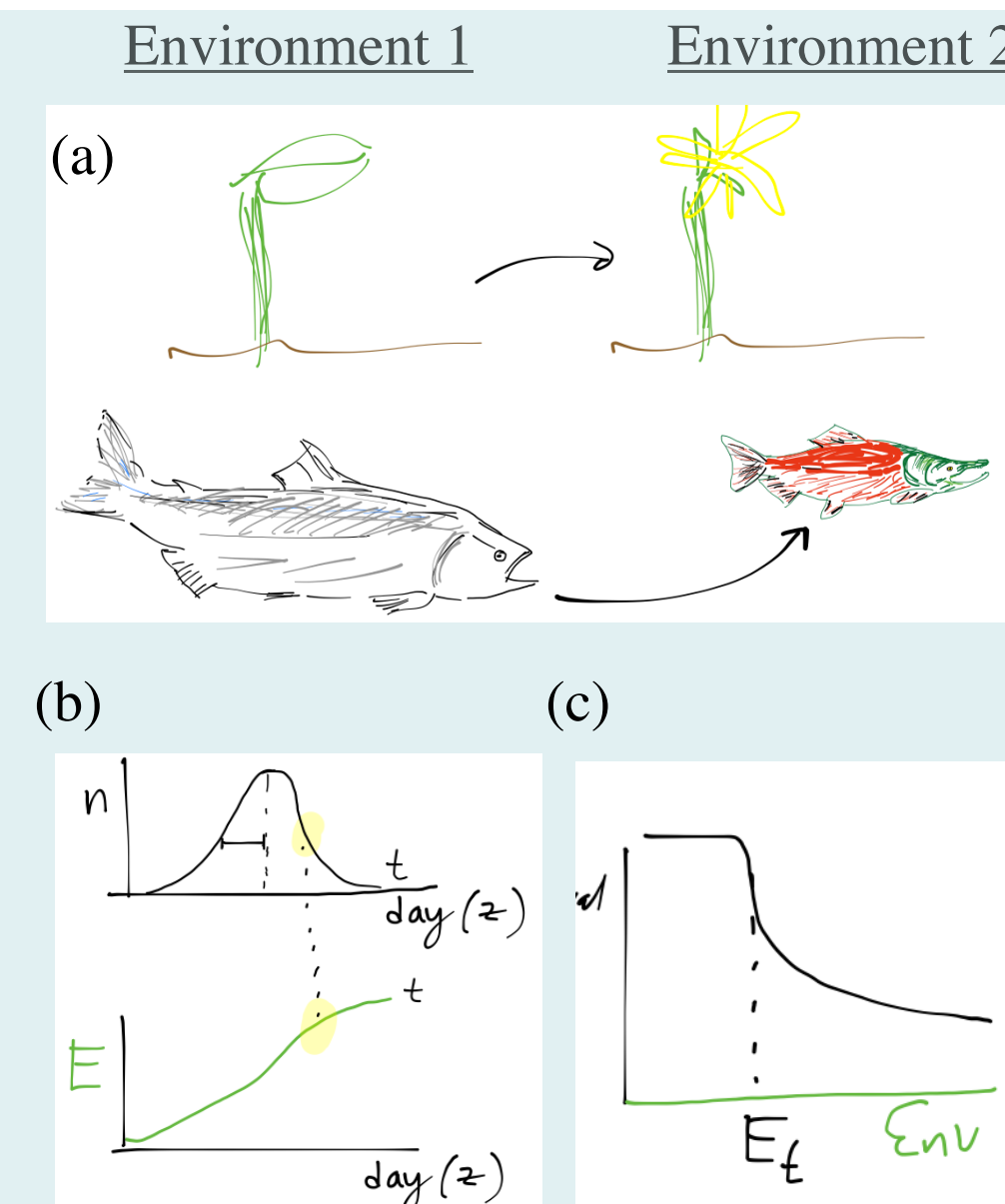


Understanding the joint effects of plastic and evolutionary change on demography from time series



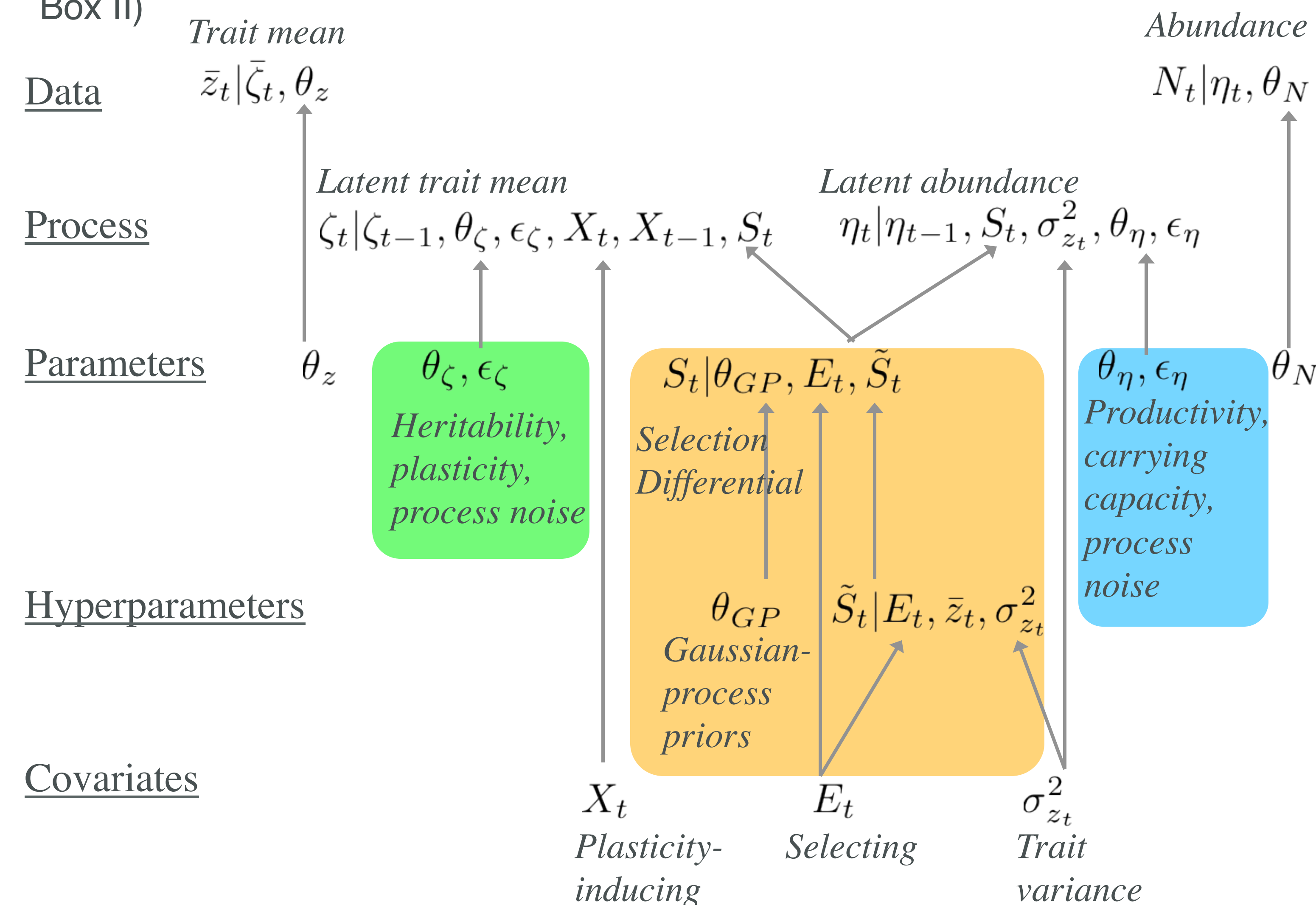
Motivation

Traits that govern life-history timing often respond plastically to aspects of the environment (a). Seasonal trends (b) mean such traits also determine the environment experienced. An environment that imposes differential mortality (c) causes selection on timing, potentially affecting demographic trajectories. To forecast responses to environmental change we wish to **predict population and trait dynamics**, and to do so we must **leverage prior studies** to help us **reduce uncertainty concerning the interaction of plasticity and selection** in our focal system.



A Bayesian approach to eco-evo dynamics

Accommodate prior estimates of **heritability**, **plasticity**, and/or **demography**, while admitting uncertainty about “backcasting” [e.g., 1,4] **selection gradients** by assuming the true gradient is some smooth function of the environment (see Box II)

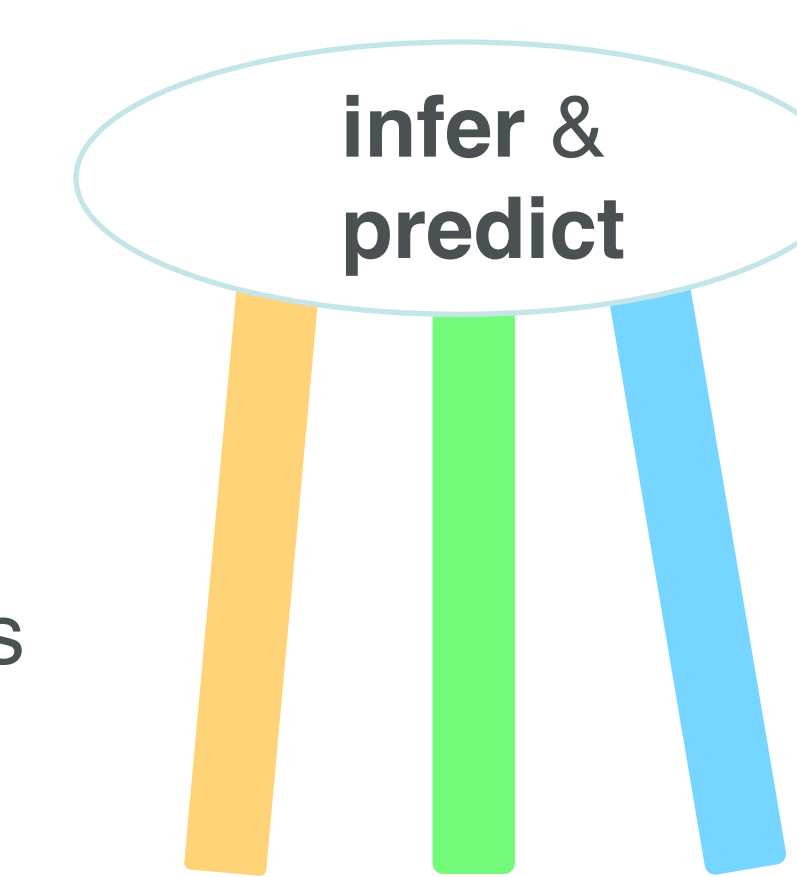


Methods

- State-space model in Bayesian framework implemented and sampled using MCMC via Hamiltonian Monte Carlo sampler in `stan` (Gelman *et al.* [2])
- Assumed Gaussian error (i.e., Kalman filter)
- Priors used
 - Heritability of focal trait h^2
 - Population dynamic parameters $\theta_\eta := \lambda, K$
 - Reconstructed selection differentials $S_t | \theta_{GP}, E_t, \tilde{S}_t$

Applying the framework

Population and trait dynamics are influenced by three interdependent processes: selection, response to selection and plastic change \rightarrow prior information on two of these can help us **infer** the action of the third, and thus **predict** the trajectories of interest.



I: heritability v plasticity

- Strong priors on population dynamics
- Assume backcast selection gradients reflect actual selection

$$\theta_\eta := \lambda, K$$

$$S_t = \tilde{S}_t \quad \forall t$$
$$\tilde{S}_t = f(E_t, \bar{z}_t, \sigma_{z_t}^2)$$

Outcomes

- Assess plasticity versus selection h^2 **b**
- Determine most-explanatory covariates X_t
- Population predictions provide additional data for predictive inference N_t

II: past selection by environment

- Strong priors on population dynamics
- Strong priors on evolutionary processes
- Variation in confidence of backcast selection

$$\theta_\eta := \lambda, K$$

$$\theta_\zeta := h^2, \mathbf{b}$$

$$\tilde{S}_t = f(E_t, \bar{z}_t, \sigma_{z_t}^2)$$

$$S_t = \tilde{S}_t \quad \text{for } t \in \Omega_N$$

$$S_t | \theta_{GP}, E_t, \tilde{S}_t \quad \text{for } t \notin \Omega_N$$

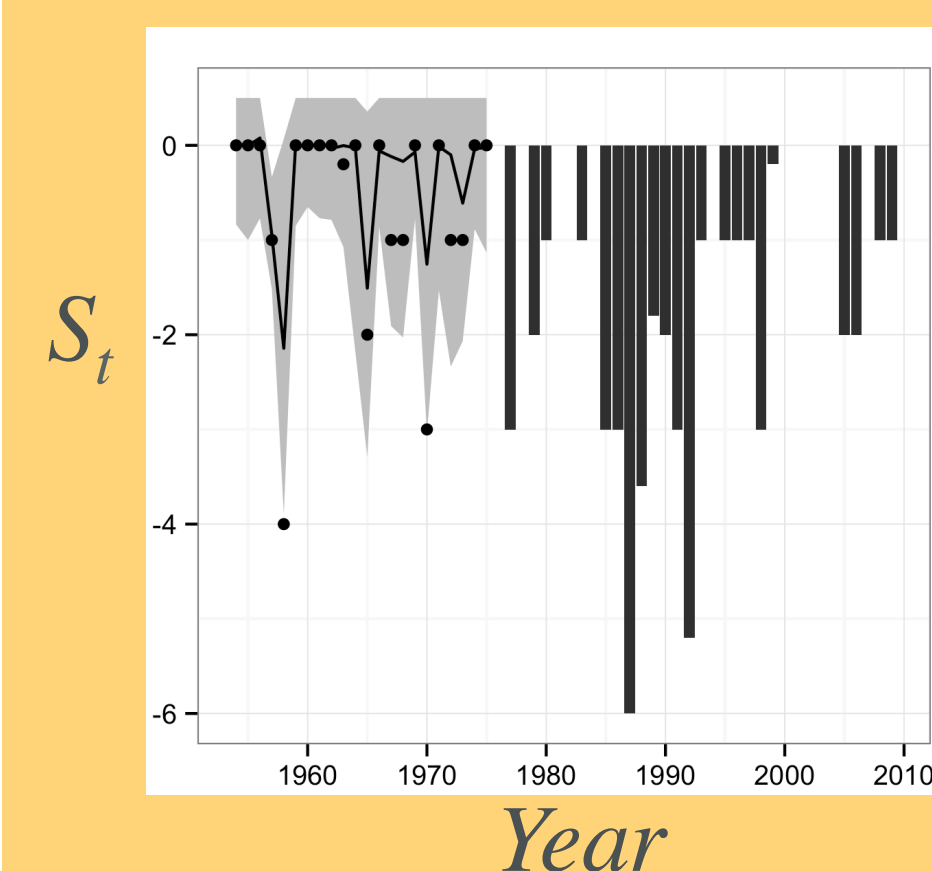


Figure: Prior to 1976 selection differential S_t is modeled as a Gaussian Process based on the relationship between selection and the environment from 1976-2009.

Outcomes

- Understand selection by the environment S_t over time
- Determine most-explanatory covariates X_t



Case study: Sockeye (*O. nerka*) in the Columbia River

- Strong prior on population dynamic parameter for growth, assuming Ricker form (Myers *et al.* [3])
- Backcast selection differentials based on temperature-survival relationship (after Crozier *et al* [1])

Outcomes

- Prediction for population dynamics given out-of-sample environment
- Re-ranking models for best predictors **b** X_t

	Model	WAIC [5]	dWAIC	Crozier et al. Rank
1	S+F4+Umay	268.29	0.00	4
2	S+F4+PDO4	284.97	16.68	1
3	S+F4+Umar	294.01	25.72	5
4	S+F4+NPI4	296.67	28.38	6
5	S+F4	303.03	34.73	3
6	S+F4+NPGO4	307.17	38.88	2
7	S+PDO4+NPGO4	316.19	47.89	7

Table: Model scores based on WAIC [5] for a selection of covariates used to predict migration date and abundance. S=selection, F=mean June flow, NPGO/PDO=oceanographic indices, U=upwelling indices. The same covariates were considered in Crozier *et al.* [1], but as shown, including population in the model results in a different ordering of the models.

Acknowledgements

- Special thanks to Lisa Crozier for encouragement. Discussions with Marissa Baskett, my PhD supervisor, have been instrumental in developing these ideas. Baskett Lab for feedback, and Richard McElreath and Michael Turelli for advice. Funding from NSF REACH IGERT at UC Davis, Center for Population Biology, ESP Travel Award to J.A. and Hellman Foundation Award to M.B. Cluster time from M.B. account on FARM.

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