



Apport des données génétiques pour modéliser la connectivité du paysage

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Cartographier la connectivité écologique : trames, données génétiques et paysages









Influence de la structure du paysage sur la connectivité des populations de passereaux forestiers

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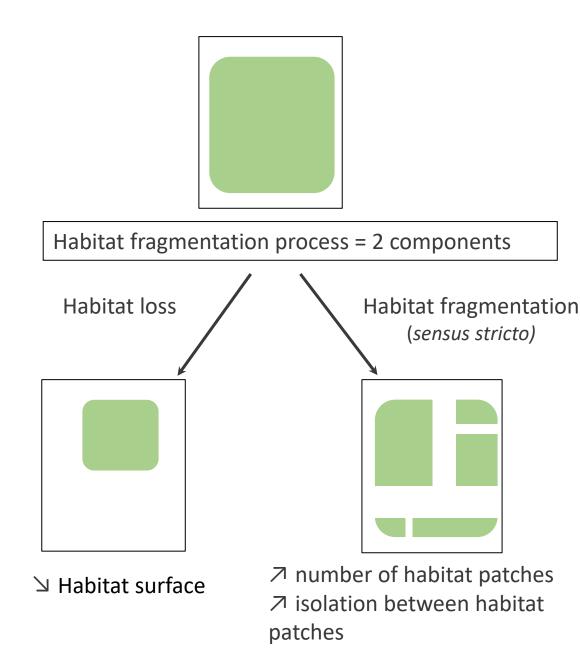
Habitat fragmentation – Forest Fragmentation

Continuous forest habitat

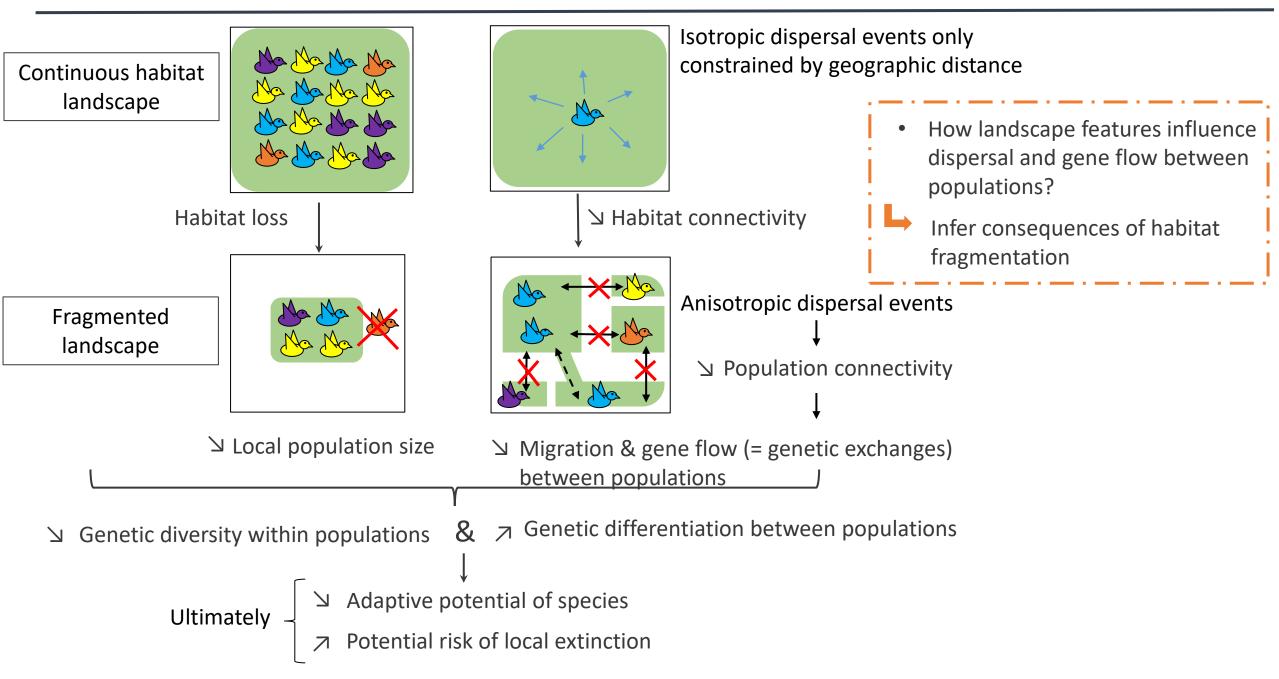


Fragmented forest habitat

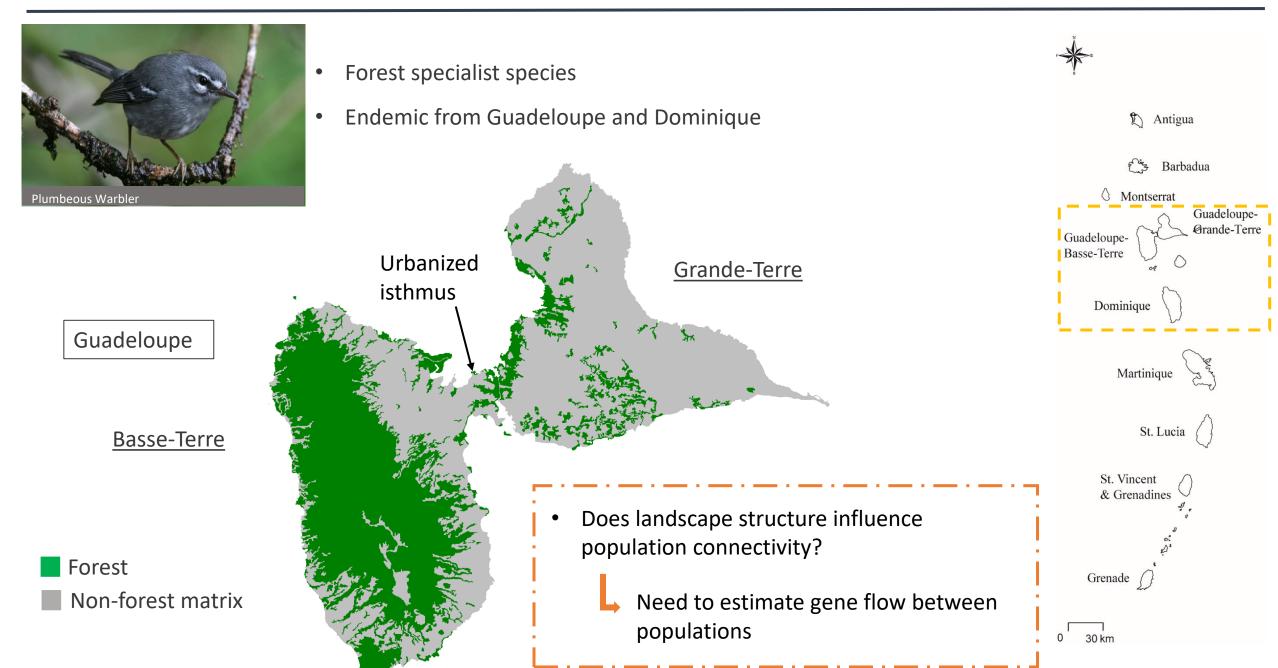




Habitat fragmentation - One of the major threats to biodiversity



A forest specialist bird species – In a fragmented landscape



Indirect estimation of gene flow – genetic data

In the field



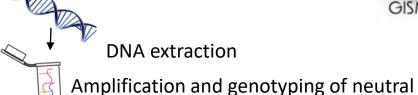
Birds are captured with mist-nets



Blood samples are gently collected for each individual

Plateforme d'écologie moléculaire





genetic regions (microsatellites)

	Pop1	Pop2	Pop3
Pop1	0		
Pop2	X1-2	0	
Pop3	X1-3	X2-3	0

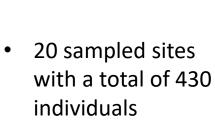
Measure of genetic differentiation between pairs of populations

- Low genetic differentiation = high level of gene flow
- Strong genetic differentiation = low level of gene flow

Genetic similarity between pairs of populations is an estimate of the landscape functional connectivity

D sampled sites Link functional connectivity & structural connectivity

Landscape genetics



Landscape genetics framework – Combining landscape ecology & population genetics

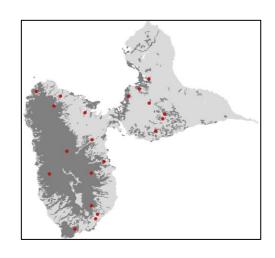
1/ 20 sampled sites

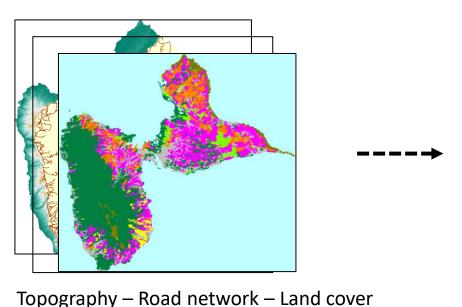
2/ Environmental surfaces (raster)

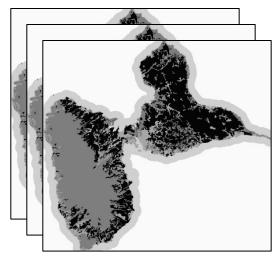
3/ Resistance surfaces (raster)

1 pixel = Altitude value/ 1 land cover class

1 pixel = Resistance value







High resistance

Low resistance

How to compute ecological distances

Genetic data $(12 \mu sats)$

> Matrix of pairwise genetic differentiation between populations **F**ST

Landscape functional connectivity

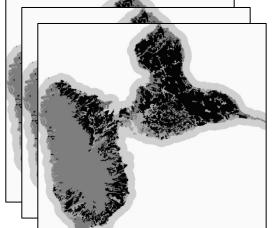
relationship

4/ Test for

Landscape structural connectivity

Matrix of pairwise ecological distances between populations

Indirect estimation of gene flow



Two classical measures of ecological distances

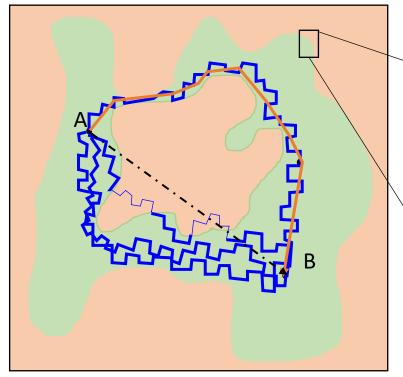
Least cost distance (LCP)

Suitable habitat (low cost value of pixels)Unsuitable matrix (high cost value of pixels)

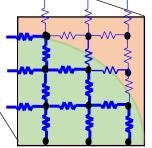
Each population pair is connected following the **optimal route** = least cost cells

Limitation

- Cannot incorporate the effects of
- corridor size
- independant parallel paths connecting A-D



- Resistance distance (McRae 2006)
- Distance metric based in circuit theory Analogy between gene flow and conductance in electronic circuits



Landscape = network of nodes connected to their neighbors by resistors

- Resistance distance integrates all possible paths connecting each population pair

Landscape genetics framework

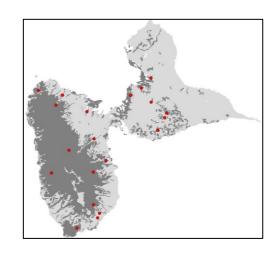
1/ 20 sampled sites

2/ Environmental surfaces (raster)

3/ Resistance surfaces (raster)

1 pixel = Altitude value/ 1 land cover class

1 pixel = Resistance value



Topography – Road network – Land cover

How to derivate resistance surfaces

High resistance

Low resistance



Genetic data $(12 \mu sats)$

> Matrix of pairwise genetic differentiation between populations

Landscape functional connectivity

relationship

4/ Test for

Landscape structural connectivity

Matrix of pairwise ecological distances between populations

FST

Indirect estimation of gene flow

Parameterization of resistance surfaces: no consensus about the best method

A priori assignation of resistance values for each landscape feature

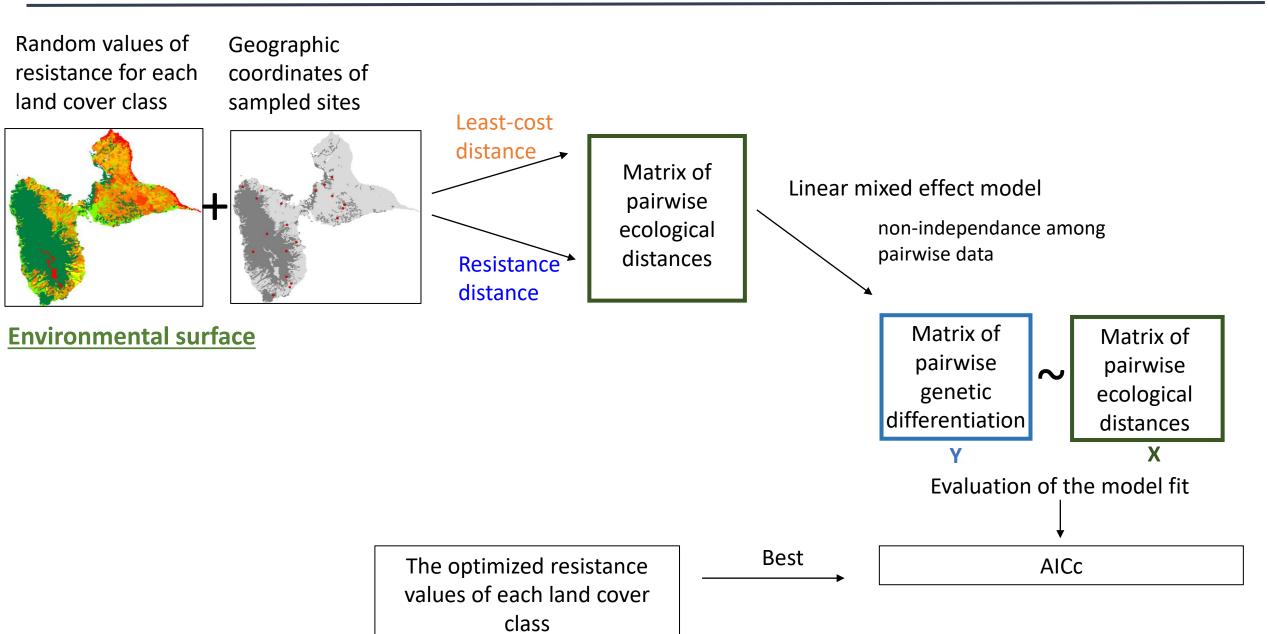
Commonly based on

- Expert optinion about species ecology -> poor performance
- Habitat suitability maps (from SDM; high suitability = low resistance value)

Less ofen based on

- Empirical movement studies
- No such available data for the Plumbeous warbler
 - Optimization procedure using Genetic Algorithms (GA); R package ResistanceGA (Peterman 2004)
 - Exploration of parameter space without any *a priori*

Framework for optimization of resistance values



Landscape genetics approach – General framework

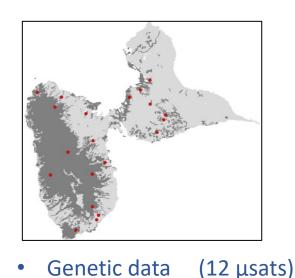
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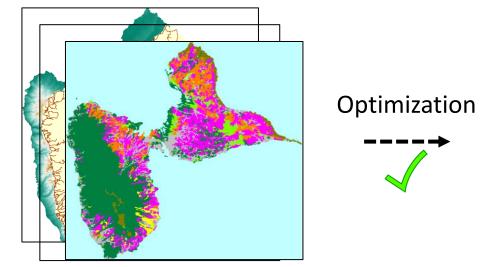
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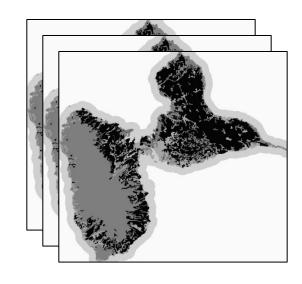
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High resistance

Low resistance

Topography – Road network – Land cover

Resistance & **CP** distances

Matrix of pairwise genetic differentiation

Landscape functional connectivity

relationship

4/ Test for

Landscape structural connectivity

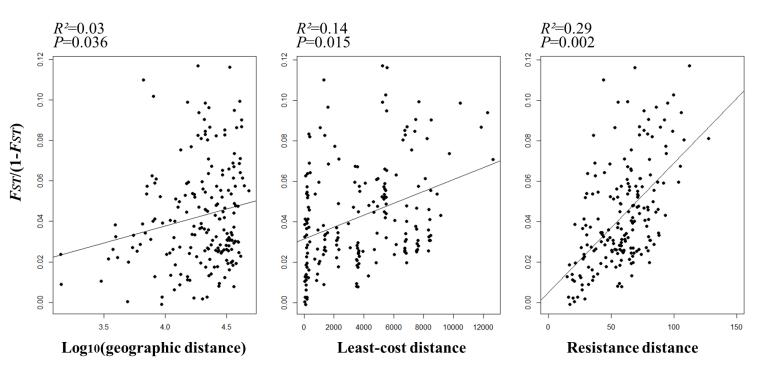
Matrix of pairwise ecological distances between populations

between populations **F**ST

Indirect estimation of gene flow

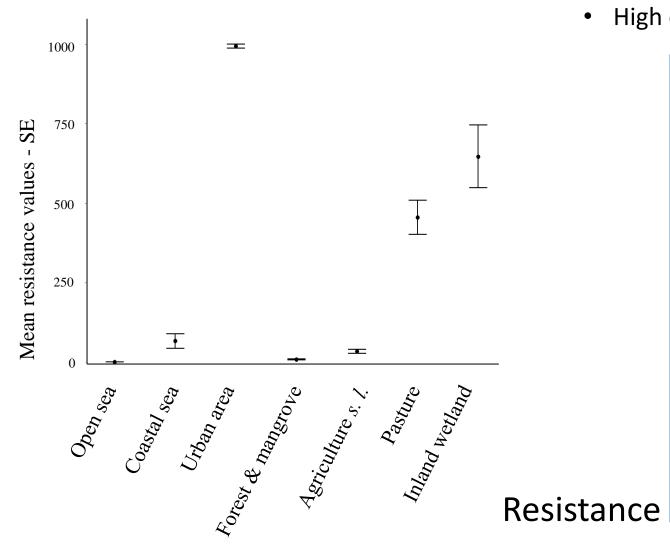
Relative performances of LCP and Resistance distances

Simple Mantel Tests



- Significant simple correlations
 Genetic dist ~ Geographic or Ecological distances
- → Higher variance explained by Resistance distances

Resistance of land cover types – Optimization from resistance distances



High consistency of optimized values across the 10 replicates

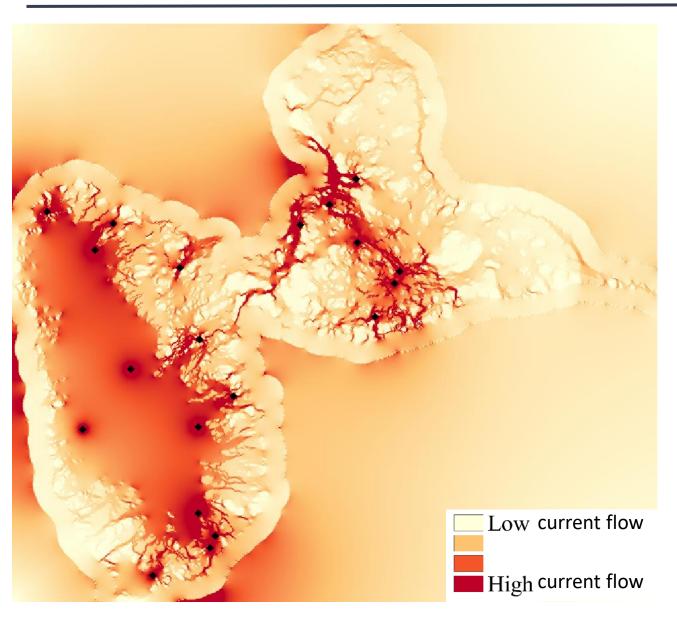
Urban area (1000X)

Pasture and Inland wetland

Agriculture s. l. (4X) and Coastal sea (7X)

Forest & Mangrove and Open sea

Current flow among samples locations



Density of current in each pixel = probability of use during dispersal

- Connectivity between BT-GT populations rely on a narrow corridor of suitable habitat
 - → Special attention in conservation effort

Next steps...

