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Supplemental Material

Predicting the Effects of Climate Change on Dengue Vectors Densities in Southeast Asia through Process-Based Modeling

Lucas Bonnin, Annelise Tran, Vincent Herbreteau, Sébastien Marcombe, Sébastien Boyer, Morgan Mangeas, and Christophe Menkes

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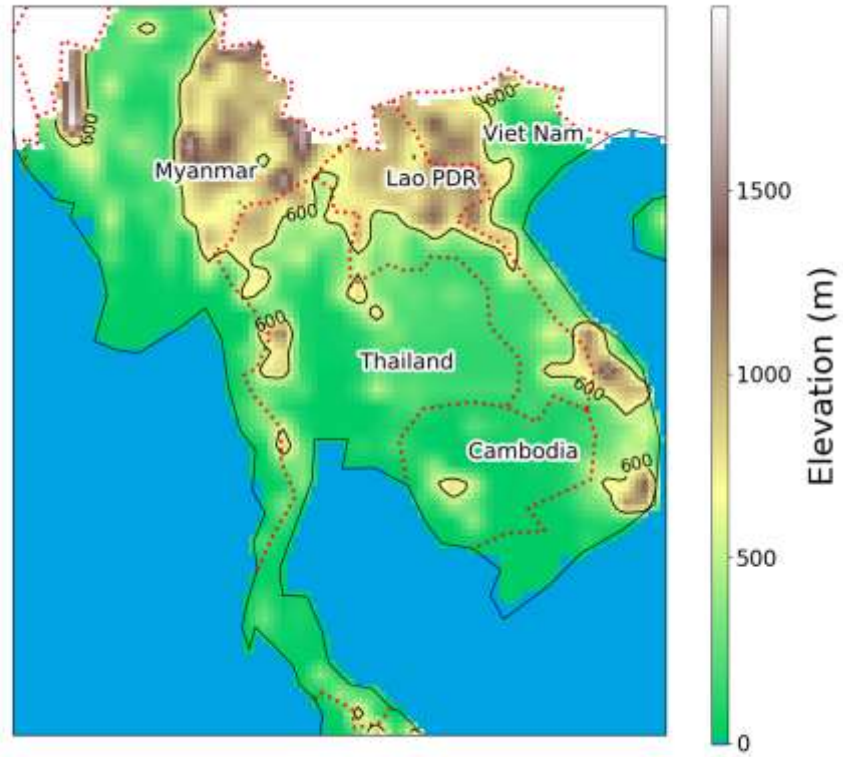


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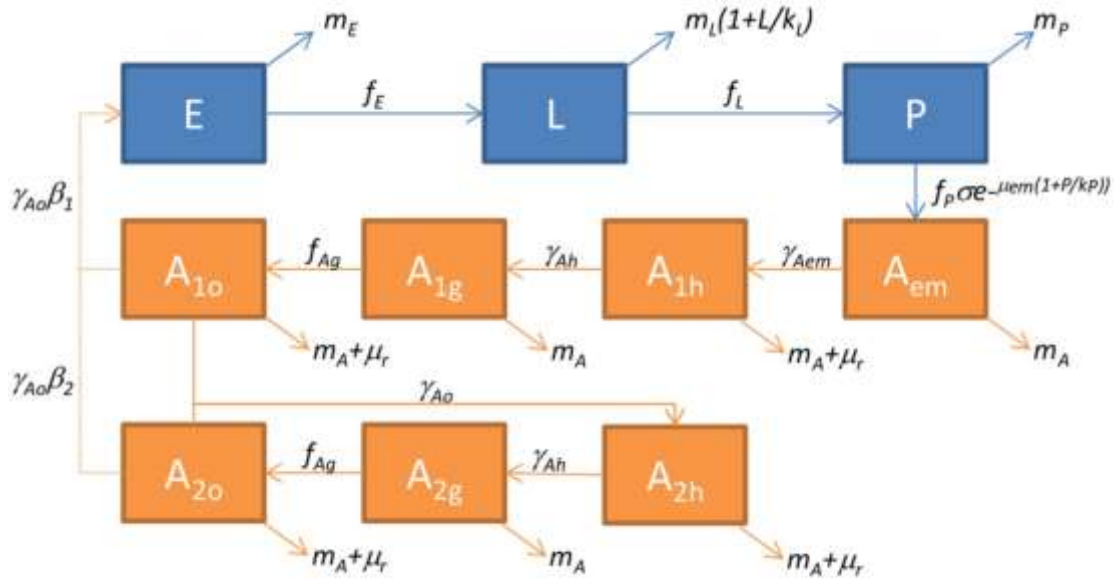


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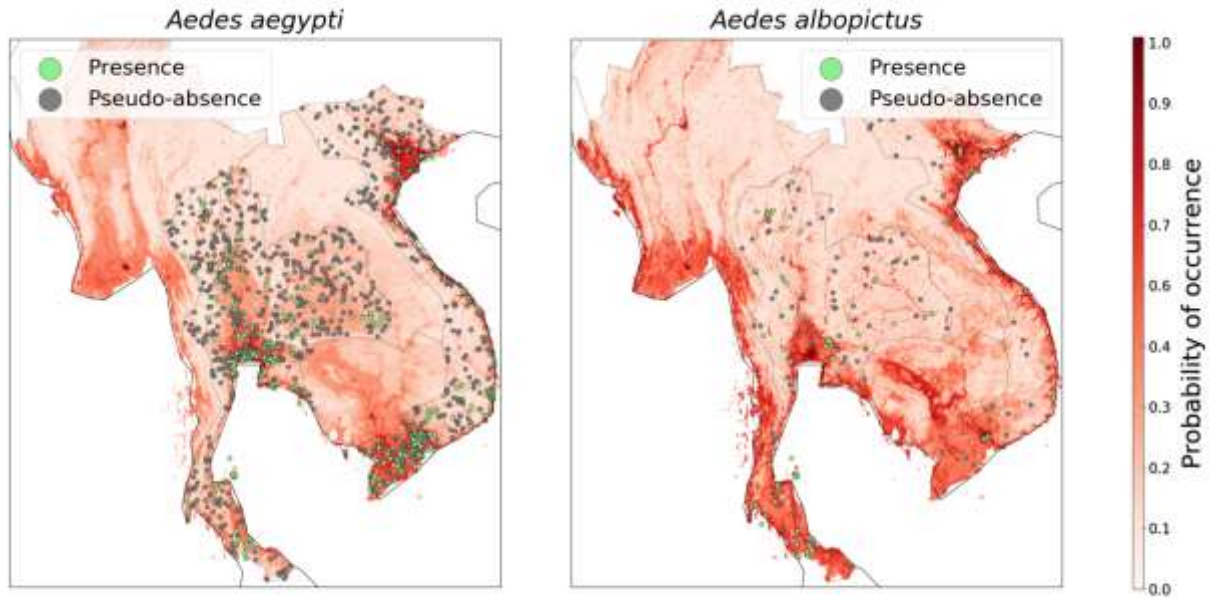
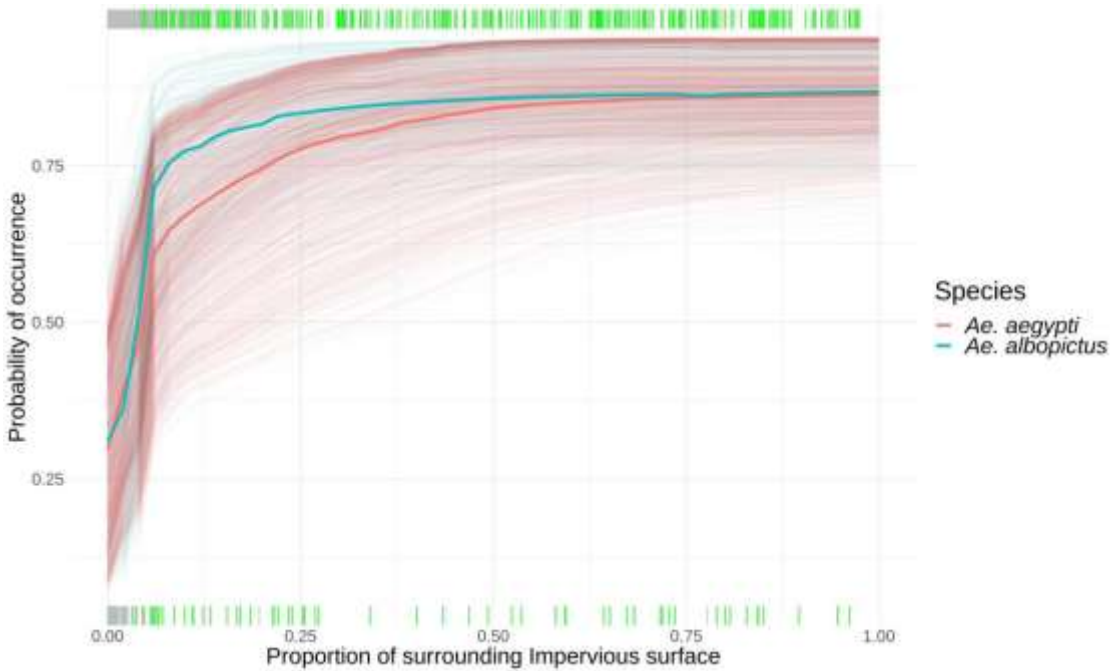


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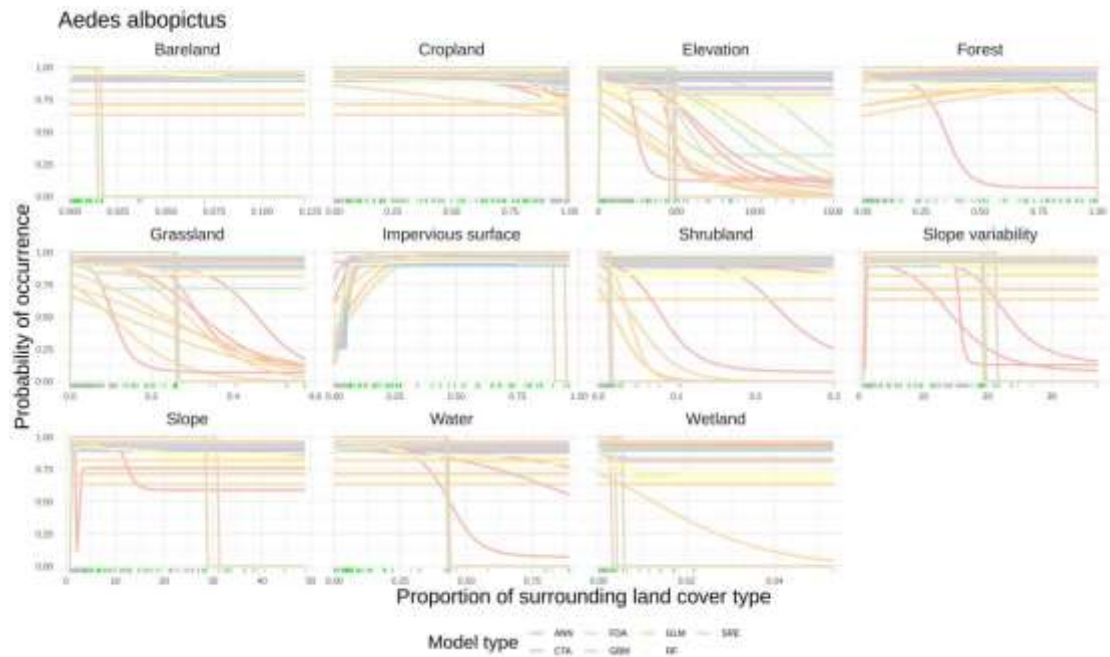
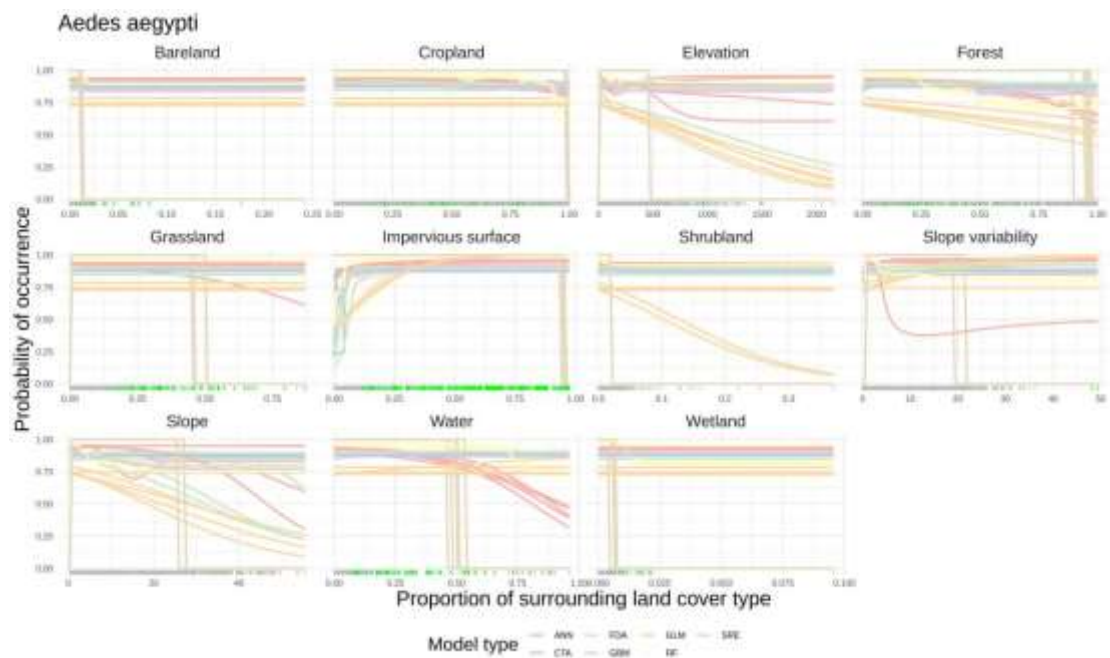


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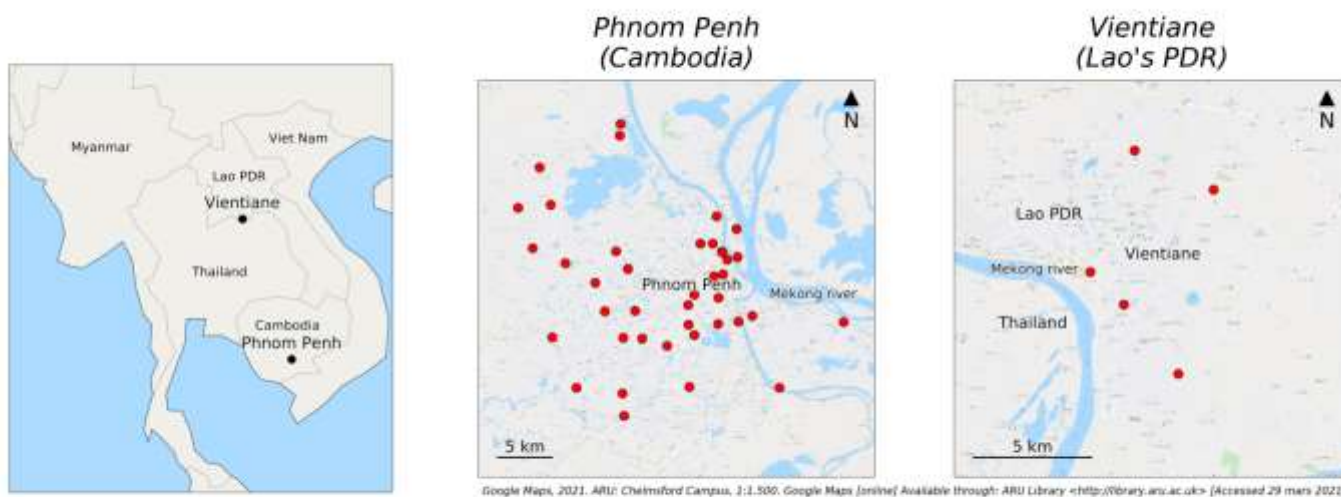


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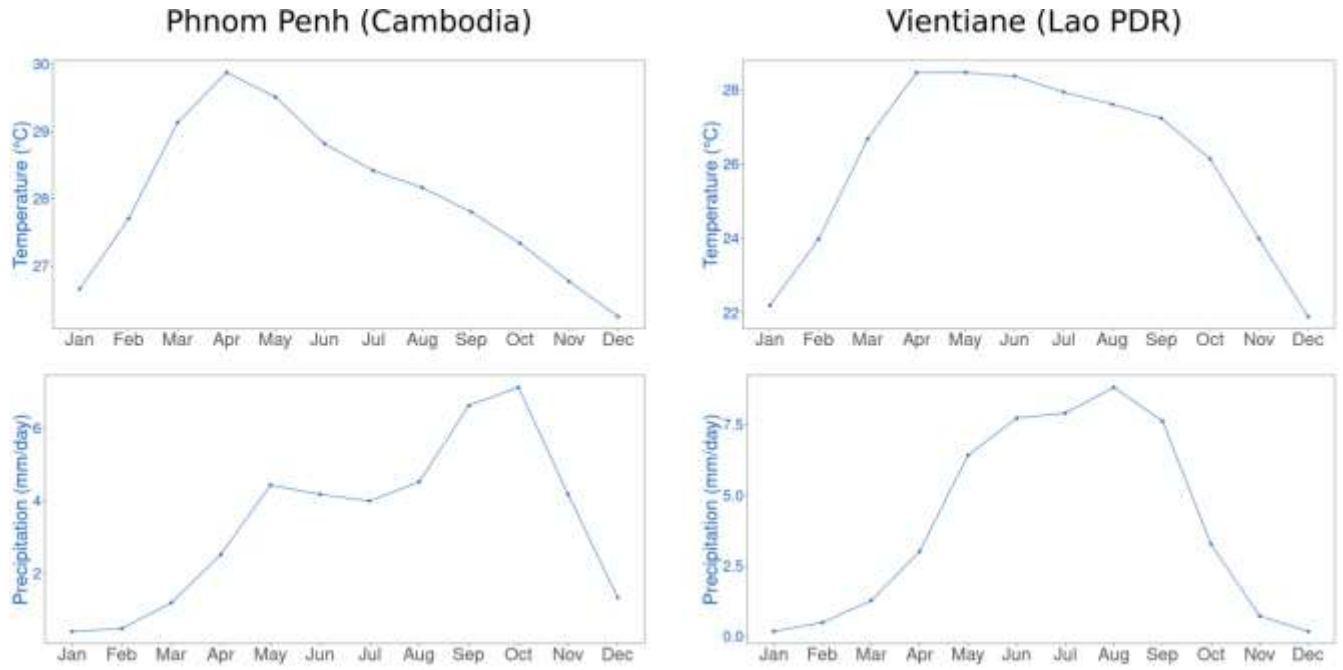


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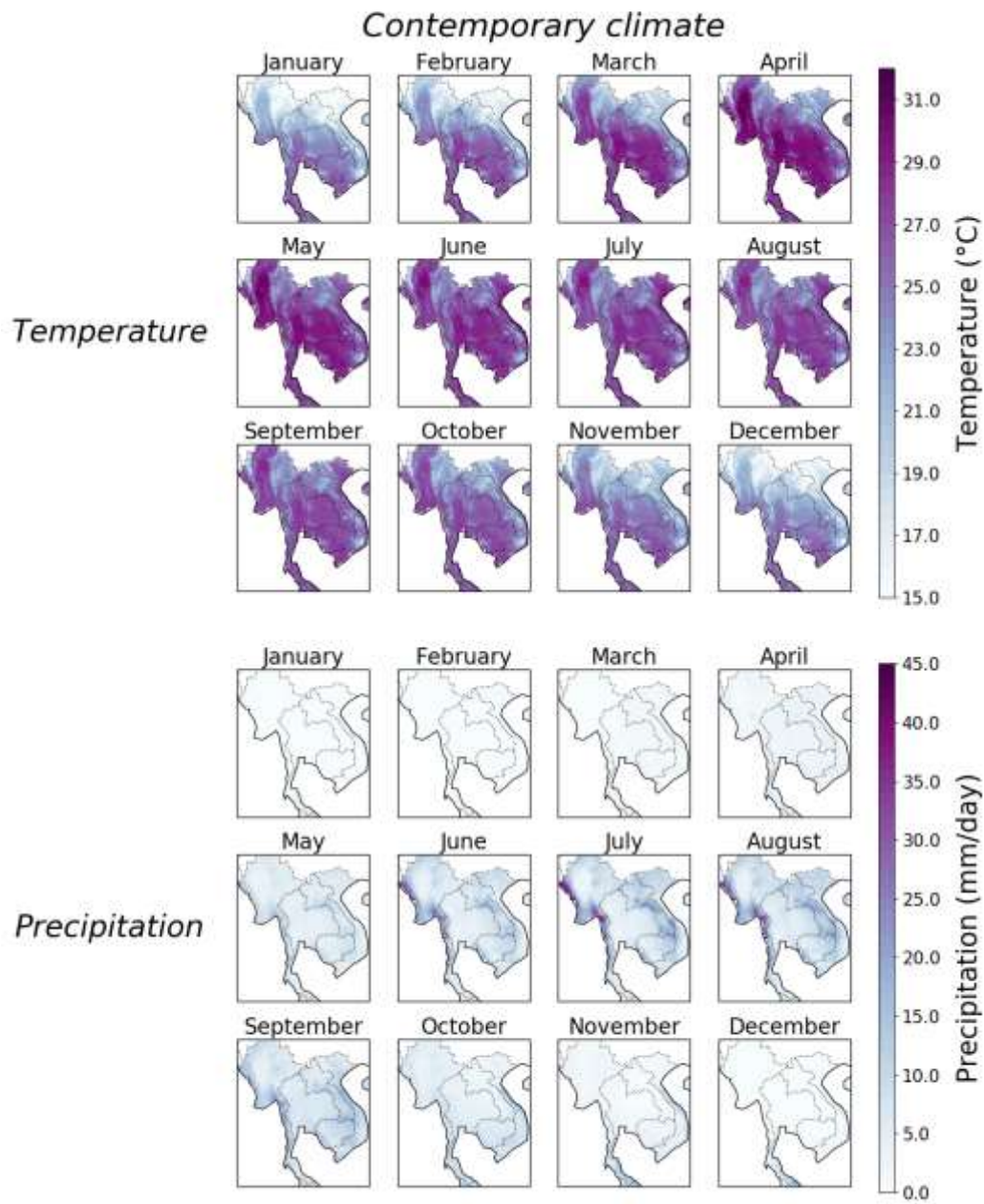


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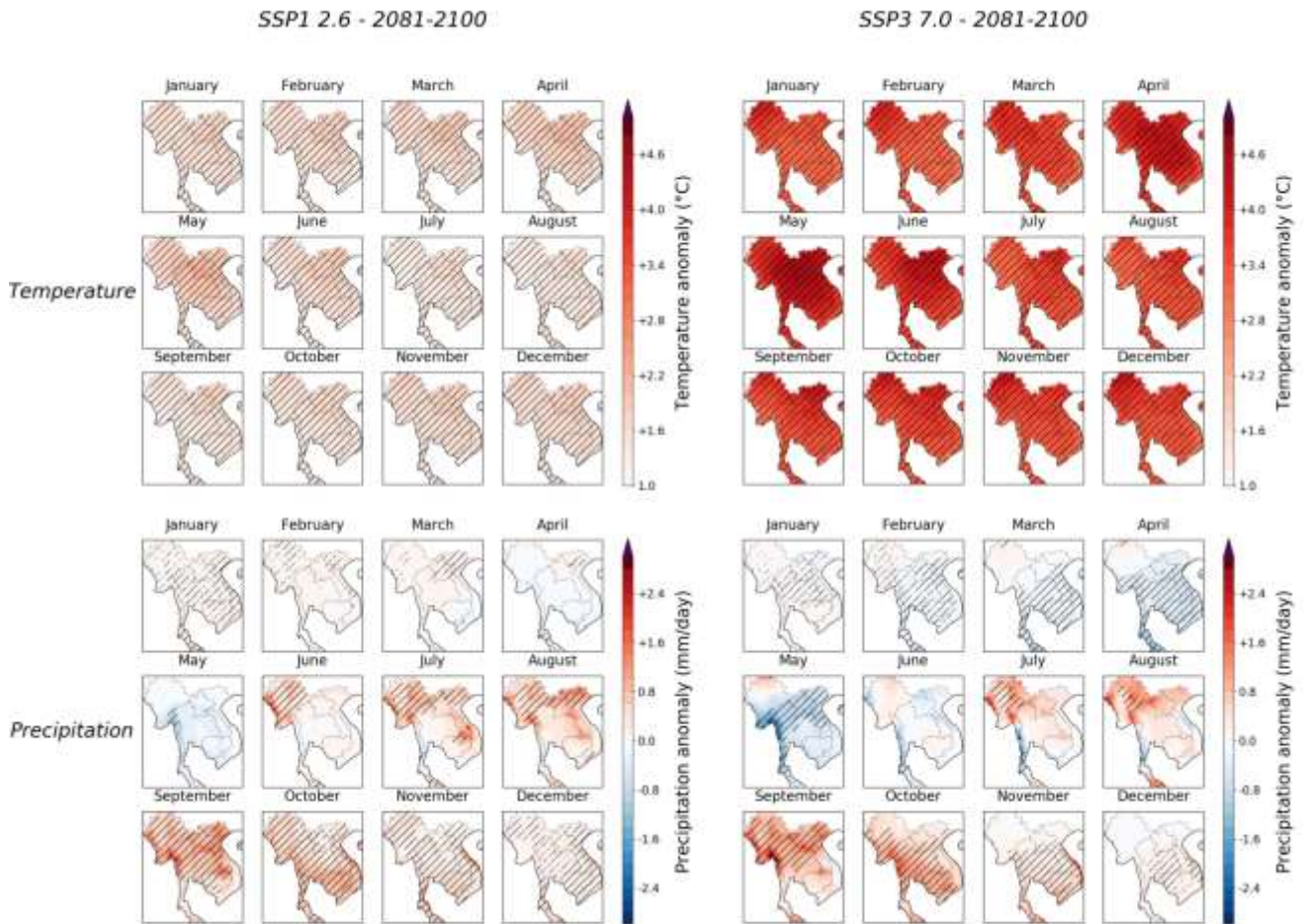


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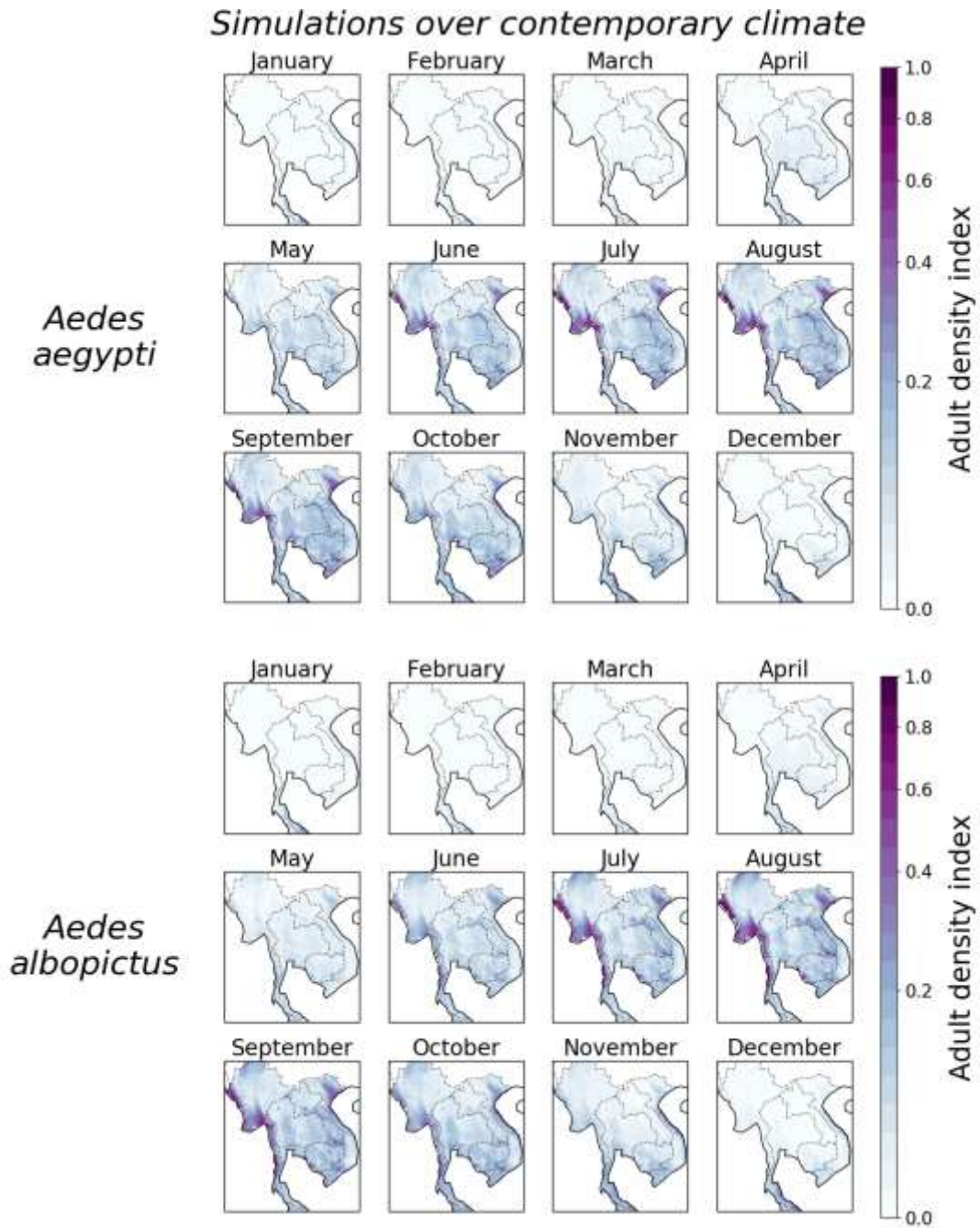


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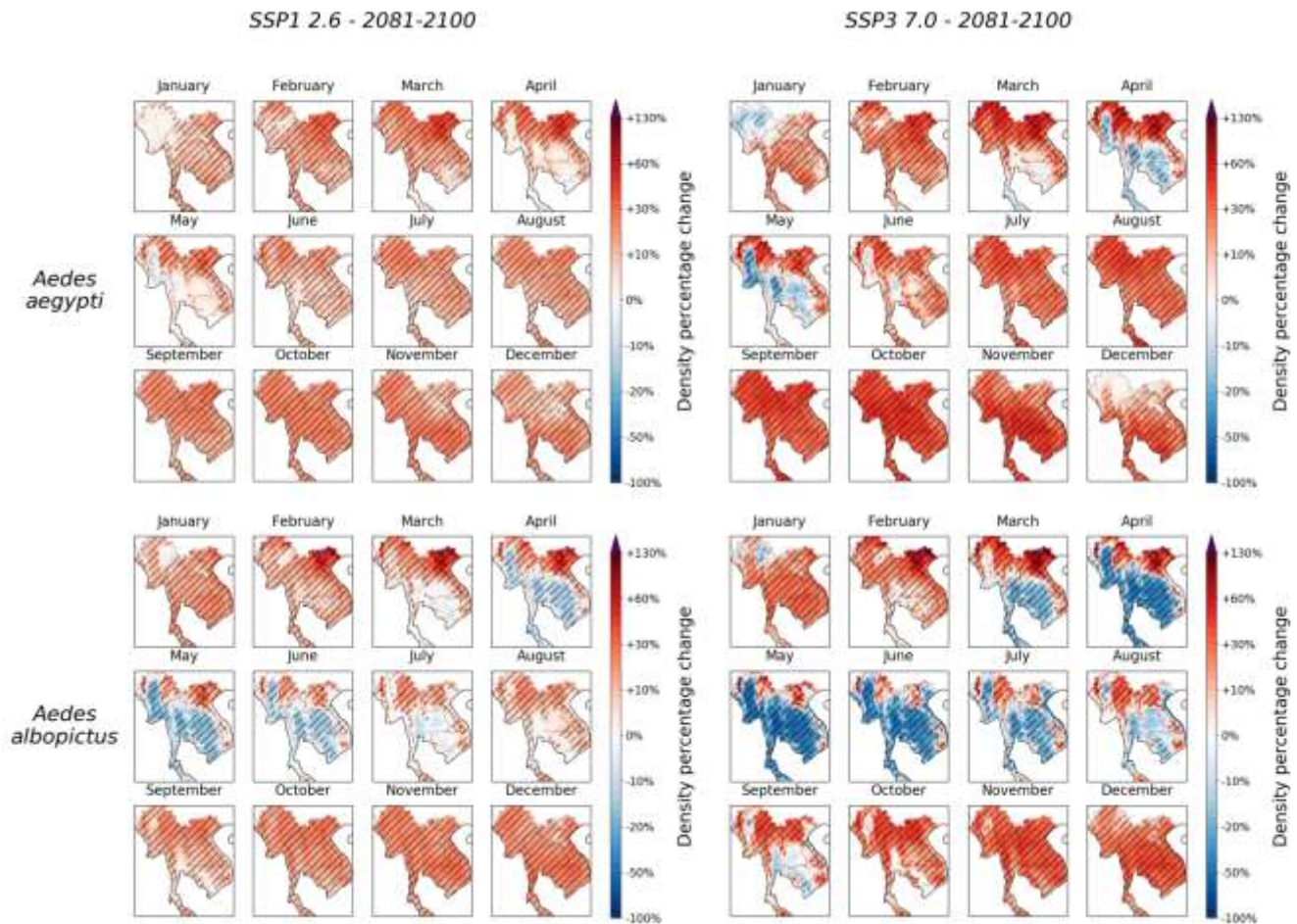


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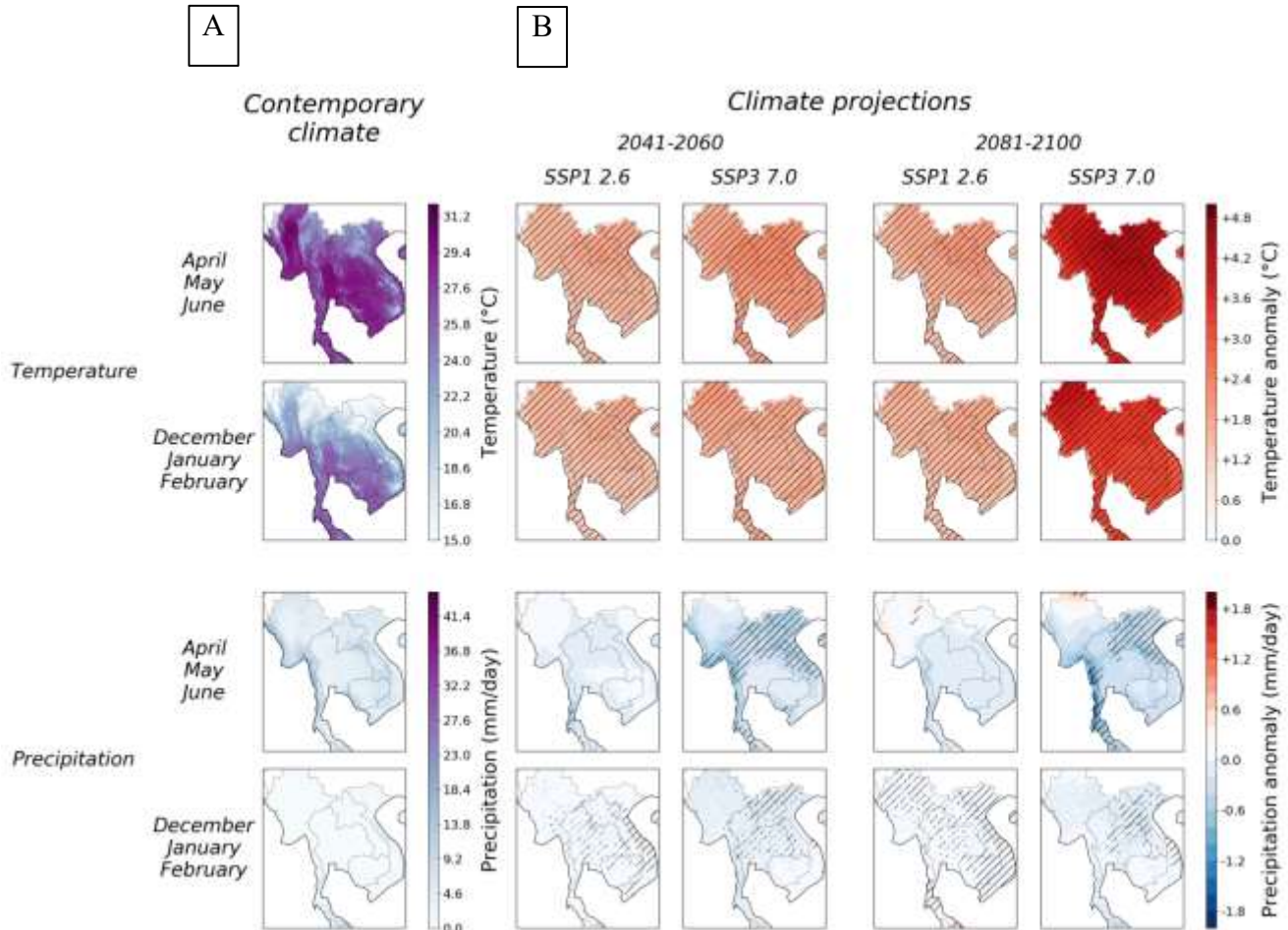
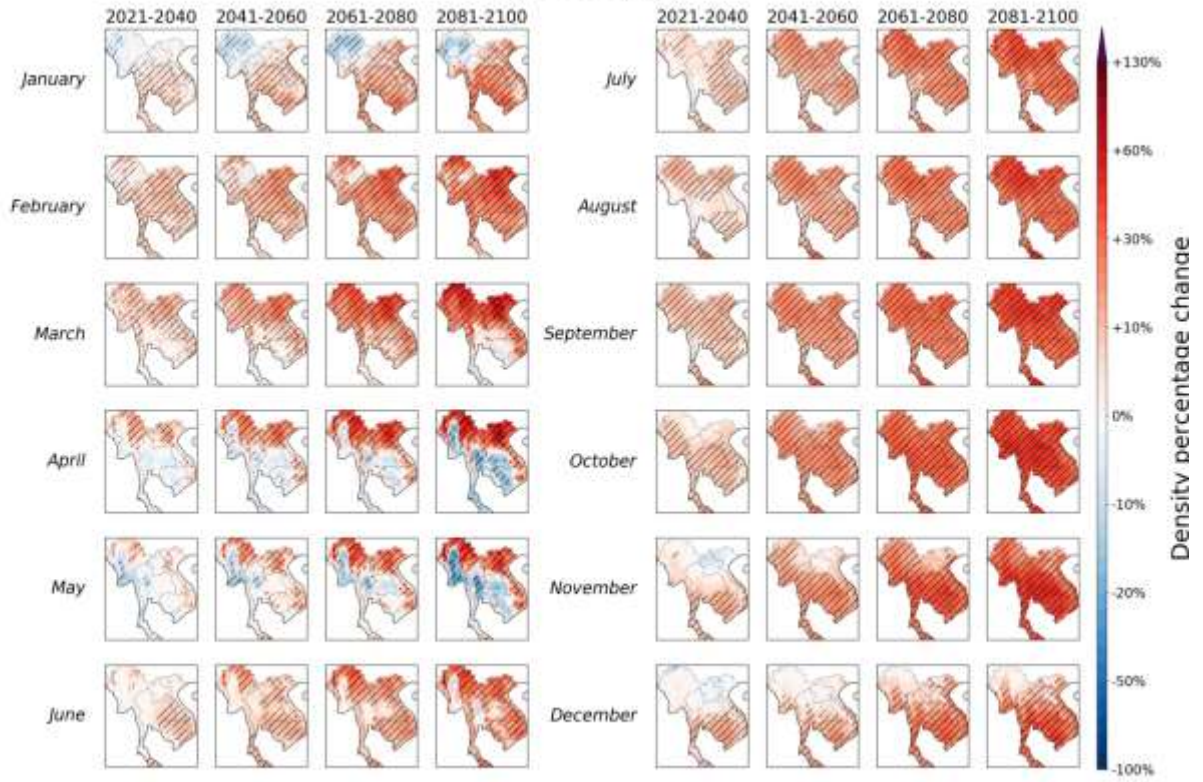


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Aedes aegypti



Aedes albopictus

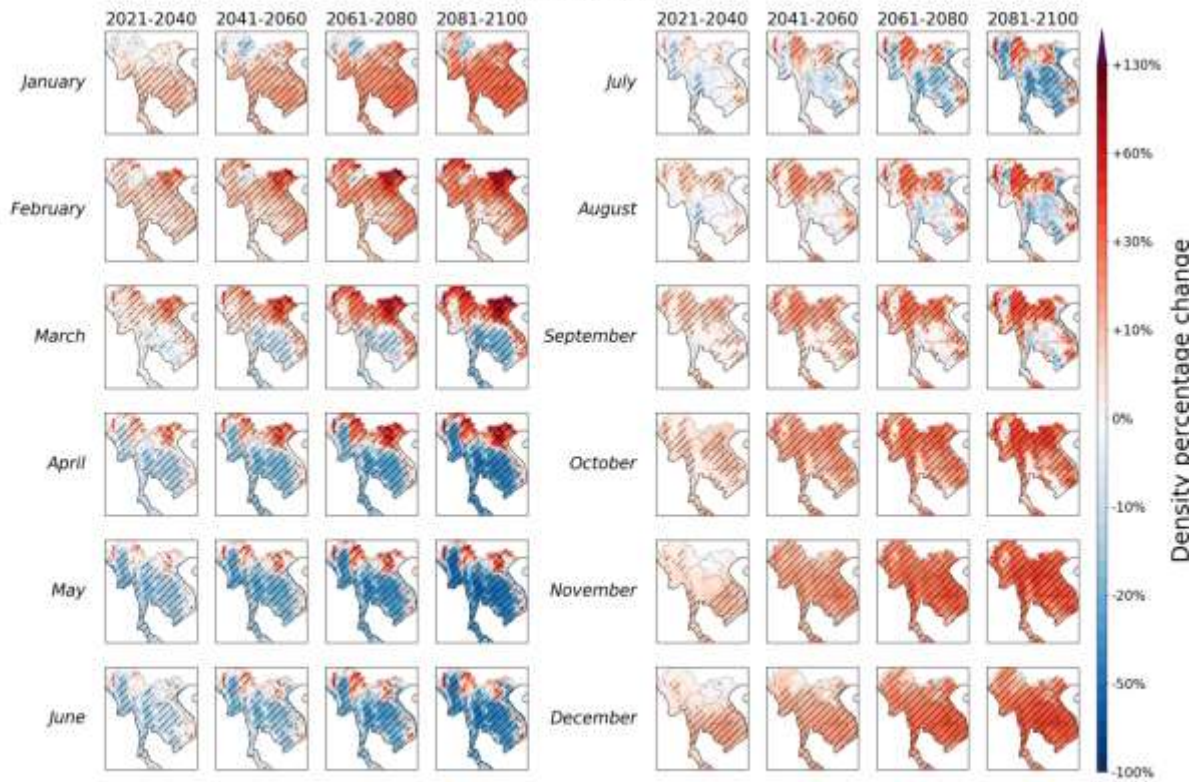
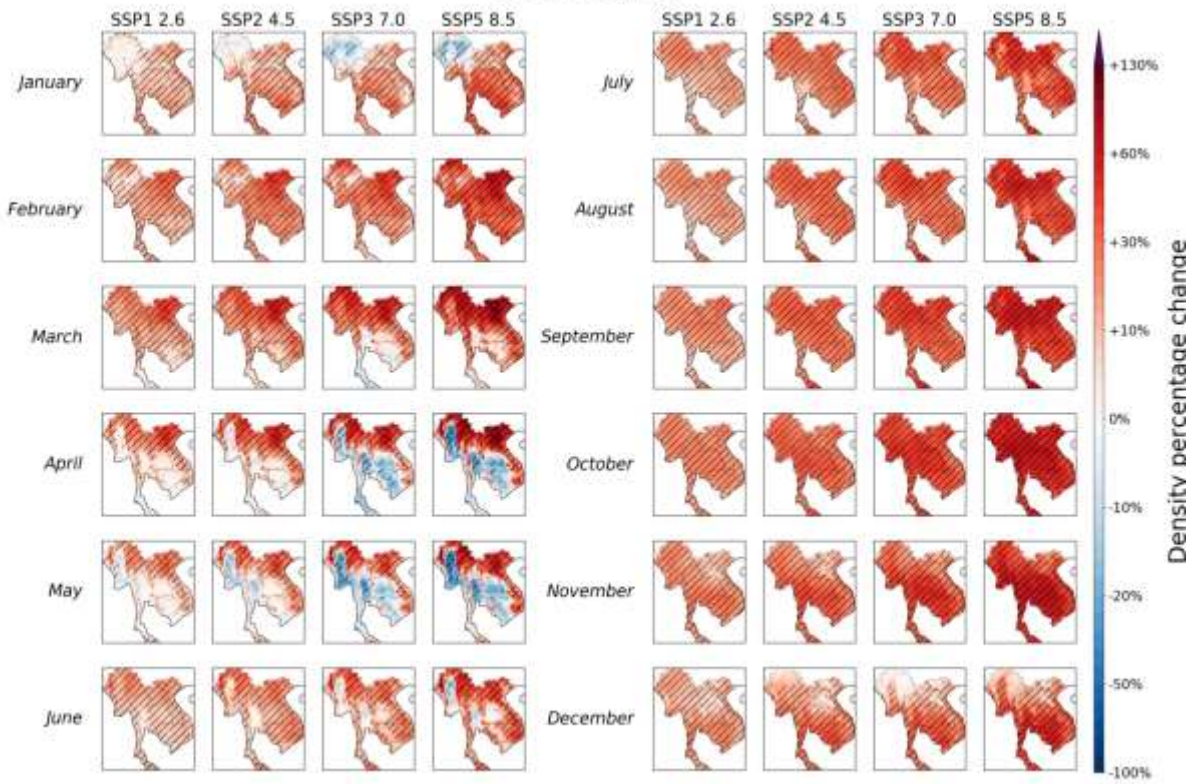


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Aedes aegypti



Aedes albopictus

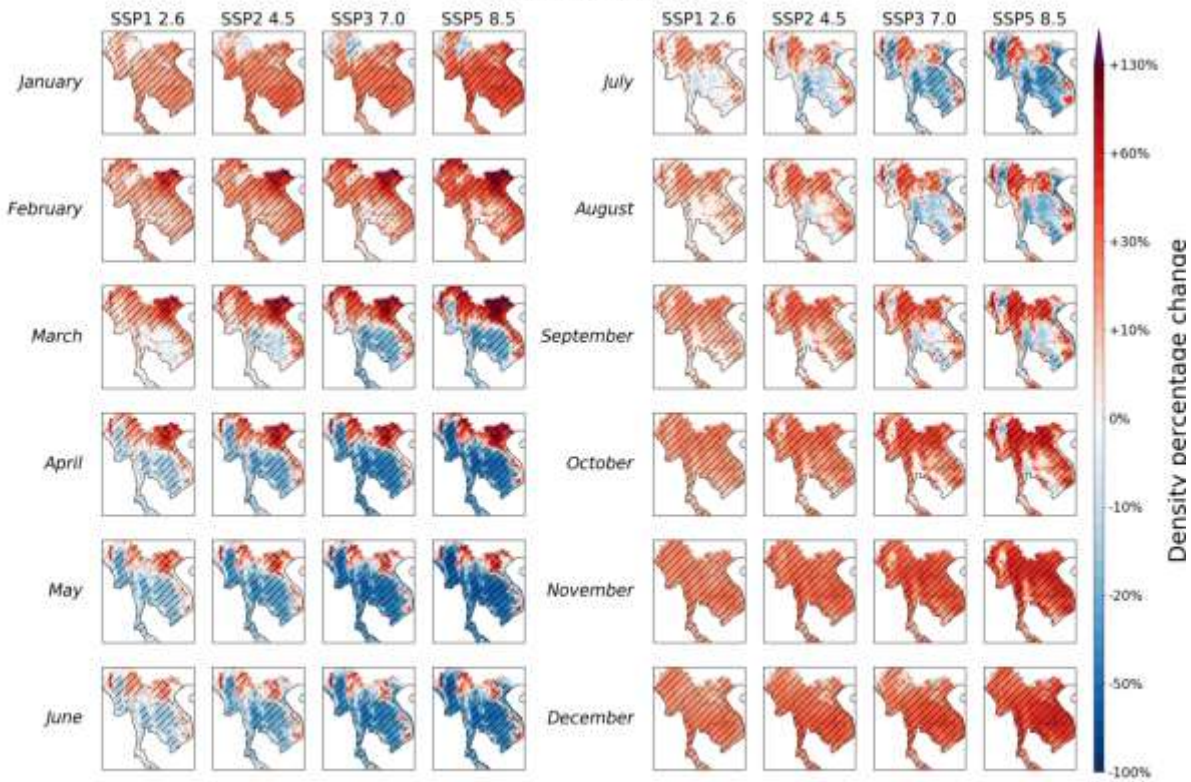


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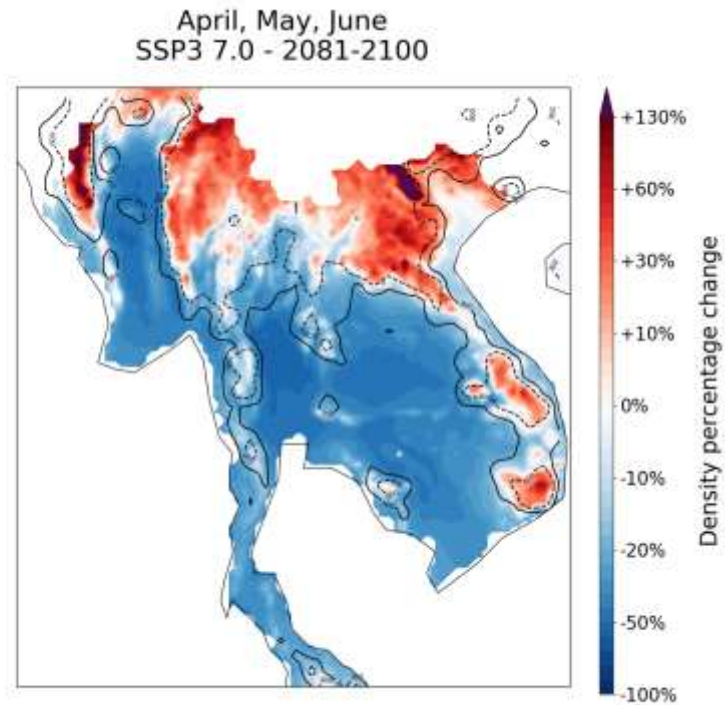


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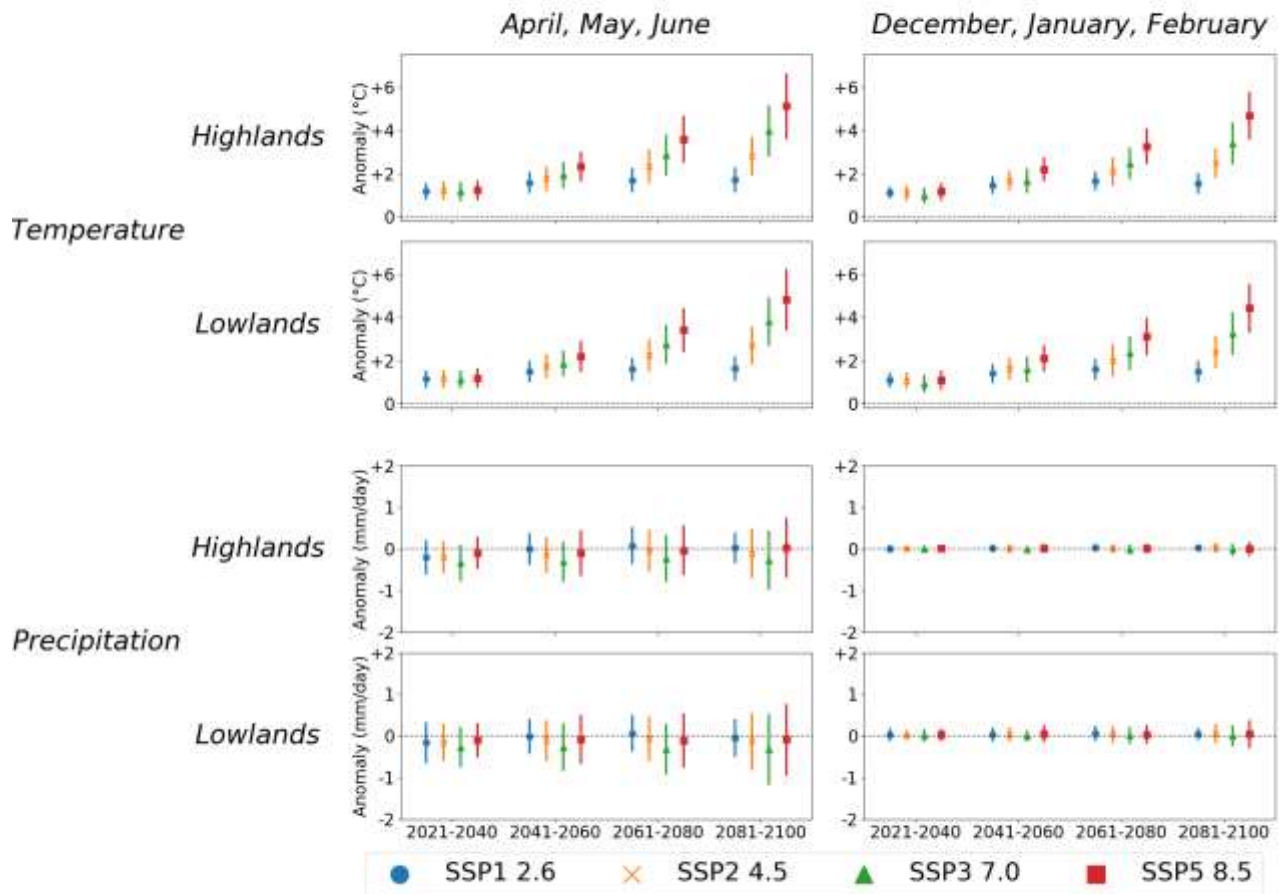


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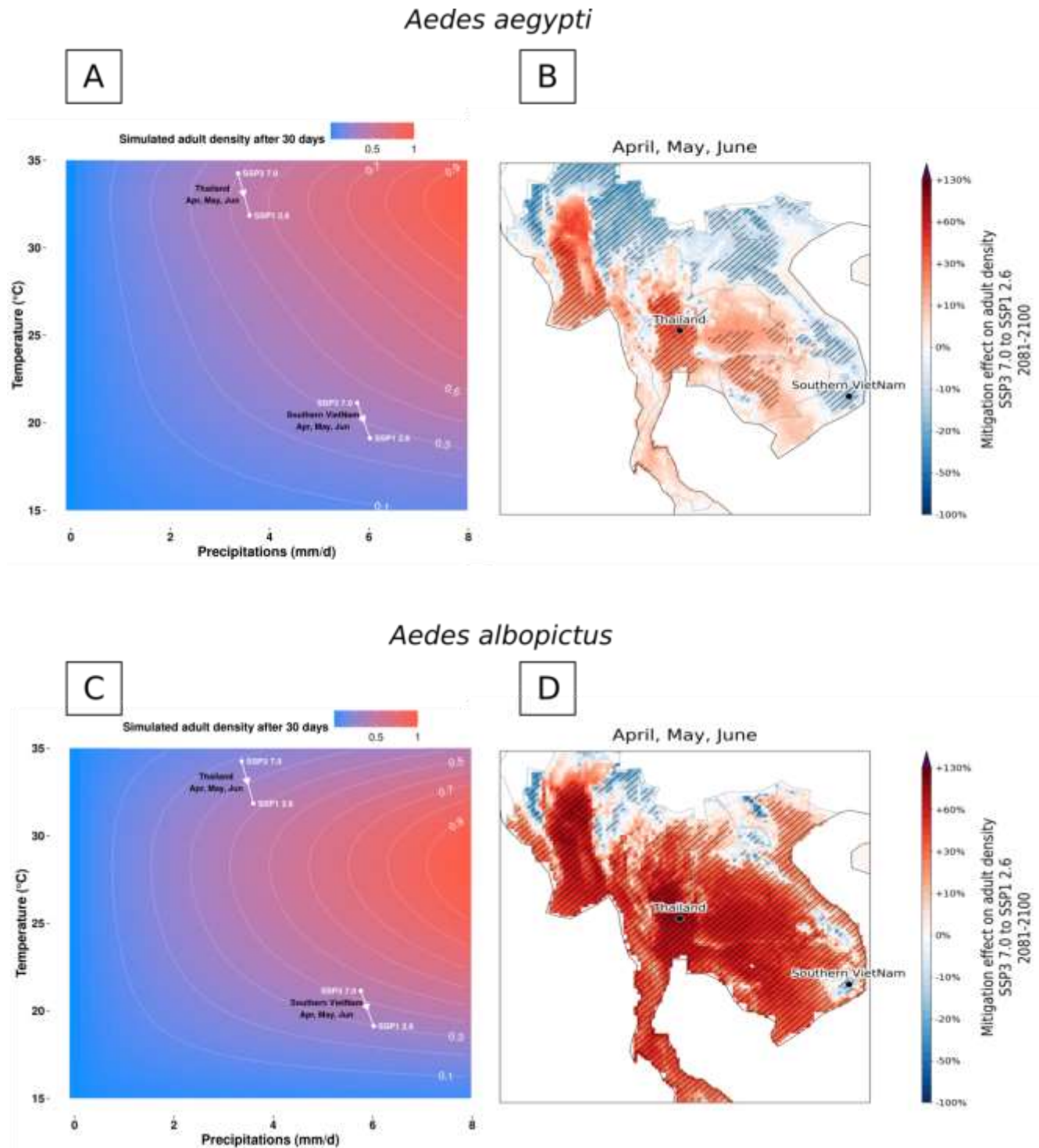


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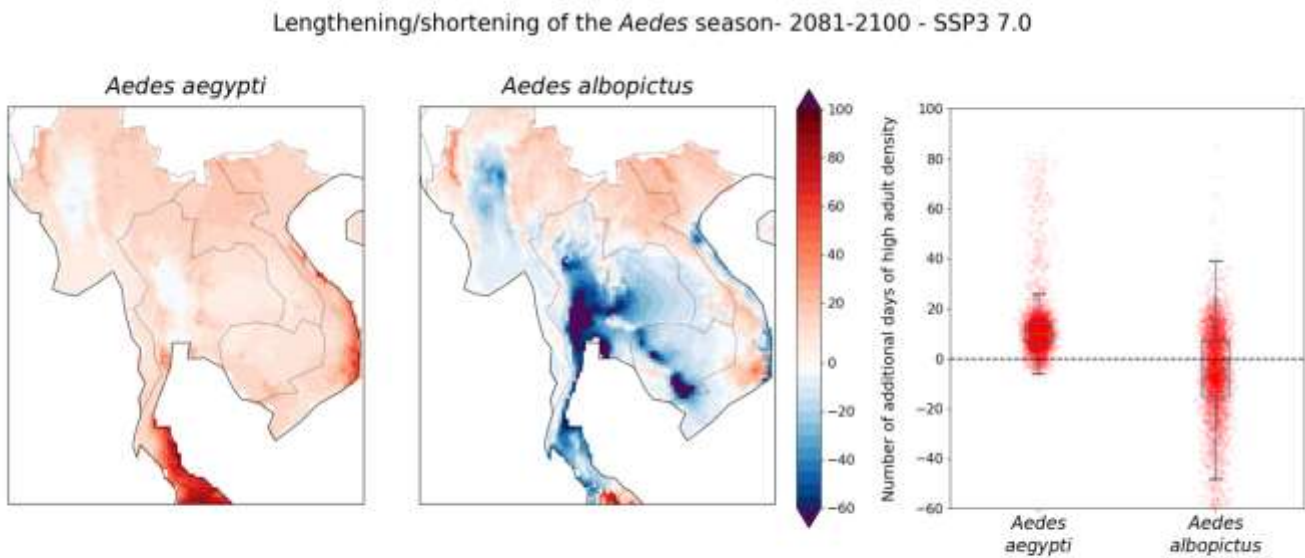
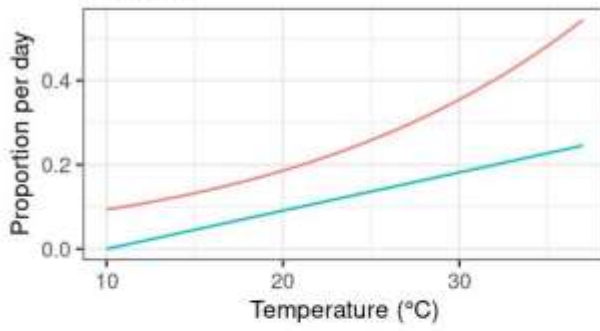
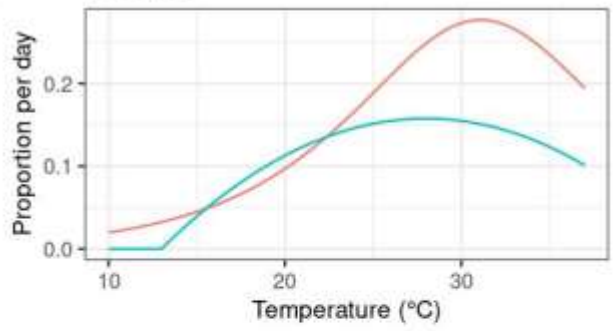


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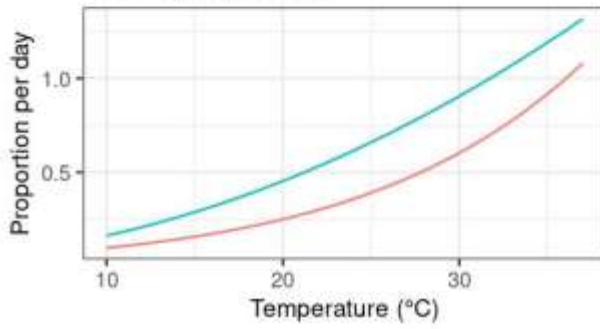
Transition function from eggs to larvae



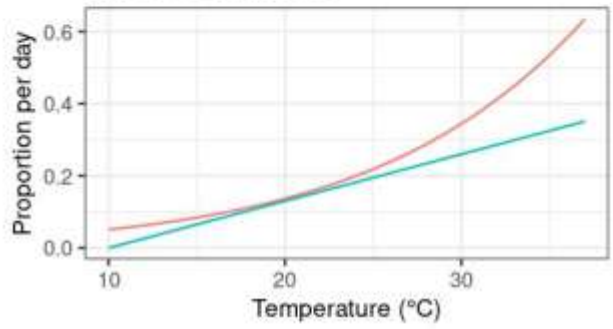
Transition function from larvae to pupae



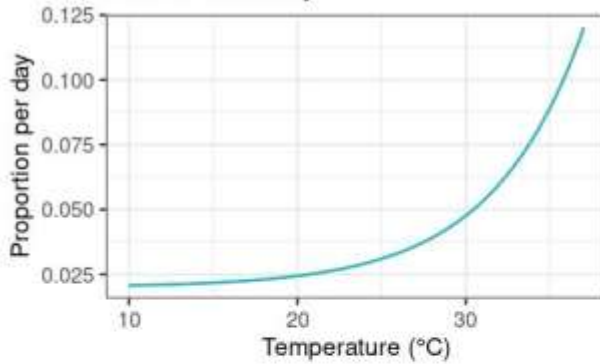
Transition function from pupae to emerging adults



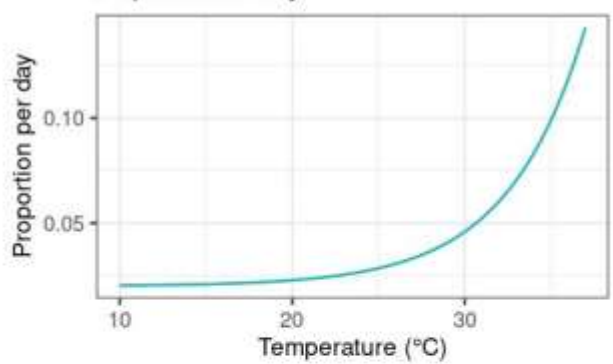
Transition function from engorged adults to oviposition



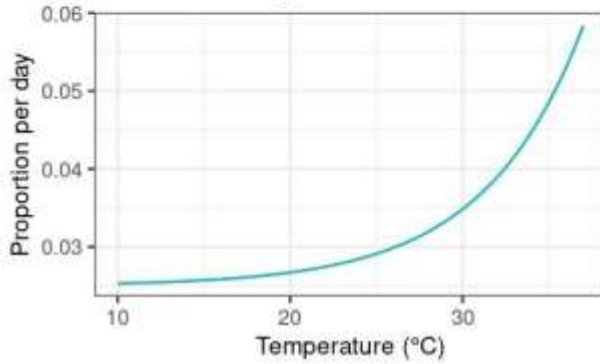
Larvae mortality



Pupae mortality



Adults mortality



Species

- *Ae. aegypti*
- *Ae. albopictus*

Figure S18. Comparison of temperature-dependent transition rates used in process-based models. Temperature-dependent transition rates used in the model developed by Liu-Helmersson et al. (2019),³ were compared to those used in the present study. Although the two models differed in several aspects, notably on the number of life stages considered, comparing the functions describing transitions rates is still pertinent.

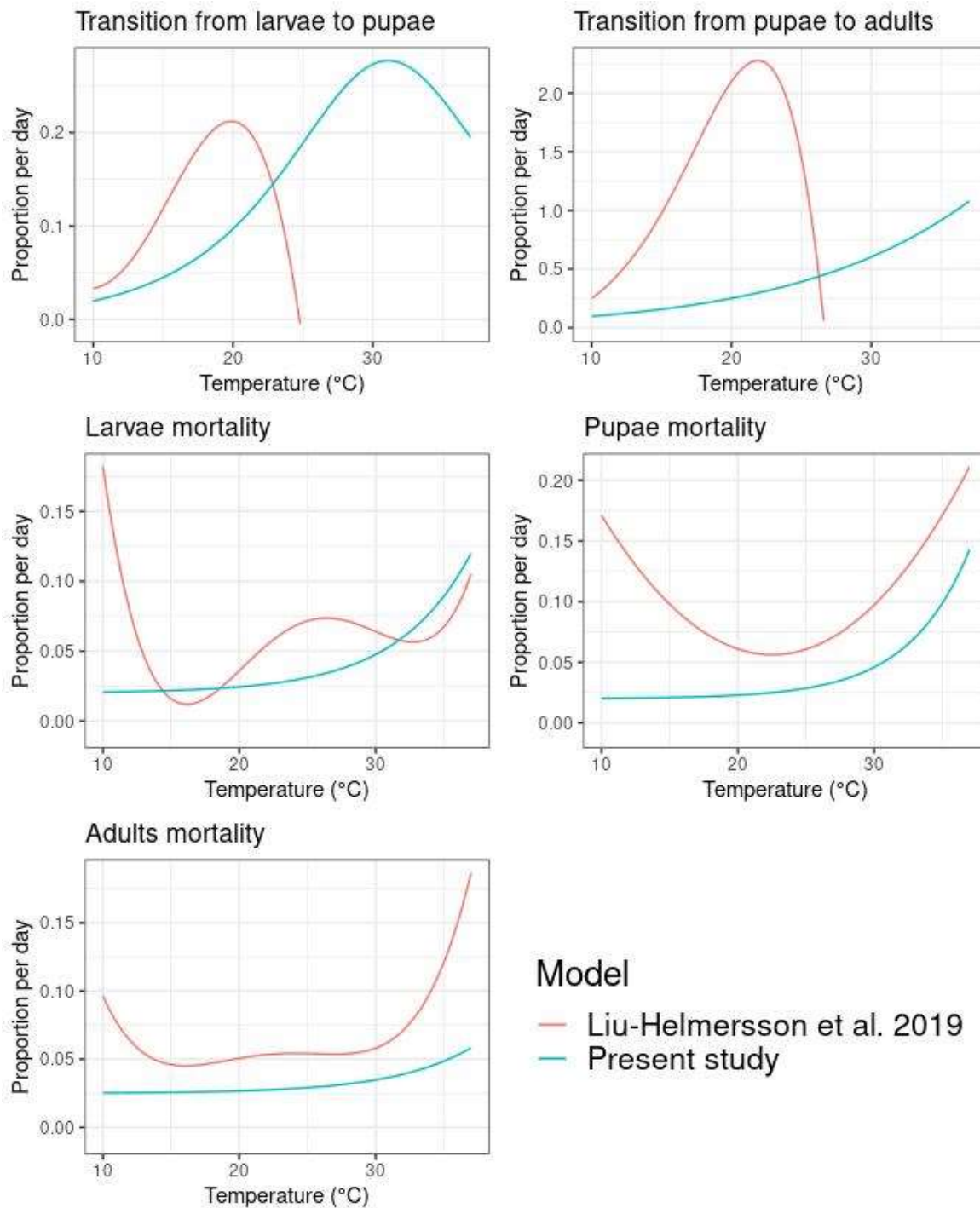


Table S1. Species Distribution Models quality metrics.

Species	Model family	Testing data AUC	
<i>Aedes aegypti</i>	GLM	0.88	
		0.893	
		0.846	
		0.86	
		0.876	
	GBM	0.885	
		0.932	
		0.886	
		0.876	
		0.89	
	CTA	0.839	
		0.864	
		0.83	
		0.787	
	ANN	0.803	
		0.88	
		0.908	
		0.872	
	FDA	0.865	
		0.876	
		0.879	
		0.9	
		0.863	
	RF	0.851	
		0.858	
		0.884	
		0.939	
		0.897	
	<i>Aedes albopictus</i>	GLM	0.883
			0.894
0.816			
0.694			
0.775			
GBM		0.866	
		0.882	
		0.938	
		0.747	
		0.838	
CTA		0.896	
		0.87	
		0.725	
		0.752	
		0.755	
ANN	0.786		
	0.775		
	0.912		
	0.608		
	0.799		
FDA	0.898		
	0.895		
	0.84		
	0.768		
	0.706		
RF	0.831		
	0.898		
	0.948		
	0.769		
	0.798		
		0.897	
		0.875	

For each species, the Ensemble modeling process was performed over 6 model families, the 5-fold cross-validation process thus resulted in building 30 models. For *Ae. aegypti*, the 30 models displayed an average AUC of 0.87 (+ 0.03 SD). Ensemble forecasting was performed using the 26 models presenting an AUC>0.85 (colored in black in the following table). For *Ae.*

albopictus, the 30 models displayed an average AUC of 0.84 (+- 0.05 SD). Ensemble forecasting was performed using the 14 models presenting an AUC>0.85 (colored in black in the following table).

Supplemental Material references:

1. Tran, A. *et al.* Complementarity of empirical and process-based approaches to modelling mosquito population dynamics with *Aedes albopictus* as an example—Application to the development of an operational mapping tool of vector populations. *PLOS ONE* **15**, e0227407 (2020).
2. Kraemer, M. U. G. *et al.* The global compendium of *Aedes aegypti* and *Ae. albopictus* occurrence. *Sci. Data* **2**, 1–8 (2015).
3. Liu-Helmersson, J., Brännström, Å., Sewe, M. O., Semenza, J. C. & Rocklöv, J. Estimating Past, Present, and Future Trends in the Global Distribution and Abundance of the Arbovirus Vector *Aedes aegypti* Under Climate Change Scenarios. *Front. Public Health* **7**, (2019).