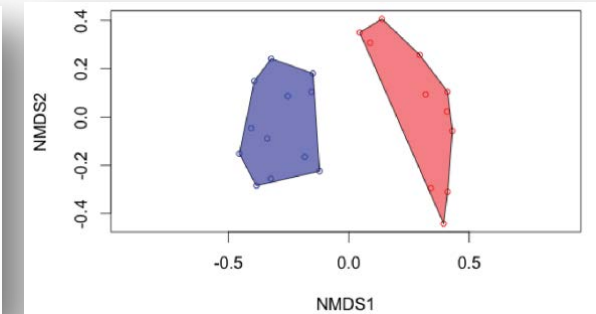
## Large-Scale Biomonitoring of Remote and Threatened Ecosystems via High-Throughput Sequencing

Joel F. Gibson<sup>1,2</sup>, Shadi Shokralla<sup>1</sup>, Colin Curry<sup>3</sup>, Donald J. Baird<sup>3</sup>, Wendy A. Monk<sup>4</sup>, Ian King<sup>1</sup>, Mehrdad Hajibabaei<sup>1\*</sup>

Morphology-based taxonomy

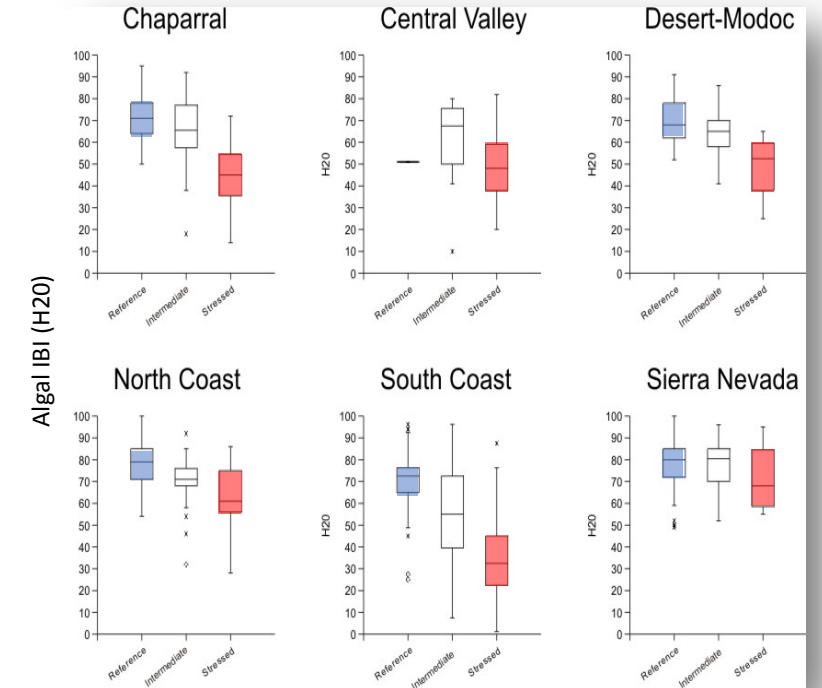


DNA – species level



# Reference screening

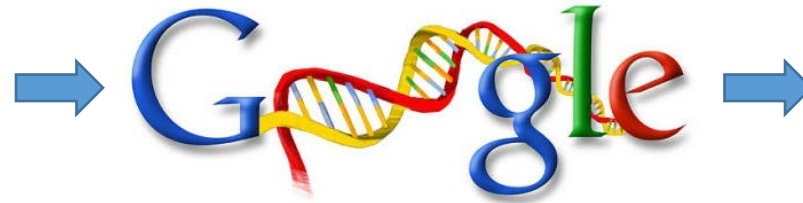
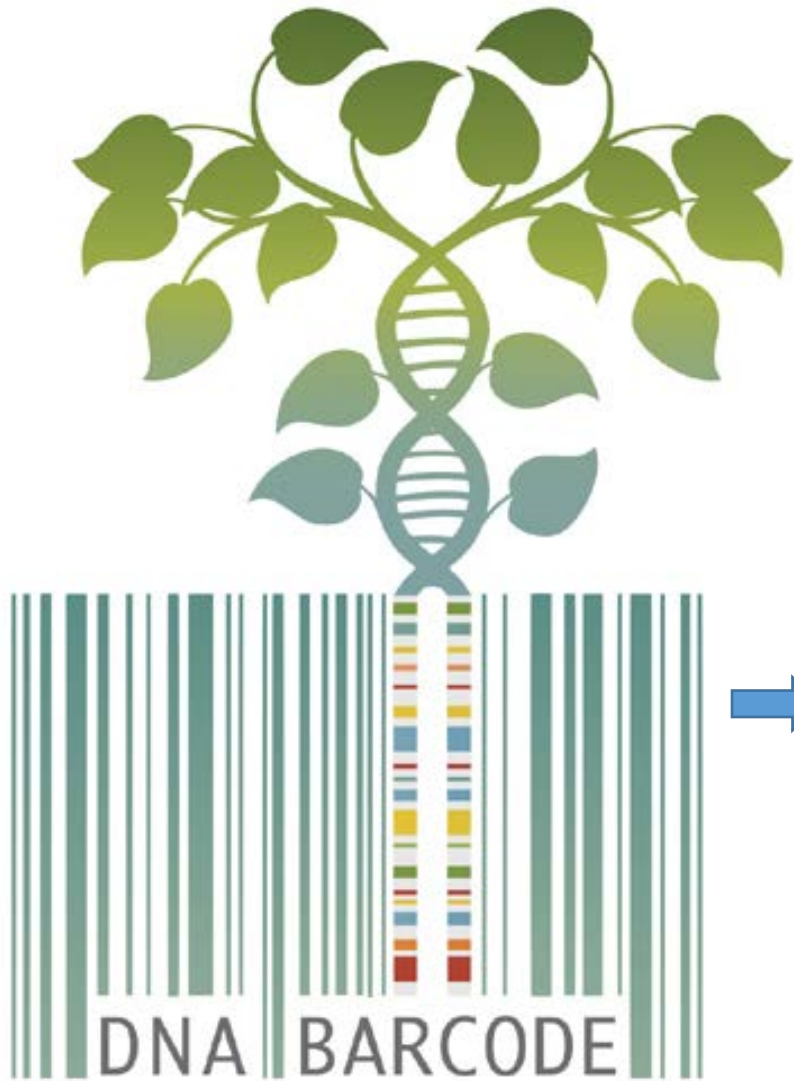
Variable	Scale	Threshold	Unit
% Agriculture	1 km, 5 km, WS	<3	%
% Urban	1 km, 5 km, WS	<3	%
% Ag + % Urban	1 km, 5 km, WS	<5	%
% Code 21	1 km and 5 km WS	<7 <10	%
Road density	1 km, 5 km, WS	<2	km/km <sup>2</sup>
Road crossings	1 km 5 km WS	<5 <10 <50	crossings/ km <sup>2</sup>
Dam distance	WS	<10	km
% Canals and pipelines	WS	<10	%
Instream gravel mines	5 km	<0.1	mines/km
Producer mines	5 km	0	mines
Specific conductance	Site	99/1**	prediction interval
W1_HALL	Sample reach	<1.5	NA



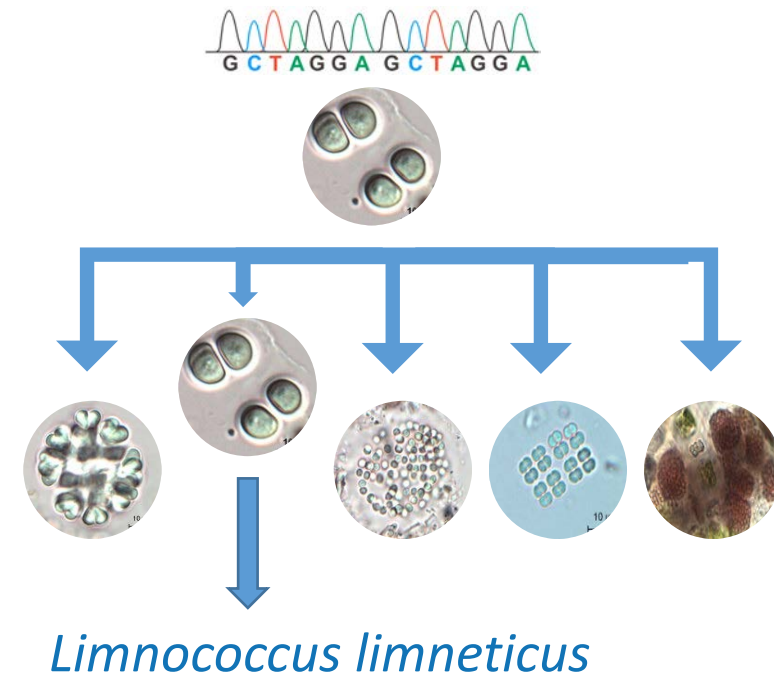
Andy Rehn



# DNA barcoding – what is it?



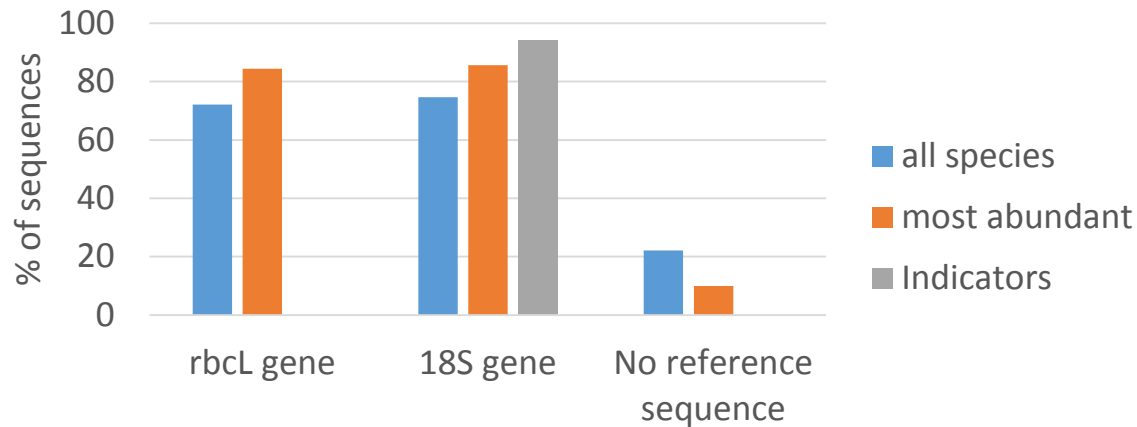
- Sequence one or two DNA regions as a “fingerprint”
- Infer species of organisms by comparing to reference databases



# DNA barcode reference database

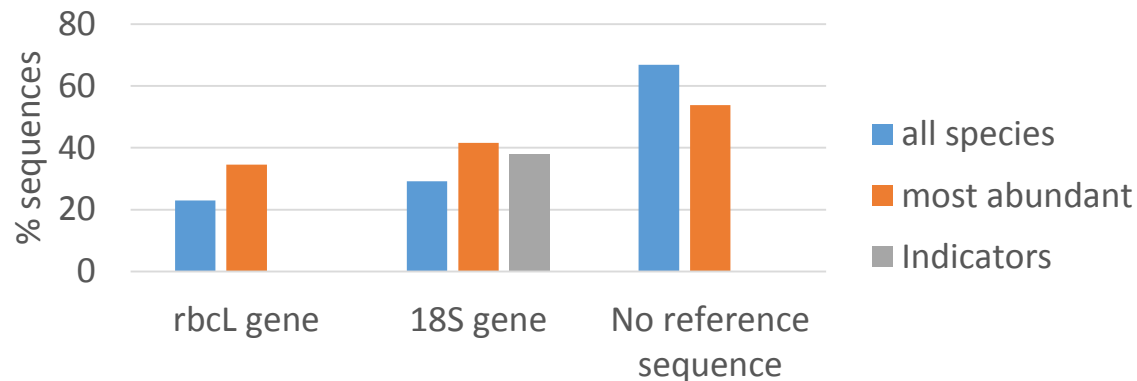
## SMC (2009-2013) Diatoms in Genbank

Genus-level



- 90% of most abundant taxa and 93% of indicator species have a sequence in Genbank database at *genus* level.

Species-level



- 50% of most abundant taxa and 38% of indicator species have a sequence in Genbank database at *species* level.



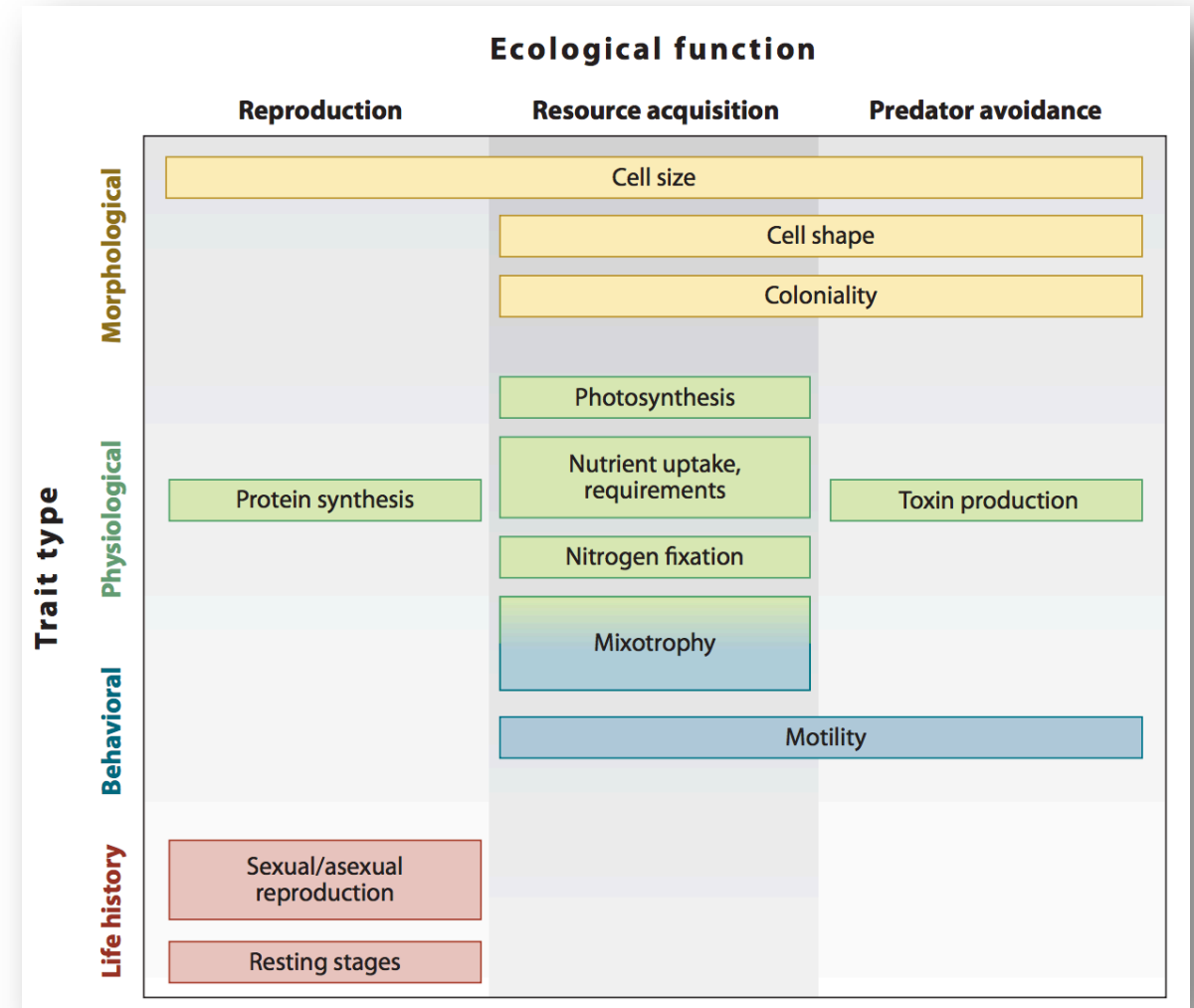
# Statewide Algal Index

## Reference-based approach

Site-specific expectations for each site based on taxa found at groups of similar reference sites

### Candidate traits

1. sedimentation tolerant (incl. highly motile)
2. low-nitrogen indicators (incl. N fixers)
3. halobiontic
4. nitrogen heterotrophs
5. requiring > 50% saturation DO
6. organic-associated
7. copper-associated
8. low-phosphorus



# Statewide Algae Plan – current efforts

- Development of a statewide scoring tool
- Capacity building
  - Development of standard taxonomic effort (STE) for algal species
  - Exploration of molecular methods

