

Climate fidelity and genetic variation in mountain landscapes

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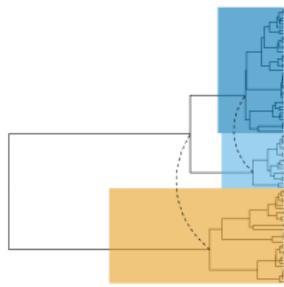
Climate structures biodiversity



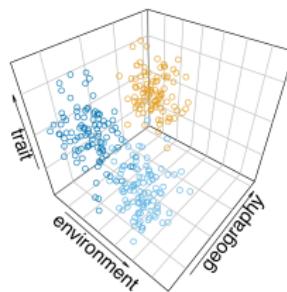
Climate fidelity: the tendency of a species or population to track its climatic niche through time (Wang et al. 2023 *PNAS*)

Integrative population genetics

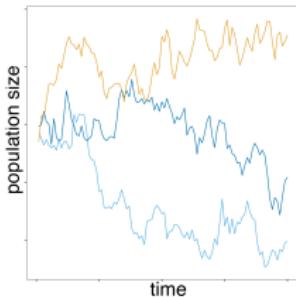
genetic differentiation



species range evolution



global change biology



Linck et al. 2020 *J. Evol. Biol.*

Linck et al. 2023 *Am. Nat.*

Linck et al. 2021 *Biol. Lett.*

Linck et al. 2019 *Syst. Biol.*

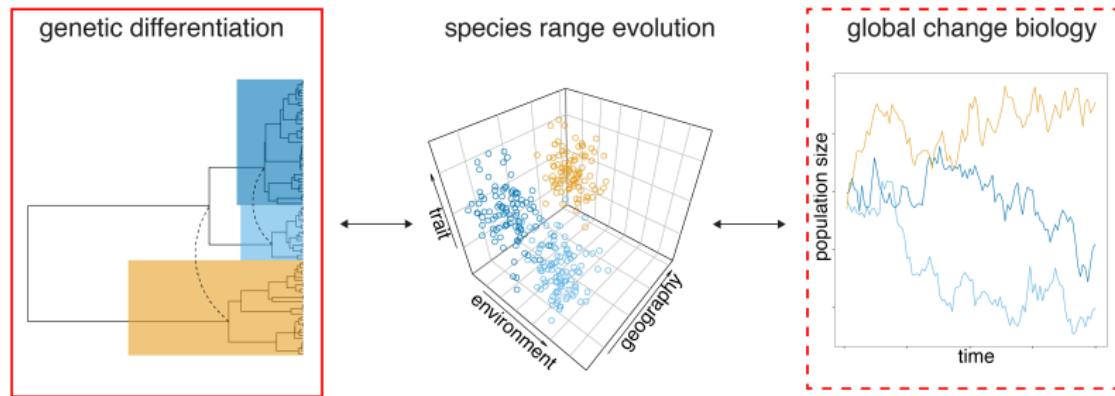
Linck et al. 2020 *Ecol. Evol.*

Mamantov et al. 2023 *Glob. Ecol. Biogeogr.*

Linck & Battey 2019 *Mol. Ecol. Res.*

Battey, Linck et al. 2018 *Am. Nat.*

Integrative population genetics



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How do mountains and climate fidelity
act to structure genetic variation?

What, if anything, can we do with this information on a burning planet?

Outline

- ▶ Q_1 : How has climate fidelity shaped genetic variation in Western *Empidonax* flycatchers?
- ▶ Q_2 : What can we learn from adding climatic complexity to population genetic models?
- ▶ Q_3 : How can genomics help us predict vulnerability to climate change?
- ▶ Q_4 : What does any of this mean for conservation?

Outline

- ▶ **Q_1 : How has climate fidelity shaped genetic variation in Western *Empidonax* flycatchers?**
- ▶ Q_2 : What can we learn from adding climatic complexity to population genetic models do?
- ▶ Q_3 : How can genomics help us predict vulnerability to climate change?
- ▶ Q_4 : What does any of this mean for conservation?

Acknowledgements:

- ▶ **Collaborators:** Kevin Epperly, Paul van Els, Garth Spellman, Rob Bryson, John McCormack, **Ricardo Canales-del-Castillo**, John Klicka, **Alex Hopping**
- ▶ **Funding:** Department of Defense NDSEG Fellowship

Climatic cycles shape biodiversity

GLACIATION, A FACTOR IN SPECIATION

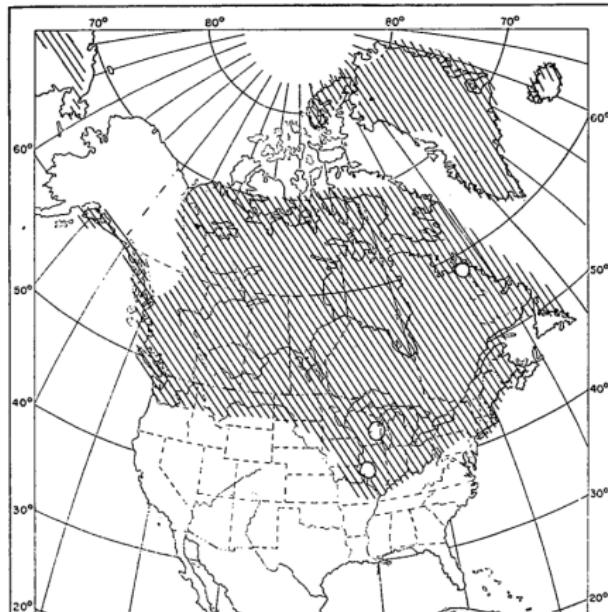
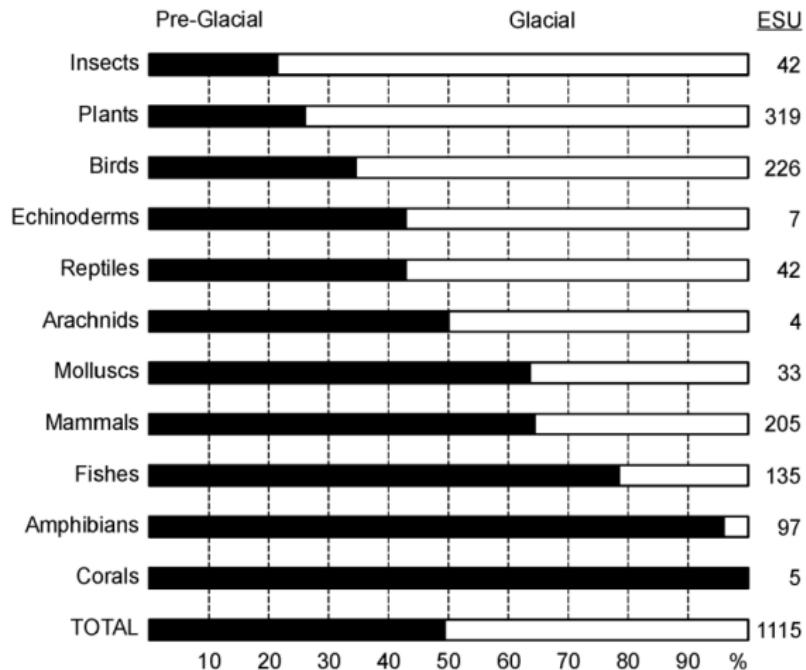


FIG. 1. Maximum extent of solid ice sheets according to Hulten, 1937; additional local glaciation also occurred. Flint (see fig. 2) believes that glaciation was more extensive than this.

(Rand 1948 *Evolution*)

Climatic cycles shape biodiversity



(Rull 2008 *Molecular Ecology*)

“Western” *Empidonax* Flycatchers



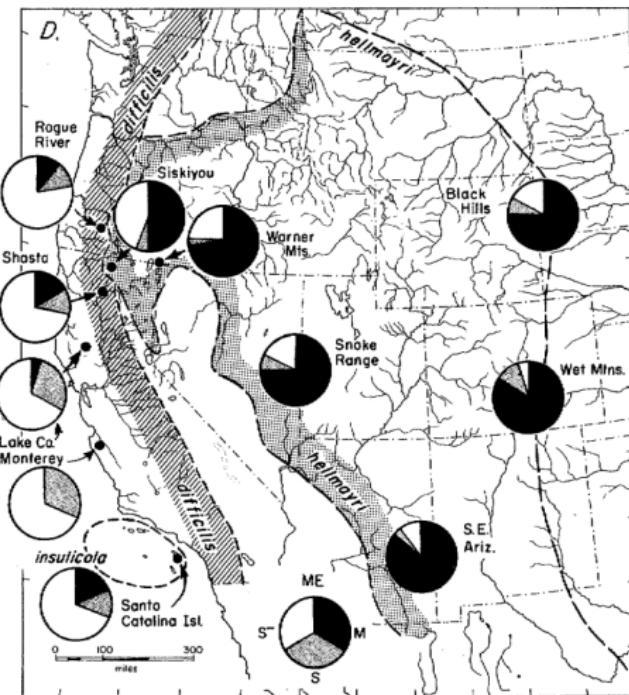
Cordilleran Flycatcher (COFL)
Empidonax occidentalis



Pacific-slope Flycatcher (PSFL)
Empidonax difficilis

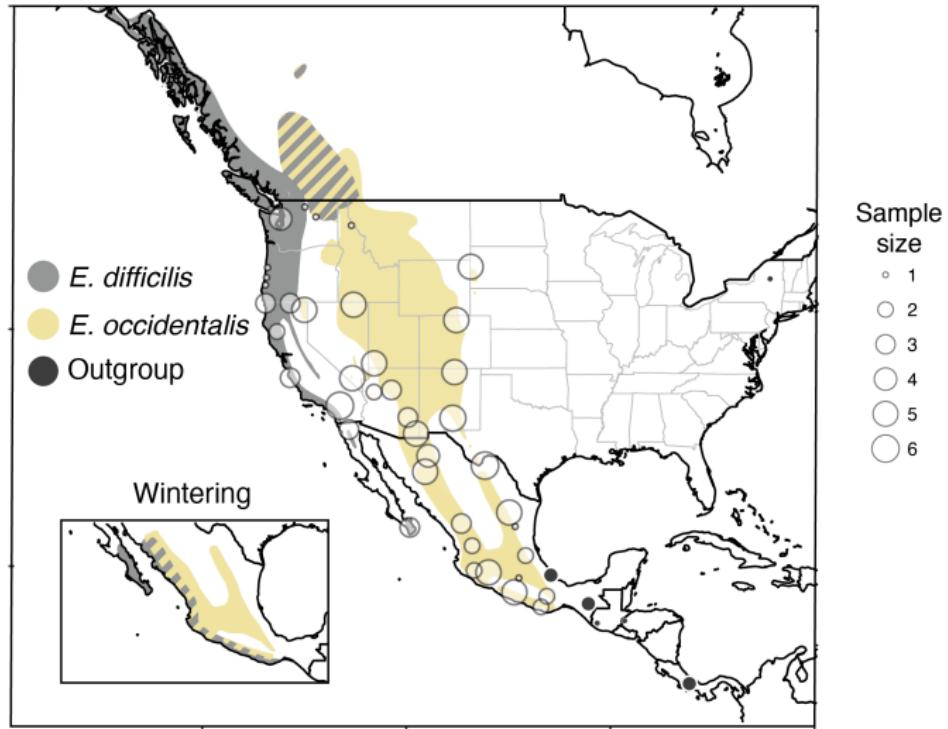
(Macaulay Library 109315481 & 236799211)

One species or two?



(Johnson & Marten 1988 *Auk*)

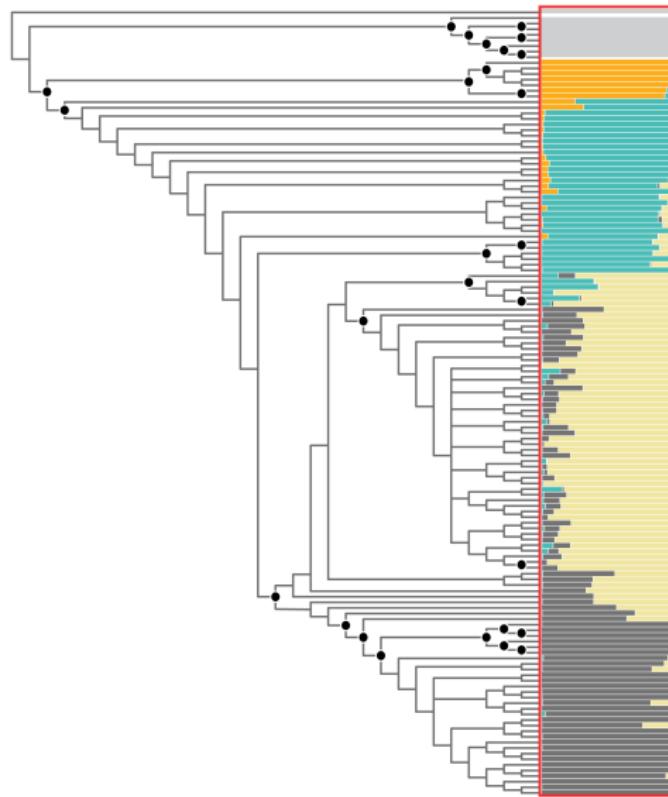
New information, more genes, more geography



(Linck et al. 2019 *Syst. Biol.*)

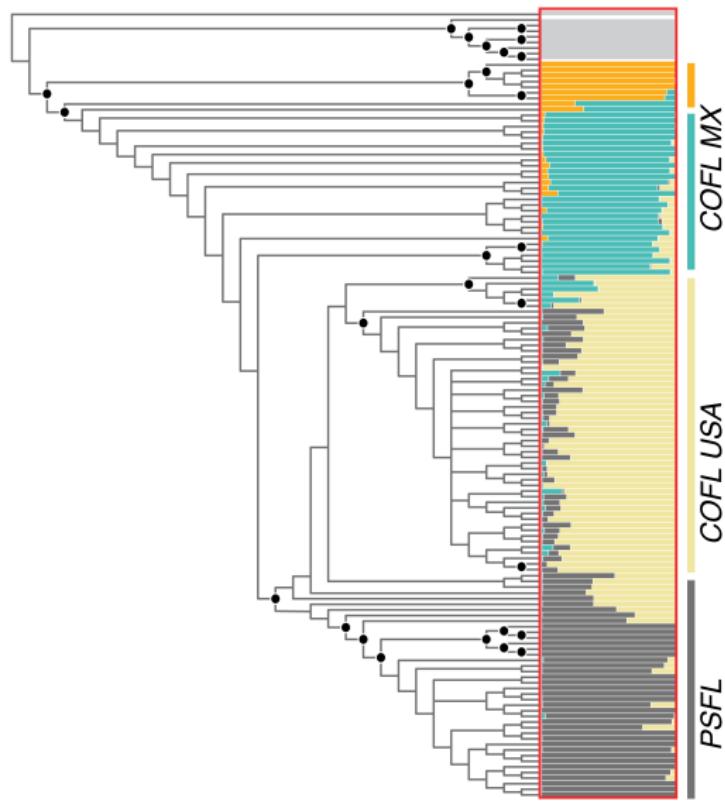
H_1 : Mountain ranges structure genetic variation in Western *Empidonax*

Geographically structured populations



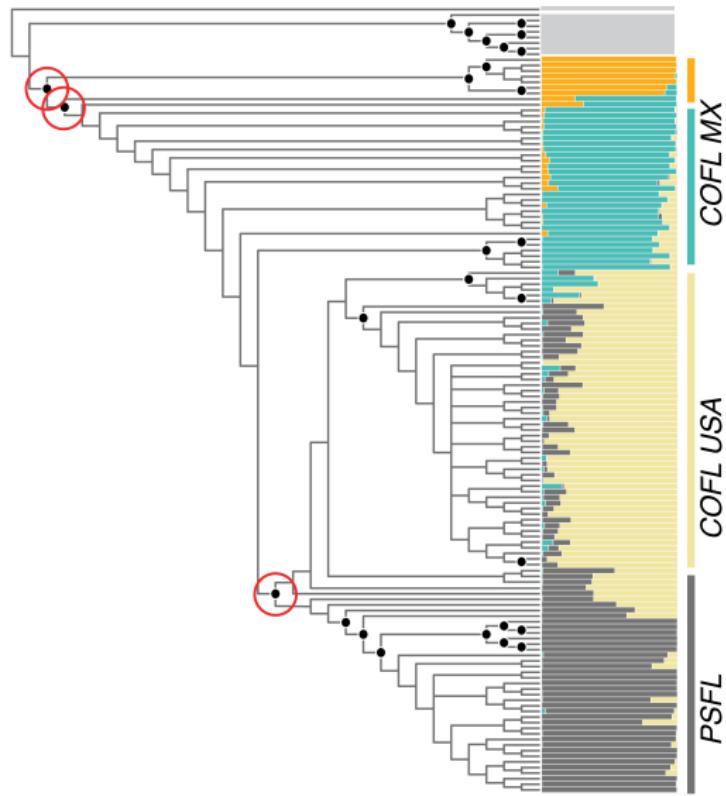
(Linck et al. 2019 *Syst. Biol.*)

Geographically structured populations



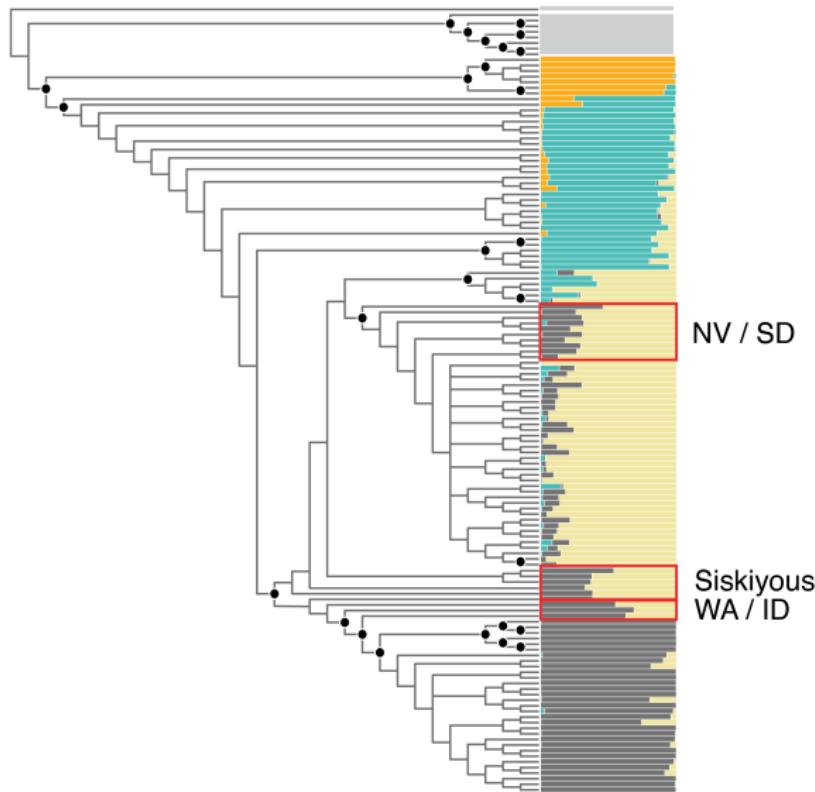
(Linck et al. 2019 *Syst. Biol.*)

COFL is paraphyletic!



(Linck et al. 2019 *Syst. Biol.*)

Hybridization at forest corridors



(Linck et al. 2019 *Syst. Biol.*)

Takeaways

- ▶ 1 species of WEFL (for now)
- ▶ Hybridization is key
- ▶ Climate fidelity and montane geography explain genetic structure in *Empidonax* flycatchers
- ▶ **Application:** There's no getting around comprehensive sampling for delimitation of management units

Outline

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- ▶ Q_2 : **What can we learn from adding climatic complexity to population genetic models?**
- ▶ Q_3 : How can genomics help us predict vulnerability to climate change?
- ▶ Q_4 : What does any of this mean for conservation?

Acknowledgements:

- ▶ **Collaborators:** CJ Battey, Ana Bedoya
- ▶ **Funding:** NSF DDIG #1701224

Climatic cycles shape biodiversity

GLACIATION, A FACTOR IN SPECIATION

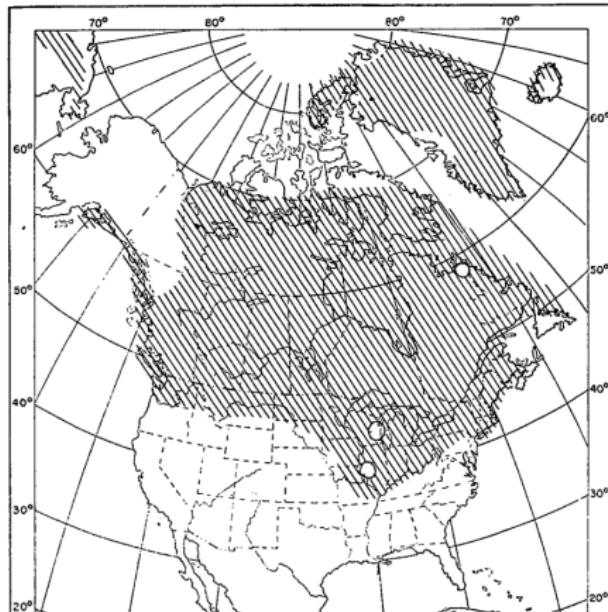
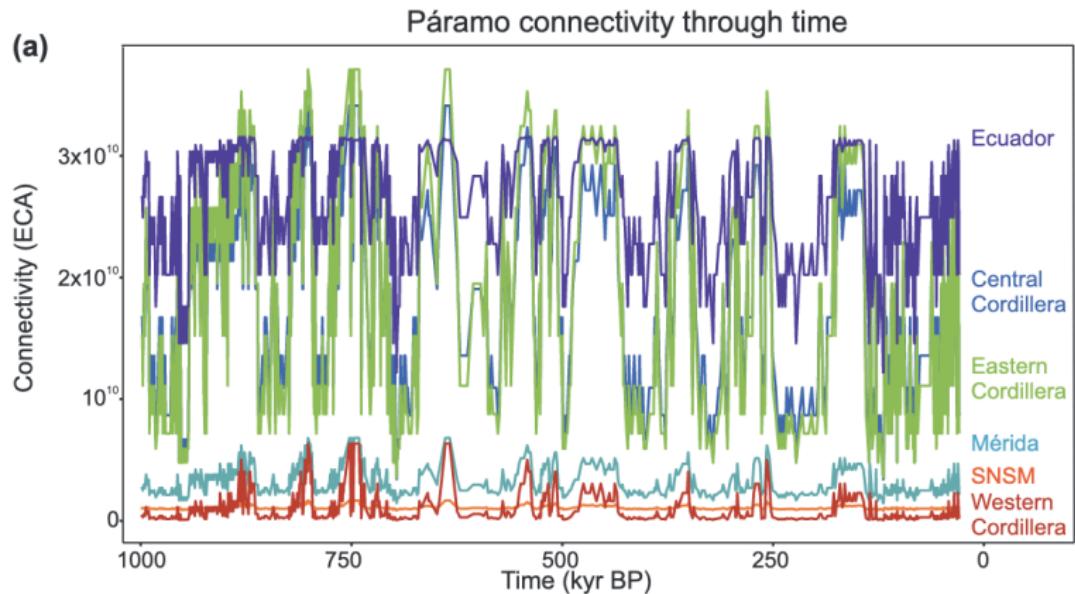


FIG. 1. Maximum extent of solid ice sheets according to Hulten, 1937; additional local glaciation also occurred. Flint (see fig. 2) believes that glaciation was more extensive than this.

(Rand 1948 *Evolution*)

Climate fidelity and cyclical contact



(Flantua et al. 2019 *J. Biogeog.*)

Simple models of divergence

Geographic modes of speciation

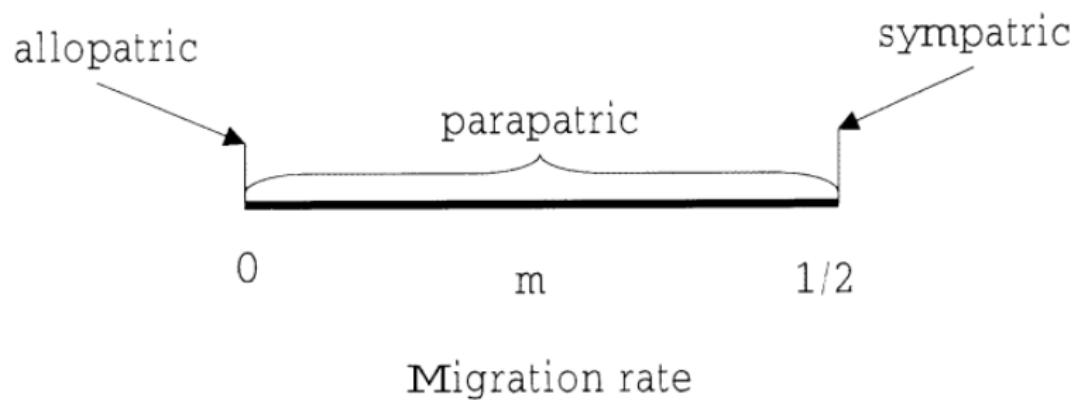


FIG. 1. Geographic modes of speciation.

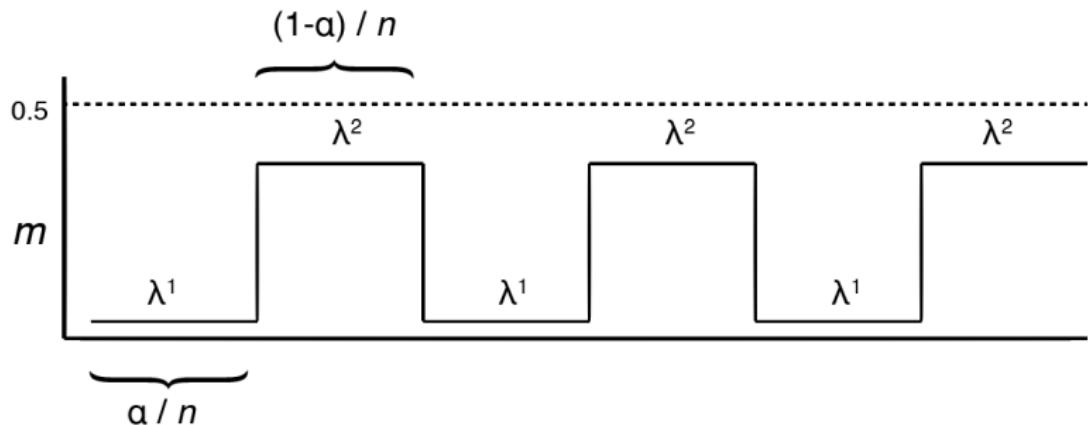
(Gavrilets 2003 *Evolution*)

For example: allopatric speciation

$$\text{time to speciation} \quad \overbrace{T^a} = \underbrace{\frac{2}{\text{no. alleles}}} * \overbrace{\frac{1}{\mu}}^{\text{substitution rate}} = \frac{2}{\mu}$$

H_1 : Cyclical contact slows divergence,
but not *that* much

Our own toy model



(Linck & Battey In revision *Am. Nat.*)

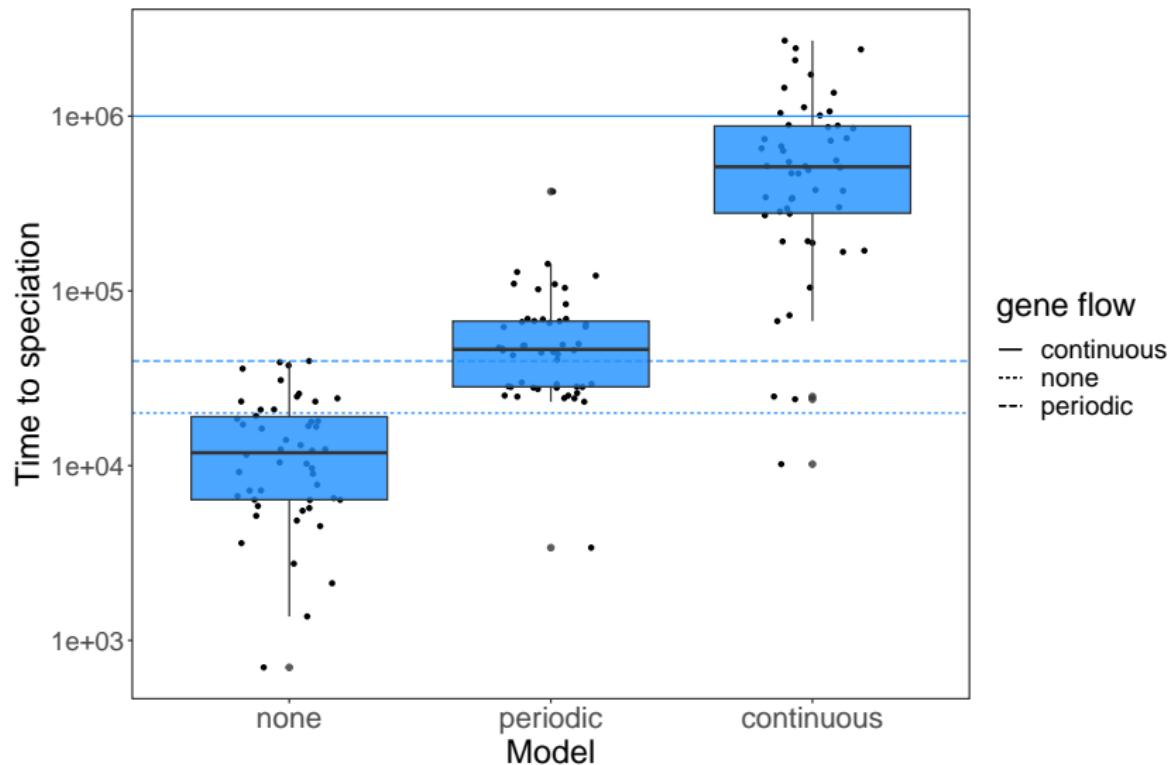
Our own toy model

$$\widehat{T}^t = \frac{\text{time to speciation}}{\sum_{i=1}^n \underbrace{\alpha}_{\text{time in allo.}} \underbrace{\left(\frac{\mu}{2}\right)}_{\text{allo. model}} + \underbrace{(1 - \alpha)}_{\text{time in para.}} \underbrace{\left(\frac{\mu^2}{m}\right)}_{\text{para. model}}}$$

number of climate cycles
 \widehat{n}

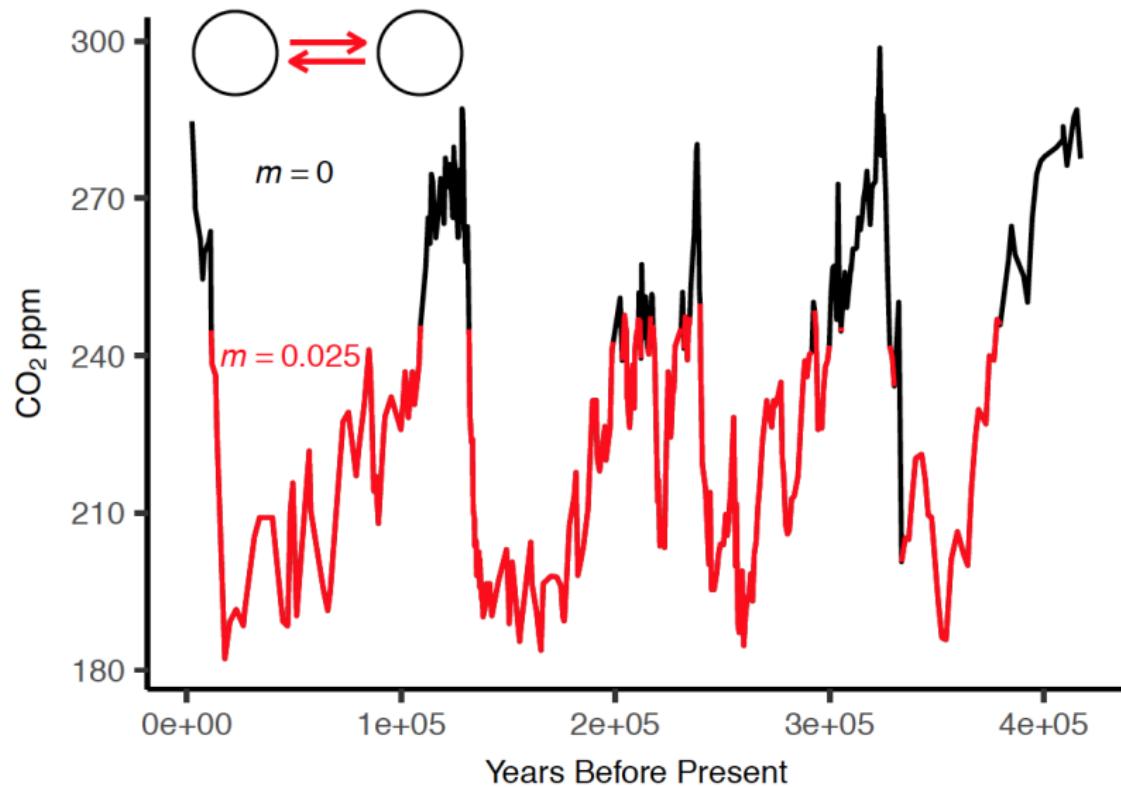
Our own toy model

$$\alpha = 0.5, \mu = 1e-4, m = 0.01$$

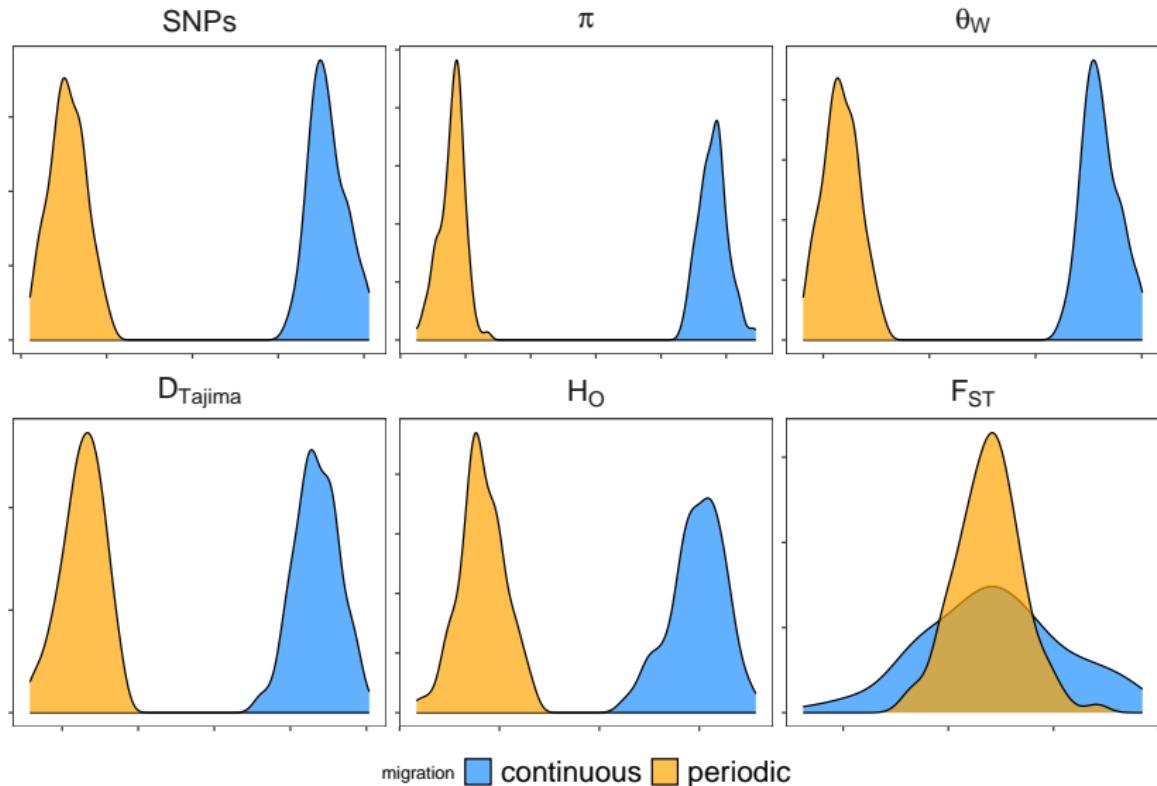


H_2 : Cyclical contact has a distinct genomic signature

Parameterizing simulations



Different summary statistics



Takeaways

- ▶ Climate fidelity involved in an efficient mode of divergence
- ▶ Potentially detectable with sequence data
- ▶ Climatic complexity adds value to population genetic models
- ▶ **Application:** Identifying corridors of intermittent connectivity on the landscape

Outline

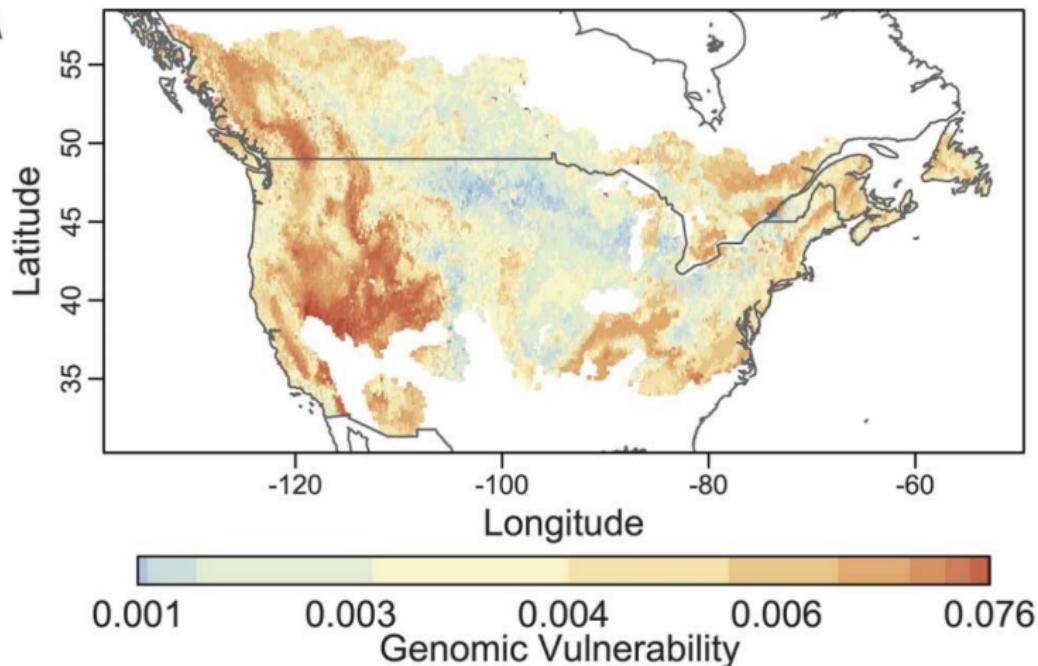
- ▶ Q_1 : How has climate fidelity shaped genetic variation in Western *Empidonax* flycatchers?
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- ▶ **Collaborators:** Daniel Cadena, Cameron Ghalambor, Ben Freeman, Gavin Jones, Laura Céspedes
- ▶ **Funding:** NSF PRFB

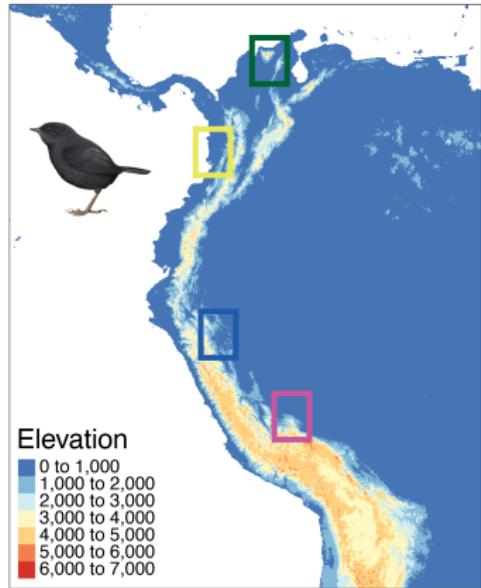
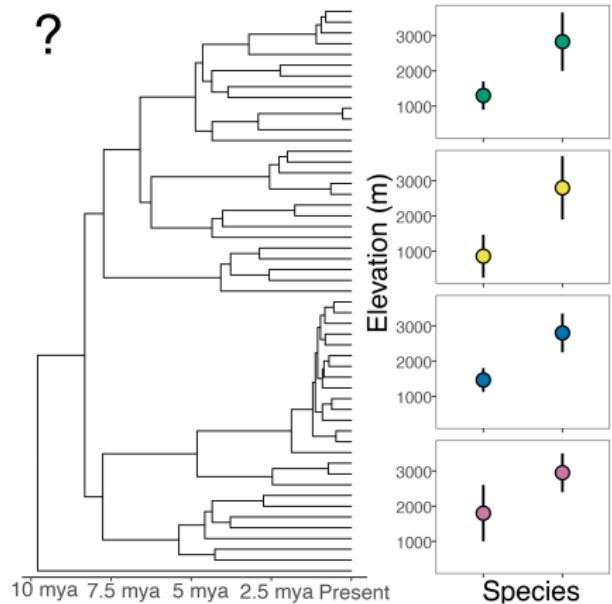
Move, **adapt** / acclimate, or die

A



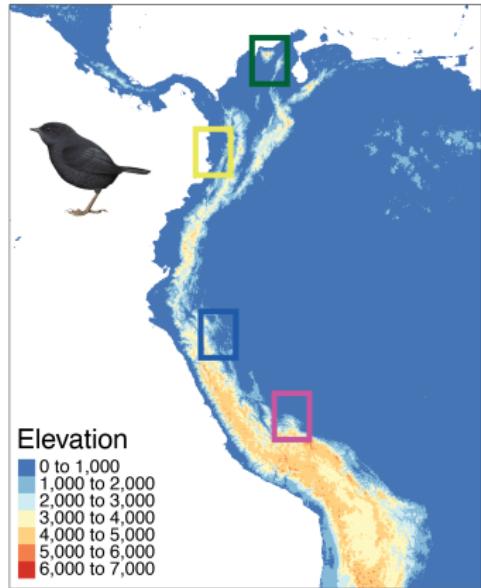
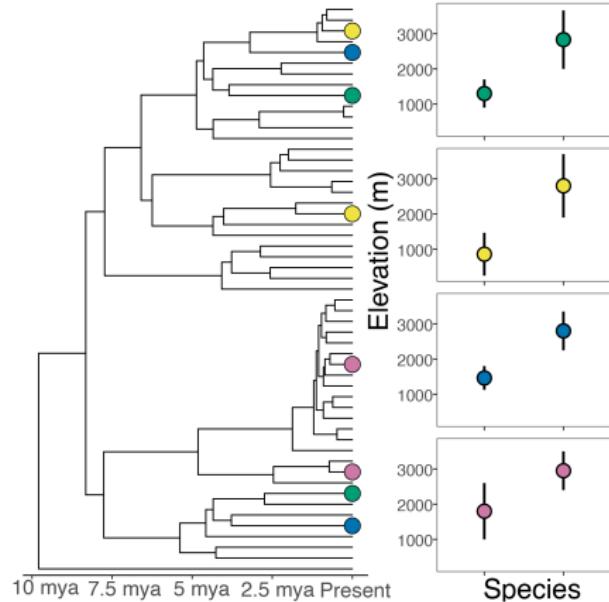
(Bay et al. 2018 *Science*)

How quickly do species adapt to new climates?



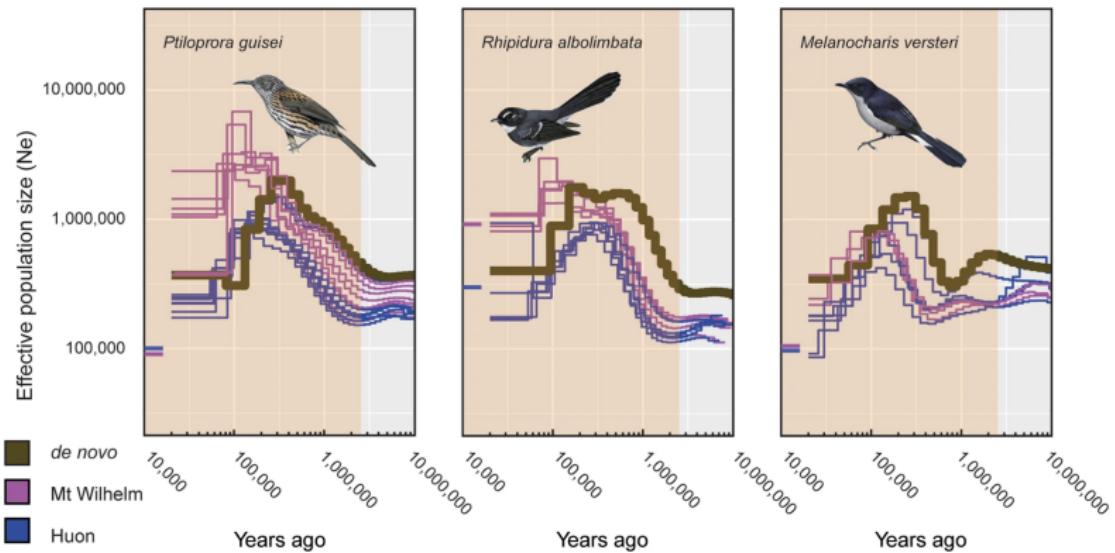
(Linck et al. 2021 *Biol. Lett.*)

Climate fidelity across phylogeny in tapaculos



(Linck et al. 2021 *Biol. Lett.*)

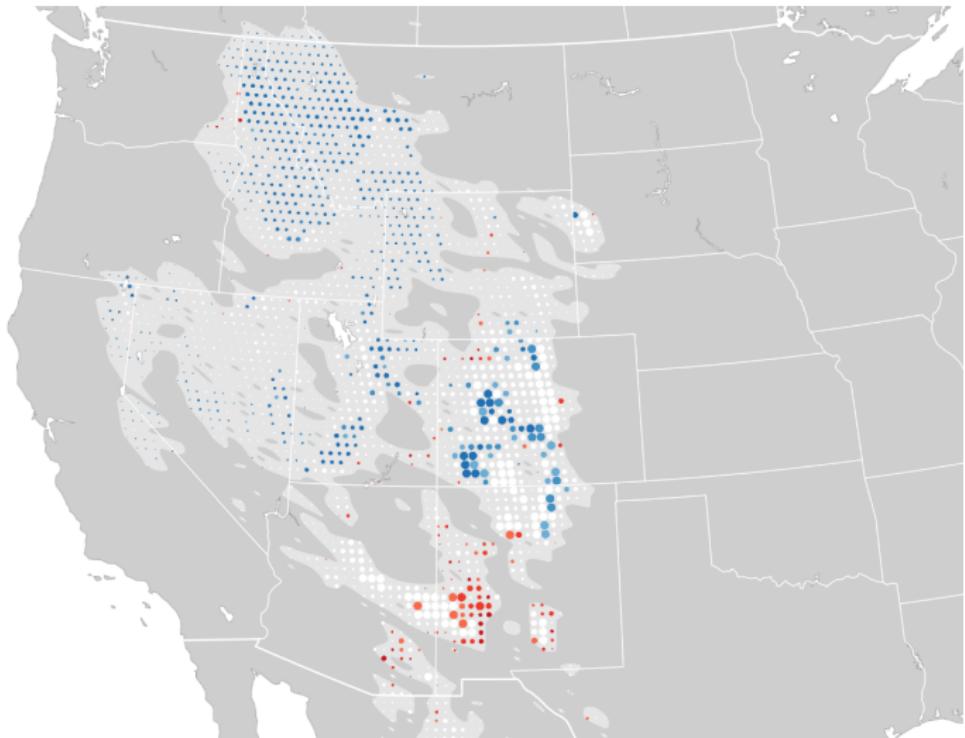
Climate fidelity shapes population trajectories



(Pujolar et al. 2022 *Nat. Comm.*)

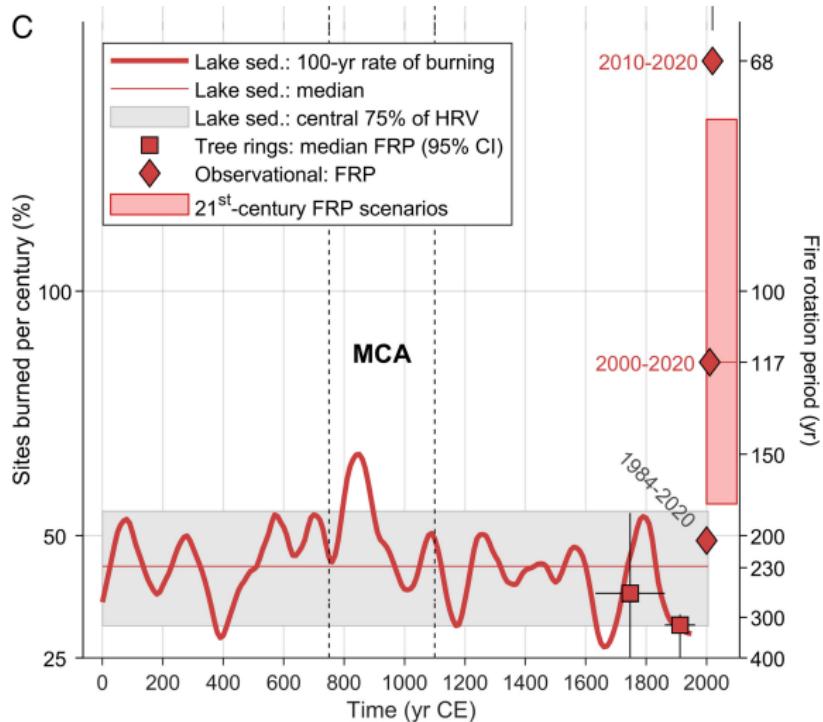
H_1 : Climate fidelity tells us about adaptive potential to future climates

Western forests host niches under threat



(eBird)

Megafires as evolutionary experiment



(Higuera et al. 2021 *PNAS*)

Future work: Demographic responses to climate change in subalpine forest taxa

- ▶ Aim 1: What do we know about animal evolution in the “Pyrocene” (Jones et al. In review *TREE*)
- ▶ Aim 2: Did the Medieval Climate Anomaly leave a genomic signature in subalpine birds and mammals?
- ▶ Aim 3: Can museum genomics reveal population impacts of recent megafires?
- ▶ Aim 4: **Application:** Do metrics of climate fidelity improve assessments of risk?

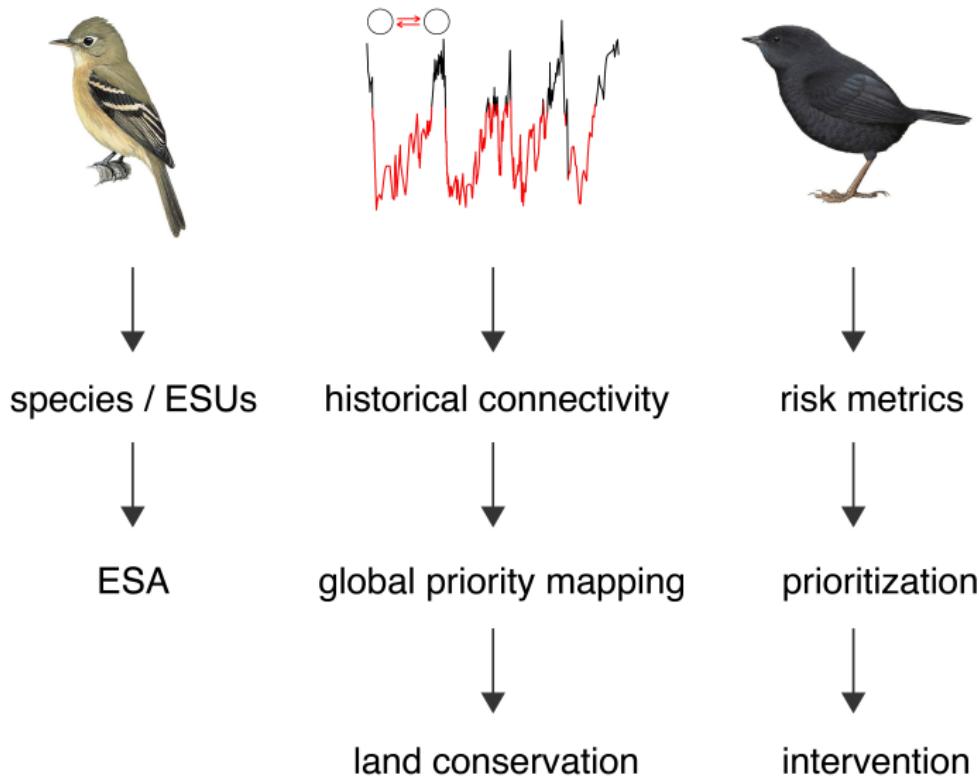
Takeaways

- ▶ Tapaculos adapt to new climates very slowly
- ▶ Climate fidelity shapes population trajectories
- ▶ Megafire is a natural experiment on genetic consequences of climate fidelity
- ▶ **Application:** The genomic signature of climate fidelity contains information for risk metrics and prioritization

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Conservation connects data to policy levers

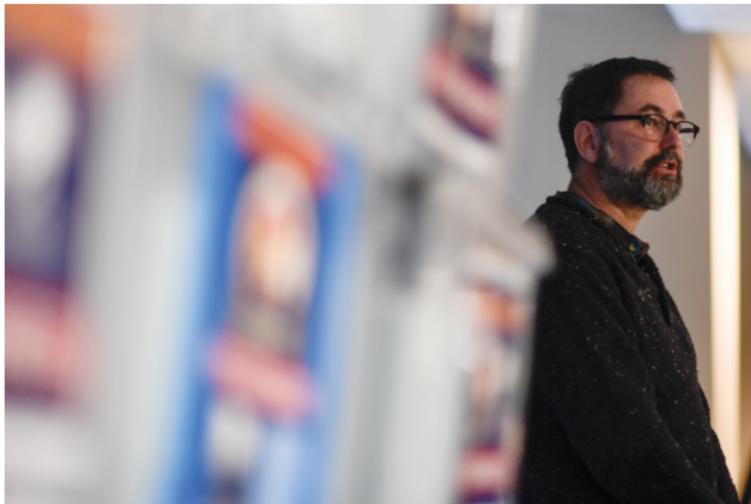


Conservation is mostly about people



(Schimel & Linck 2019 *bioGraphic*)

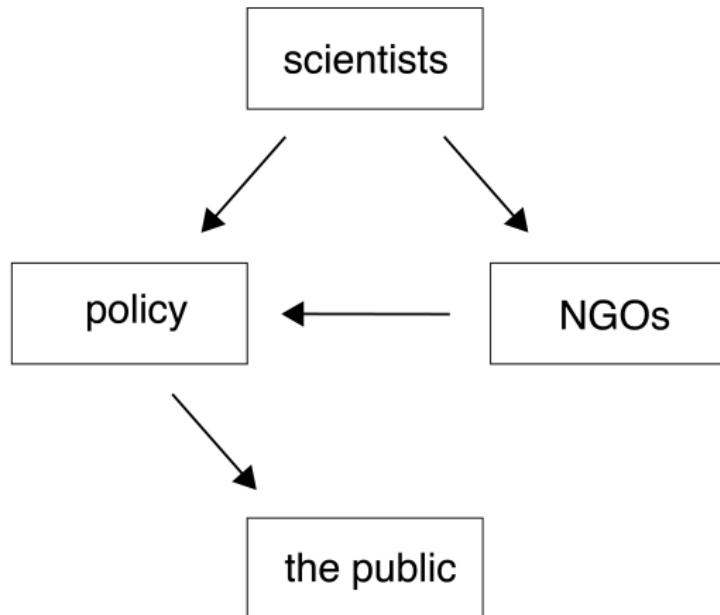
Values influence the interpretation of data



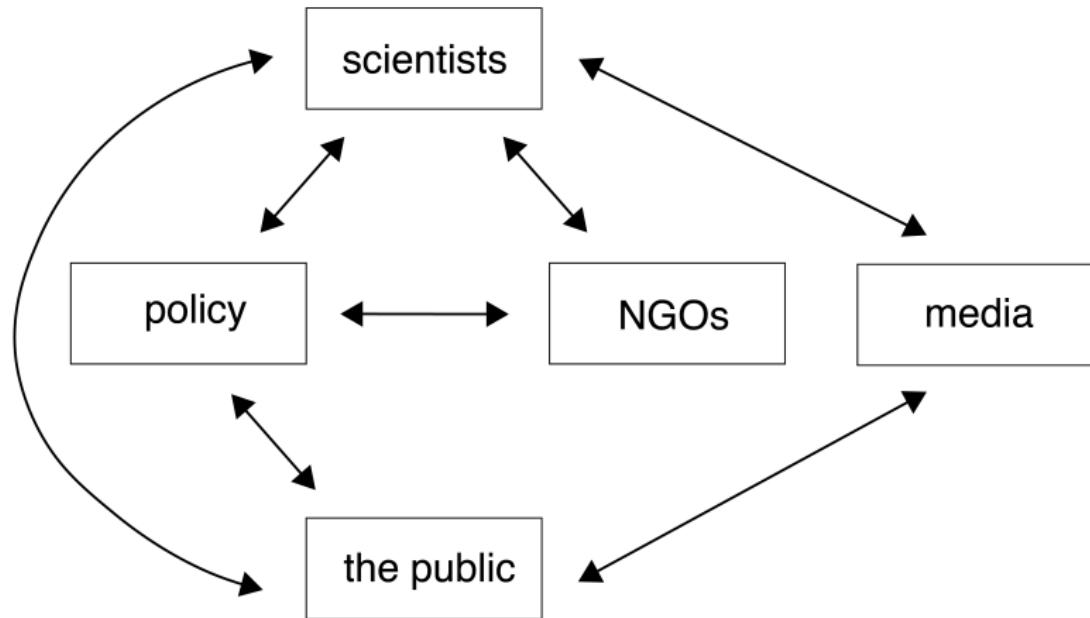
"They have wolves on the Gaza Strip. They have wolves in Italy. They have wolves in Northern-freaking-California. Why can't we have wolves here?"

(Rob Edwards of RMWP in Linck 2020, *High Country News*)

Conservation as a chain



Conservation as a network



Resources

- ▶ slides and code: https://github.com/elinck/misc_talks/
- ▶ website: <https://elinck.org/>
- ▶ twitter: @ethanblinck

References

- ▶ Rand 1948. *Evolution*, 2(4), 314-321.
- ▶ Rull 2008. *Molecular Ecology*, 17(11), 2722-2729.
- ▶ Johnson & Marten 1988. *The Auk*, 105(1), 177-191.
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- ▶ Bay et al. 2018. *Science*, 359(6371), 83-86.
- ▶ Linck et al. 2021. *Biology Letters*, 17(10), 20210363.
- ▶ Pujolar et al. 2022. *Nature Communications*, 13(1), 268.
- ▶ Higuera et al. 2021. *PNAS*, 118(25), e2103135118.
- ▶ Schimel & Linck 2020. *bioGraphic*.
- ▶ Linck 2020. *High Country News*.

Thanks!

- ▶ Q_1 : Climate fidelity and montane geography explain genetic structure in *Empidonax* flycatchers
- ▶ Q_2 : Climatic complexity adds value to population genetic models
- ▶ Q_3 : The genomic signature of past climate fidelity may contain information about future risk
- ▶ Q_4 : Conservation is a network, not a chain