Weighted Mean Occurrence From an Ensemble of Species Distribution Models

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Collaborators

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 - Jason Tack (HAPET)
 - Neal Niemuth (HAPET)
 - Kevin Barnes (HAPET)
- BCR
 - Chris Latimer
- ECCC
 - Barry Robinson
- PHJV
 - Jim Devries











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Why are spatial models important?

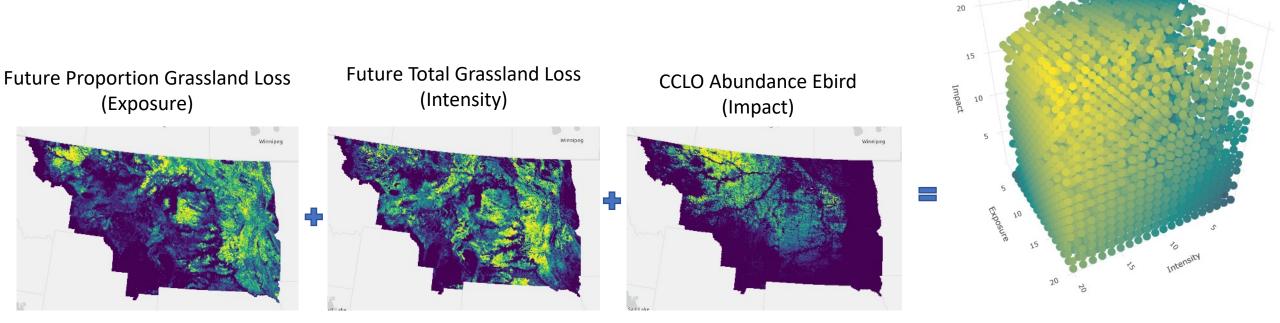
- Spatial information on biological value, threats, and cost are useful for prioritizing landscapes for conservation action
- Relationships identified by models provide direction for specific conservation actions such as <u>protection, restoration, and</u> <u>enhancement</u>



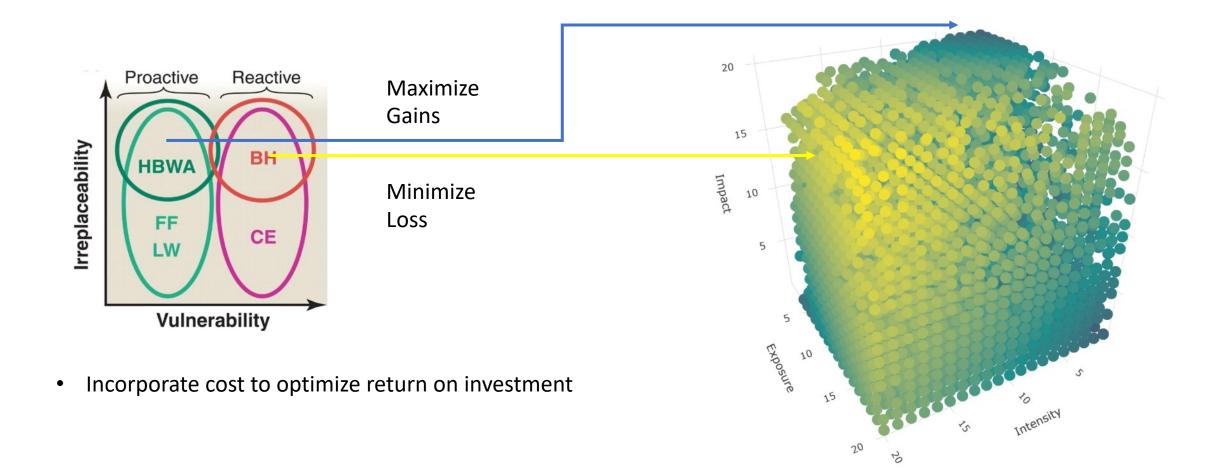


Example

- Wilson et al. (2005) defined vulnerability as having 3 dimensions:
 - Exposure= probability of threatening process
 - Intensity = magnitude threatening process
 - Impact = potential biological loss



Prioritize to meet your strategy!



Conservation planning

U.S. Fish & Wildlife Service and Prairie Pothole Joint Venture

A Full Annual-Cycle Conservation Strategy for Sprague's Pipit, Chestnut-collared and McCown's Longspurs, and Baird's Sparrow



Spraque's Pipit (top, left); Chestnut-collared Longspur (bottom, left); McCown's Longspur (top, right); Baird's Sparrow (bottom, right)

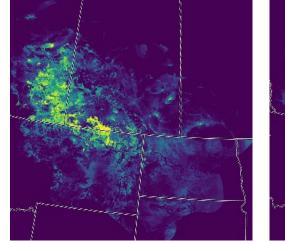
Conservation planning

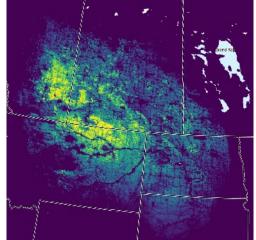
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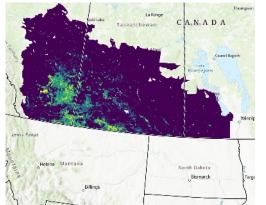
A Full Annual-Cycle Conservation Strategy for Sprague's Pipit, Chestnut-collared and McCown's Longspurs, and Baird's Sparrow



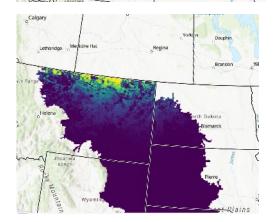
- Do we have models of biological value?
 - Sather et al.
 - Fink et al.
 - Robinson et al.
 - Niemuth et al.
 - Pavlacky et al.
 - Fields et al.
 - Fedy et al.
 - Etc.

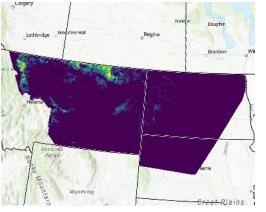






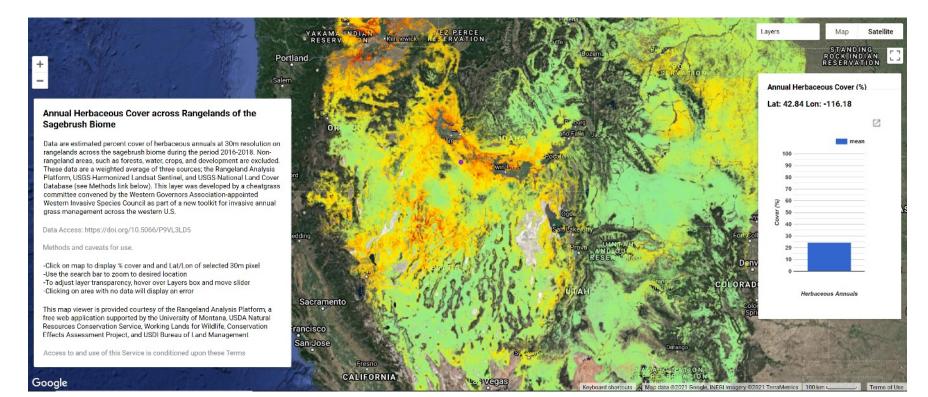






What have other done?

 Western Invasive Species Council developed a weighted average of three cheatgrass models to serve as a threat layer to better prioritize conservation actions.

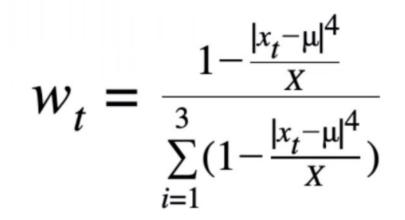


How did they do it?

- Assign weights to pixel values.
- A model's pixel is weighted less if it is further from the ensemble's mean pixel value, and weighted more if closer.

$$w_{t} = \frac{1 - \frac{|x_{t} - \mu|^{4}}{X}}{\sum_{i=1}^{3} (1 - \frac{|x_{t} - \mu|^{4}}{X})}$$

Weighting Expression:



$$X = \sum_{t=1}^{3} |x_t - \mu|^4$$

weighted mean $= \sum_{t=1}^{3} w_t x_t$

Example:

0.3				0.1436094675
Mean Pixel Value	Sum x-Mu ^4 (X)	Sum 1- x-Mu ^4 /		weighted Mean
0.7	8 0.04100625	0.1137390147	0.05686950737	0.04435821575
0.1	0.00194481	0.957967158	0.478983579	0.05747802948
0.0	0.00331776	0.9282938272	0.4641469136	0.04177322223
Pixel Value (Occurrence)	xt-Mu ^4	1- x-Mu ^4 / X	Weight	wtxt

Potential Issues?

What about different scales and extents?

Solution: resample to minimum mapping unit, use arithmetic mean when 2 available, use any prediction when only 1 available

What about varying state variables? (i.e. not true occurrence?) Solution: Could rescale values to common metric (std normal distribution? Scale to largest value observed across datasets?) Abundance to Occurence via Royle-Nichols equation.

*0 vs NA (e.g. BBS model NA in CAN)

Proof of concept: Sprague's Pipit



Source	Model	Prediction	Y~ Source	Landcover data	Landscape scales	Resolution	Spatial Extent
Sather et al.	Random Forest	Occurrence	Multiple (2007-2012)	ACI/NLCD	1-10000 m	160 m	Breeding Range
Fink et al.	adaSTEM	Occurrence	eBird (2019)	FAO	1500 m	~3 km	Breeding Range
Robinson et al.	Boosted Regression Tree	Density	Multiple (2009-2019)	ACI	800 m- 4000 m	800 m	PPR - Canada
Niemuth et al.	GLMM	Occurrence	BBS Stop (2005-2011)	NLCD	1200 m	30 m	<mt, nd,="" sd<="" td=""></mt,>
Fields et al.	GLMM	Density	BBS Stop (2008-2016)	NLCD	1600 m	30 m	<mt, nd,="" sd<="" td=""></mt,>

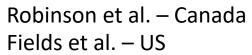
Some decisions we had to make...

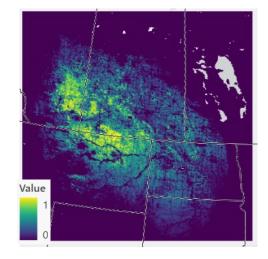
- Common projection & snap
- Transform density to binomial : 1-Exp(density*-1)
- If occurrence < 1; divide by max value
- Process each model at native resolution/extent
- Process composites using maximum extent and minimum resolution

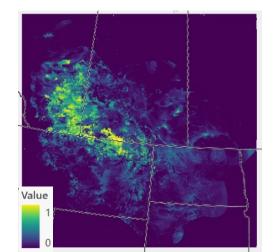
Fink et al. (eBird)

Sather et al.

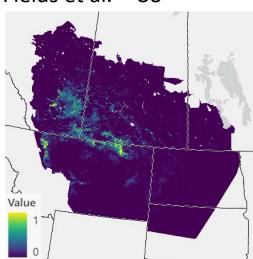
Niemuth et al.

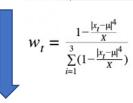


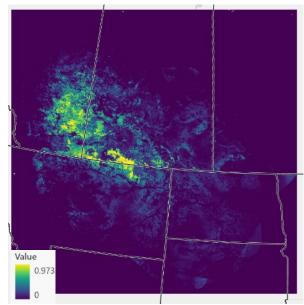






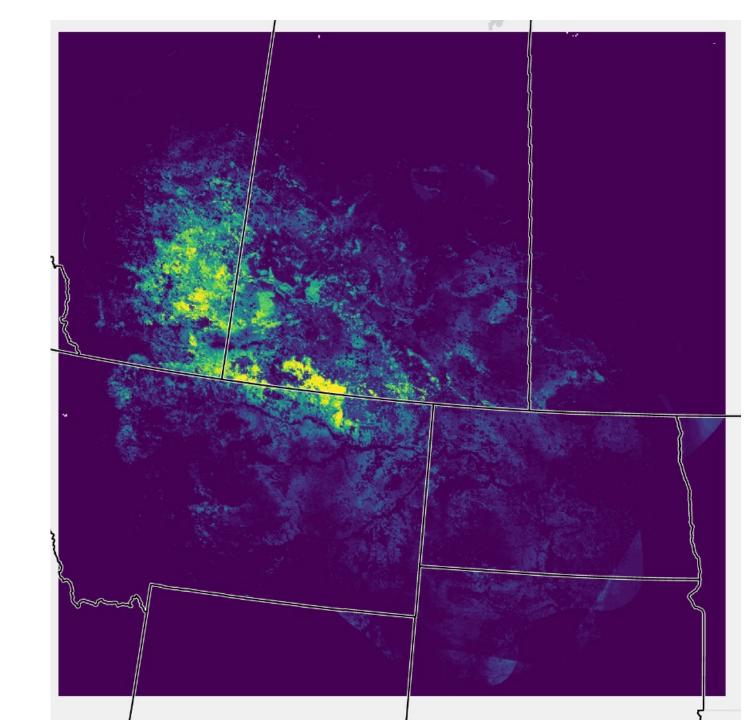






Next steps

- Incorporate models from Fedy et al. and BCR.
- Rerun analyses for SPPI, BAIS, CCLO, TBLO.
- Make outputs & processing scripts publicly available
- Update when necessary



Thank you!

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