



PRIMER-e
empowering research



Modelling species' responses to climate change using a flexible non-linear Bayesian approach

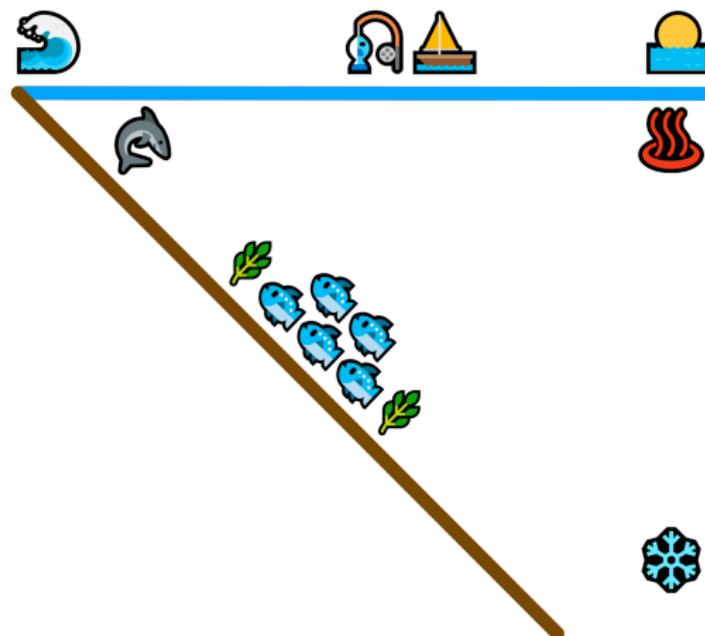
Hayden Rabel - Massey University & PRIMER-e

June 27, 2022 - ISBA World Meeting Montréal, Canada

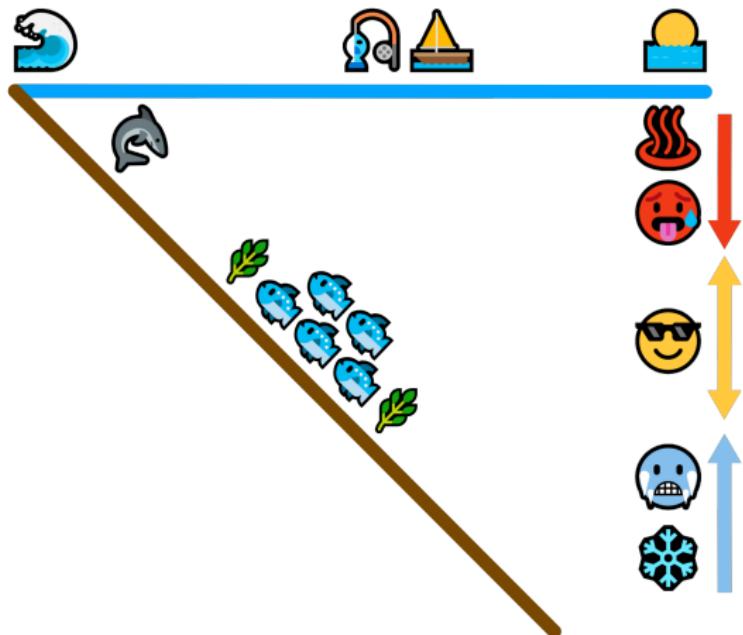
New Zealand
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 MASSEY UNIVERSITY
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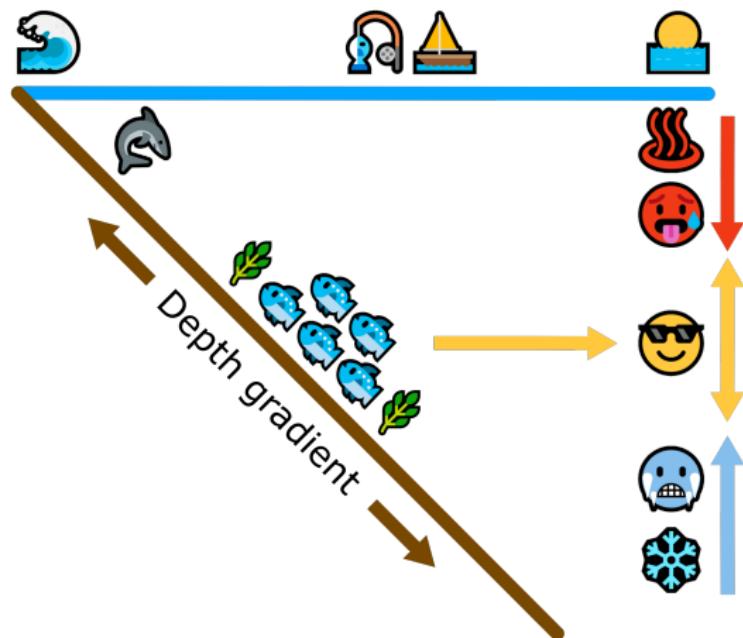
Species' responses



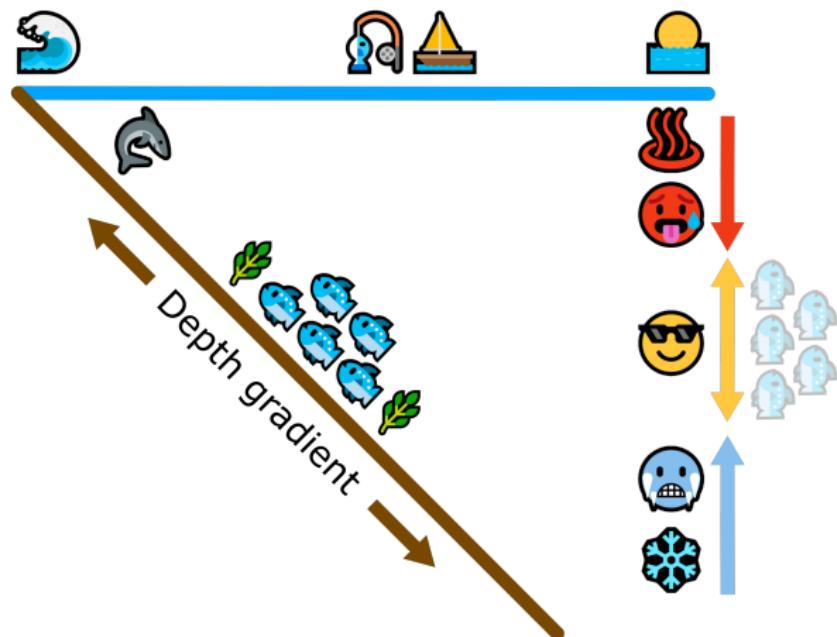
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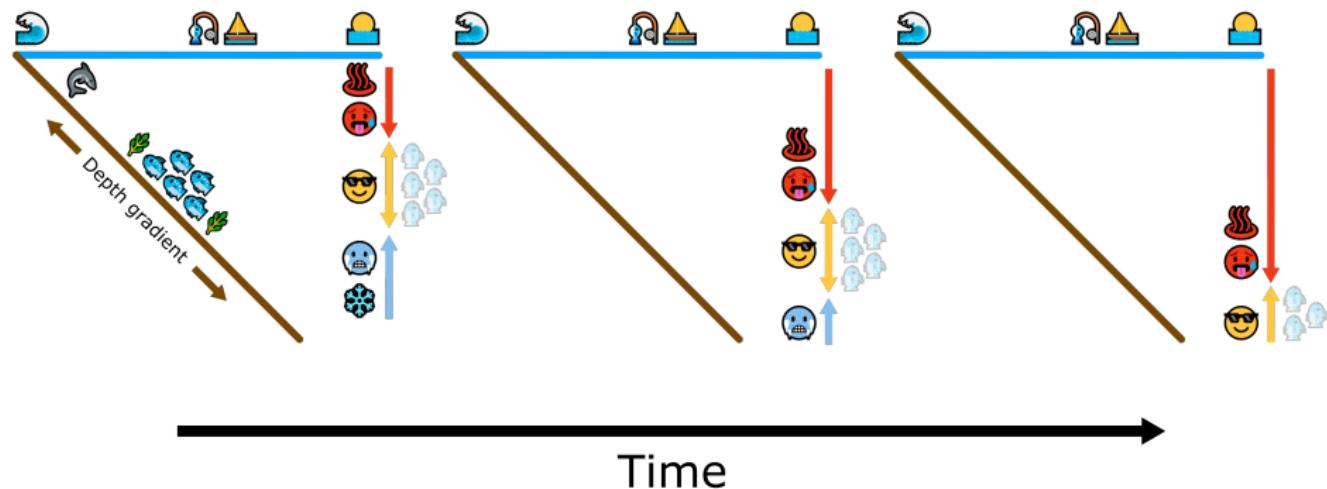
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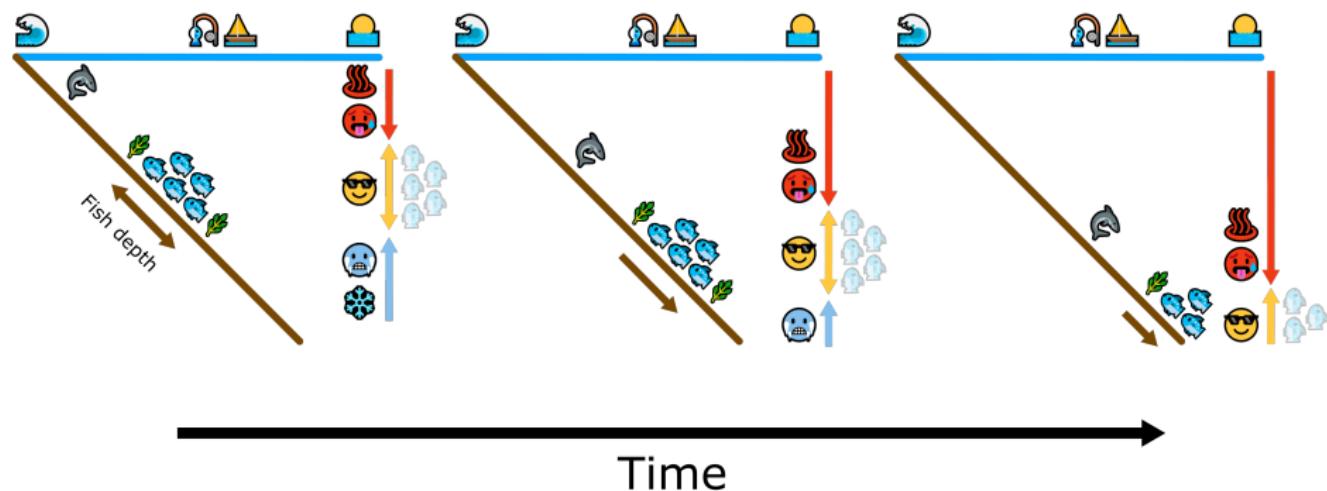
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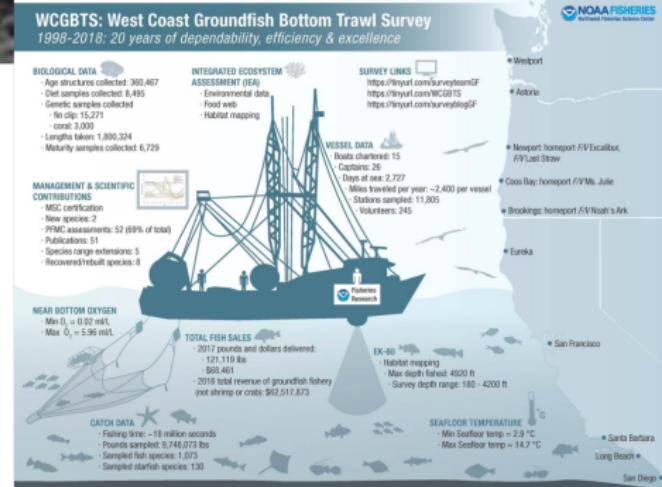
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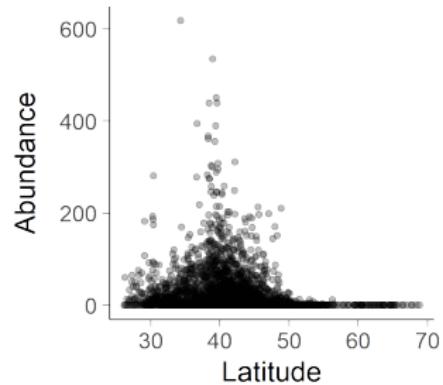


We need broad-scale data

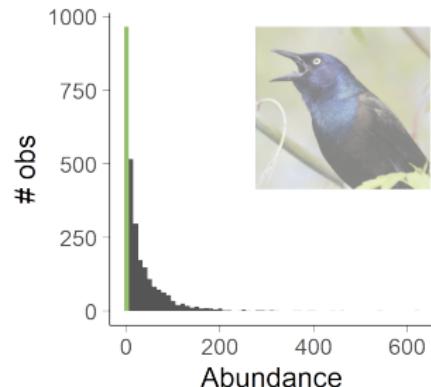


Species distributions along gradients

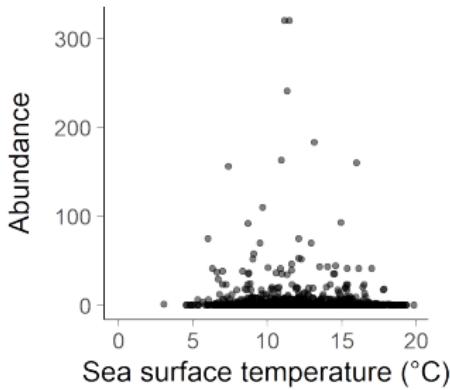
Quiscalus quiscula



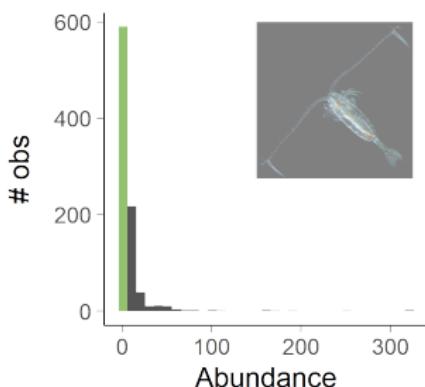
35% zero + bins of 10



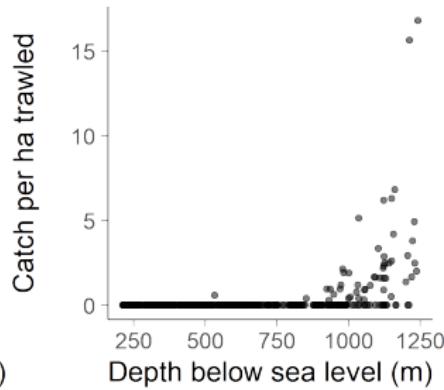
Calanus finmarchicus



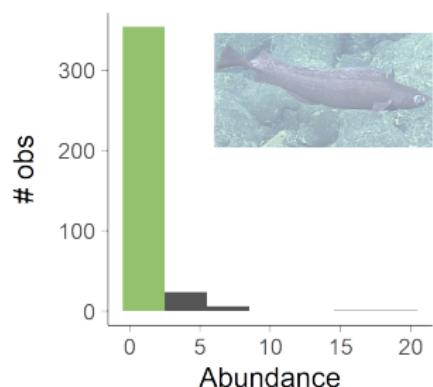
66% zero + bins of 10



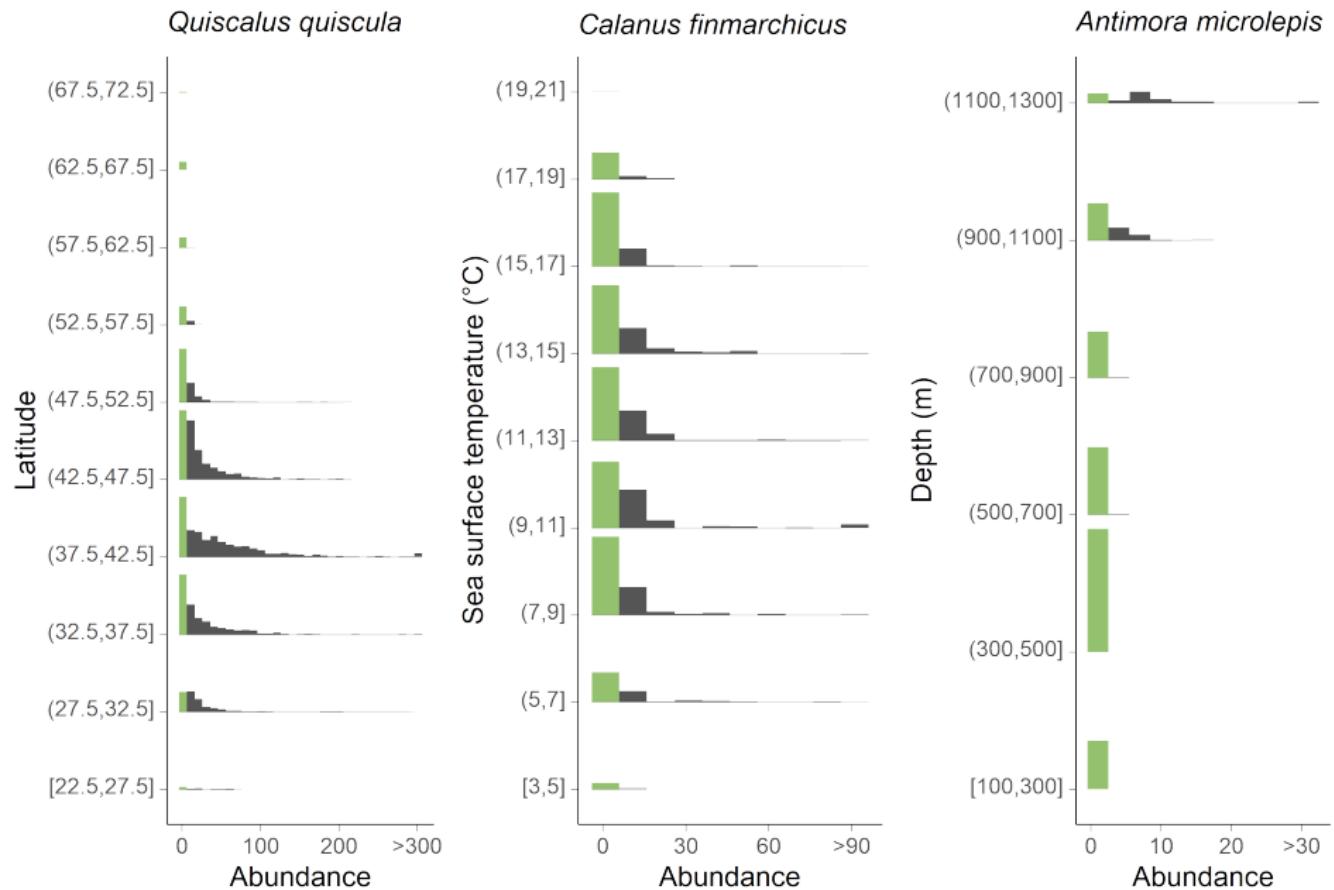
Antimora microlepis



87.6% zero + bins of 5



Species distributions along gradients



Some distributions for count data

Variance = mean:

$$y_i \sim \text{Poisson}(\eta_i = \lambda_i, \theta = \{\})$$

$$\mathbb{E}[y_i] = \eta_i$$

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Zero Inflation (constant prob. excess zeros):

$$y_i \sim \text{ZID}(\eta_i, \theta, \pi) \quad E[y_i] = (1 - \pi)\eta_i$$

$$= \begin{cases} \pi + (1 - \pi) \times D(0, \theta), & \text{if } y_i = 0, \\ (1 - \pi) \times D(\eta_i, \theta), & \text{if } y_n > 0 \end{cases}$$

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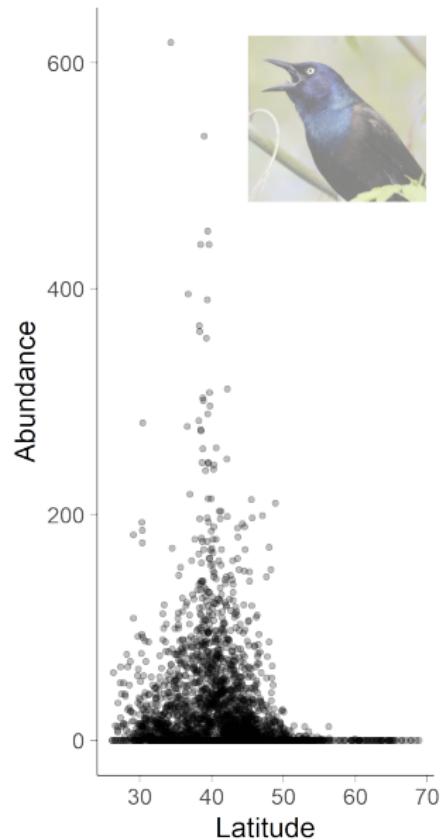
ZI (prob. excess zeros Linked to η):

$$y_i \sim \text{ZIDL}(\eta_i, \theta, \pi_i, \gamma_0, \gamma_1) \quad E[y_i] = (1 - \pi)\eta_i$$

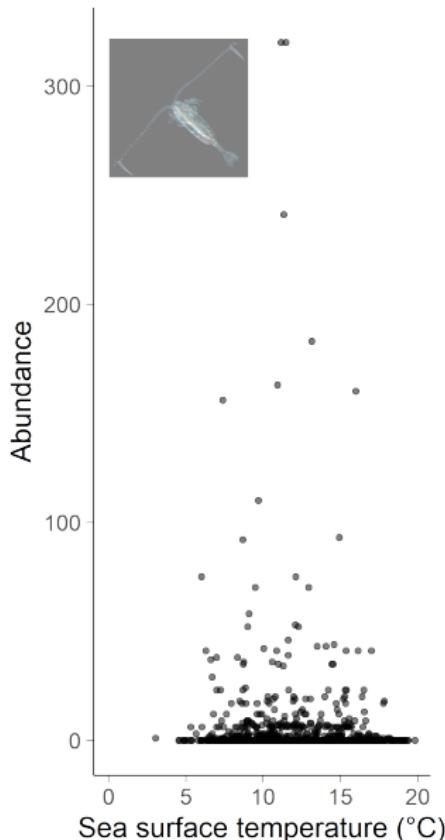
$$\text{logit } \pi_i = \gamma_0 - \gamma_1 \log(\eta_i)$$

Species-environment relationships aren't quite linear

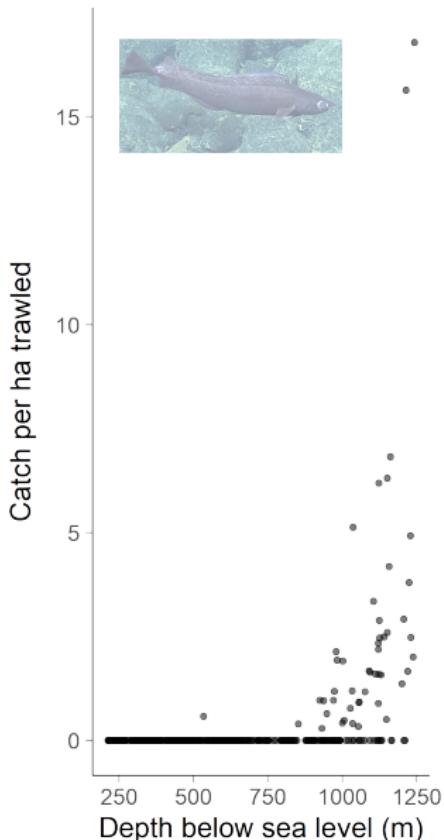
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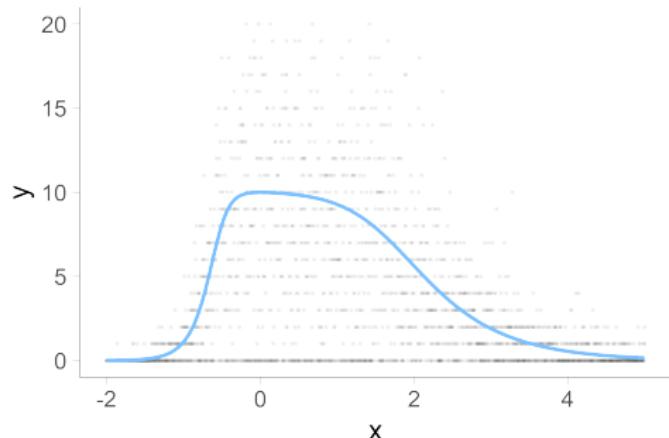


Antimora microlepis



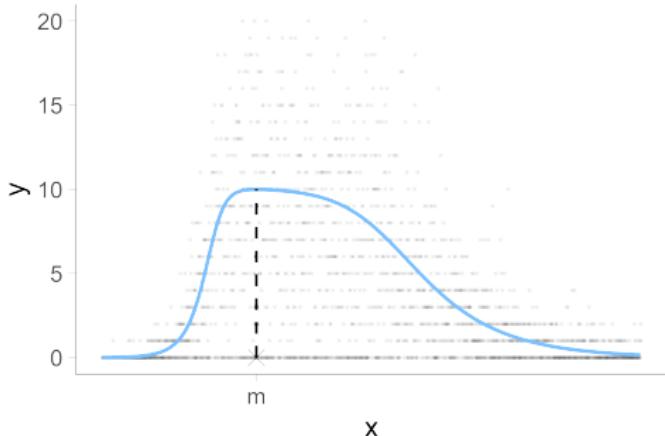
A non-linear predictor

- ▶ $\eta_i = f(x_i)$



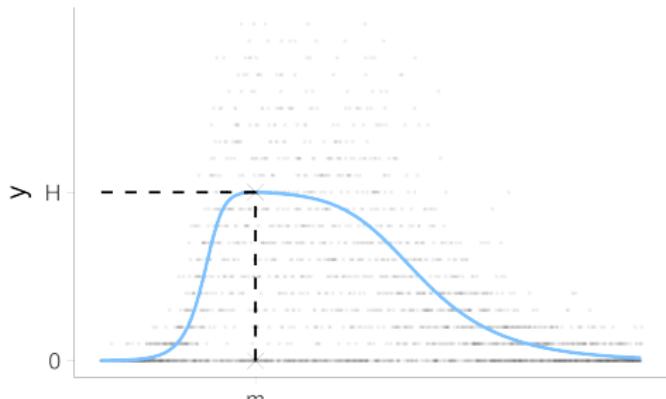
A non-linear predictor

- ▶ $\eta_i = f(x_i, m)$
- ▶ $m \in \mathbb{R}$: **mode**, optimum x



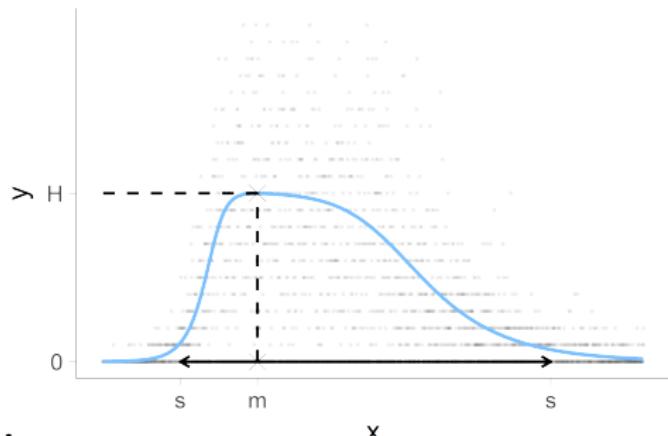
A non-linear predictor

- ▶ $\eta_i = f(x_i, H, m)$
- ▶ $m \in \mathbb{R}$: **mode**, optimum x
- ▶ $H \in \mathbb{R}_{>0}$: maximum **Height**



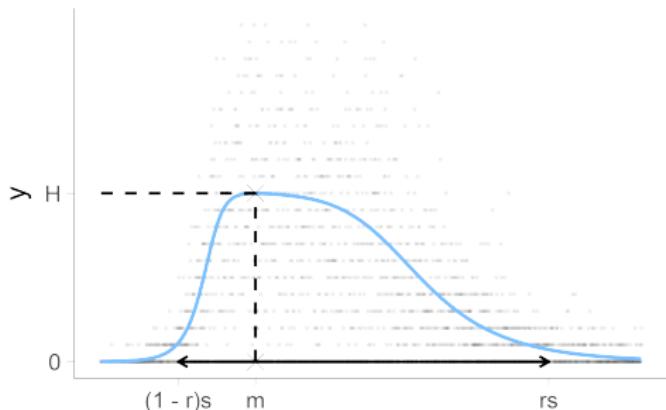
A non-linear predictor

- ▶ $\eta_i = f(x_i, H, m, s)$
- ▶ $m \in \mathbb{R}$: **mode**, optimum x
- ▶ $H \in \mathbb{R}_{>0}$: maximum **Height**
- ▶ $s \in \mathbb{R}_{>0}$: **spread**, tolerable range



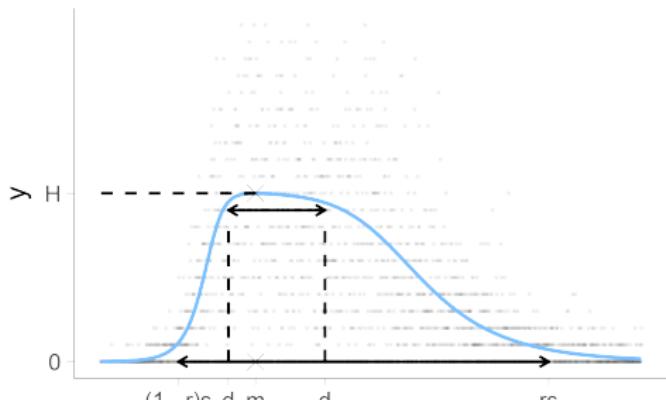
A non-linear predictor

- ▶ $\eta_i = f(x_i, H, m, s, r)$
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- ▶ $s \in \mathbb{R}_{>0}$: **spread**, tolerable range
- ▶ $r \in (0, 1)$: **asymmetry**



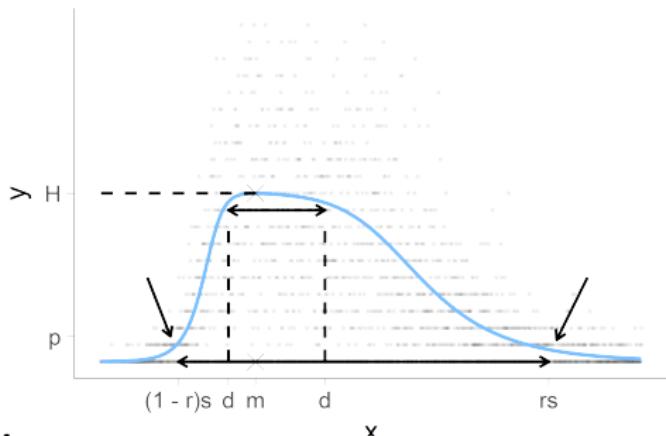
A non-linear predictor

- ▶ $\eta_i = f(x_i, H, m, s, r, d)$
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- ▶ $d \in \mathbb{R}$: **broadness** or peakedness around the optimum



A non-linear predictor

- ▶ $\eta_i = f(x_i, H, m, s, r, d, p)$
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- ▶ $s \in \mathbb{R}_{>0}$: **spread**, tolerable range
- ▶ $r \in (0, 1)$: **asymmetry**
- ▶ $d \in \mathbb{R}$: **broadness** or peakedness around the optimum
- ▶ $p \in \mathbb{R}_{>0}$: **exaggeration** or pinching of tails



The modskurt family

$$\eta_i = f(x_i)$$

$$f(x_i|\psi) = H \left(r e^{\frac{1}{p}(\frac{z_i}{r}-d)} + (1-r)e^{-\frac{1}{p}(\frac{z_i}{1-r}+d)} - e^{-\frac{d}{p}} + 1 \right)^{-p}$$

$$z_i = \frac{x_i - m}{s}$$

$$\psi = \{H, m, s, r, d, p\}$$

- ▶ All parameters for complex species-env relationships
- ▶ $p \rightarrow 1$ removes tail exaggeration
- ▶ $d \rightarrow 0$ removes peak flatness
- ▶ $r \rightarrow 0.5$ removes asymmetry
- ▶ primer-e.github.io/senlm/articles/modskurt.html

What we have so far

For obs $i \in 1, \dots, N$

$$y_i \sim \text{ZIDL}(\eta_i, \theta = \{\phi, \pi, \gamma_0, \gamma_1\})$$

$$\eta_i = f(x_i, H, m, s, r, d, p)$$

$$H \sim \text{Beta}(2, 4) \cdot \max y \quad H \in (0, \max y)$$

$$m \sim \text{Beta}(1, 1) \cdot (u - l) + l \quad \approx \text{Unif}[l = \min x, u = \max x]$$

$$s \sim \text{HalfNormal}(1, \sigma_s) \cdot \bar{\sigma}_x \quad \text{standardise over } x$$

$$r \sim \text{Beta}(2, 2)$$

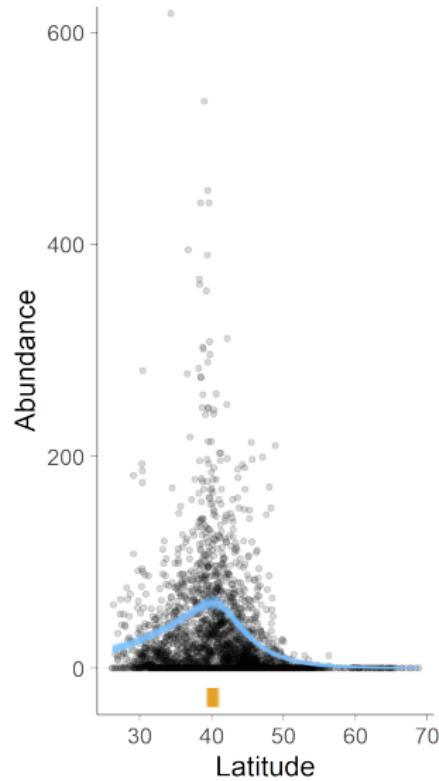
$$d \sim \text{HalfNormal}(0, \sigma_d)$$

$$p \sim \text{HalfNormal}(1, \sigma_p)$$

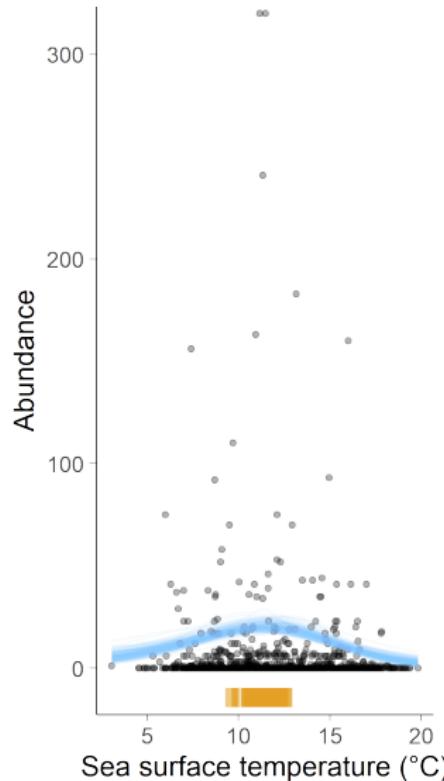
$$\theta \sim P(\mu_\theta, \sigma_\theta) \quad \text{indep. dist priors}$$

Posterior predictive checks

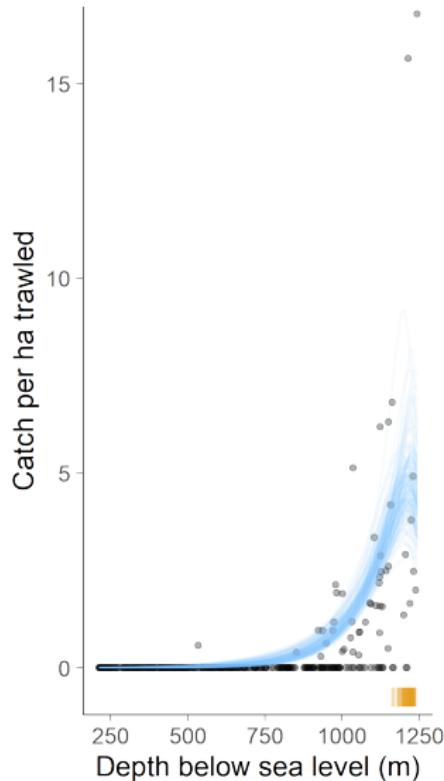
Quiscalus quiscula



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— NB mean (η) ■ Mode, optimal x (m)

Responses need time

For year $t \in 1, \dots, T$, each with observations $i \in 1, \dots, N_t$

$$y_{t[i]} \sim \text{ZIDL}(\eta_{t[i]}, \theta = \{\phi, \pi, \gamma_0, \gamma_1\})$$

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Parameters for the error distribution consistent across years

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Different mean curves fit for each year

$$\eta_{t[i]} = f(x_{t[i]}, H_t, m_t, s_t, r_t, d_t, p_t)$$

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Modal position as a function of time

$$m_t = g(t) + \epsilon_t$$

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Remaining modskurt parameters are partially pooled across years

$$\psi = \{H, s, r, d, p\}$$

$$\bar{\psi}_j \sim P(\mu_\psi, \sigma_{\bar{\psi}}) \leftarrow \text{average priors the same as earlier}$$

$$\psi_t \sim \text{Normal}(\bar{\psi}, \sigma_\psi) \leftarrow \text{varying each year}$$

$$\sigma_\psi \sim \text{HalfNormal}(0, \tau_\psi) \leftarrow \text{with some regularisation}$$

Technical issues

```
functions {
  vector zinbl(vector pooled, vector pars_t, array[] real x_t, array[] int y_t) {
    real H = pars_t[1]; real m = pars_t[2]; real s = pars_t[3];
    real r = pars_t[4]; real d = pars_t[5]; real p = pars_t[6];

    real phi = pooled[1]; real gamma_0 = pooled[2]; real gamma_1 = pooled[3];

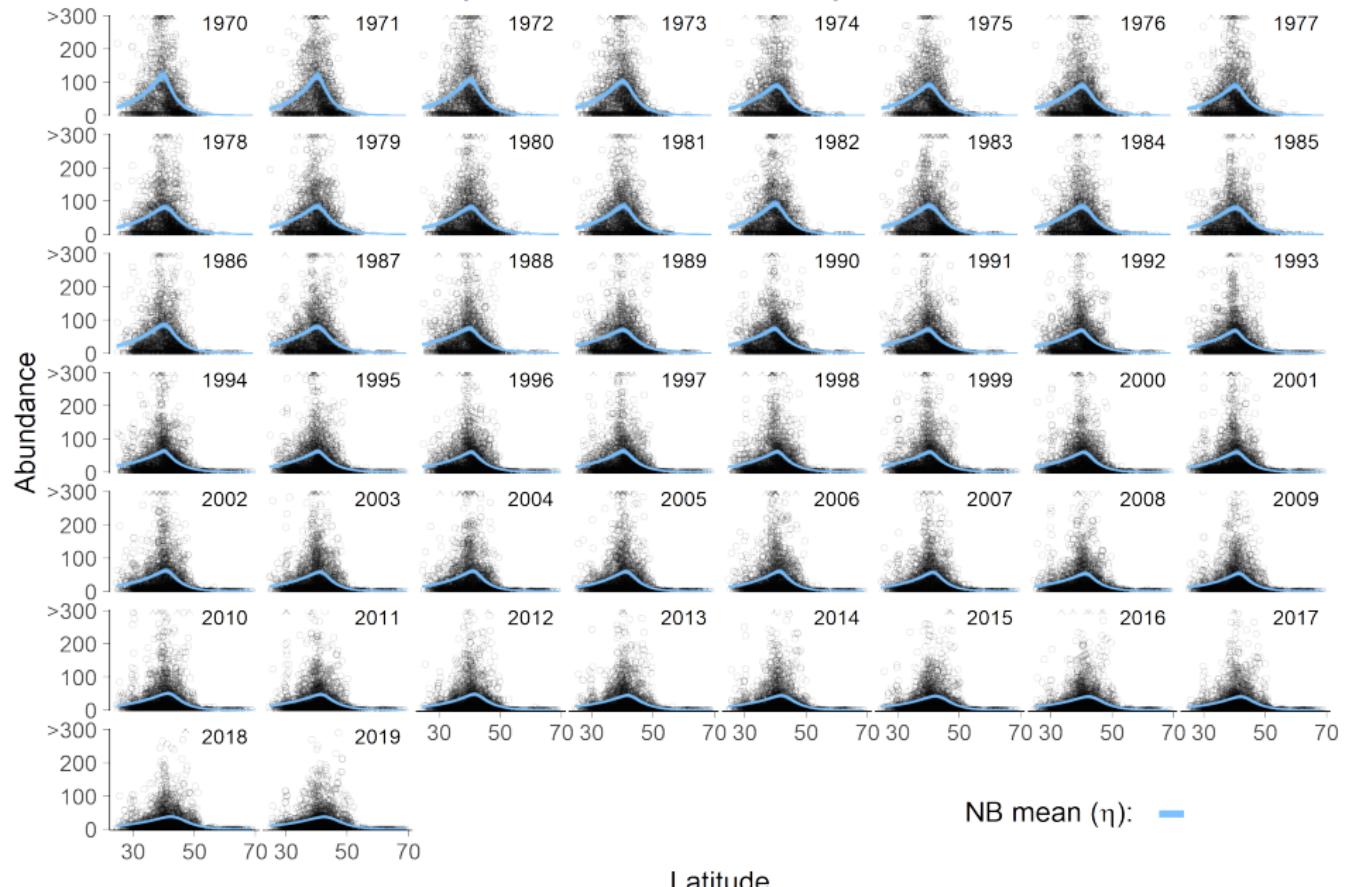
    int N_t = y_t[1]; int Nz_t = y_t[2]; int Npos = N_t - Nz_t;

    vector[N_t] x = to_vector(x_t[3:(N_t + 2)]);
    array[Npos] int y = y_t[(Nz_t + 3):(N_t + 2)];

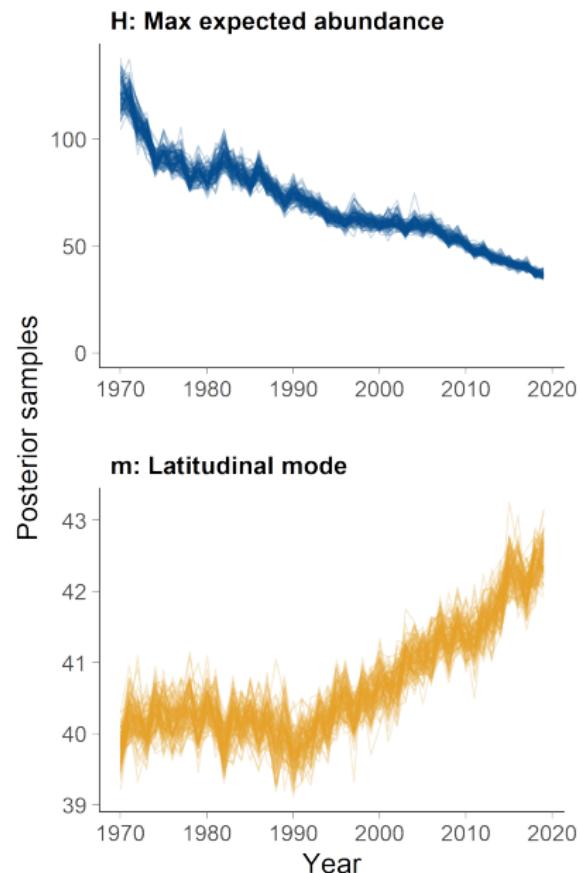
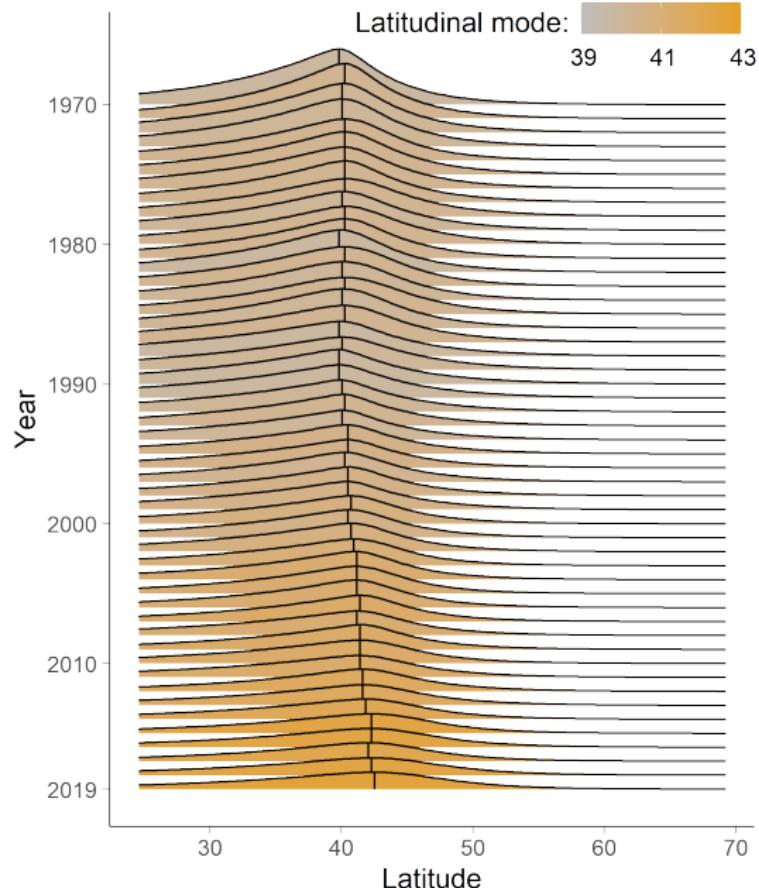
    vector[N_t] log_eta = log_modskurt(x, H, m, s, r, d, p);
    vector[N_t] logit_pi = fma(-gamma_1, log_eta, gamma_0);

    real lp = 0;
    if (Nz_t > 0) {
      lp += log_sum_exp(bernoulli_logit_lpmf(1 | logit_pi[1:Nz_t]),
                        bernoulli_logit_lpmf(0 | logit_pi[1:Nz_t]) + neg_binomial_2_log_lpmf(0 | log_eta[1:Nz_t], phi));
    }
    lp += bernoulli_logit_lpmf(0 | logit_pi[(Nz_t + 1):N_t]) + neg_binomial_2_log_lpmf(y | log_eta[(Nz_t + 1):N_t], phi);
    return [lp]';
  }
}
model {
  target += sum(map_rect(zinbl, pars_pooled, pars_t, x, y));
}
```

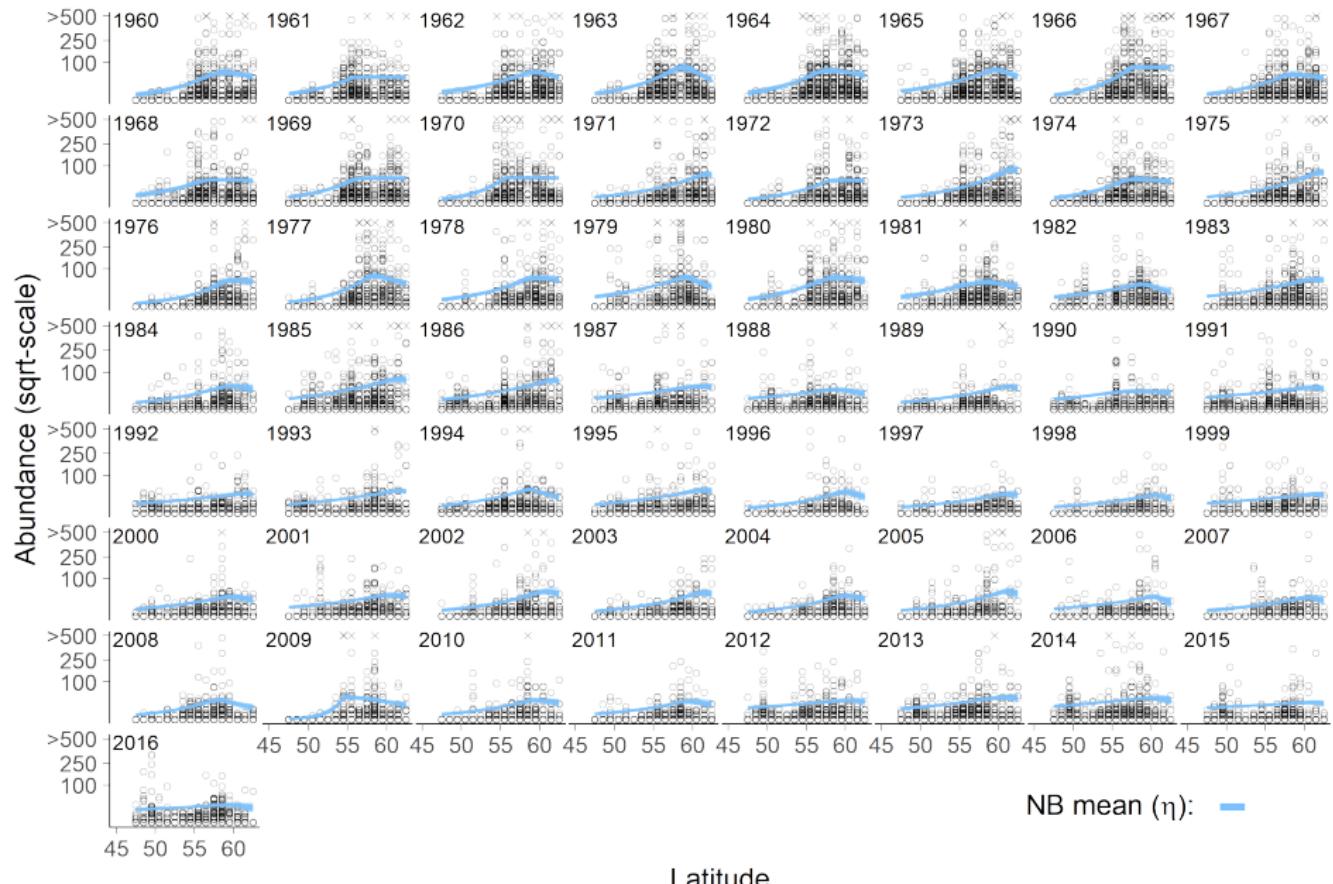
Quiscalus quiscula (common grackle) + lat



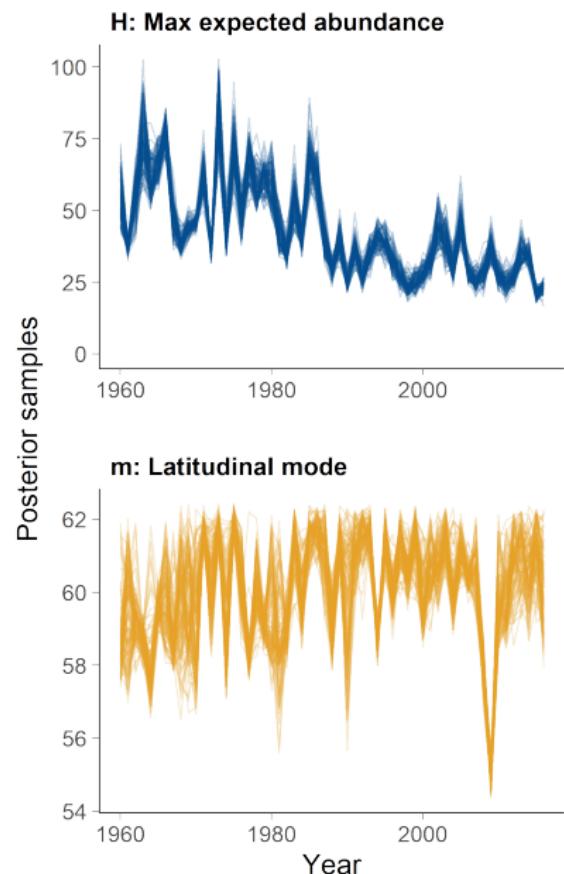
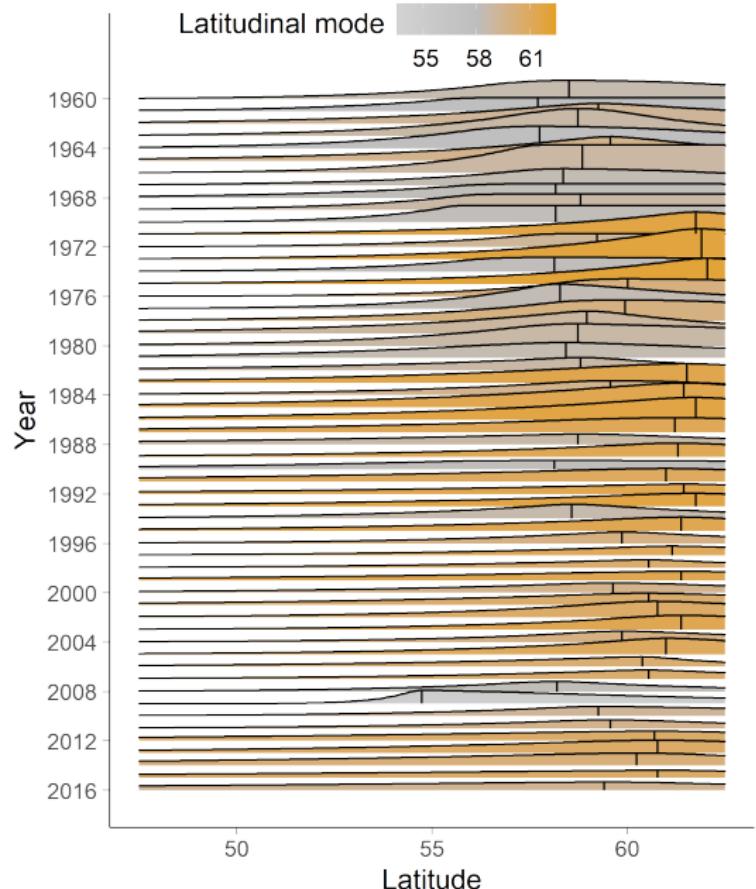
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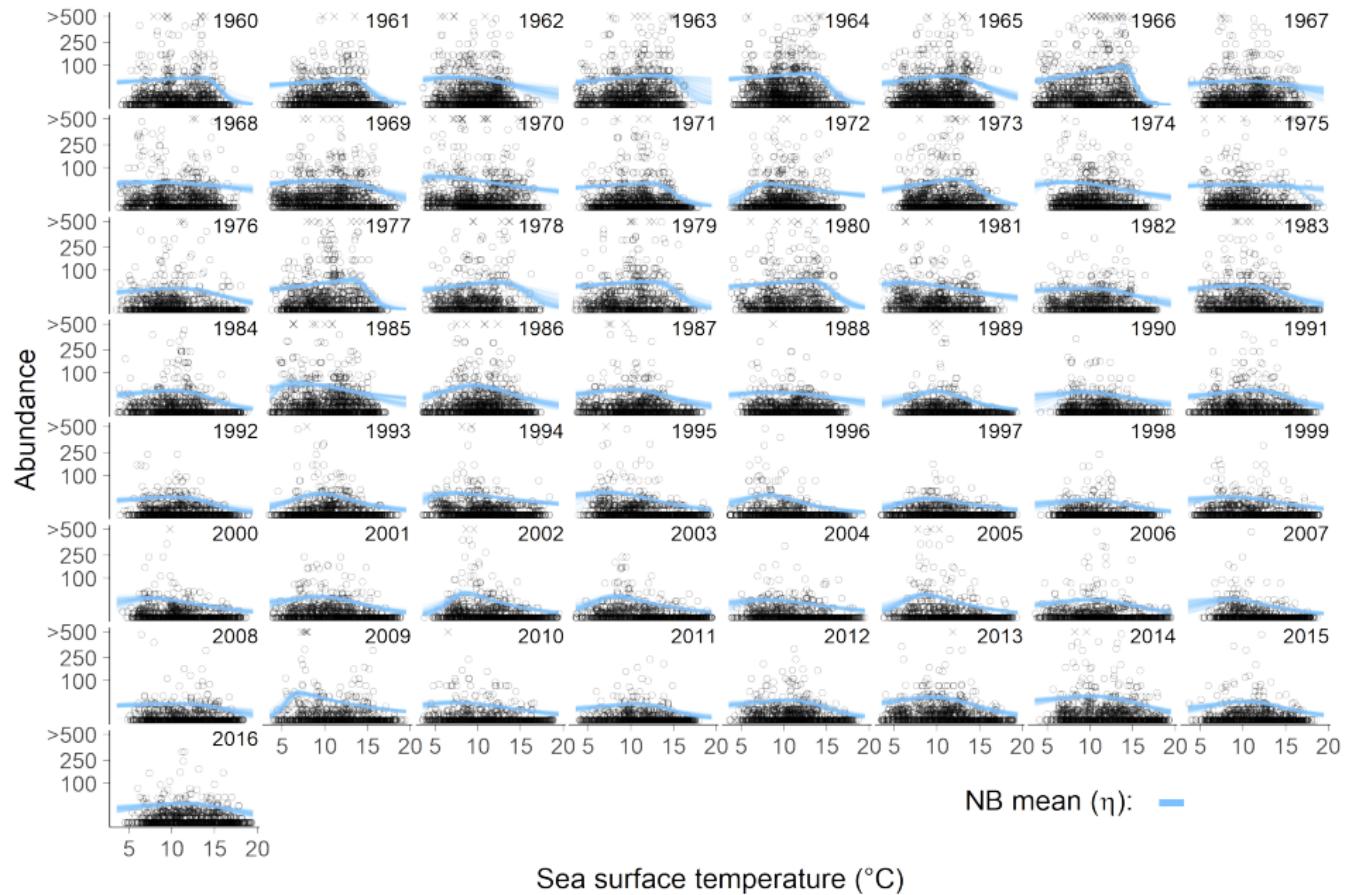
Calanus finmarchicus (cold water copepod) + lat



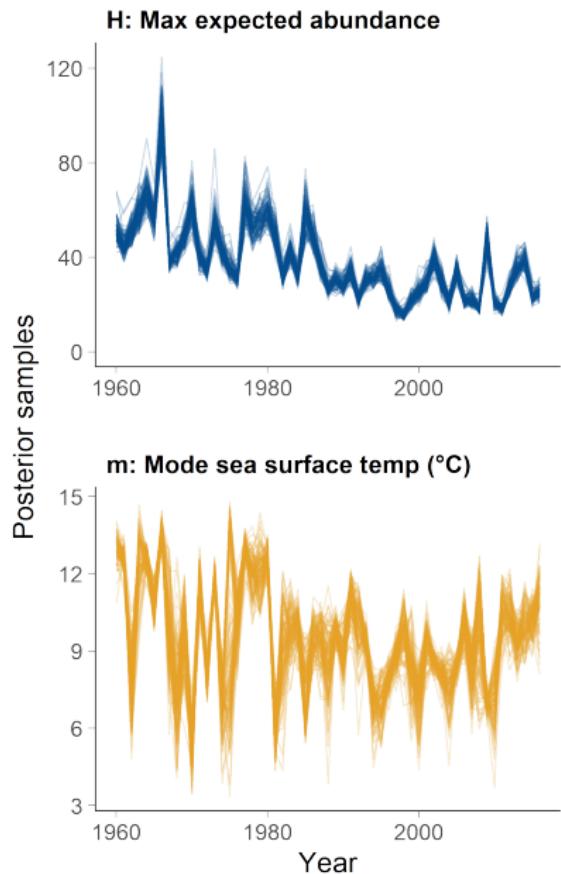
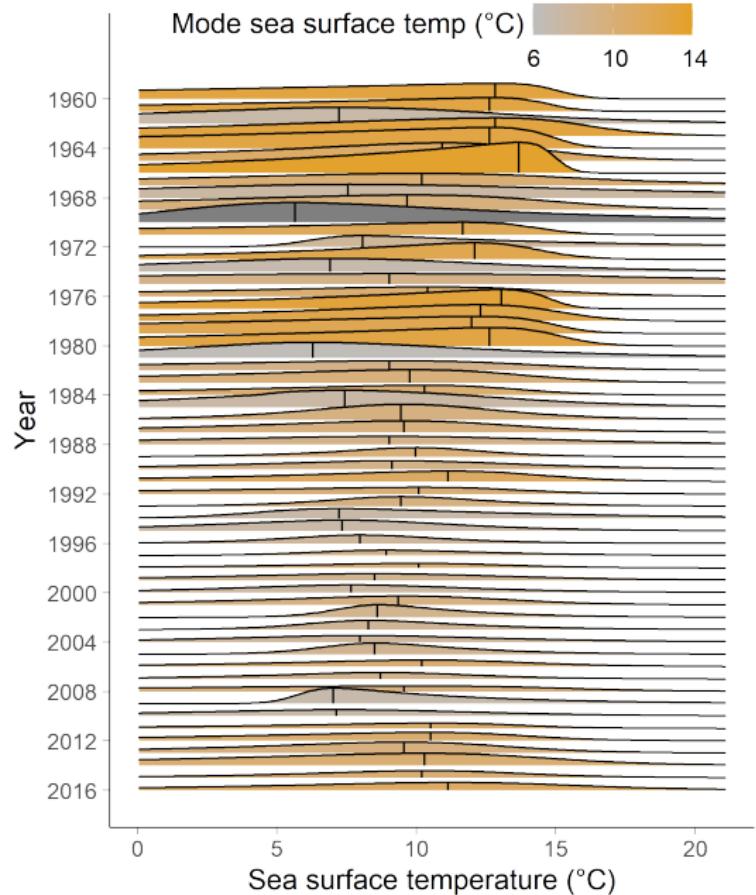
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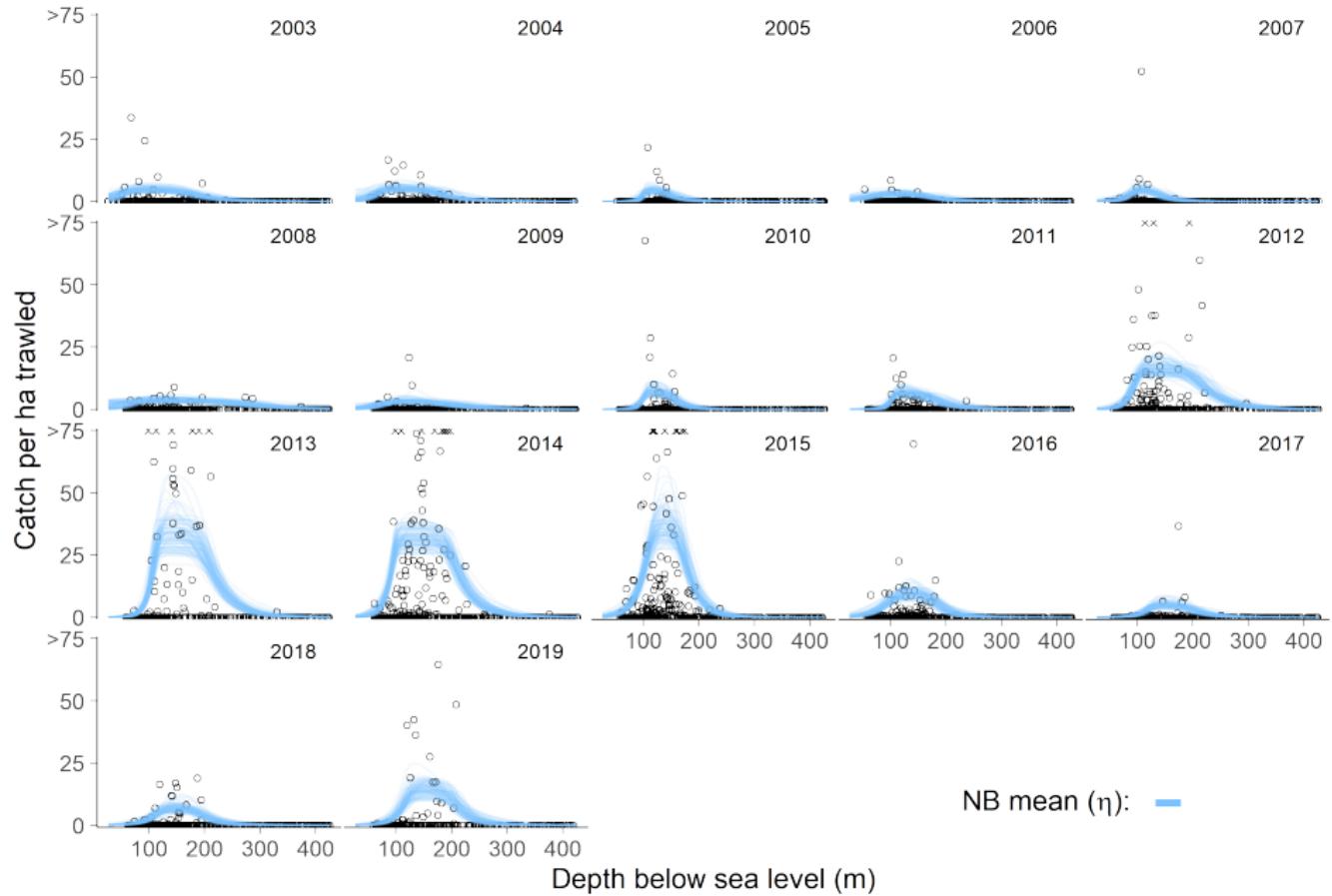
Calanus finmarchicus (cold water copepod) + temp



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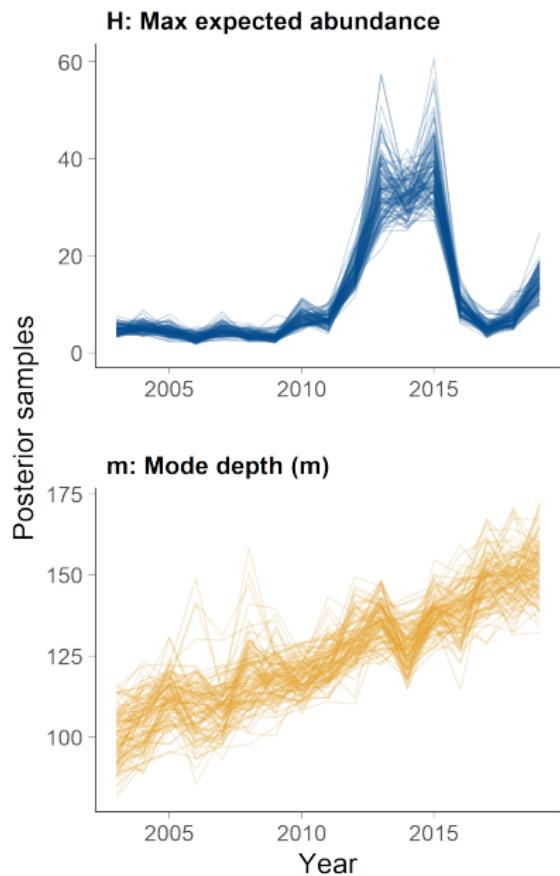
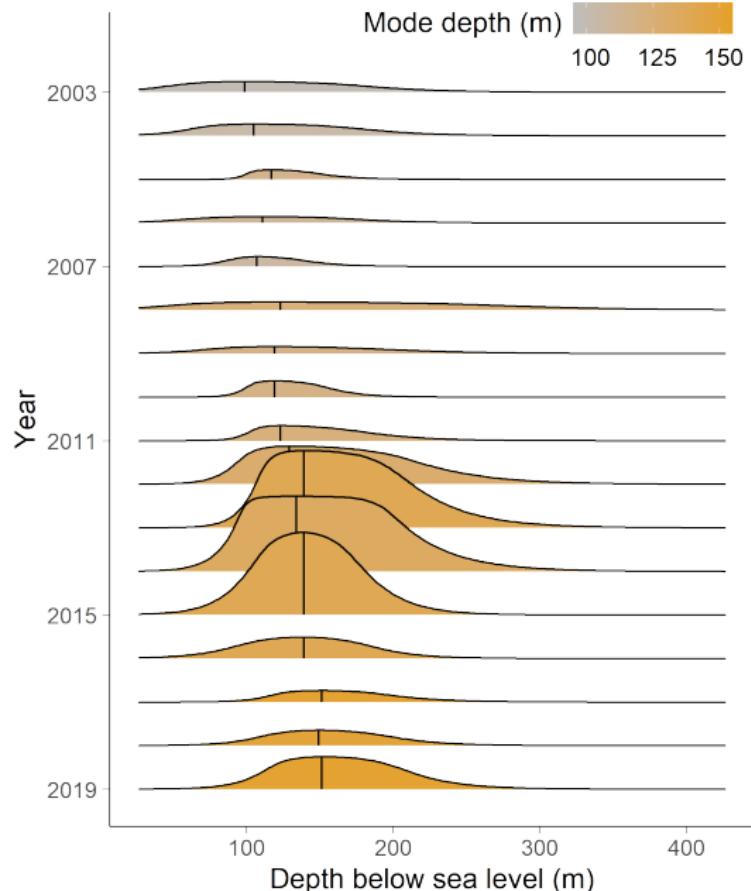


Thaleichthys pacificus (eulachon) + depth

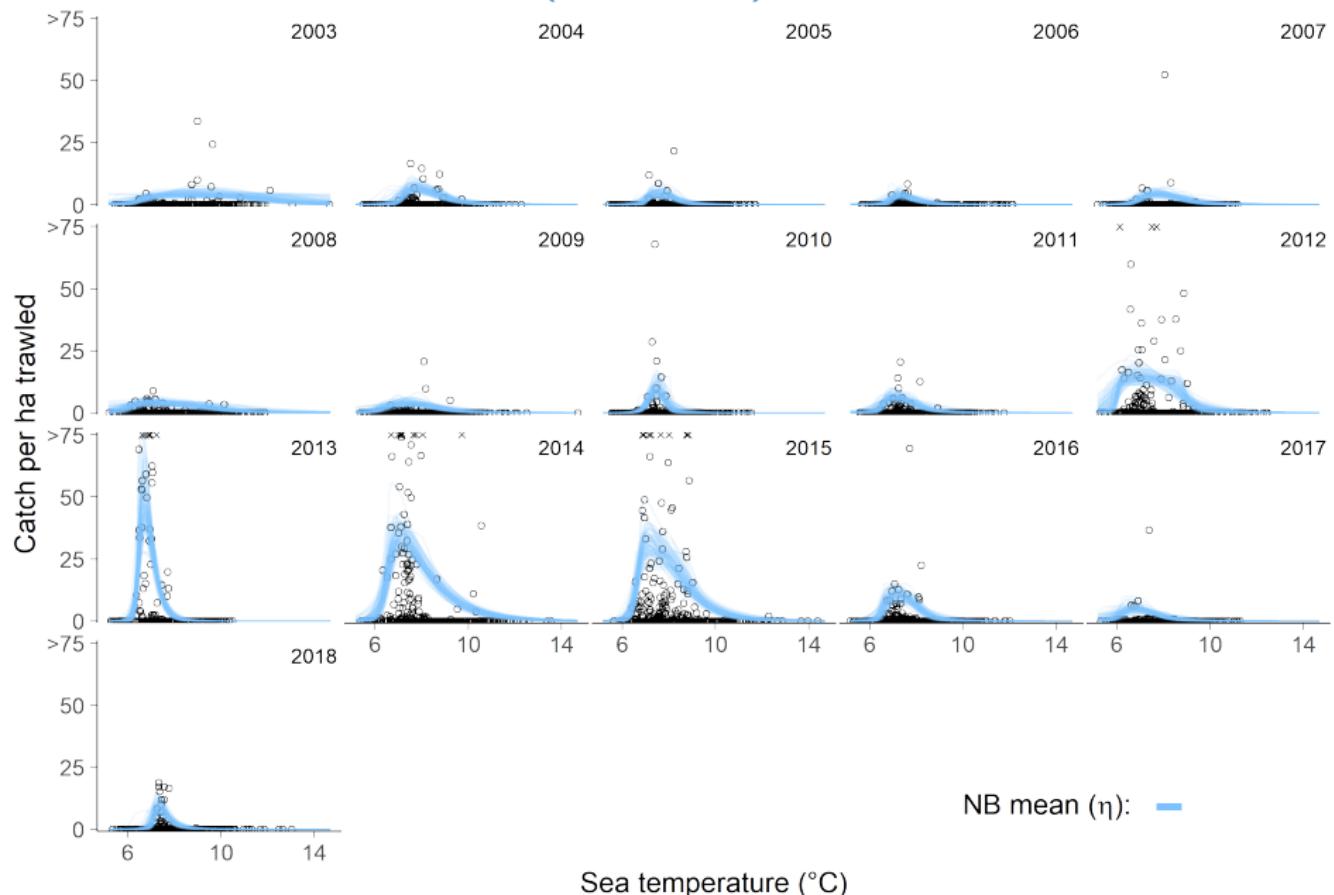


NB mean (η): —

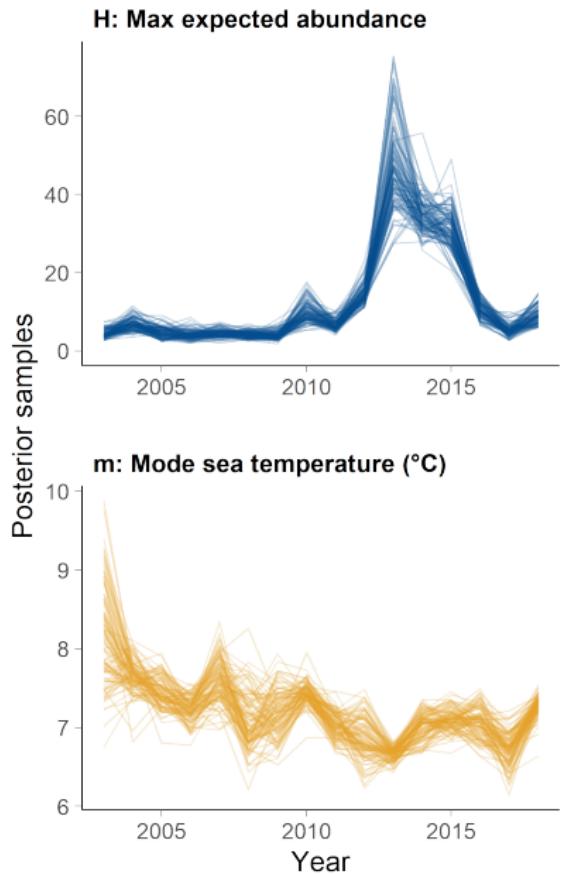
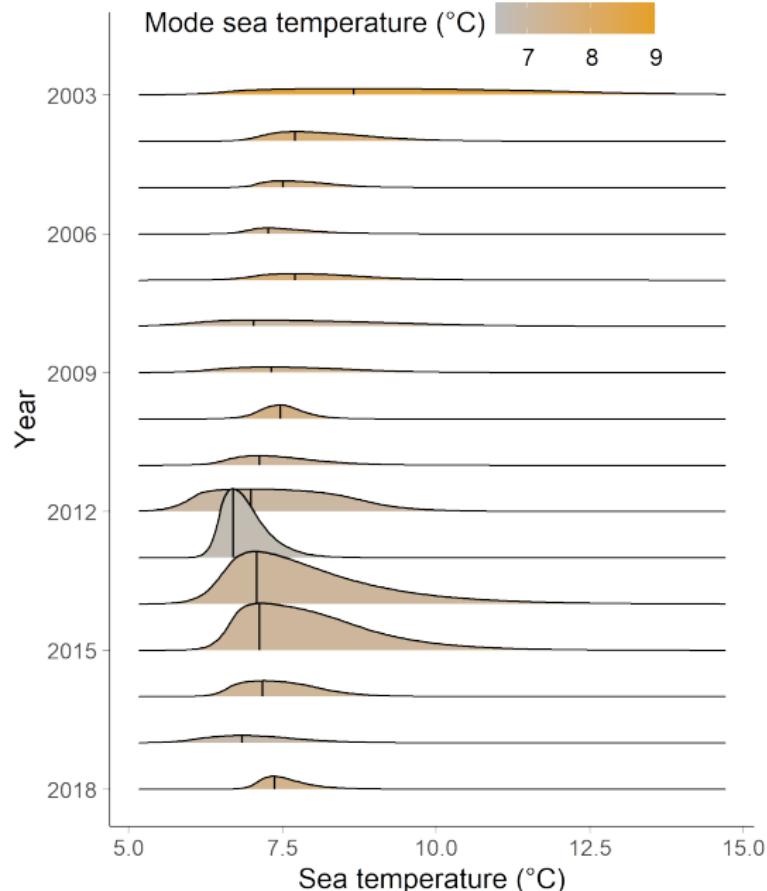
Thaleichthys pacificus (eulachon) + depth



Thaleichthys pacificus (eulachon) + temp



Thaleichthys pacificus (eulachon) + temp



In summary

- ▶ Flexible and interpretable model for exploring species-environment data over time (some things to iron out)
- ▶ Bayesian approach propagates uncertainty across multiple levels, and benefits from regularising prior model
- ▶ Building from principles may offer inferential advantages over more predictive approaches

Next steps

- ▶ Do ironing
- ▶ Formal models for causality
- ▶ Publish R package to extend `brms`

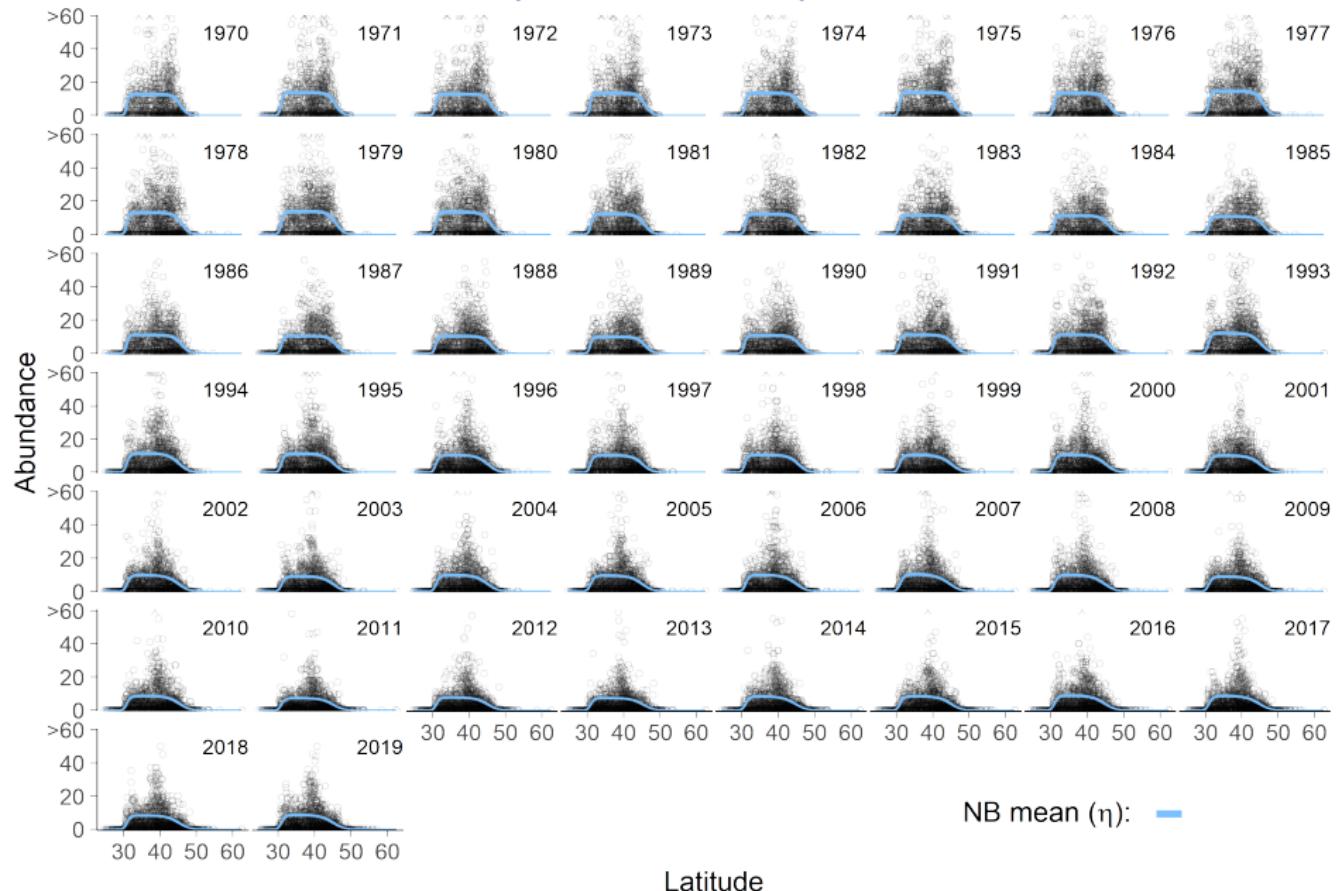
Acknowledgements

- ▶ My supervisors and team: Marti Anderson, Adam Smith, Winston Sweatman, Andrew Punnet, Paulo Mateus Martins
- ▶ Massey University Masters Research Scholarship
- ▶ PRIMER-e Inaugural Masterate Scholarship
- ▶ New Zealand Institute for Advanced Study - Center for Theoretical Chemistry and Physics HPC cluster.
- ▶ Royal Society of New Zealand Marsden Grant (19-466 MAU-145)
- ▶ Canadian Wildlife Service & United States Geological Survey, UK Marine Biological Association, National Oceanic and Atmospheric Administration Fisheries

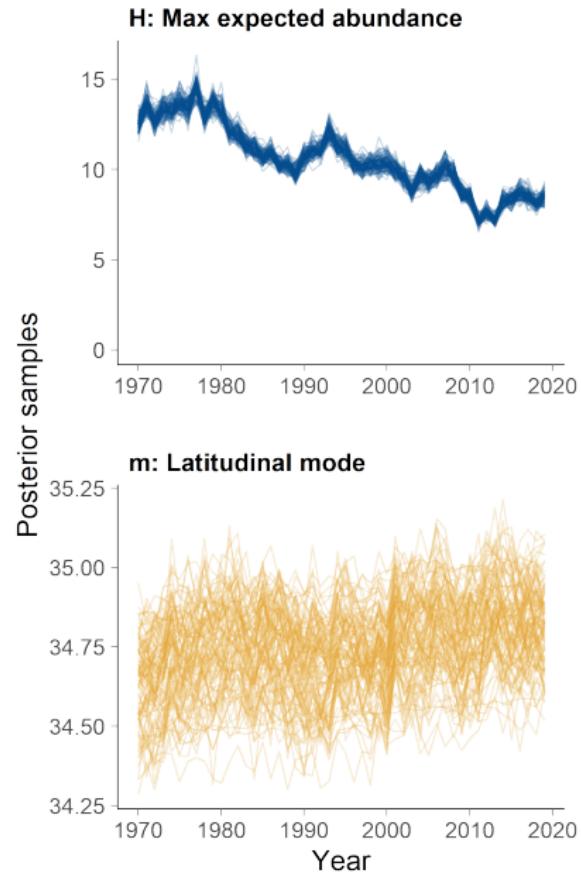
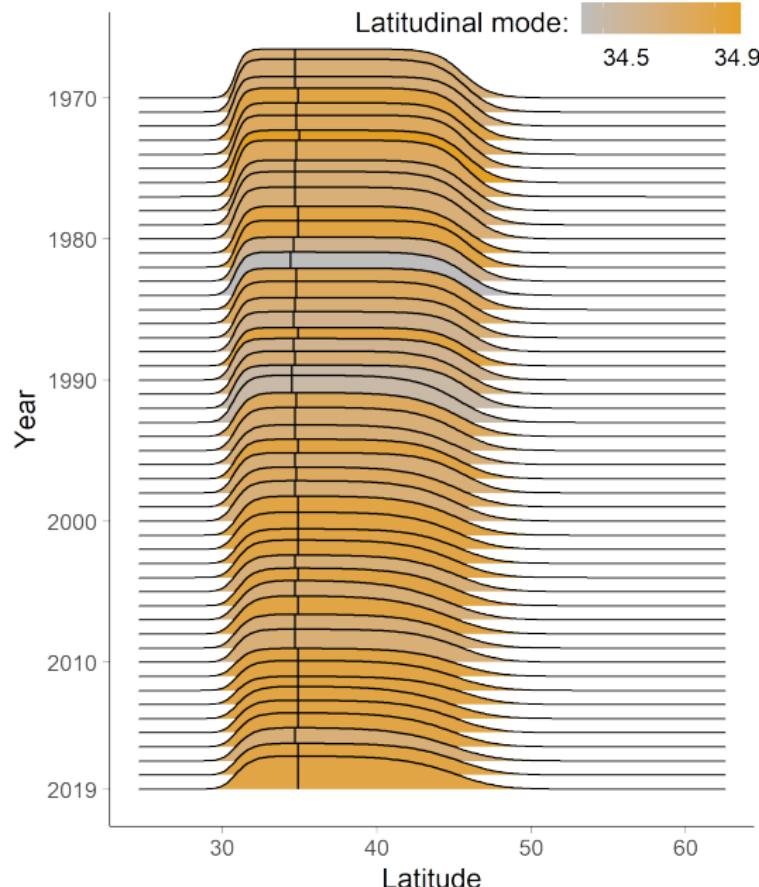
Extras

...

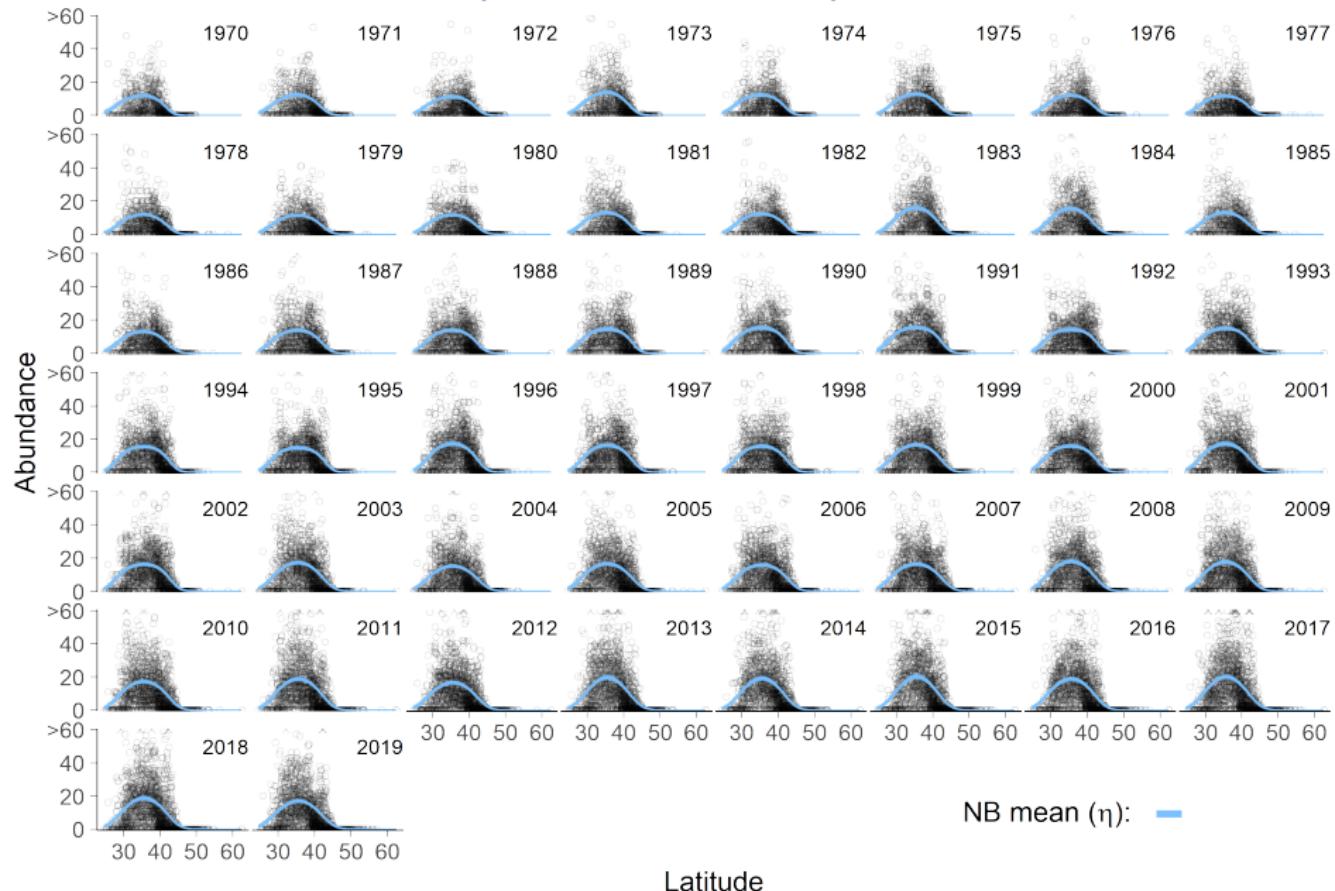
Hylocichla mustelina (wood thrush) + lat



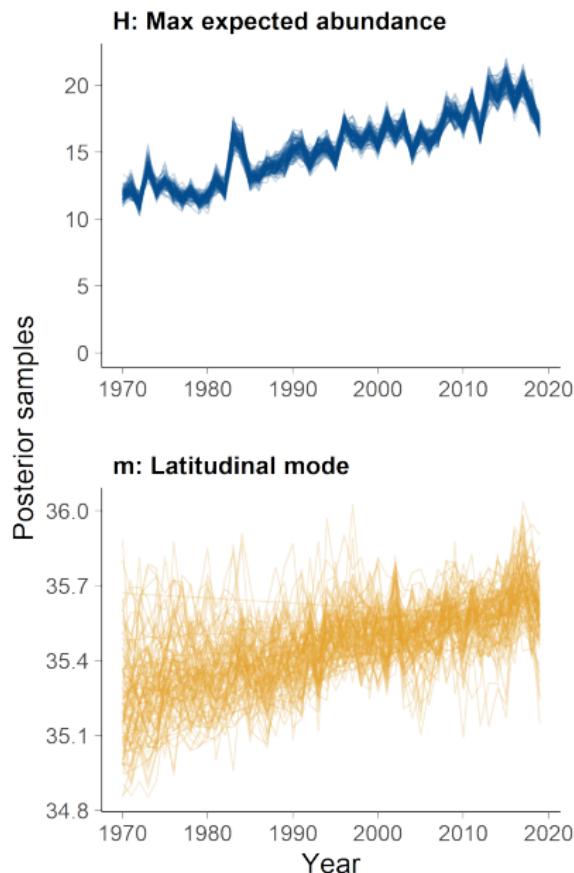
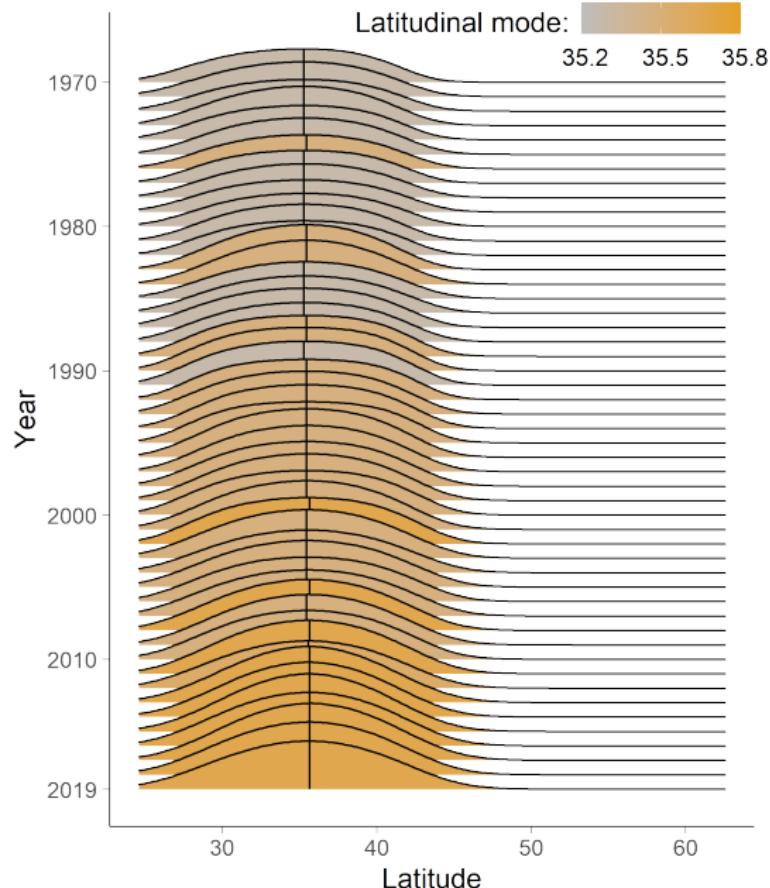
Hylocichla mustelina (wood thrush) + lat



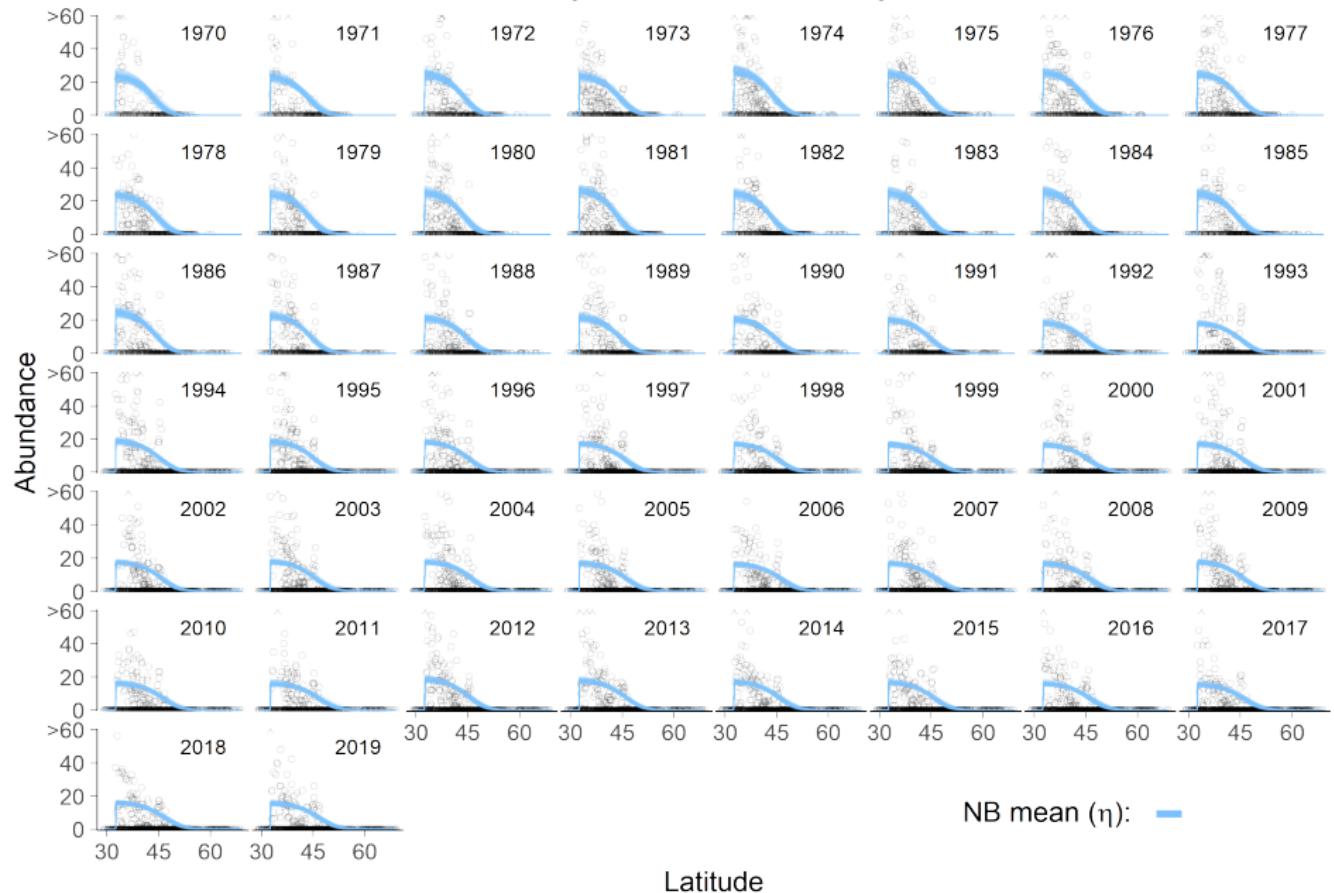
Baeolophus bicolor (tufted titmouse) + lat



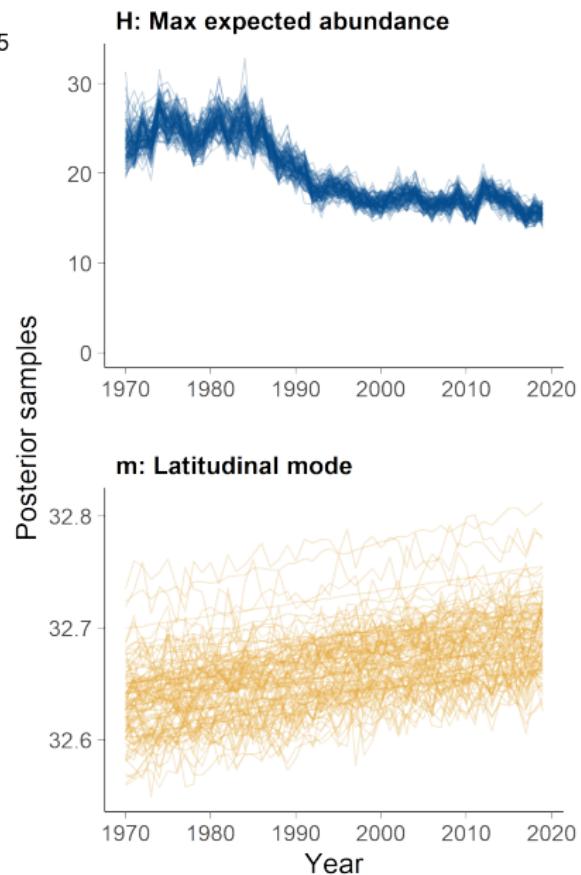
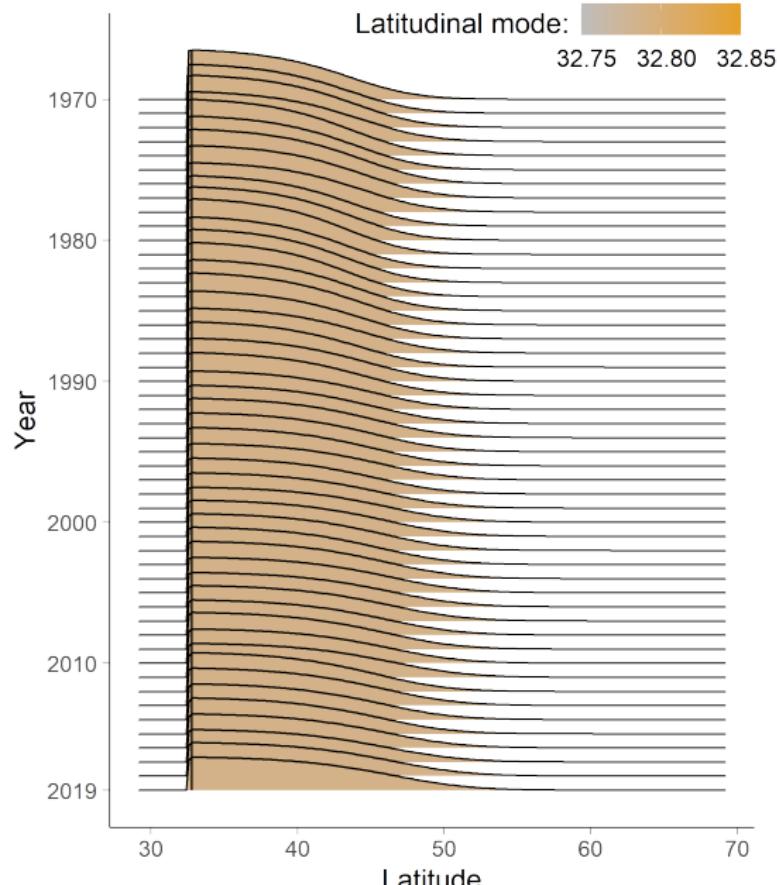
Baeolophus bicolor (tufted titmouse) + lat



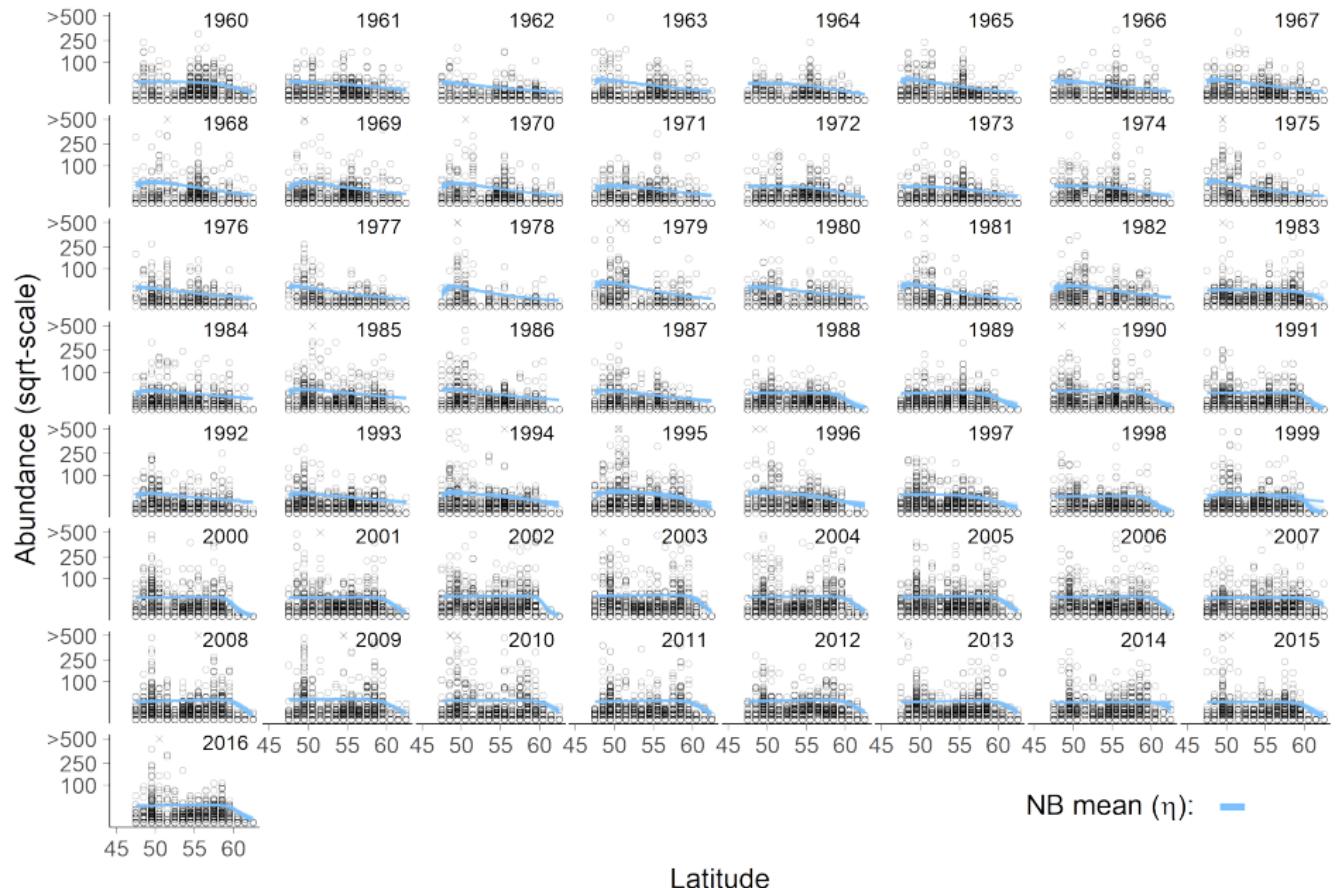
Aphelocoma californica (Cali scrub jay) + lat



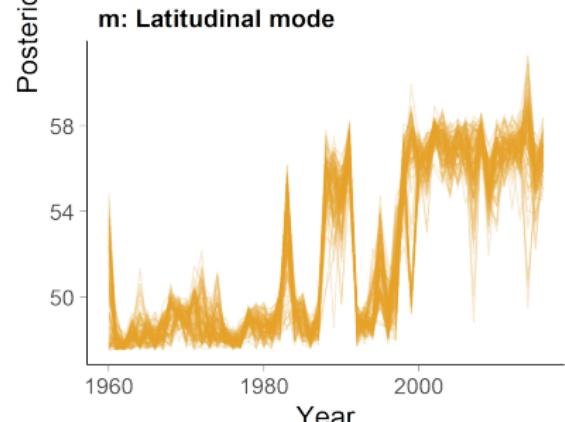
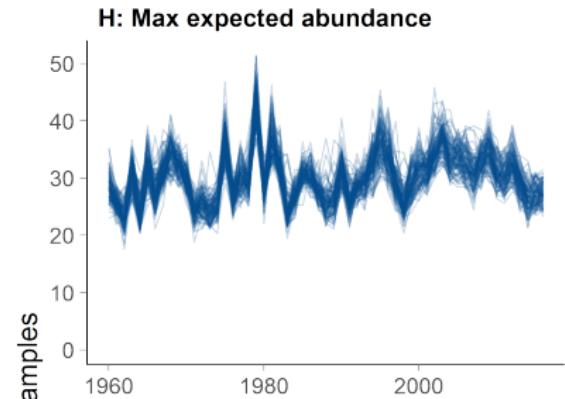
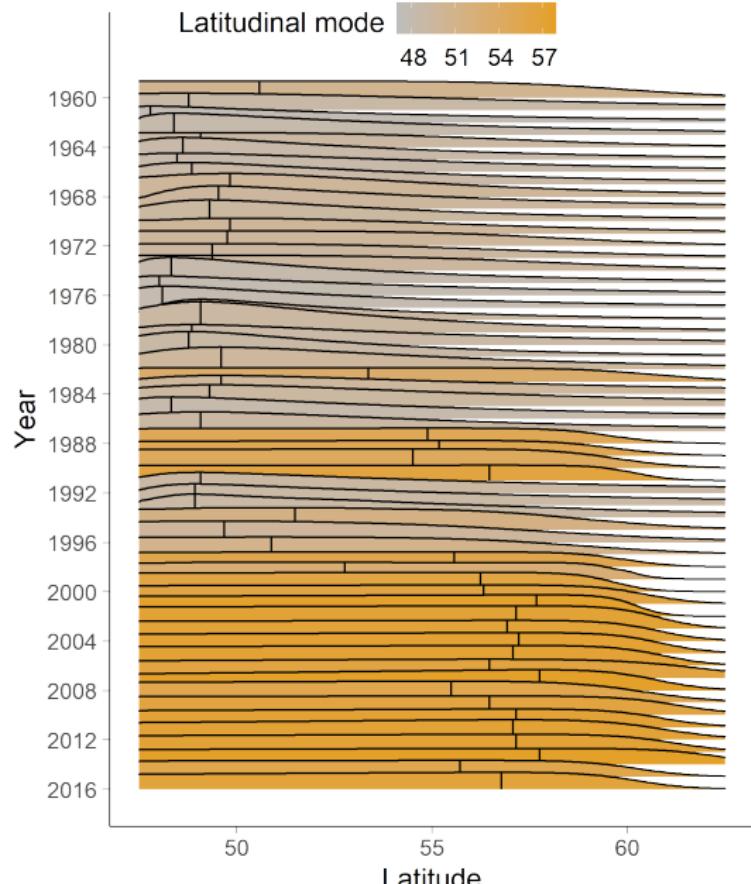
Aphelocoma californica (Cali scrub jay) + lat



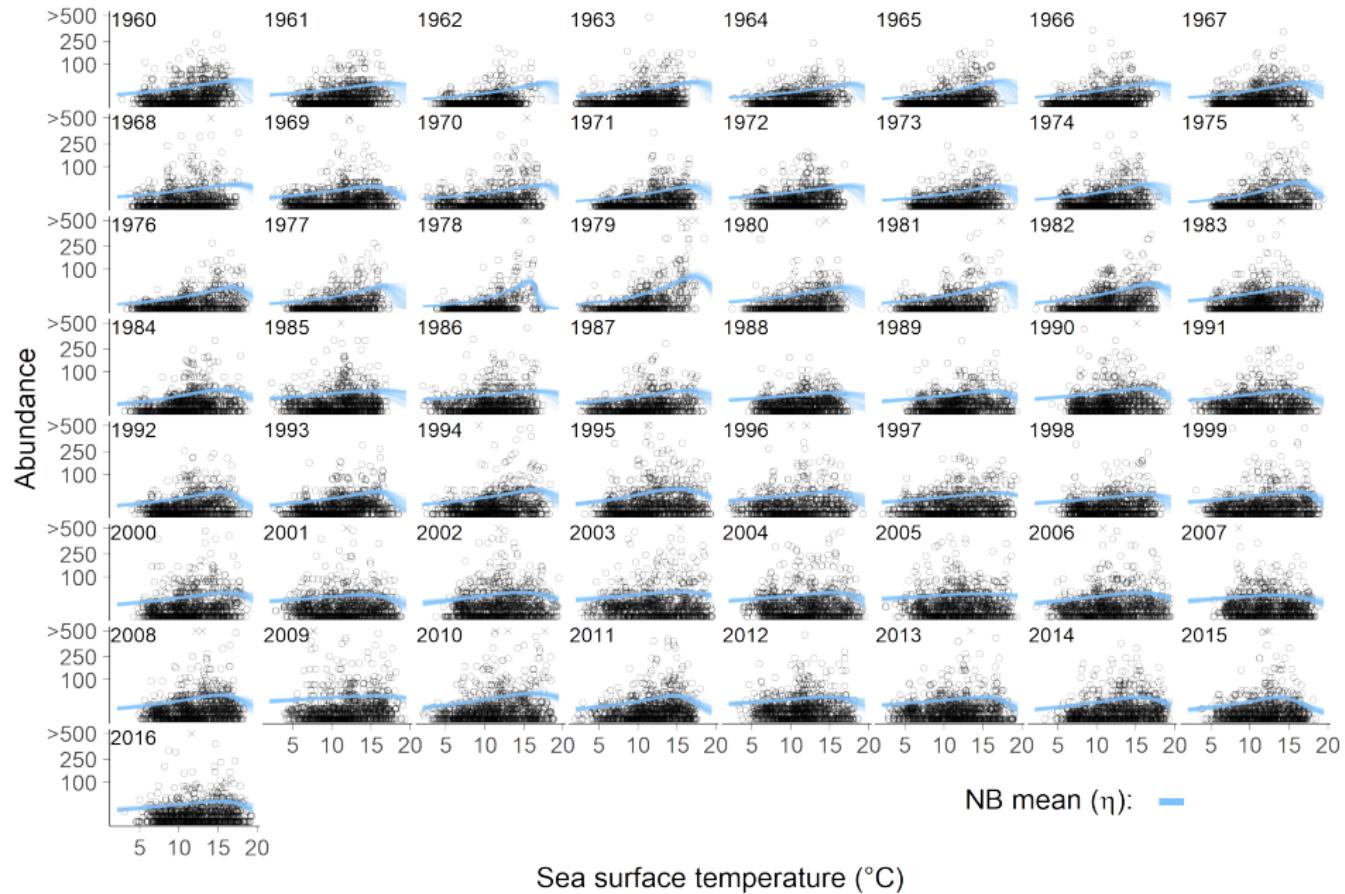
Calanus helgolandicus (warm water copepod) + lat



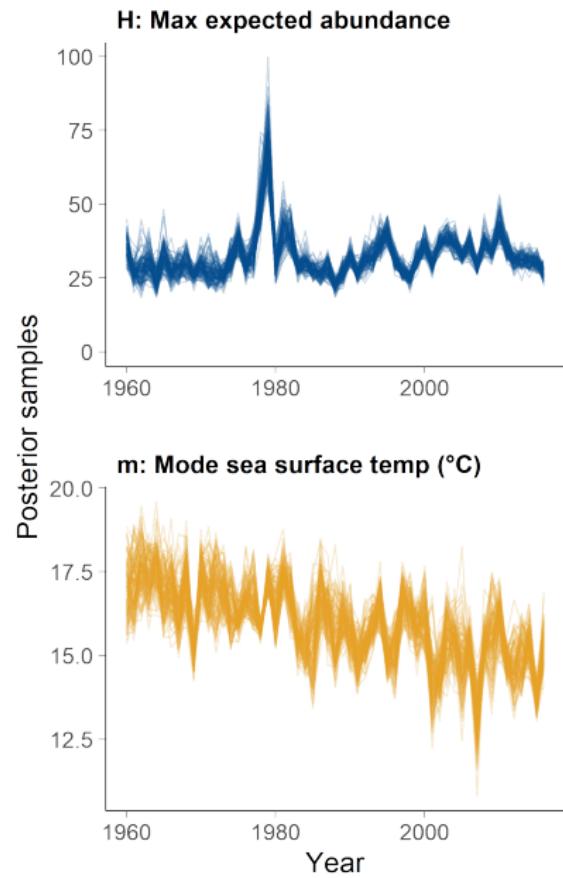
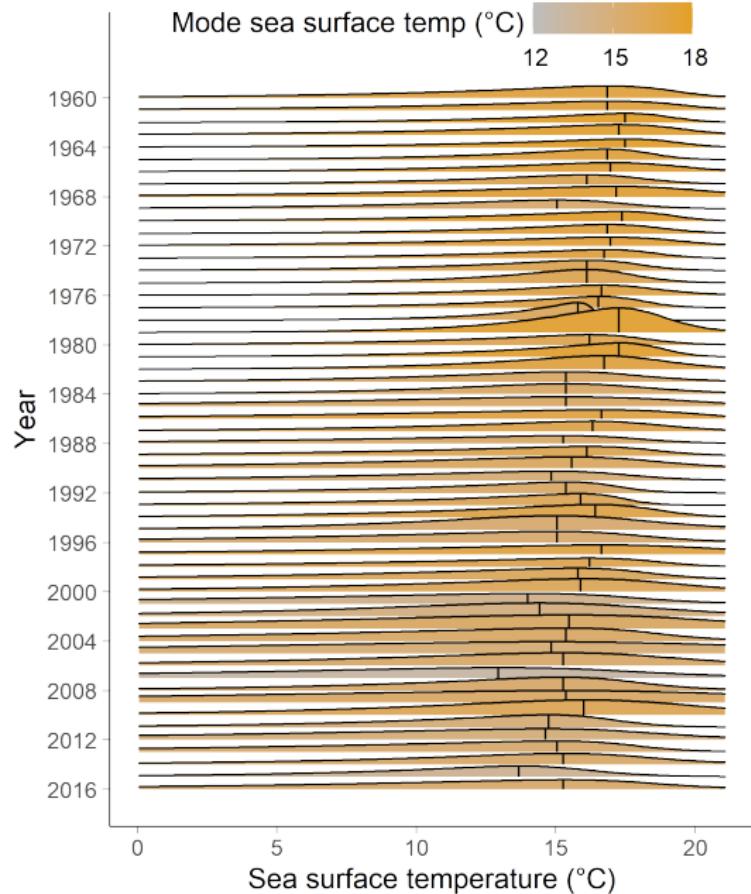
Calanus helgolandicus (warm water copepod) + lat



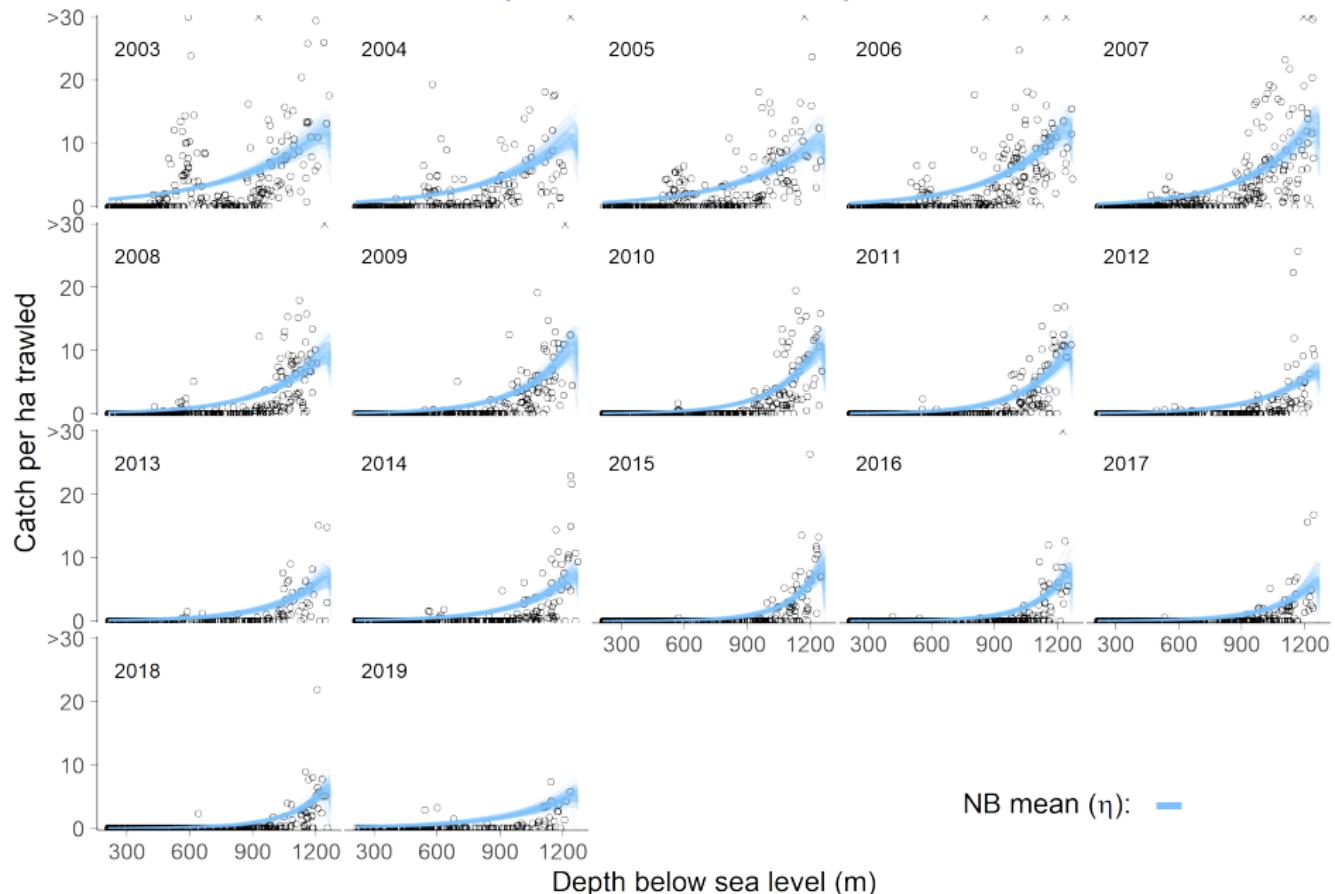
Calanus helgolandicus (warm water copepod) + temp



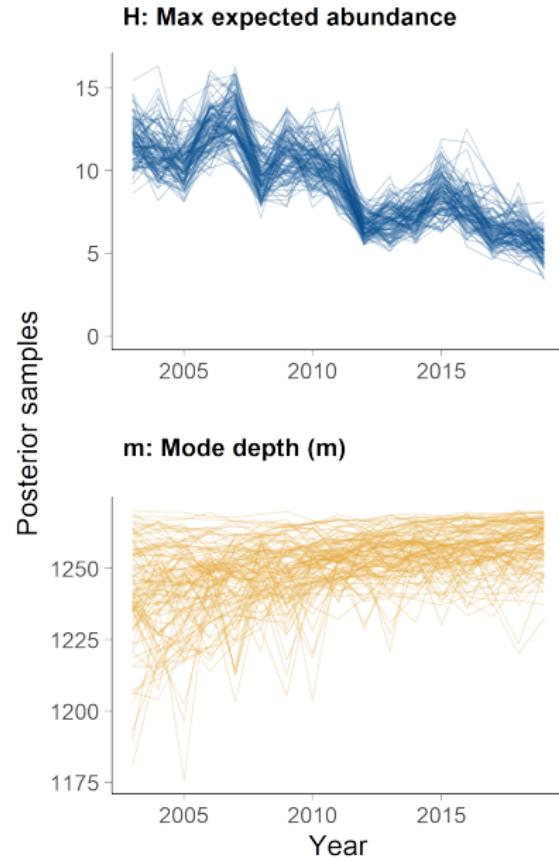
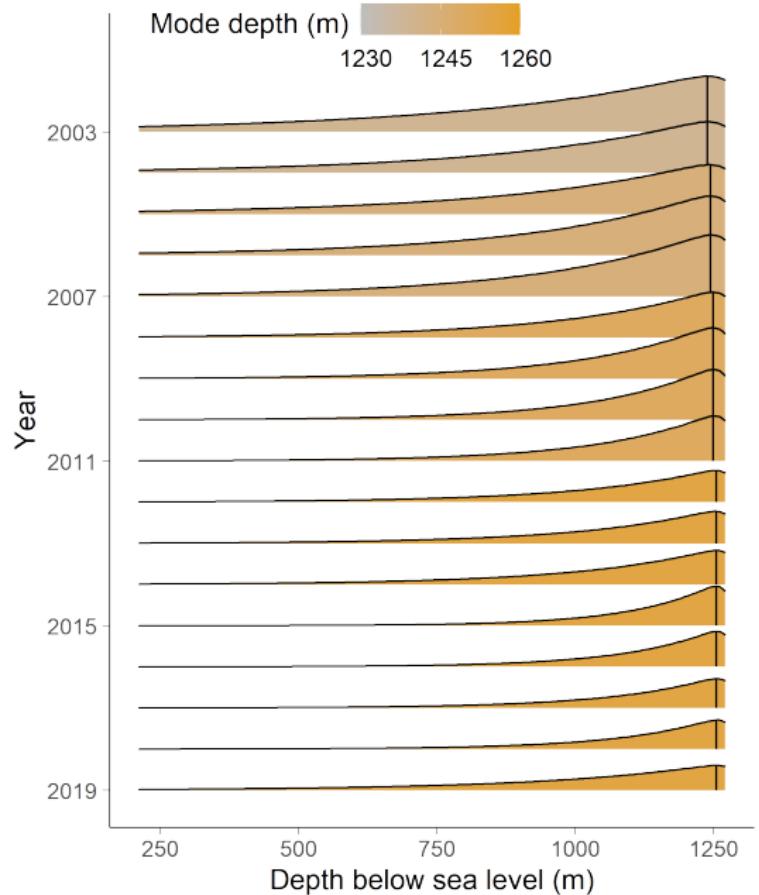
Calanus helgolandicus (warm water copepod) + temp



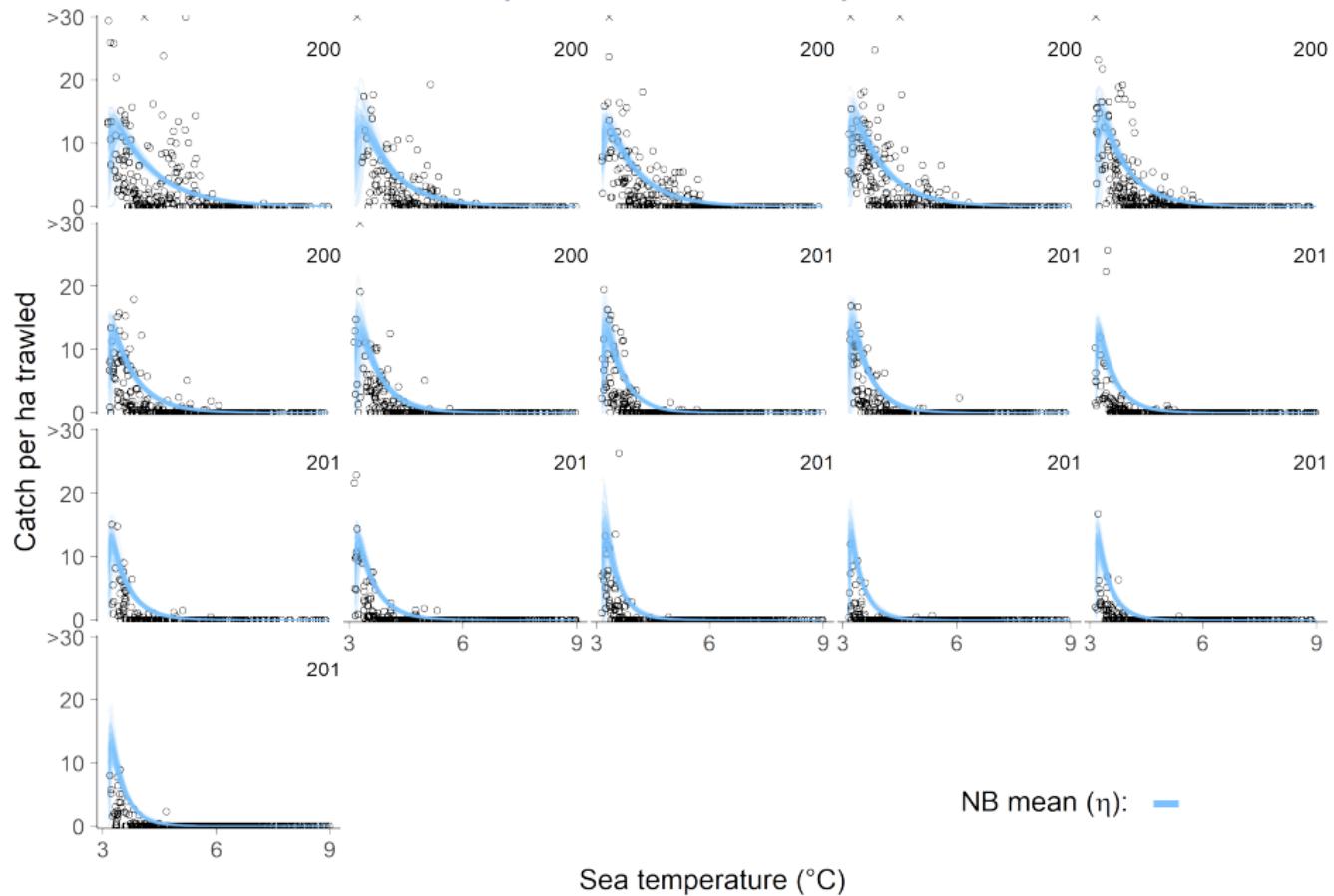
Antimora microlepis (pacific flatnose) + depth



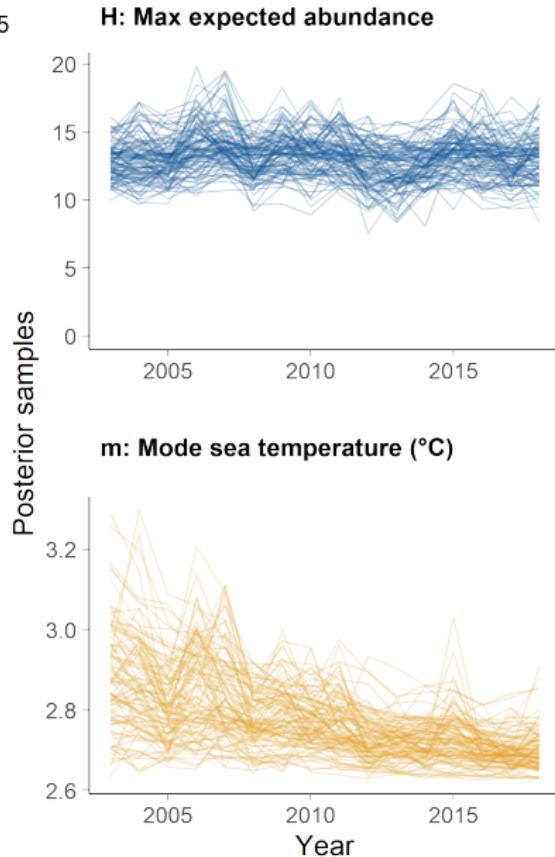
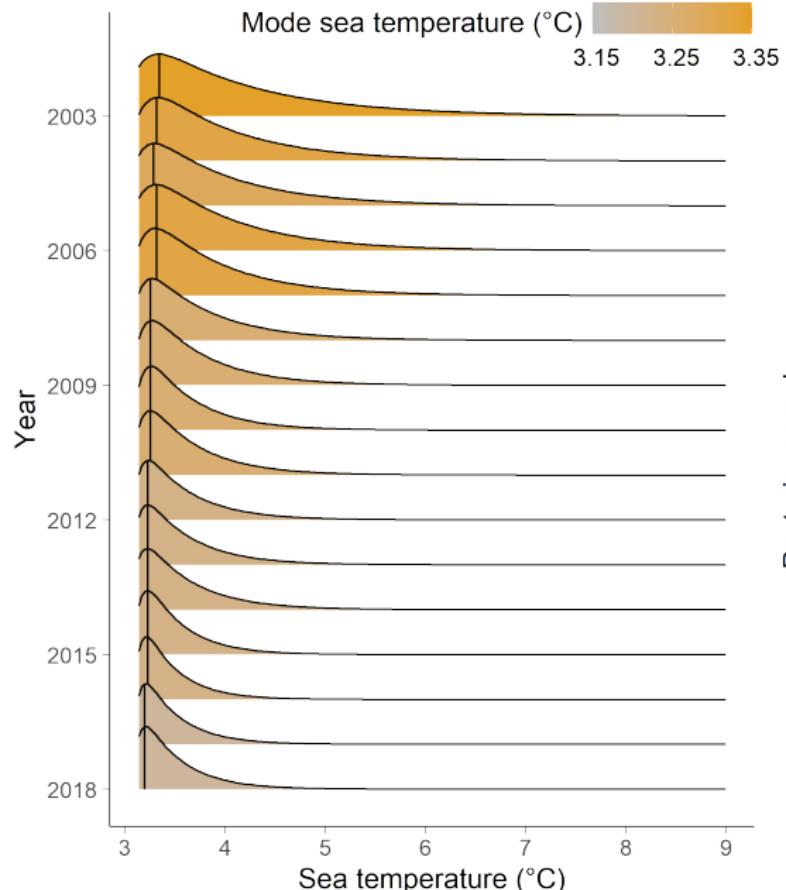
Antimora microlepis (pacific flatnose) + depth



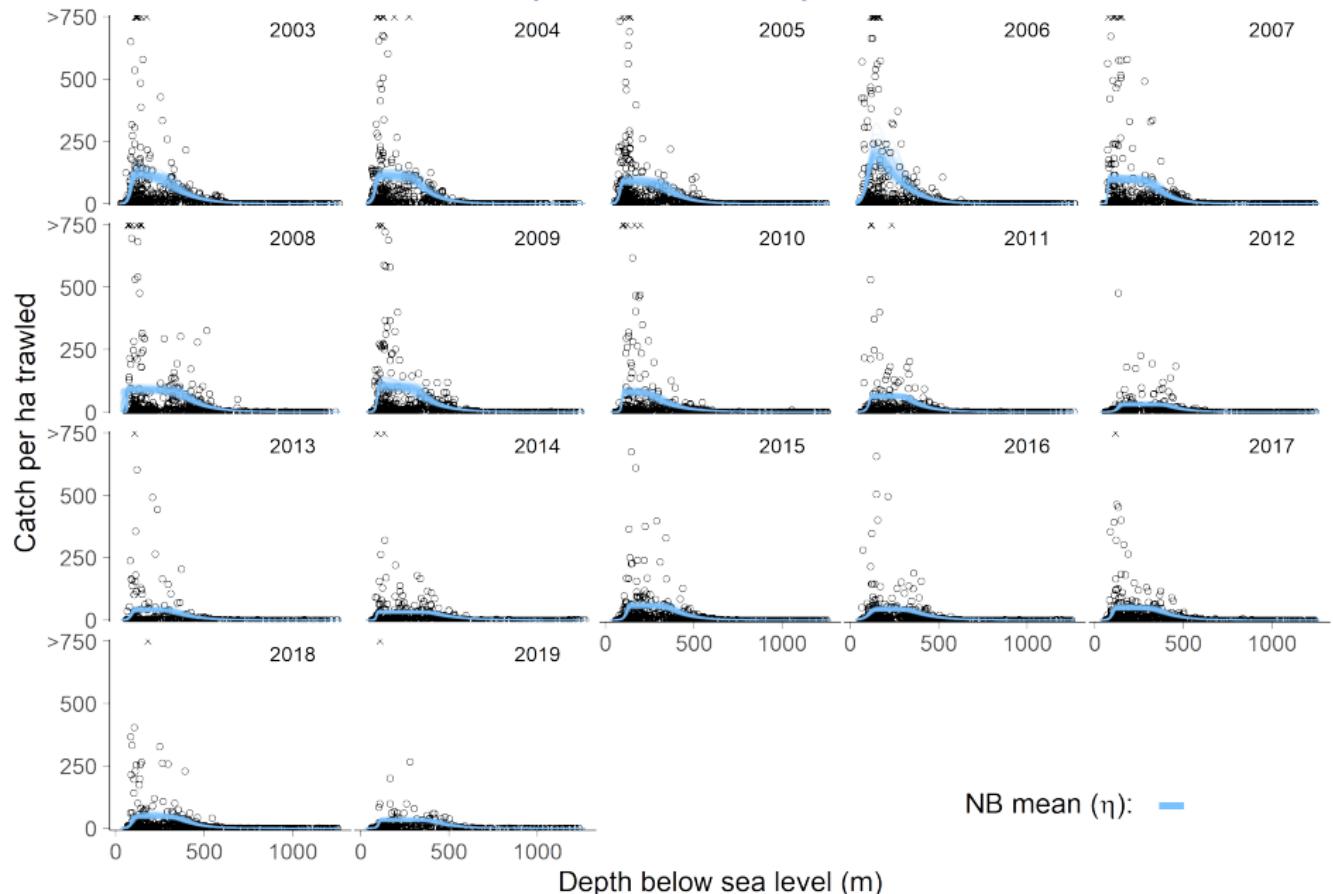
Antimora microlepis (pacific flatnose) + temp



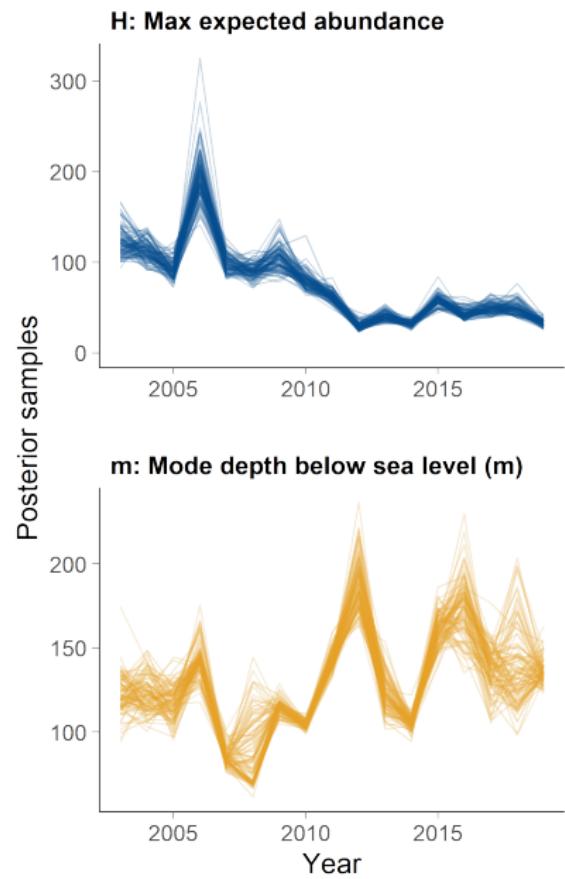
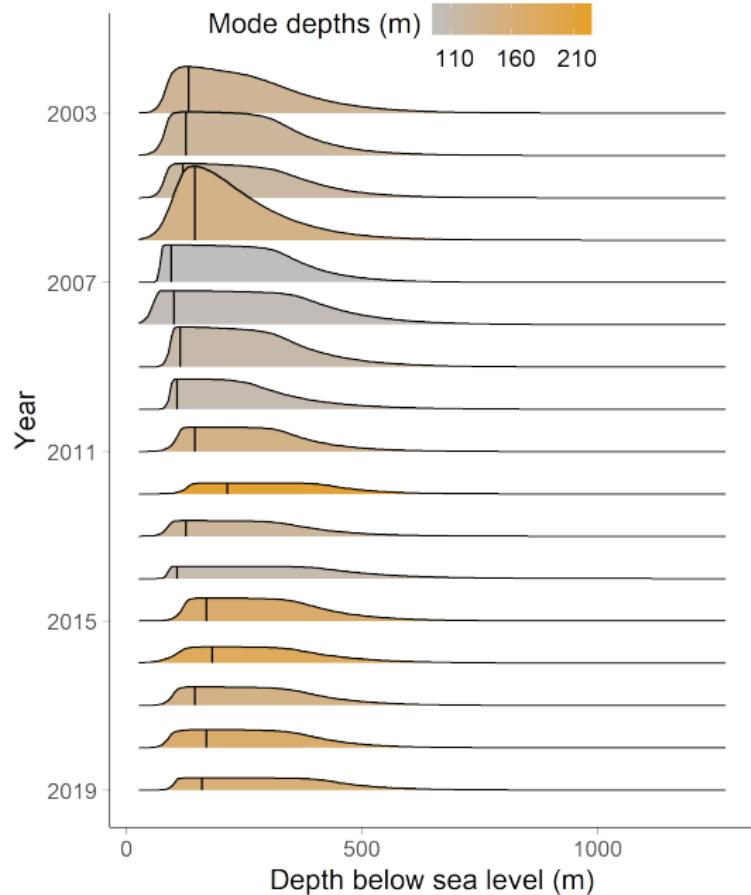
Antimora microlepis (pacific flatnose) + temp



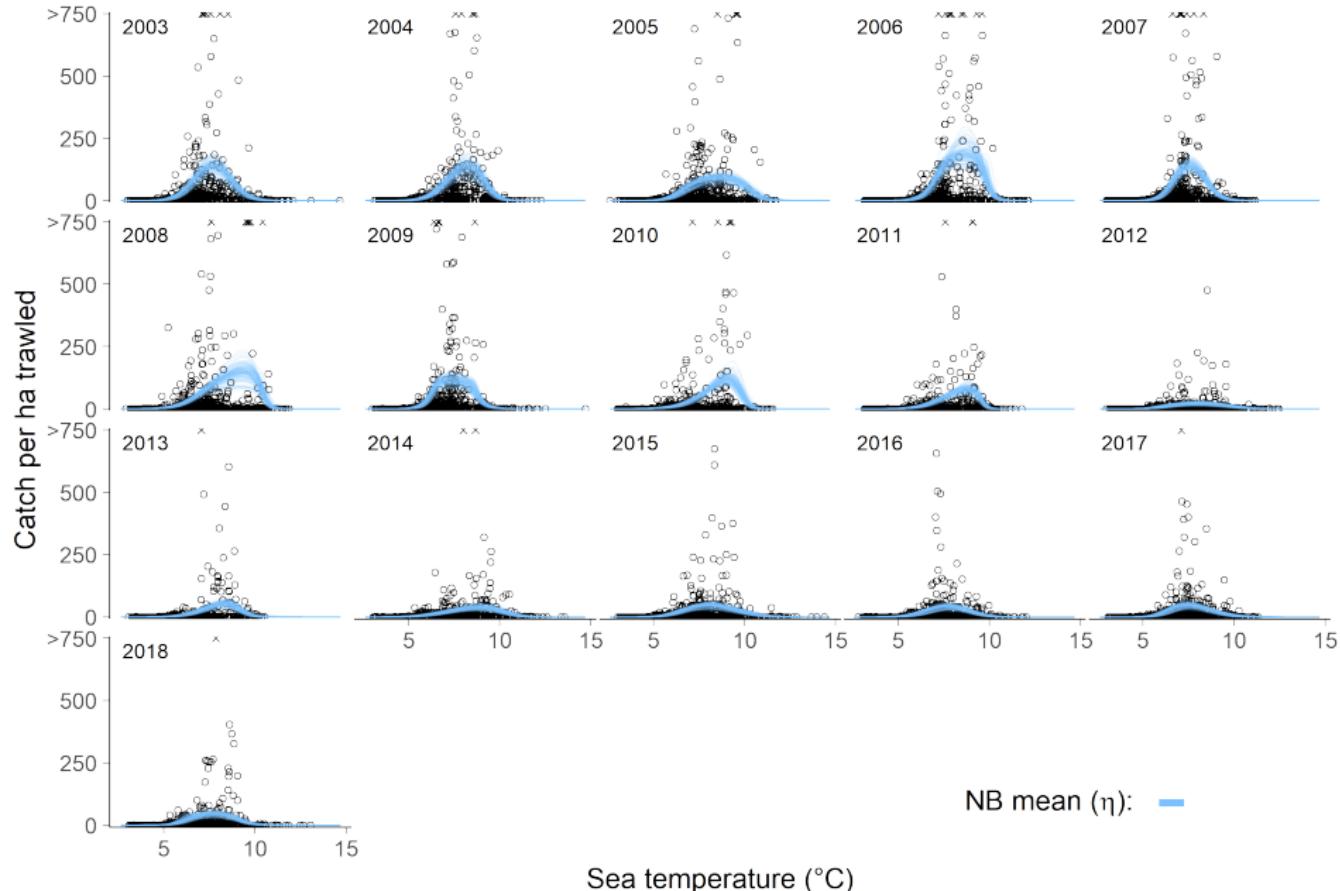
Merluccius productus (pacific hake) + lat



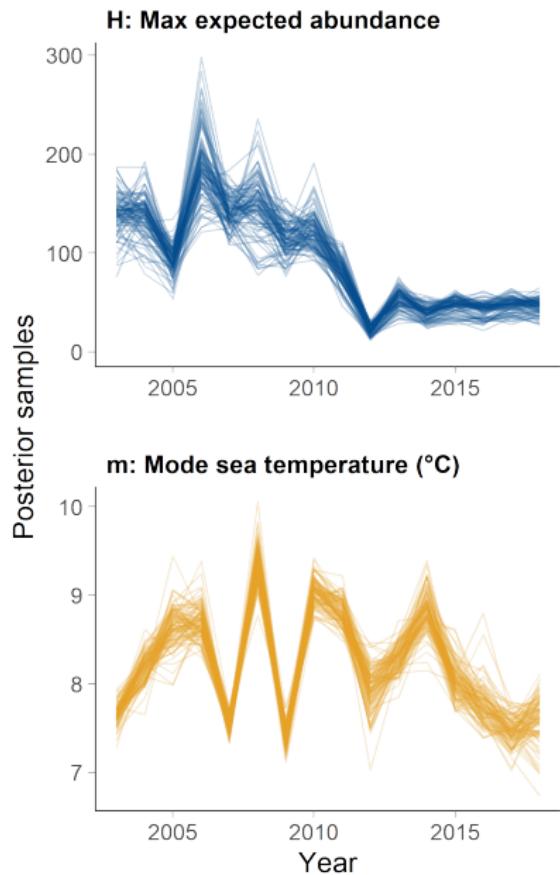
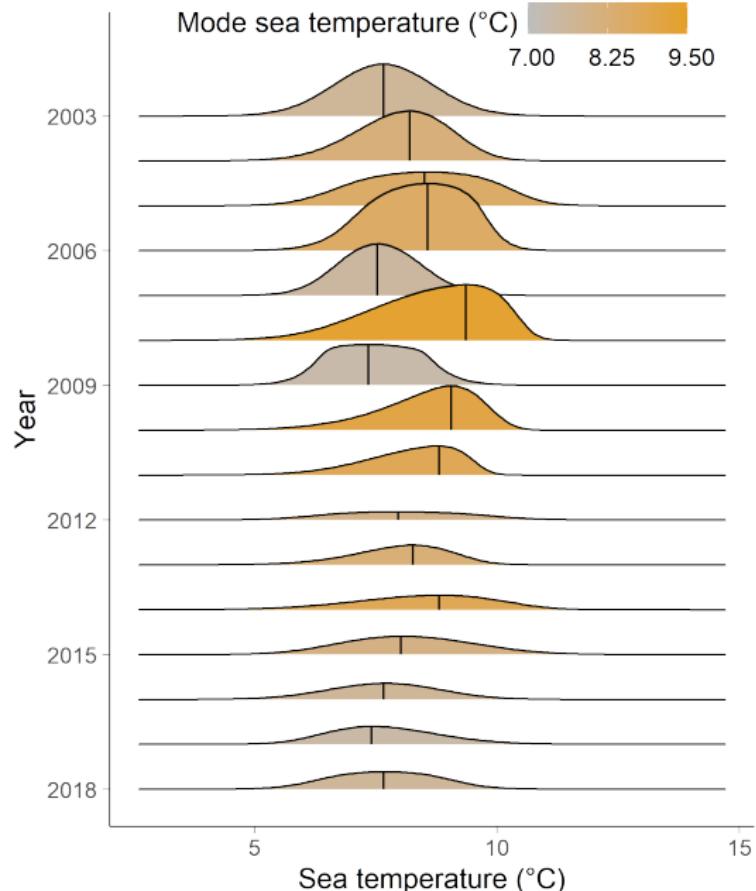
Merluccius productus (pacific hake) + lat



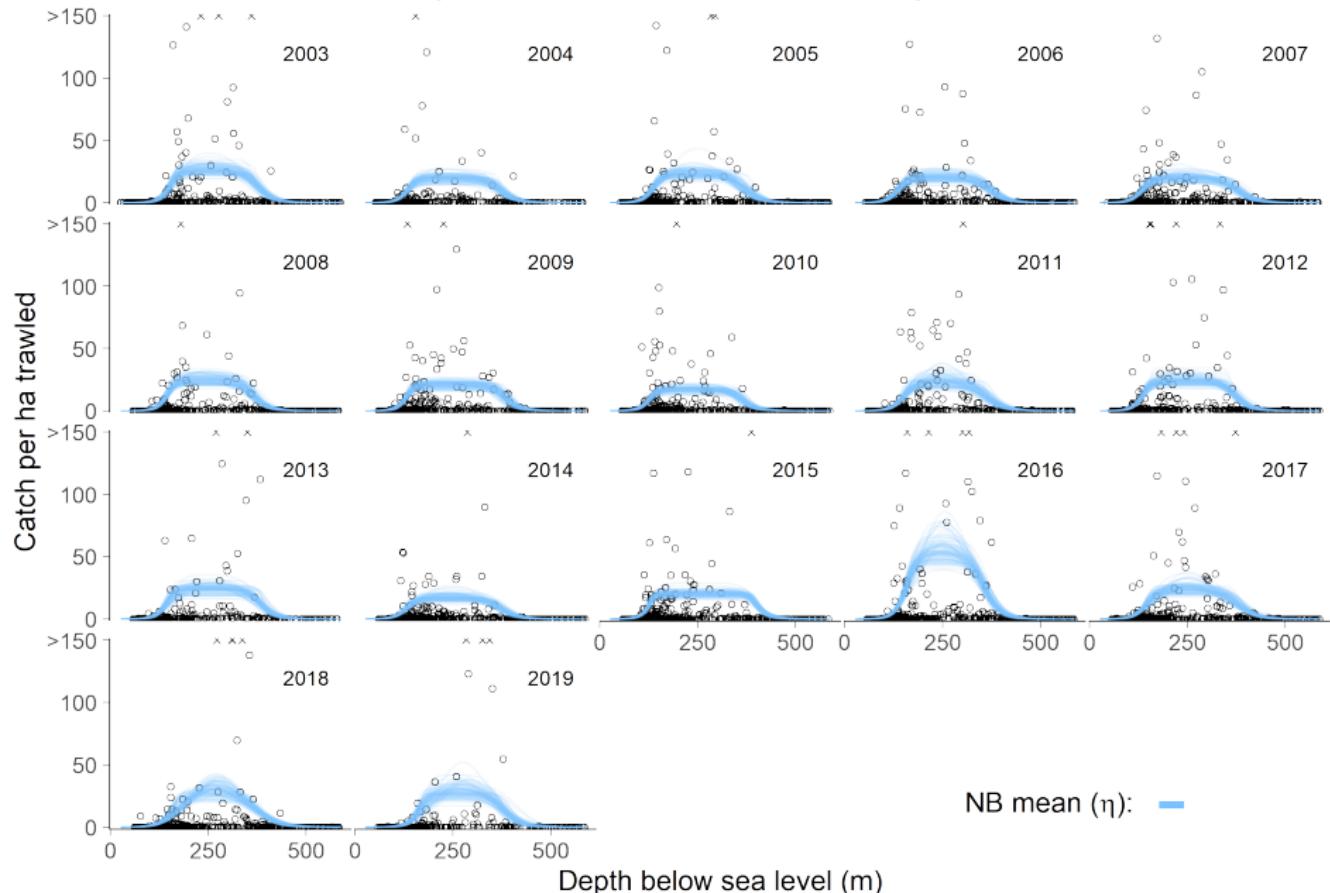
Merluccius productus (pacific hake) + temp



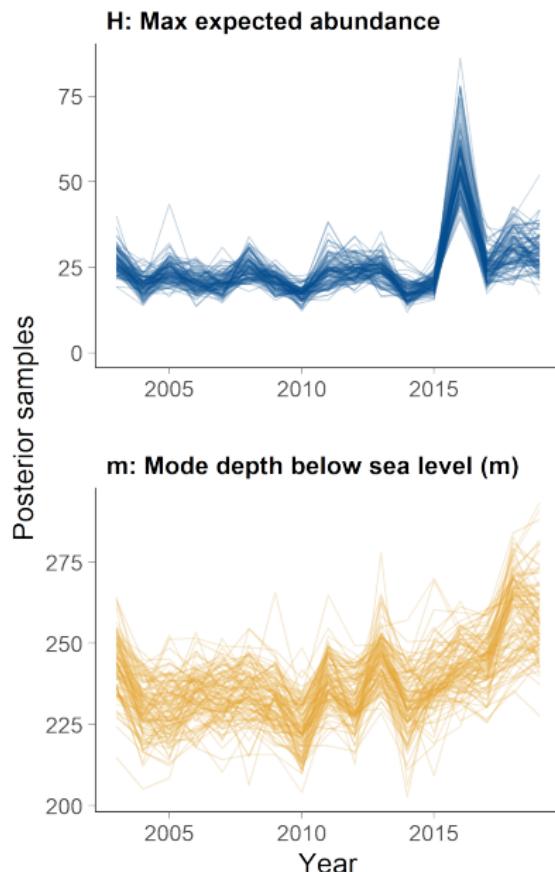
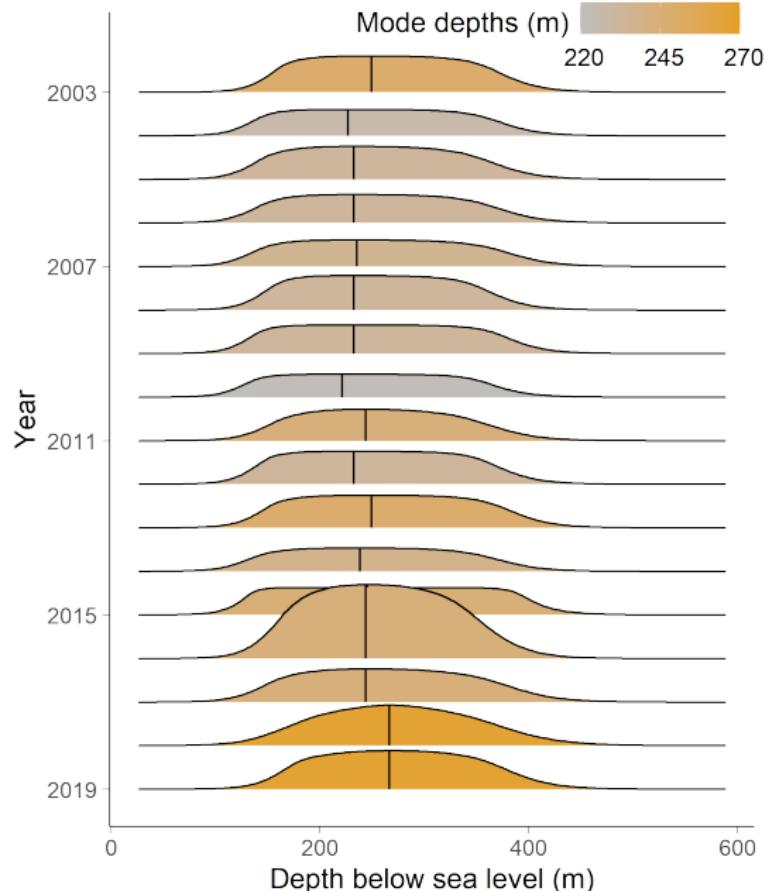
Merluccius productus (pacific hake) + temp



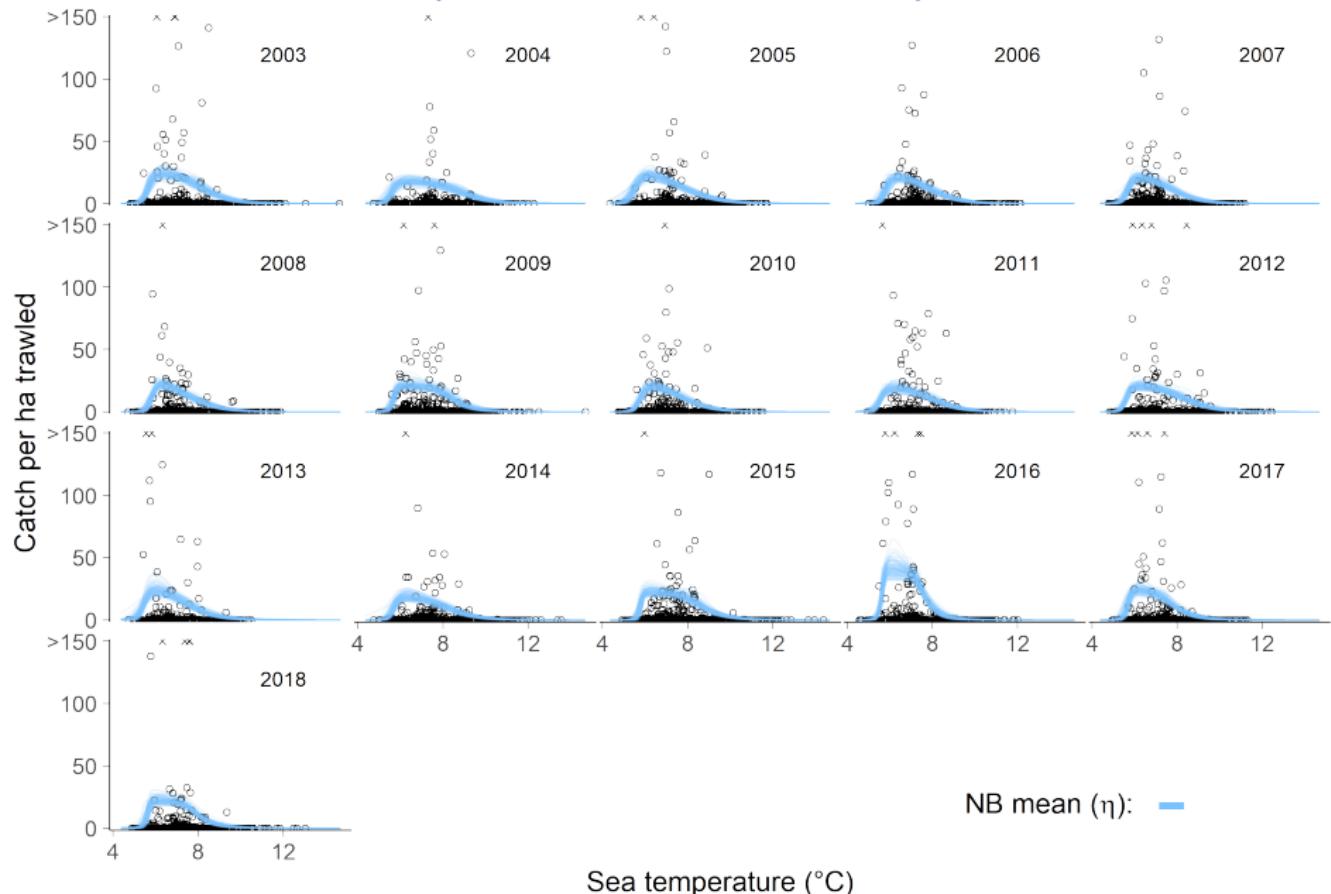
Sebastes crameri (darkblotched rockfish) + lat



Sebastodes crameri (darkblotched rockfish) + lat



Sebastodes crameri (darkblotched rockfish) + temp



Sebastodes crameri (darkblotched rockfish) + temp

