

From oceanography to top predator habitat in TOPP: Synthesis of tag data and modeling

Elliott Hazen, UH-JIMAR / NOAA-SWCSC-PFEL

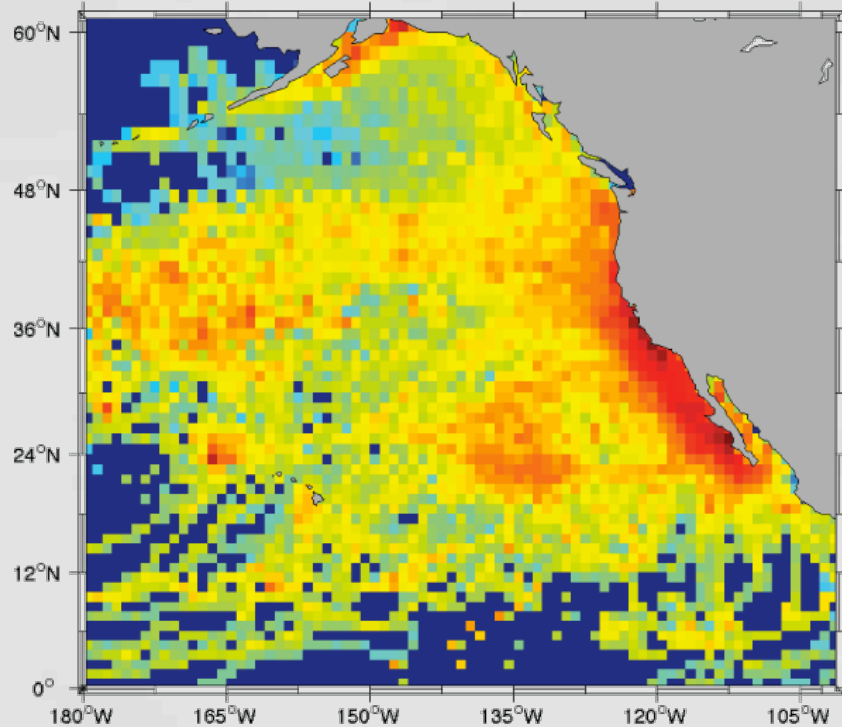
B. A. Block¹, I. D. Jonsen², S. J. Jorgensen¹, A. J. Winship², S. A. Shaffer³, S. J. Bograd⁴, E. L. Hazen⁴, D. G. Foley⁴, G. A. Breed^{2,5}, A.-L. Harrison⁵, J. E. Ganong¹, A. Swithenbank¹, M. Castleton¹, H. Dewar⁶, B. R. Mate⁷, G. L. Shillinger¹, K. M. Schaefer⁸, S. R. Benson⁹, M. J. Weise⁵, R. W. Henry⁵ & D. P. Costa⁵

Outline of Presentation

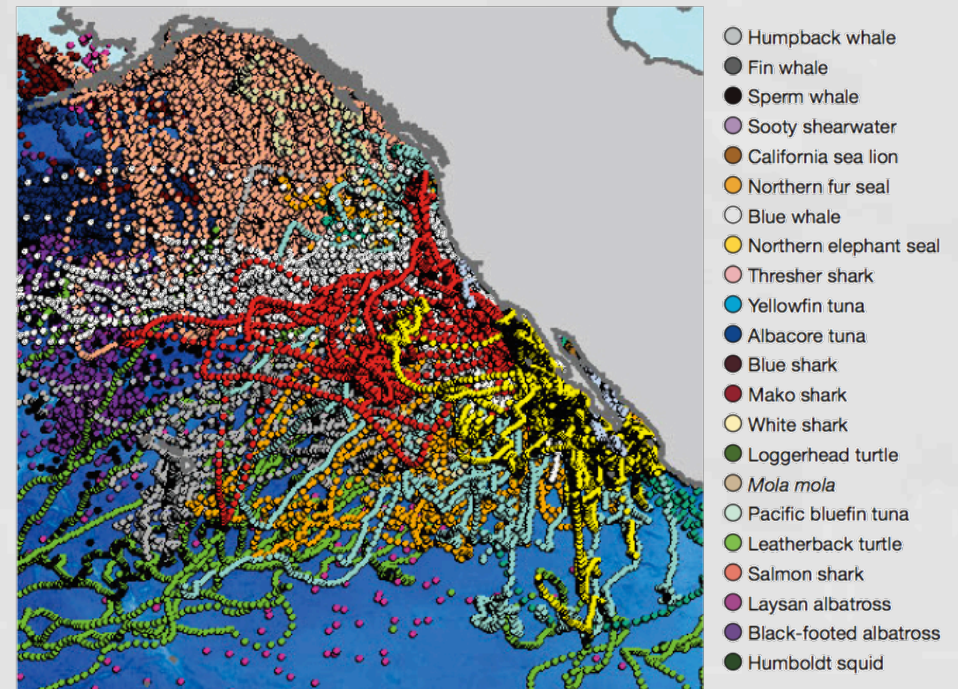
- Introduction to the data
- Exploratory analyses & environmental sampling
- Choosing & running statistical models
- Where to go from here?

TOPP dataset

ALL SPECIES NORMALIZED LOG SCALE



All Species All Positions



- From 2000-2010: 23 species; 4,300 tags; >1.1 Million profiles
- Tracking, conservation, ocean observation

Goals and Questions

- How do TOPP species use the ocean?
- Are there predictable variables that correspond to important habitat?
- How do seasonal patterns in the variables correspond to seasonal changes in top predator habitat use?

Dataset challenges

- Tag data are highly correlated spatially and temporally
 - Include spatial autocorrelation explicitly – correlation structure in the error term
 - Bin data at scales broader than correlation (e.g. 1° & ~90 days)
- Long time series can span multiple satellites e.g. SEAWiFS and MODIS for ocean color
- > 1.1 million points

Relative use maps

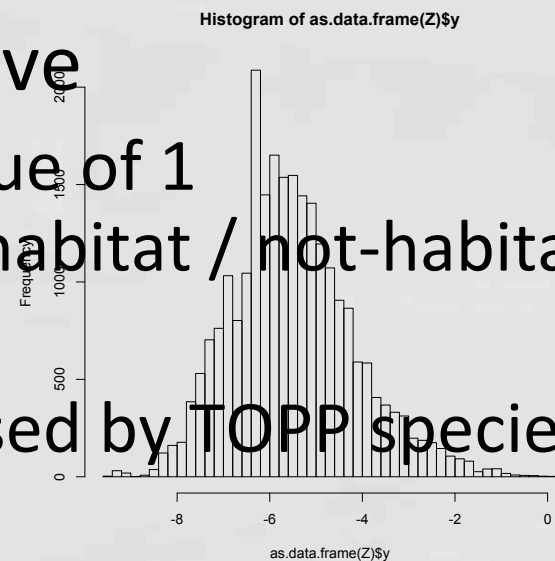
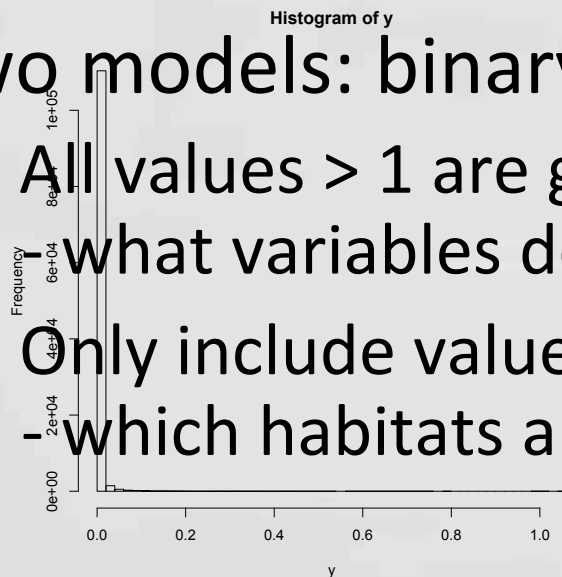
- How do you convert tag data from multiple species & tag types into useful products?
 - State space models to account for multiple tag types and corresponding positional error
 - Weight track by length - longer tracks get higher weighting to minimize tag bias
 - Normalize by species – 1000 elephant seal tracks = 100 blue whale tracks
 - Relative use $\sim f(\# \text{ of species, } \# \text{ of individuals})$

TOPP modeling framework

- R for statistics (mgcv, ape, spdep, ncf, AED)
- 5 environmental variables available for entire tracking dataset and spatial extent
- Binned data 1° & quarterly from 2000-2010

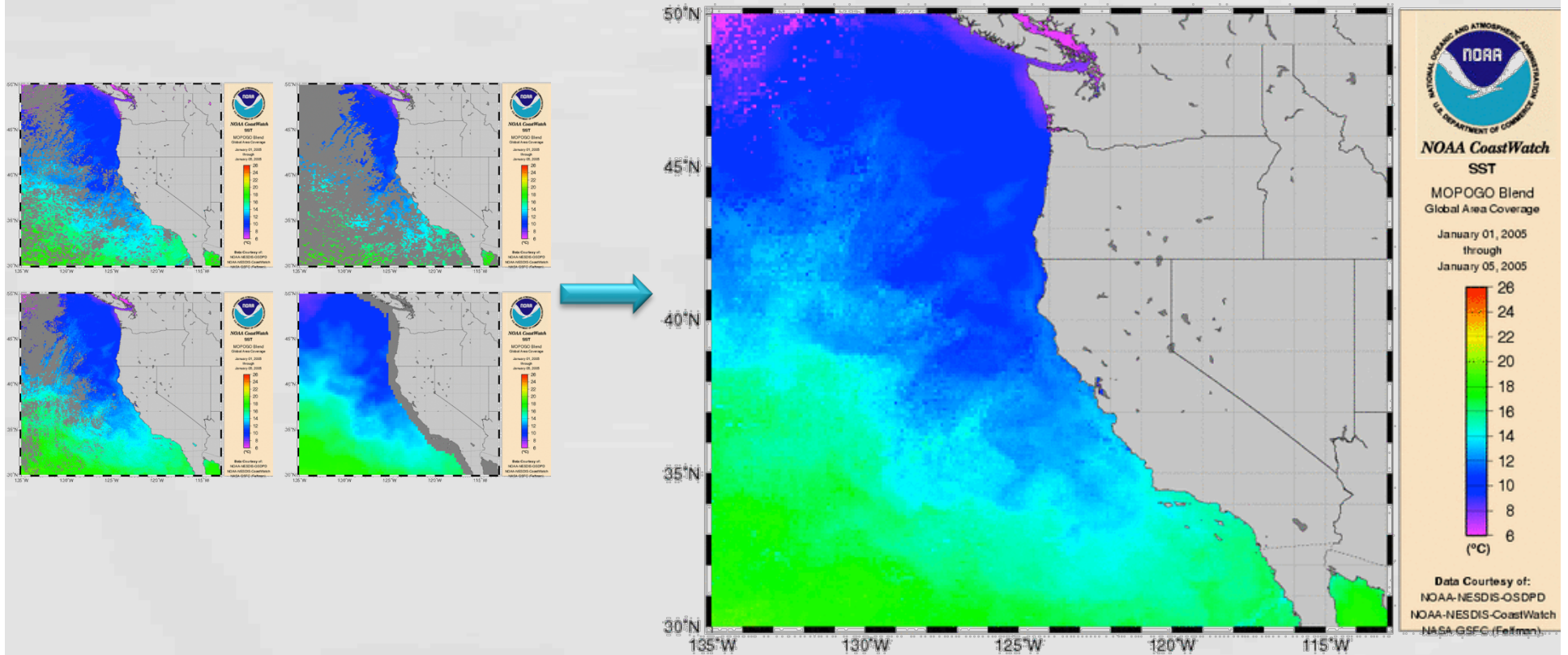
- Two models: binary & positive

- All values > 1 are given a value of 1
- what variables determine habitat / not-habitat
- Only include values > 1
- which habitats are more used by TOPP species



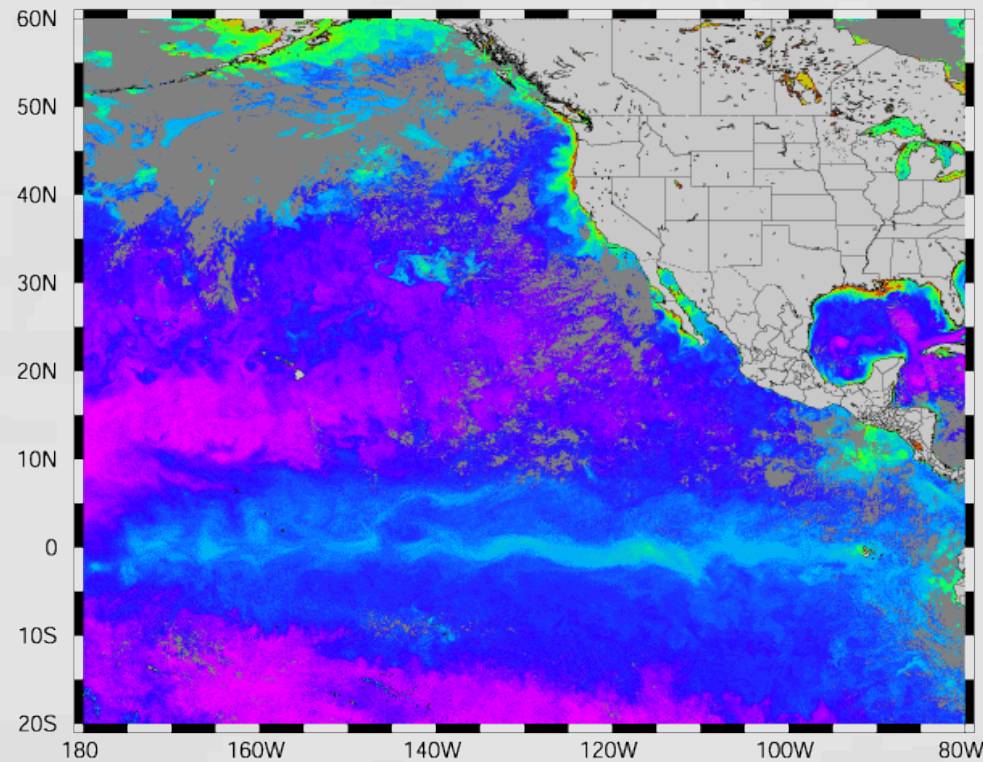
Environmental data

- Blended SST – AVHRR & GOES



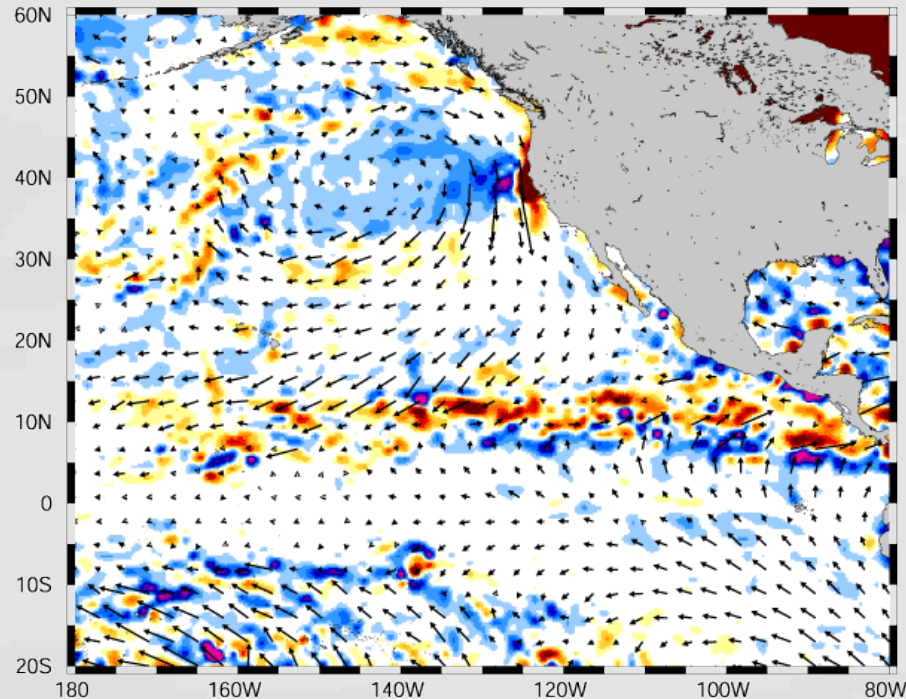
Environmental data

- Blended SST – AVHRR & GOES
- Chl – SeaWiFS (1997-2009) & MODIS (2002-)



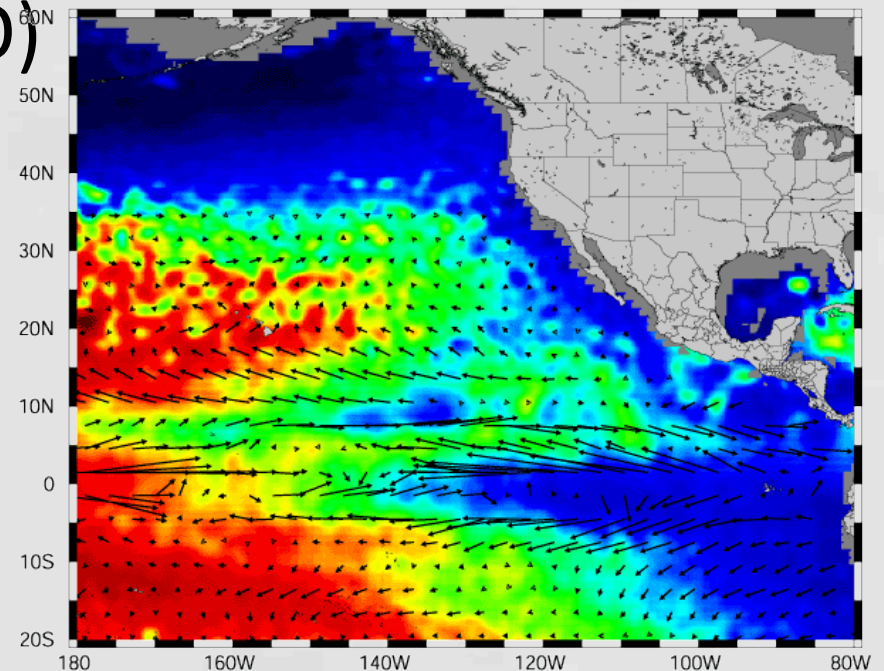
Environmental data

- Blended SST – AVHRR & GOES
- Chl – SeaWiFS (1997-2010) & MODIS (2002-)
- Wind stress curl (QuikSCAT, 1999-2009)



Environmental data

- Blended SST – AVHRR & GOES
- Chl – SeaWiFS (1997-2010) & MODIS (2002-)
- Wind stress curl (QuikSCAT)
- SSHa and SSHrms (AVISO)

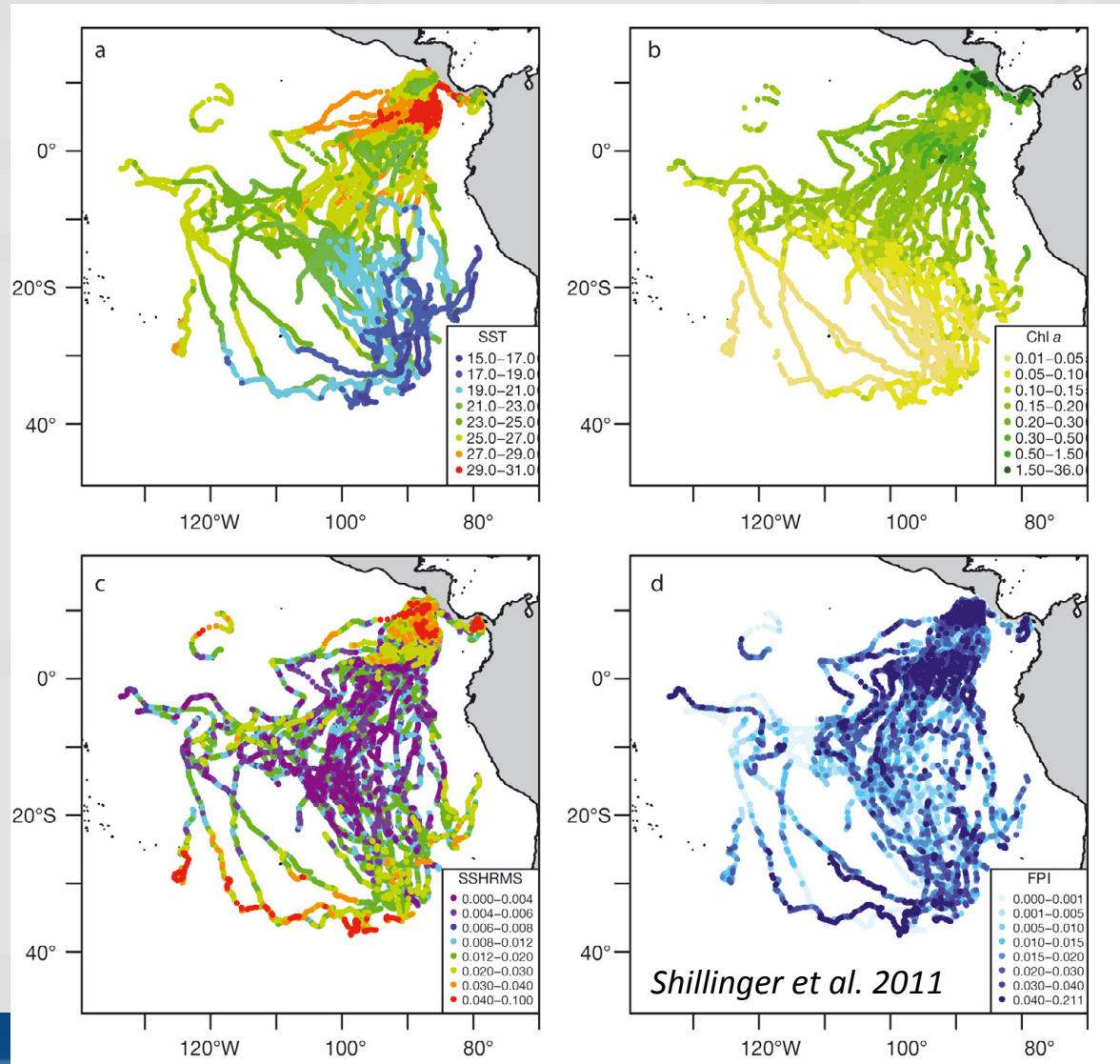


Sampling RS environmental data

- Near Real Time
 - Live access server
las.pfel.noaa.gov/oceanwatch.html
 - New and improved browser
*coastwatch.pfel.noaa.gov/coastwatch/
CWBrowser.jsp*
 - In ARCGIS – Environmental Data Connector
http://www.pfeg.noaa.gov/products/edc/
- For tagging or other similar point data
 - Xtractomatic – in R and Matlab
http://coastwatch.pfel.noaa.gov/xtracto/

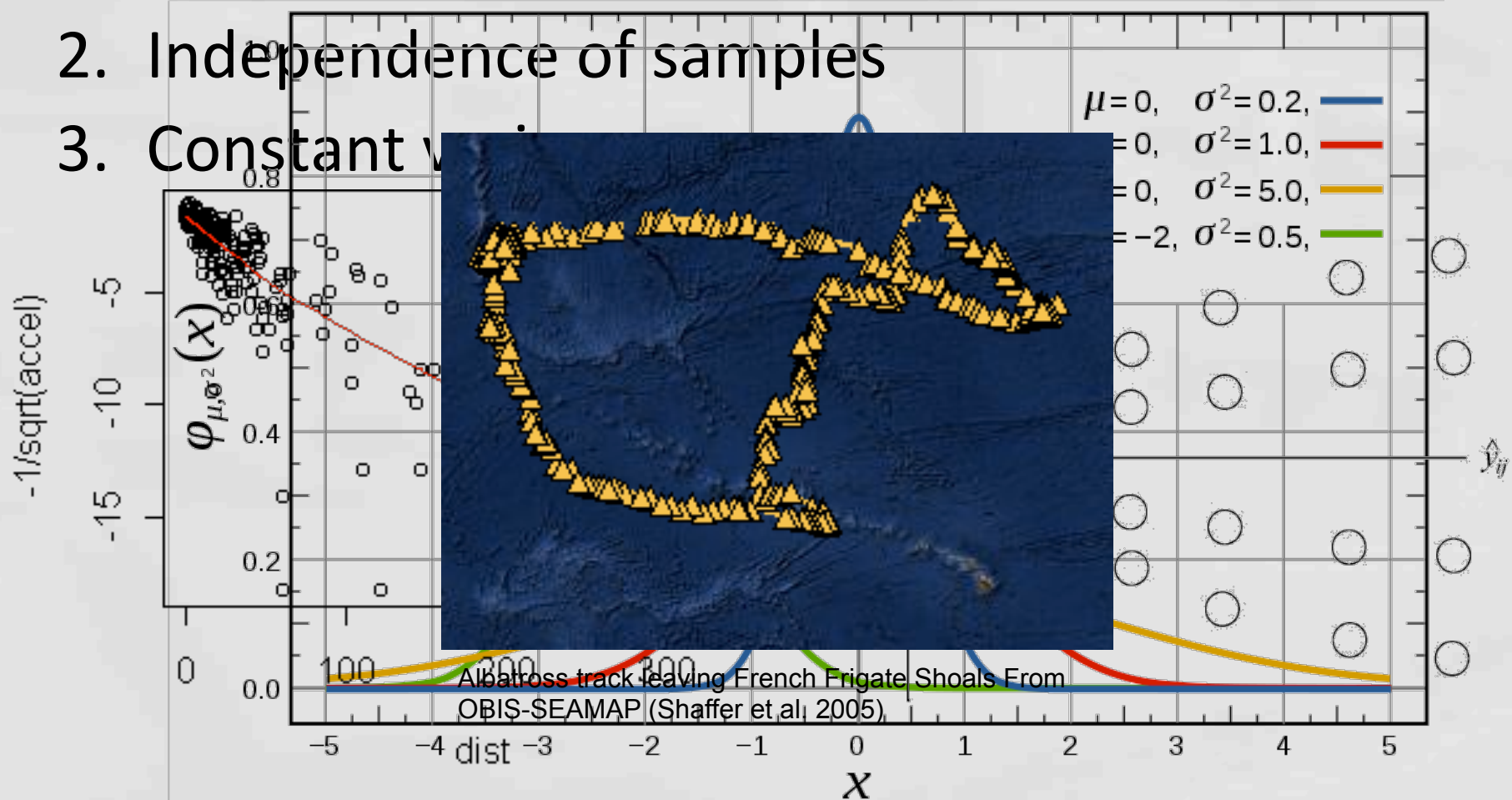
For more info: <http://www.pfel.noaa.gov/events/workshops/NOAASatCourse2012/>

Sampling RS environmental data



Common modeling assumptions

1. Normally distributed data
2. Independence of samples
3. Constant variance



Spatial autocorrelation

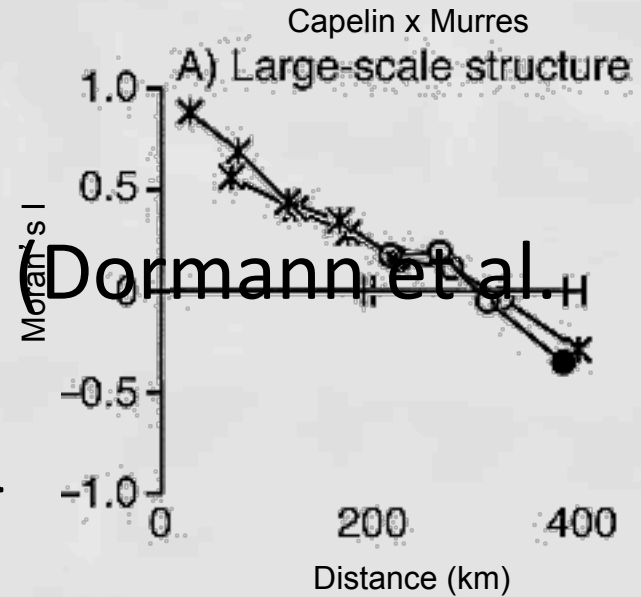
- Why do we care?
- Spatial autocorrelation can bias model signif. (see Segurado et al. 2006)
- Spatial autocorrelation is a natural component of spatial data
 - For model residuals, there may be unmeasured spatial variables affecting data



Albatross track leaving French Frigate Shoals
From OBIS-SEAMAP (Shaffer et al. 2005)

Spatial autocorrelation

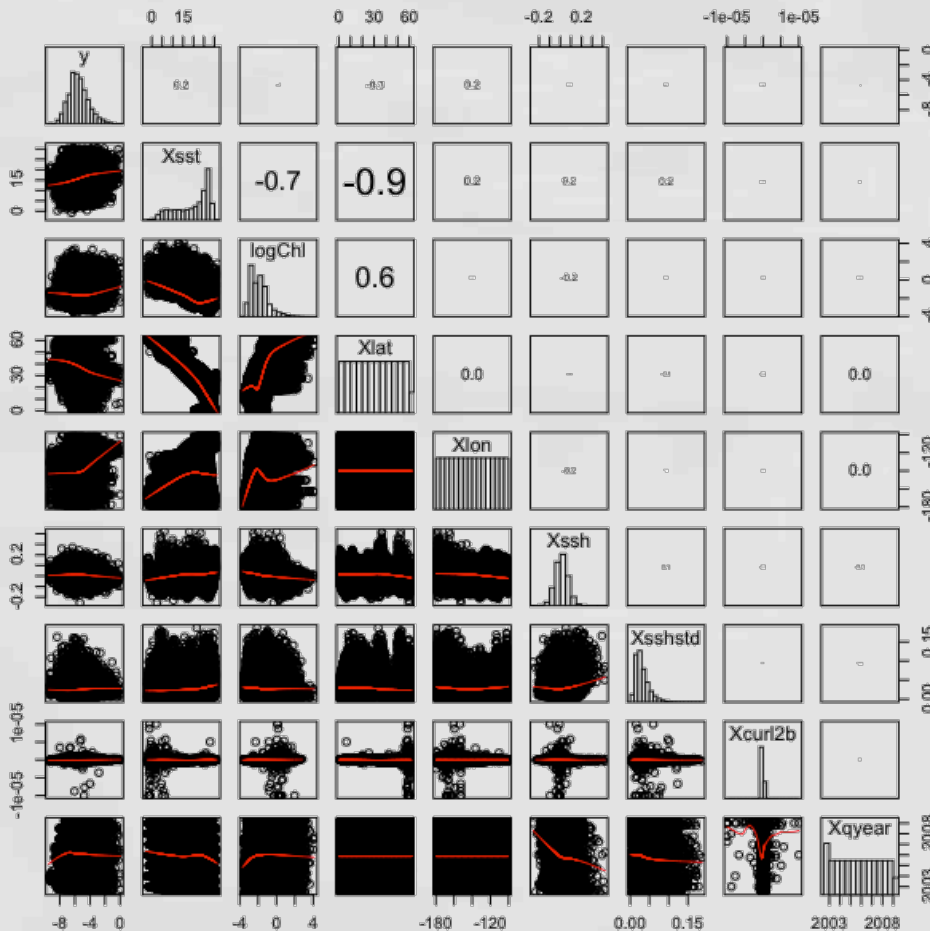
- Ways to test for it
 - Geary's C (0 to 2)
 - Moran's I (-1 to 1)
- Many methods to model it (Dormann et al. 2007)
 - Autocovariate regression & spatial eigenvectors
 - Generalized least squares (GLS), GLMMs, GEEs
 - Partial Mantel tests (Legendre and Legendre 1998)



From Fauchald et al 2000

TOPP - Exploratory Analysis

- Examined cross-correlations & normalcy



(Zuur et al. 2009)

- Chl was log transformed
- other variables were not improved via transformation

Types of statistical models

- There are many, and constantly changing / growing
- Correlation/Regression techniques – GLMs, GAMs (Austin 2002), Mixed models (Wood 2006), regression trees & random forests (Breiman 2001)
- Ordination –Multivariate dimensional scaling, e.g. CCAs (Guisan et al. 1999)
- Maximum Entropy models – species distributions “closest to uniform” (Phillips et al. 2006)
- Recent reviews of modeling approaches (Redfern et al. 2006, Elith et al. 2006, Dormann et al. 2007, Aarts et al. 2008, Elith et al. 2009)

Generalized Additive Models

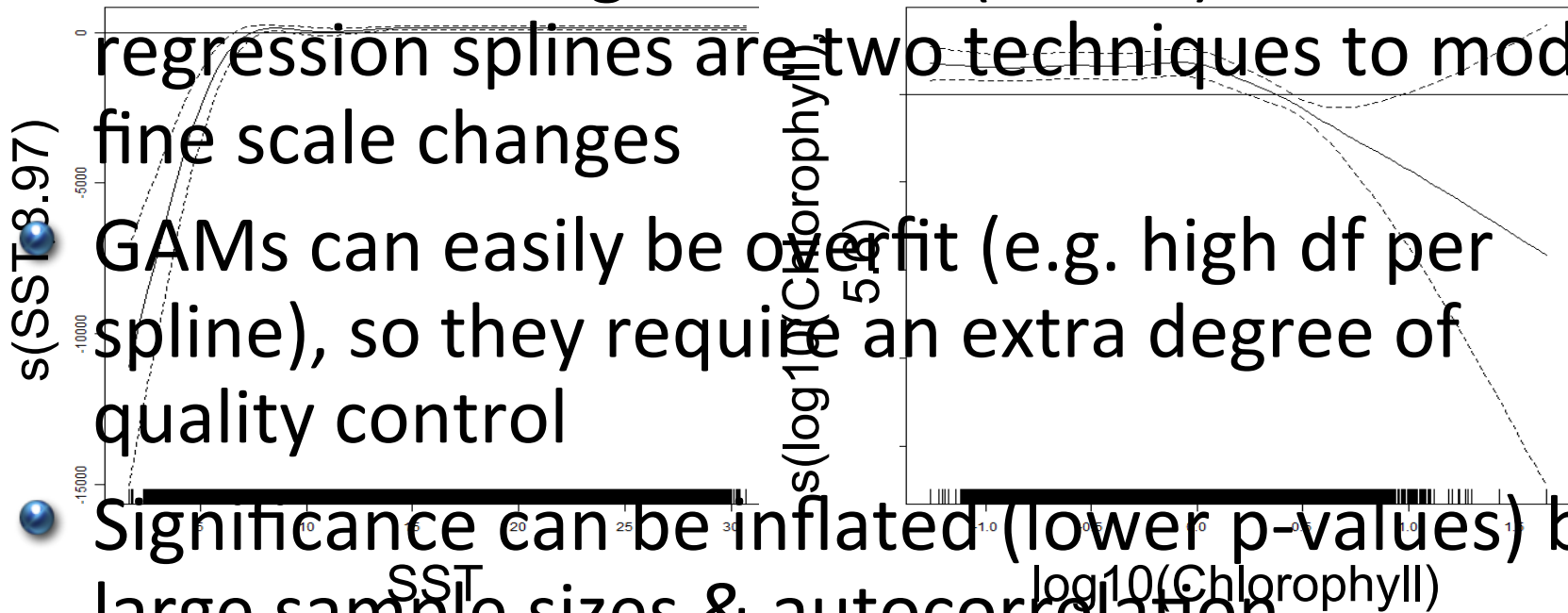
- GAMs can use a combination of parametric and non-parametric functions:

$$(y \sim A + f(x_1) + f(x_2) \dots + f(x_n) + \epsilon)$$

- Local smoothing functions (LOESS) and regression splines are two techniques to model fine scale changes

- GAMs can easily be overfit (e.g. high df per spline), so they require an extra degree of quality control

- Significance can be inflated (lower p-values) by large sample sizes & autocorrelation

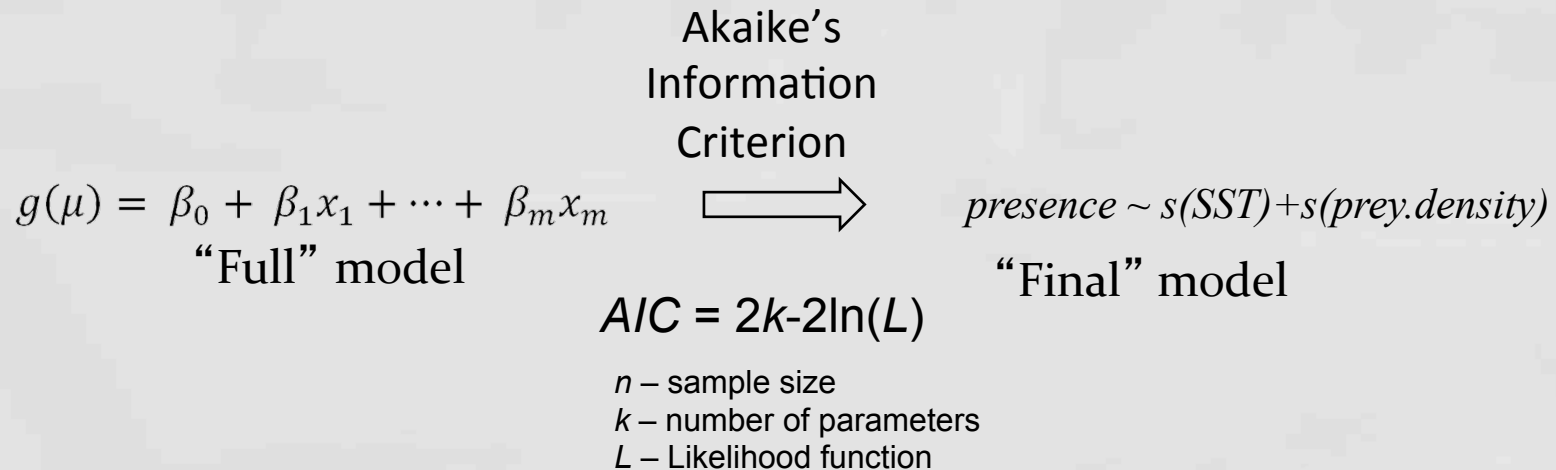


TOPP GAMM Models

- Full models

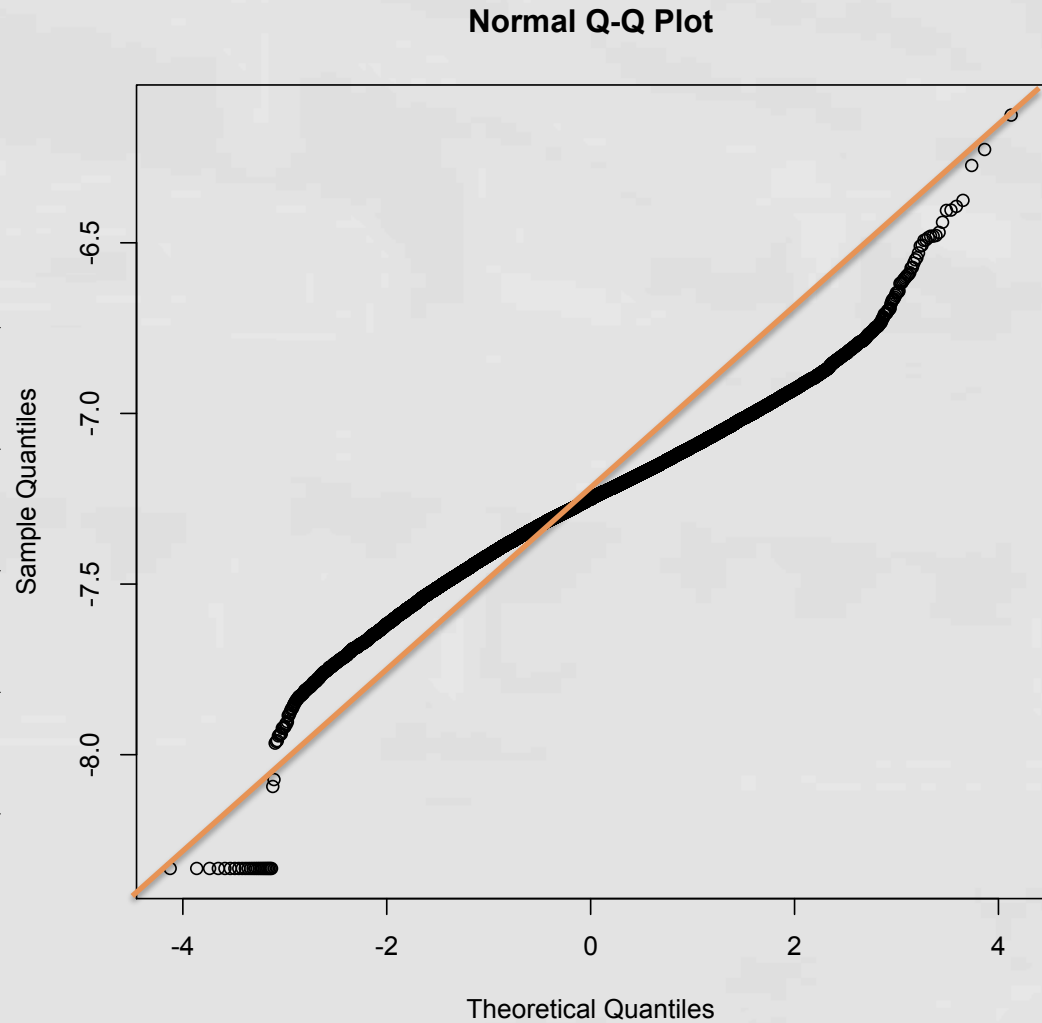
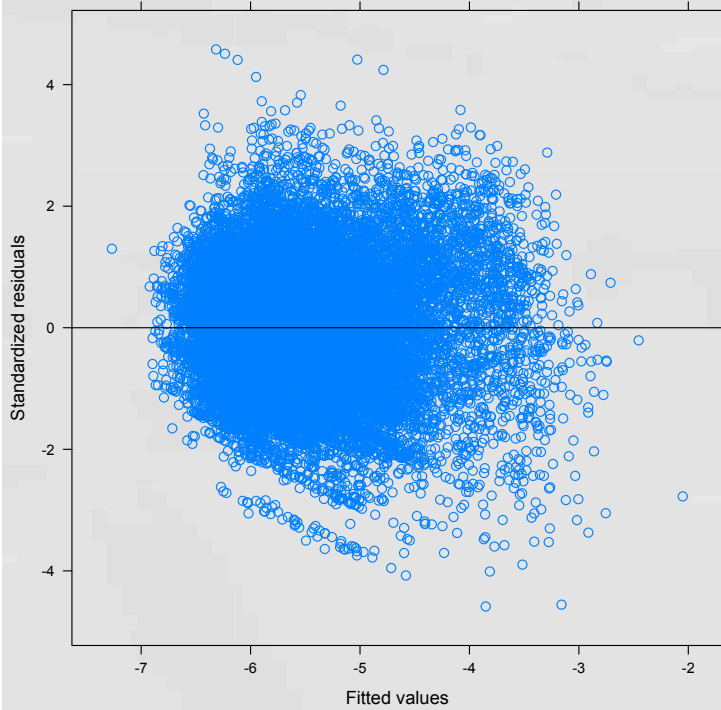
- RelUse ~ SST + log(chl) + SSH + SSHrms + Curl + Lat + Lon, family = gaussian, random = quarter+year
- PresAbs ~ SST + log(chl) + SSH + SSHrms + Curl + Lat + Lon, family = binomial, random = quarter+year

- Model selection:



TOPP GAMM residuals

- Normalcy tests



Final GAMM model results

Supplementary Table 7: Summary statistics of fitted GAMMs. Morans I statistic was calculated from the residuals of the fitted models and can range from -1 to 1 with values > 0 indicating spatial clustering and values < 0 indicating dispersion. Morans I statistic p -values indicate that there was residual but non-significant spatial clustering in the presence/absence model and in the relative density model.

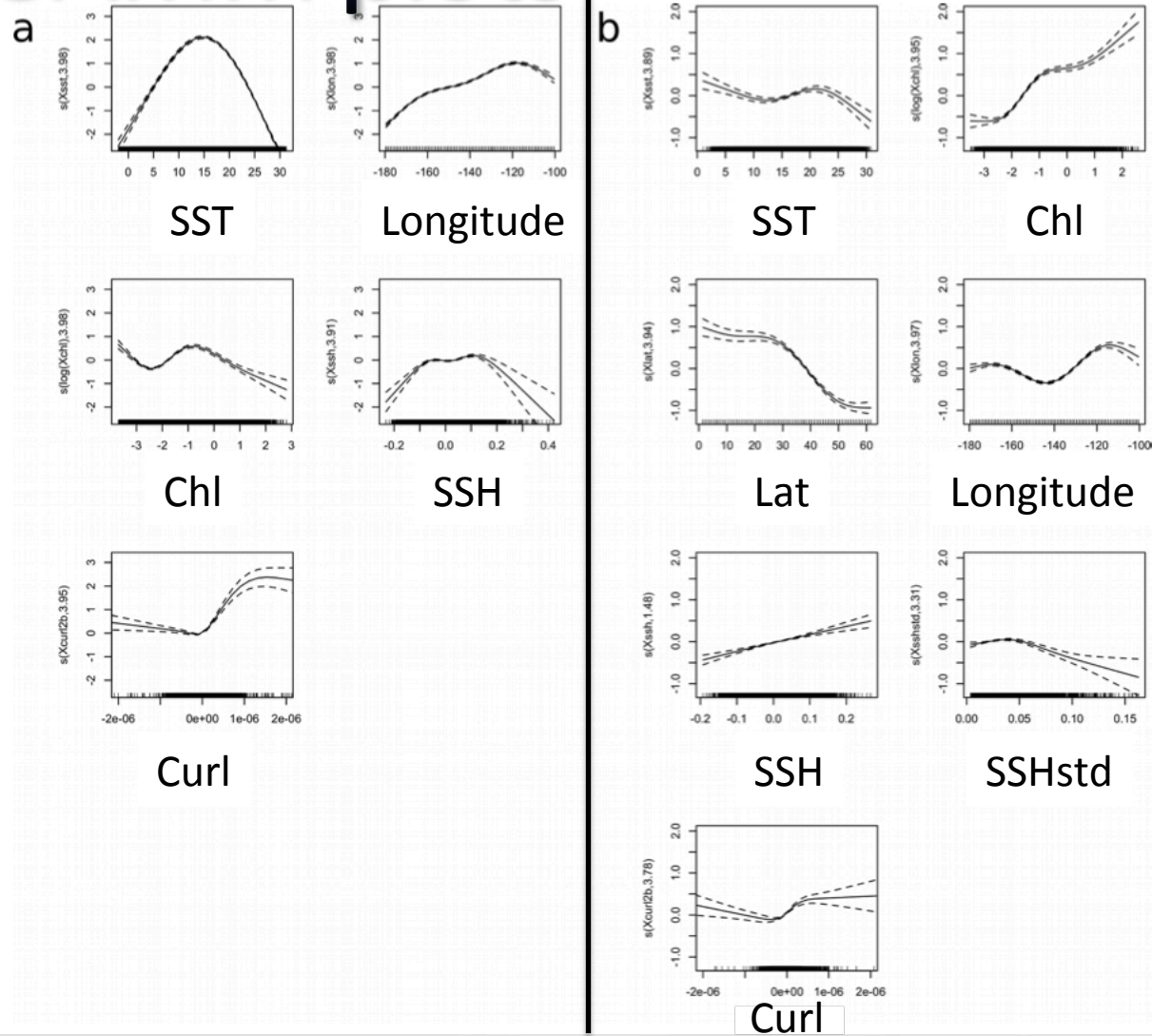
	Presence/absence	Relative density
Proportion of deviance explained	0.303	0.232
AIC	598375	74780.45
Moran's I statistic	0.013	0.050
Moran's I p -value	0.053	0.062

Final GAMM parameters

Supplementary Table 8: Model variables in final fitted presence/absence and relative density GAMMs.

Explanatory variable	Estimated DF	F	p -value
<i>Presence/absence</i>			
s(Xsst)	3.986	2506.99	< 0.001
s(log(Xchl))	3.945	1077.67	< 0.001
s(Xlon)	3.979	131.35	< 0.001
s(Xssh)	3.867	20.3	< 0.001
s(Xcurl2b)	3.935	58.6	< 0.001
<i>Relative density</i>			
s(Xsst)	3.90	31.01	< 0.001
s(log(Xchl))	3.95	226.84	< 0.001
s(Xlat)	3.94	122.57	< 0.001
s(Xlon)	3.97	147.62	< 0.001
s(Xssh)	1.48	72.25	< 0.001
s(Xsshstd)	3.31	20.11	< 0.001
s(Xcurl2b)	3.78	26.63	< 0.001

Final GAMM plots

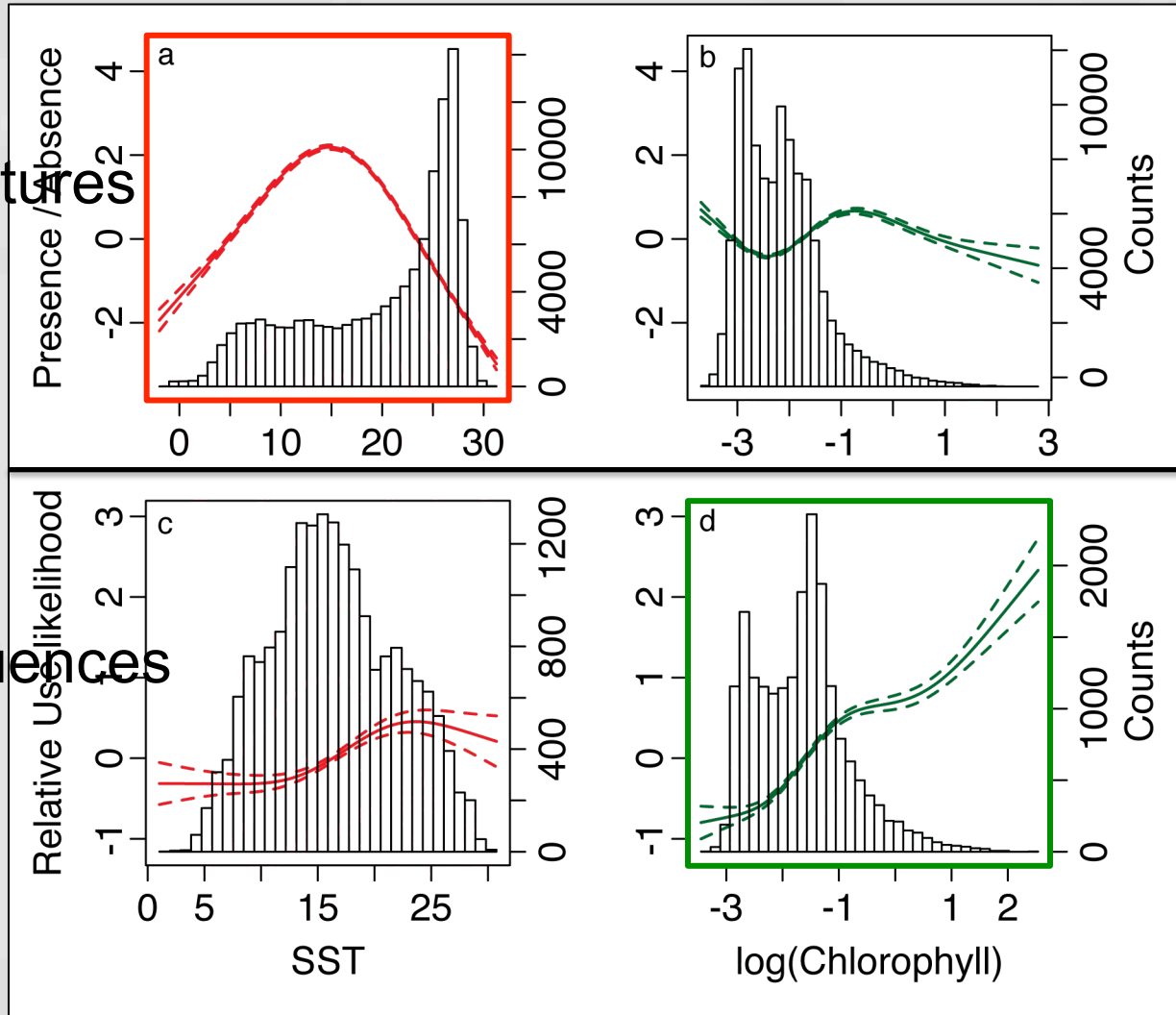


Presence /
Absence

Relative
Density

TOPP GAMM model

- SST structures habitat



- Chl-a influences use

Aknowledgements

