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Performance of Genetic Distance Metrics in Gravity and General Mixed Effects Models

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Performance of genetic distance metrics in gravity and general mixed effects models

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What are the genetic responses to fragmentation?

- Connectivity can help maintain occupied patches, genetic structure, and metapopulations
- What drives connectivity?
 - Connectivity can be influenced by at-site and between-site characteristics
- How do we determine connectivity?
 - Gene flow correlated to resistance surfaces

Issues with Genetics in Landscape Genetics

- Currently used genetic metrics were not developed for Landscape Genetics
- How long does it take to see a response to fragmentation with genetics?
- Are we selecting the correct connectivity model given the system?

Issues with Genetics in Landscape



Genetic Metrics

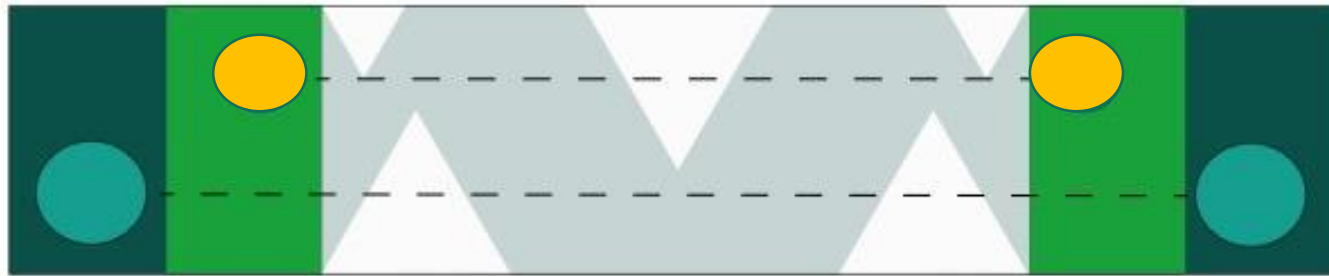
- Dps, Bray-Curtis, and PCA perform well for individual genetic metrics¹
- How do population genetic metrics perform?
 - Dps – Allele
 - Fst – Heterozygosity
 - PCA (2-axes) – Ordinal
 - Cavalli-Sforza Kinship Coefficient (Dkf) – kinship
 - Nei's D – differentiation from drift and mutation

Ecological Drivers

Between Site
Isolation by resistance



At Site
Isolation by environment



At Site and between site
Gravity

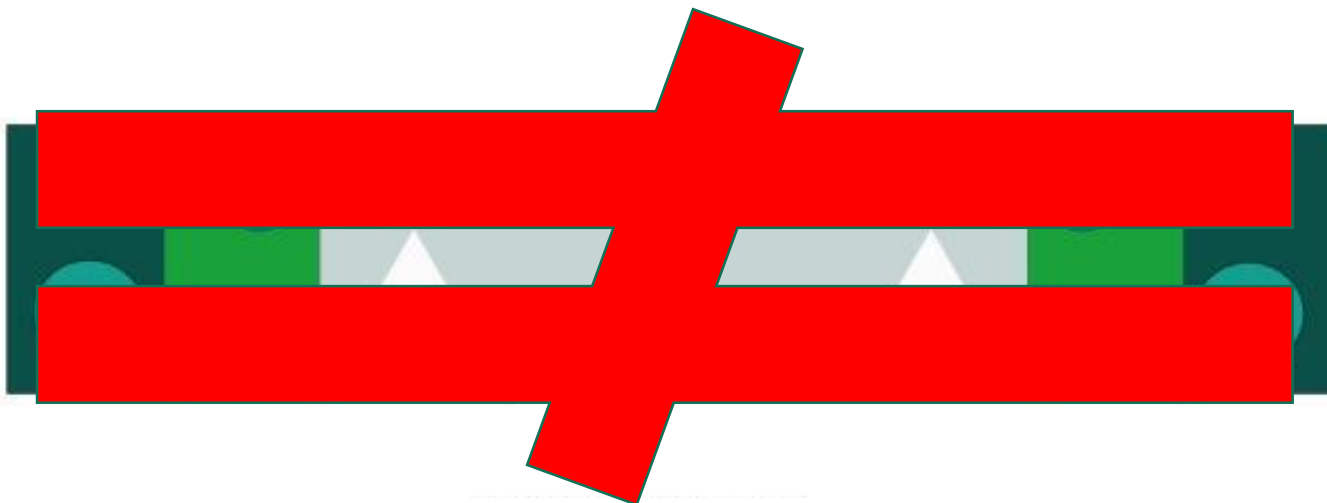


Ecological Drivers

Between Site
Isolation by resistance



At Site
Isolation by environment

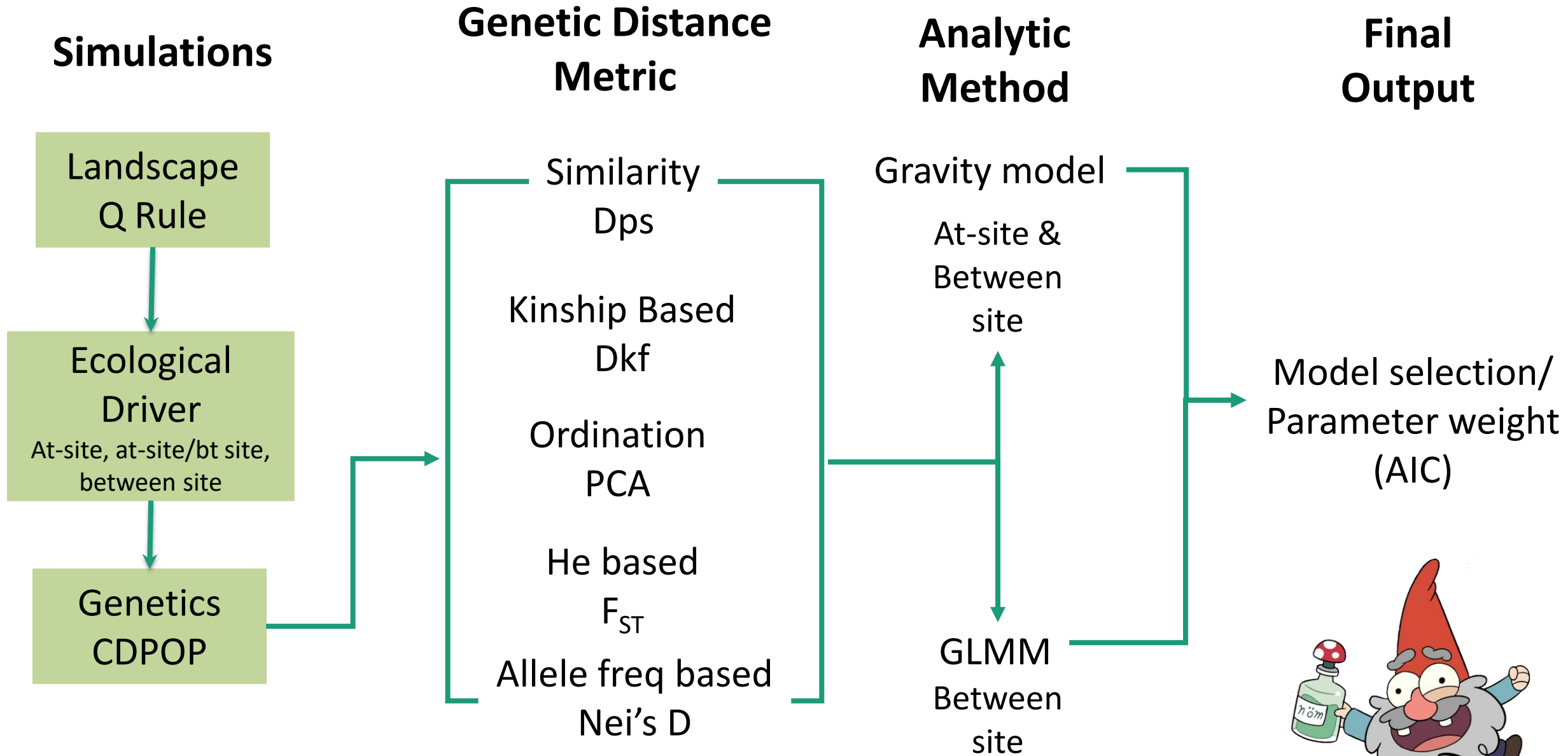


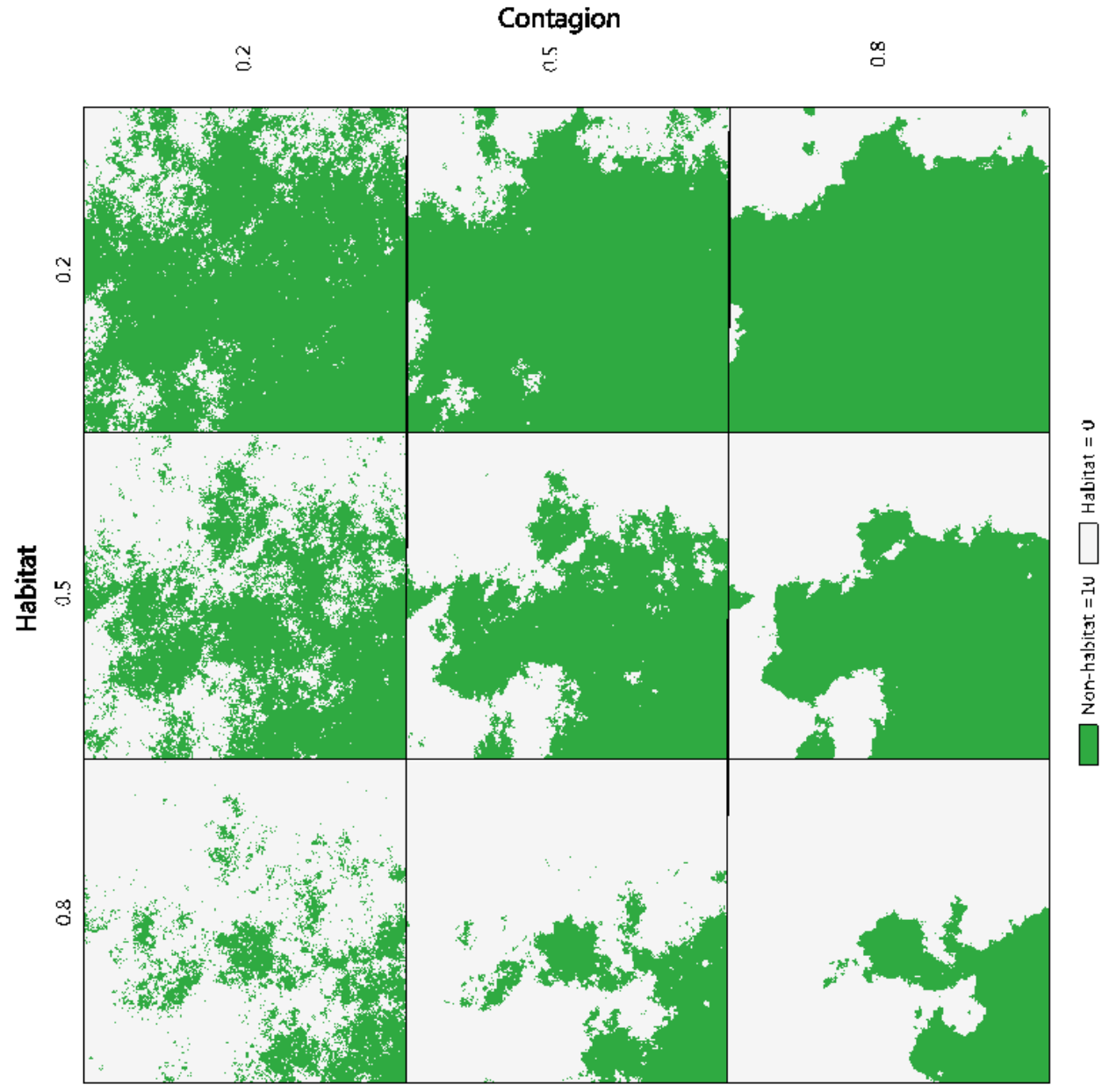
At Site and between site
Gravity

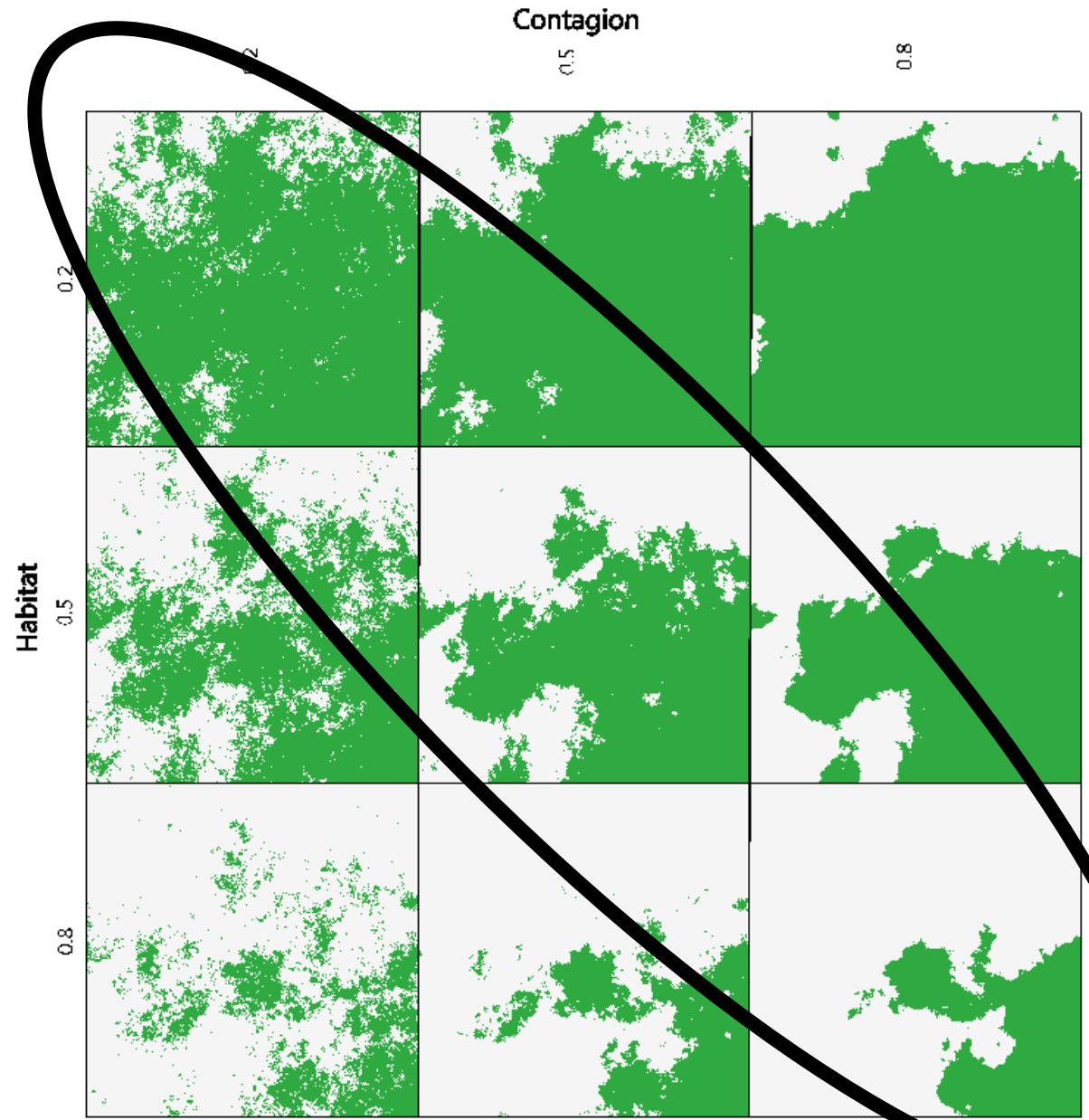


Hypotheses

- Time
 - Short duration: Allele frequency based
 - Long duration: Heterozygosity based
- Landscape
 - More landscape structure: H_e based GD metrics
 - Less landscape structure: Allele frequency based GD metrics







GLMM vs Gravity

- Gravity allows incorporation of at-site predictors impacting geneflow
- GLMM only incorporates between-site environmental costs
- **Gravity prediction:**
 - Increased genetic signature if population size is impacted by habitat metrics
- **GLMM prediction:**
 - Increased model performance if equal population sizes and only between site govern genetic exchange

Analyses

GLMM

$$Y = \underbrace{\beta X}_{\text{Fixed effects}} + \underbrace{\gamma U}_{\text{Random effects}} + \epsilon$$

Fixed effects Random effects

- Fixed effects:
 - landscape resistance cost, Euclidean distance
- Random effect: population

Analyses

GLMM

$$Y = \underbrace{\beta x}_{\text{Fixed effects}} + \underbrace{\gamma u}_{\text{Random effects}} + \varepsilon$$

Fixed effects Random effects

- Fixed effects:
 - landscape resistance cost, Euclidean distance
- Random effect: population

Gravity model

$$Y = \underbrace{kx^\alpha a^\beta e^{-\gamma}}_{\text{Fixed effects}} + \underbrace{\gamma u}_{\text{Random effects}} + \varepsilon$$

Fixed effects Random effects

- Fixed effects:
 - x = Euclidean distance between populations,
 - a = at site characteristic affecting production of migrants,
 - e = landscape resistance cost distance matrix
- Random effect: population

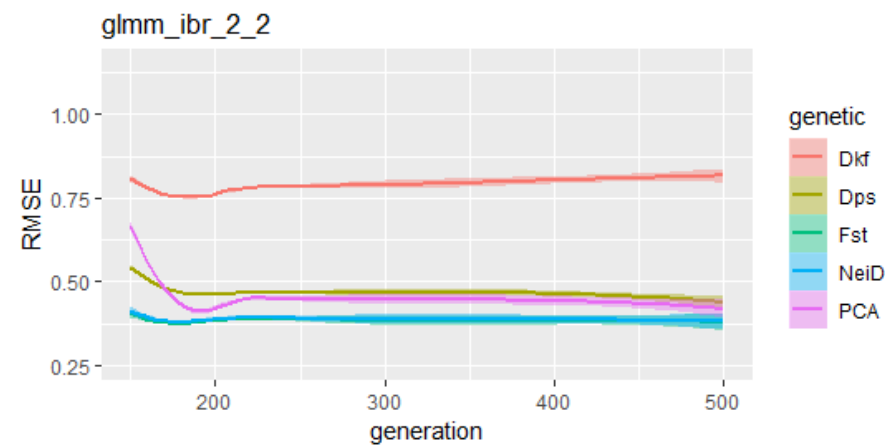
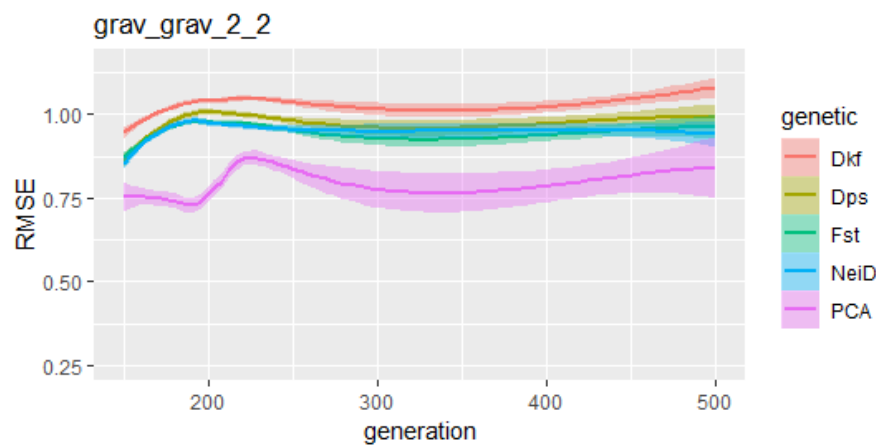
Results -RMSE con .2, hab .2



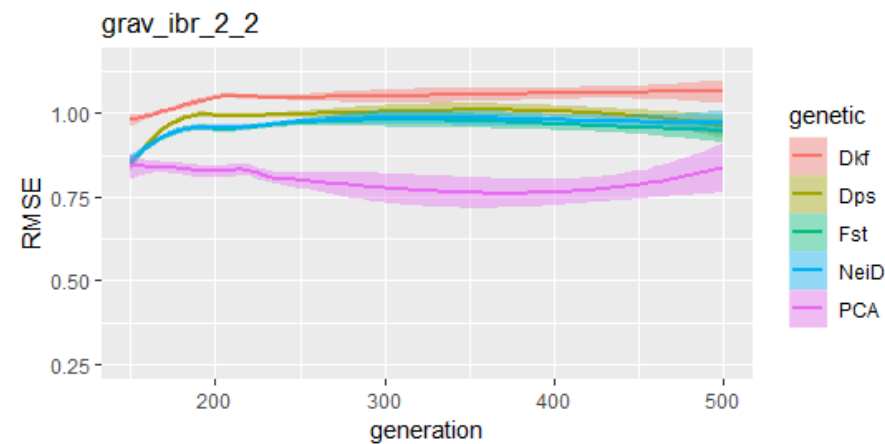
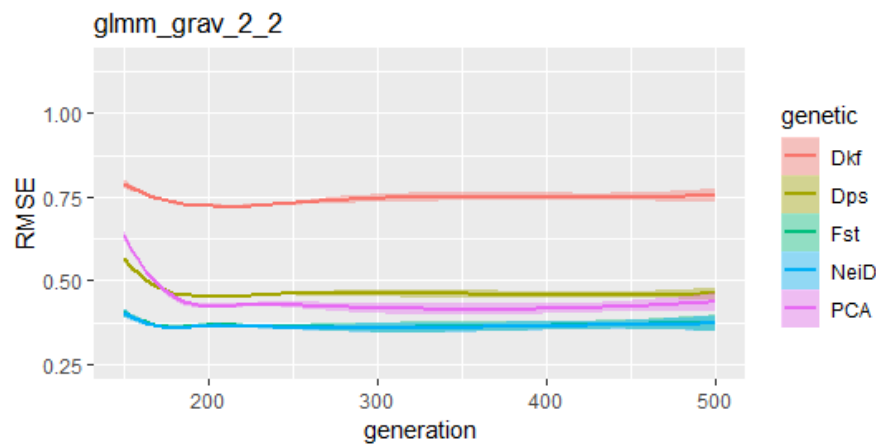
Gravity Driver

IBR Driver

Correct Model



Wrong Model



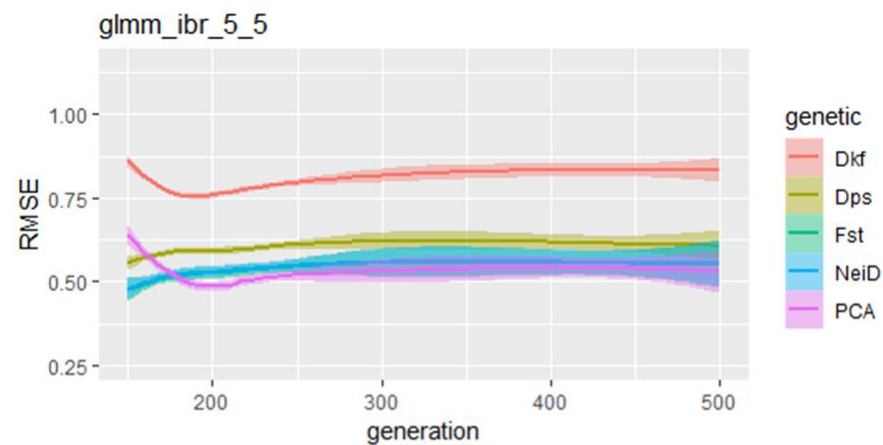
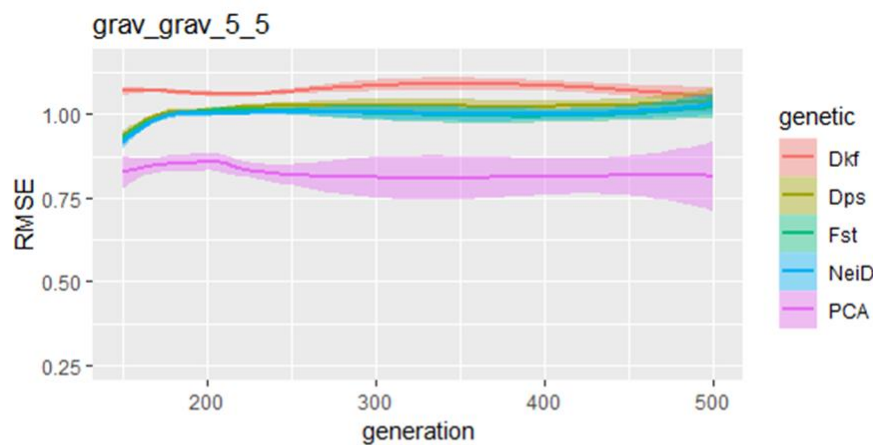
Results -RMSE con .5, hab .5



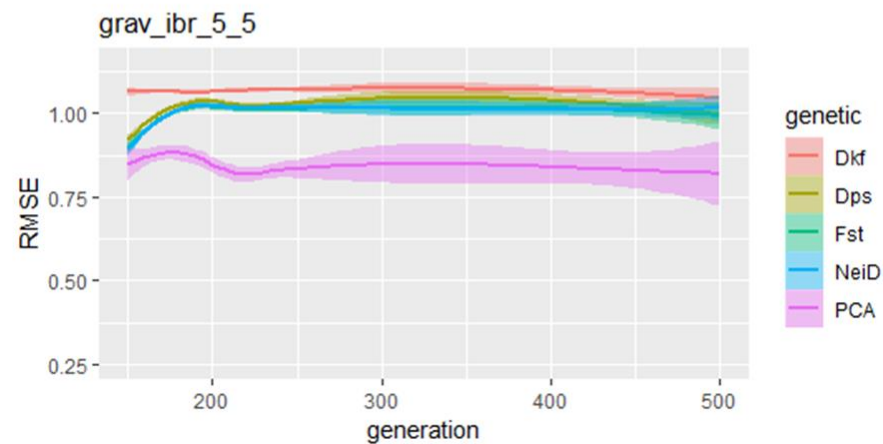
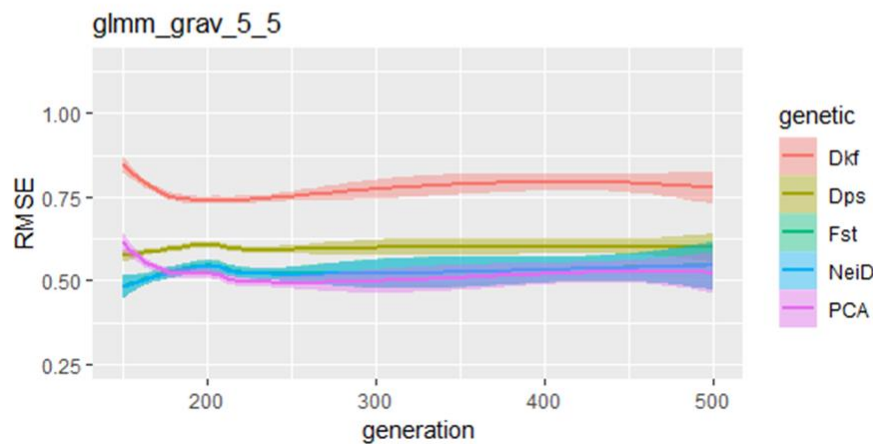
Gravity Driver

IBR Driver

Correct Model



Wrong Model



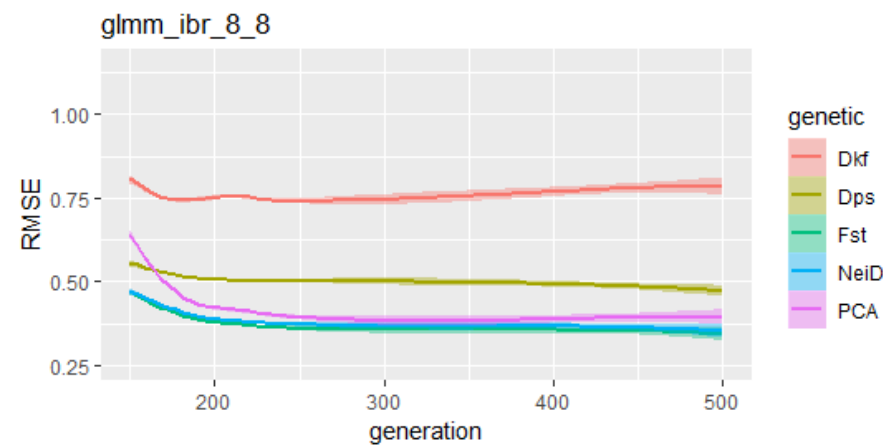
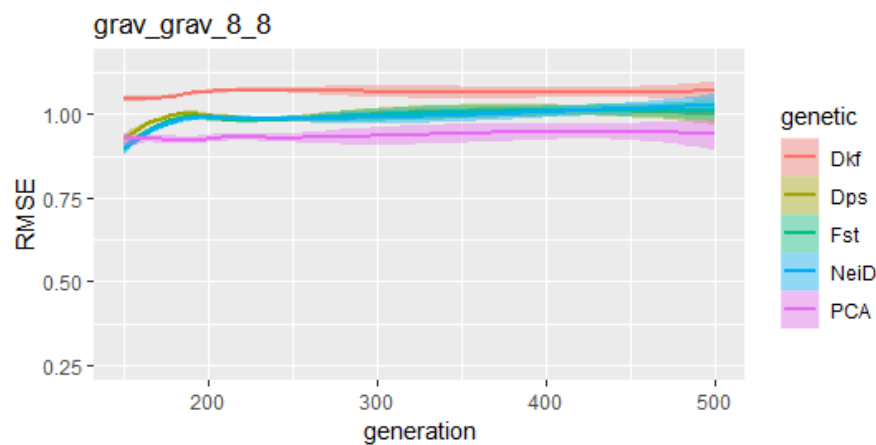
Results -RMSE con .8, hab .8



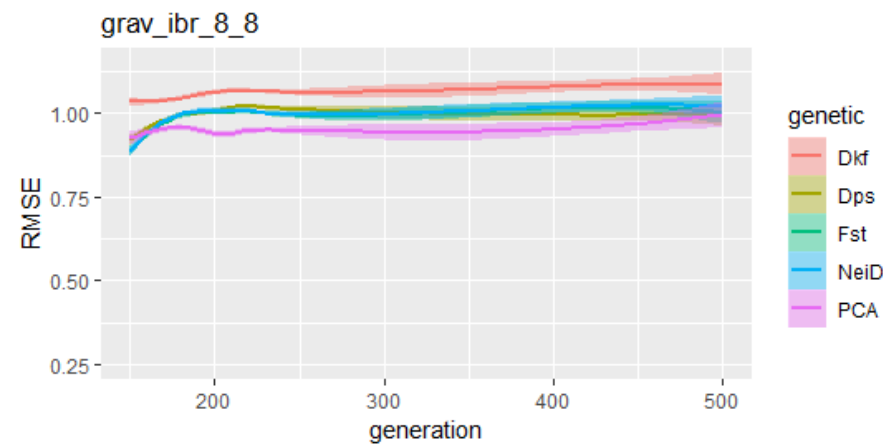
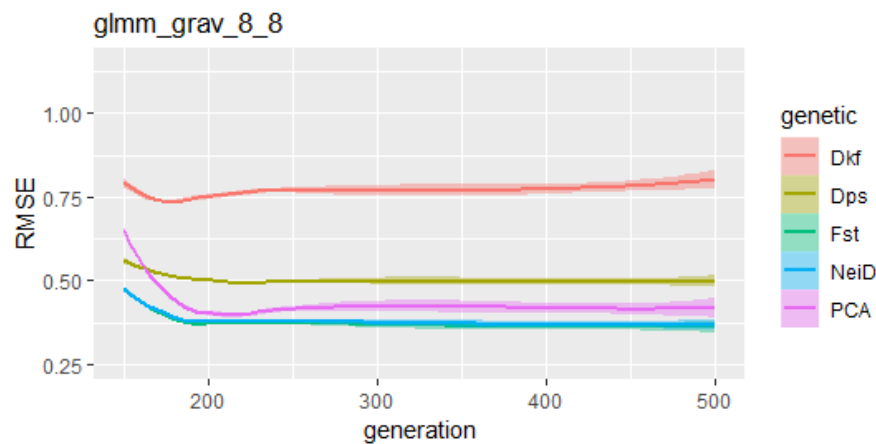
Gravity Driver

IBR Driver

Correct Model



Wrong Model



Discussion

- PCA has limited assumptions around use
- Fst assumes Hardy-Weinberg Equilibrium
- GLMM had lower RMSE than gravity models, even for our predicted correct models
- Expand habitat contagion/habitat combinations and time window

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Questions?

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