# CoLa - Connecting landscape

App demo

Founded by



Developed by









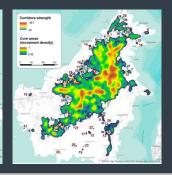
# **Tutorial**

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## Intro





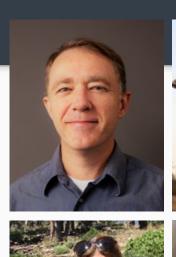


Increasing loss and fragmentation of habitats has resulted in an urgent need to identify areas for conservation that also maintain and enhance ecological connectivity of protected areas. End-users lack easy to use tools to objectively identify connectivity priorities.

**Outcome:** User friendly software to automatically generate and visualize wildlife connectivity priorities and scenarios using habitat suitability layers derived from NASA Earth observations.

**Value:** end-users can generate actionable connectivity priorities, enhancing interagency planning, improving protected area technical cooperation programs, improving sustainable forest management, and informing conservation crime assessments related to poaching and illegal land use.

# The (great!) team



























## Investigators



Patrick Jantz, PI, Northern Arizona University



Scott Goetz, Co-I, Developed by Northern Arizona University



Zaneta Kaszta, Co-I, Northern Arizona University



David Macdonald, Collaborator, Founder, WildCru, U. Oxford



Andrew Loveridge, Collaborator, Deputy Director, WildCru, TKPP, U. Oxford

Sam Cushman, Collaborator, Senjor Fellow, WildCru, U. Oxford

## Students



Ivan Gonzalez, Ph.D. student, Northern Arizona University

### **End Users**





Robynne Kotze, Research Coordinator, WildCru, TKPP, U. Oxford



Nyambe Nyambe, Executive Director, KAZA Secretariat



Kathy Zeller, Research Biologist, Bocky Mountain Research Station, USFS



Andrew Loveridge, Collaborator, Deputy Director, WildCru, TKPP, U. Oxford









## Agenda

0 - 10 min: Welcome

10 - 20: The tool. Main directions

20 - 30: Suitability to resistance

30 - 40: Edit scenarios

40 - 50: Simulate/load points

50 - 60: Corridors

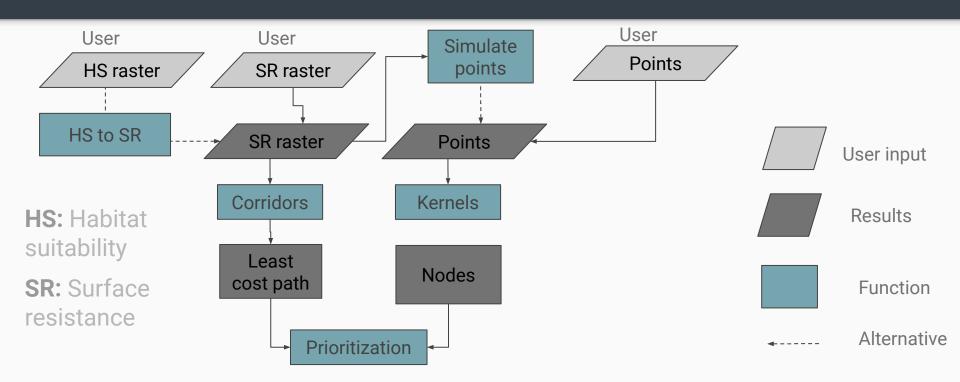
60 - 70: Kernels

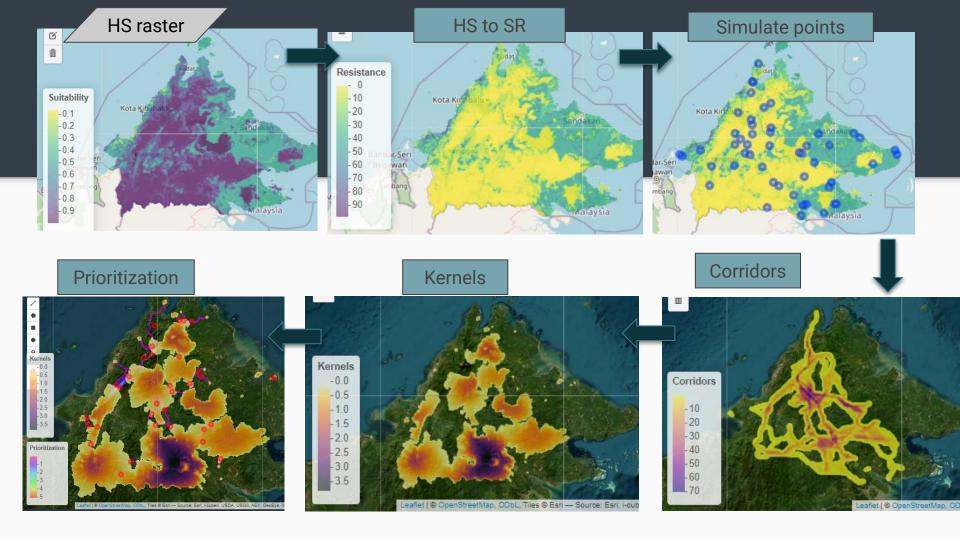
70 - 80: Prioritize

80 - 100: Debug + known issues

100 - 120m: Concluding remarks

## Complete workflow





## Overview

#### ConnectingLandscapes

- **♠** Home
- Customize resistance surface
- ? Create source points
- ⊞ Cost distance matrix

CDPOP

- ده Connectivity corridors
- ?" Connectivity dispersal kernels
- ▼ Connectivity prioritization
- Assign coords
- 77 2745F
- ♥ PDF
- ¥ Run locally

- 10	71		7	15
Home	How it works	Performance	Showcase	ShowcasePriv

#### **Toolkit Overview**

Increasing loss and fragmentation of habitats has resulted in an urgent need to identify areas for conservation that also maintain and enhance ecological connectivity of protected areas. The Connecting Landscapes (CoLa) toolkit integrates and enhances numerous landscape genetics and habitat connectivity tools (Landguth and Cushman 2010, Landguth et al. 2012) to aid in identifying connectivity conservation priorities and areas of potential human wildlife conflict.

The toolkit is supported by NASA Biological Diversity & Ecological Conservation program Grant No. 80NSSC21K1942 - Strengthening Natural Resource Management with New Protected Area Connectivity Tools, and was co-developed with the Wildlife Conservation Research Unit (WildCRU) of University of Oxford and the United States Forest Service (USFS) Rocky Mountain Research Station. USFS International Programs.

Primary inputs to the tools include habitat suitability layers or spatially explicit estimates of animal movement potential across different land use types (resistance layers), and population source points. When properly parameterized, the toolkit can generate estimates of population genetic structure, population density, core movement areas, and long-distance dispersal corridors.

For recent examples of how the individual tools have been used for wildlife research and conservation assessments see Kaszta et al. (2020a), Kaszta et al. (2020b), Zeller et al. (2021), Ash et al. (2023), Makwana et al. (2023).

#### >> References

Ash, E., Cushman, S., Kaszta, Ž., Landguth, E., Redford, T. and Macdonald, D.W., 2023. Female-biased introductions produce higher predicted population size and genetic diversity in simulations of a small, isolated tiger (Panthera tigris) population. Scientific Reports, 13(1), p.11199.

Kaszta, Ž., Cushman, S.A. and Macdonald, D.W., 2020a. Prioritizing habitat core areas and corridors for a large carnivore across its range. Animal Conservation, 23(5), pp.607-616.

Kaszta, Ż., Cushman, S.A., Htun, S., Naing, H., Burnham, D. and Macdonald, D.W., 2020b. Simulating the impact of Belt and Road initiative and other major developments in Myanmar on an ambassador felid, the clouded leopard. Neofelis nebulosa. Landscape Ecology, 35, pp.727-746.

Landguth, E.L. and Cushman, S., 2010. CDPOP: a spatially explicit cost distance population genetics program. Molecular ecology resources, 10(1), pp.156-161.

Landguth, E.L., Hand, B.K., Glassy, J., Cushman, S.A. and Sawaya, M.A., 2012. UNICOR: a species connectivity and corridor network simulator. Ecography, 35(1), pp.9-14.

Makwana, M., Vasudeva, V., Cushman, S.A. and Krishnamurthy, R., 2023. Modelling landscape permeability for dispersal and colonization of tigers (Panthera tigris) in the Greater Panna Landscape, Central India. Landscape Ecology, 38(3), pp.797-819.

Zeller, K.A., Schroeder, C.A., Wan, H.Y., Collins, G., Denryter, K., Jakes, A.F. and Cushman, S.A., 2021. Forecasting habitat and connectivity for pronghorn across the Great Basin ecoregion. *Diversity and Distributions*, 27(12), pp.2315-2329.

>> Contact

Write Patrick Jantz@nau.edu or ig299@nau.edu (Ivan Gonzalez) for questions and suggestions

## Overview

Home How it works Performance Showcase ShowcasePriv

#### User guide

Depending on the function you want to run you will need different set of inputs.

Each tab have the following goal:

#### Home:

- Project description
- Performance
- · Examples and showcase

Habitat suitability (HS) to surface resistance (SR): Converts HS into SR using user parameters Inputs:

- HS georreferenced raster
- User parameters

Outputs: SR raster (TIF)

Create source points: Inputs:

- SR georreferenced raster
- User parameters

Outputs: Points (SHP)

#### Cost distance matrix: Inputs:

- SR georreferenced raster
- · Coordinates or spatial points

User parameters

Outputs: Matrix (CSV)

CDPOP. Inputs: SR raster Coordinates or spatial points Distance matrix User parameters

Outputs: Population data (xy)

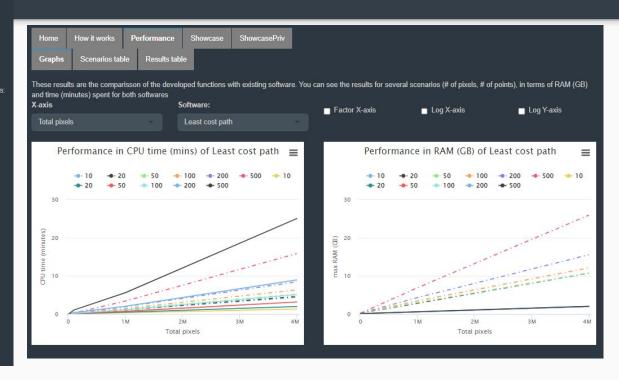
Landscape genetics: Inputs: Outputs: SR raster (TIF)

#### Corridors: Inputs:

- · SR georreferenced raster
- Coordinates or spatial points
- User parameters

Outputs: Corridors (TIF)

## Performance

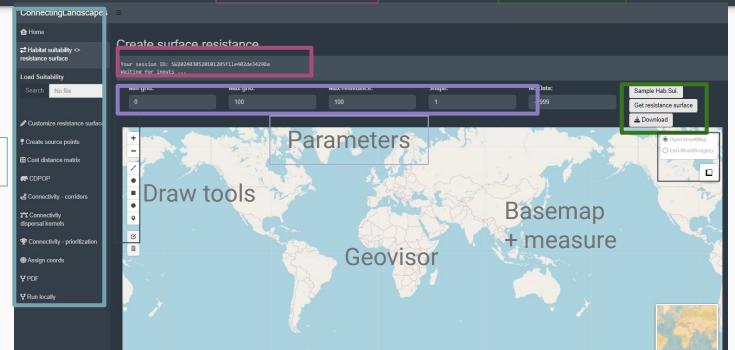


## The visuals

**Functions** 

Log box + session ID

Run / download

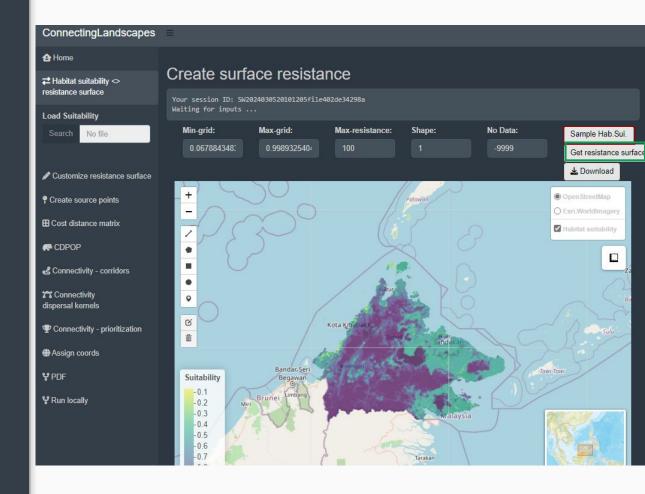


Save your session ID for questions or bugs

# 1. Suitability to resistance

#### For running:

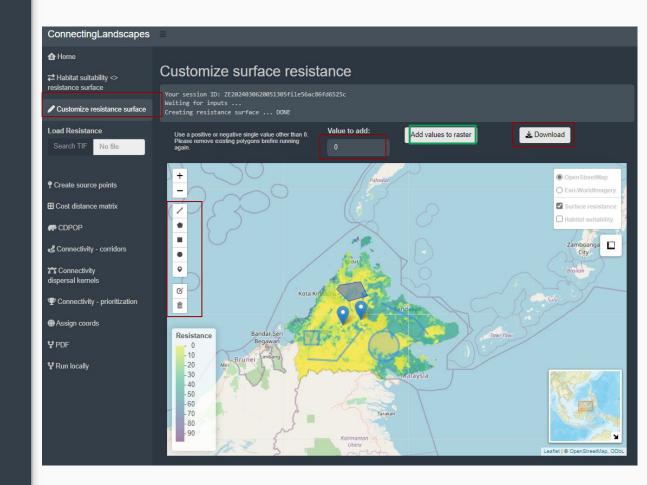
- Load a valid TIF raster with continuous values
- 2. Define the parameters. They are gess forum your raster as a prior. Modify as you consider
- 3. Click on "Get resistance surface"
- 4. Download the resulting raster in TIFF format



No data? Load the sample dataset

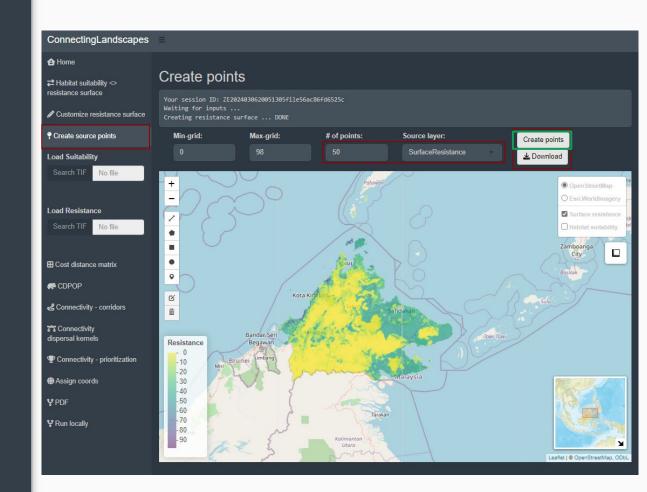
# 2. Customize resistance surface

- Load a valid TIF raster with continuous values if skipping last step. No extra SR required
- Draw figures you want to "burn".
  Multiple geometry types allowed
- Provide a "value to add".Number positive or negative
- 4. Click on "Add values to raster"
- 5. Download the resulting raster in TIFF format



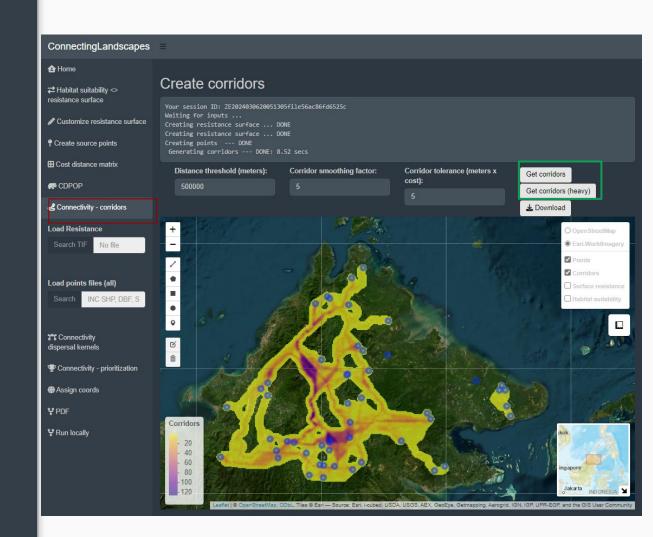
# 3. Create source points

- Load a valid TIF raster with continuous values if skipping previous steps. Can use HS or SR
- Select the min-max range on the raster to simulate the points, and the total number of points
- 3. Select which (available) layer to base the analysis on.
- 4. Click on "Create points"
- Download the resulting raster in TIFF format



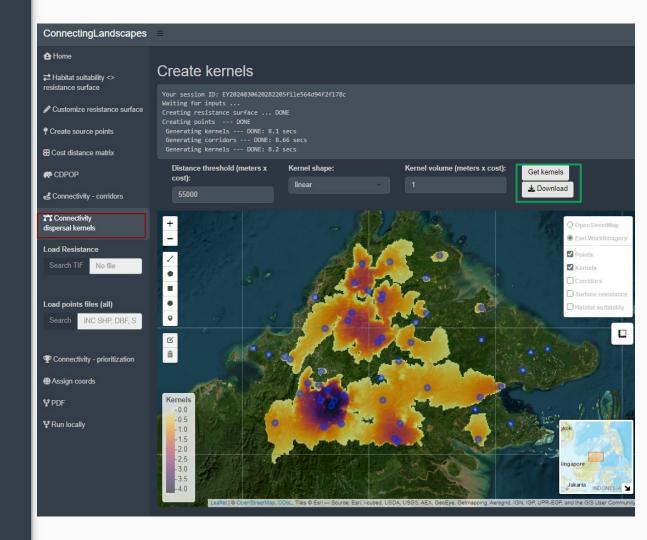
## 4. Corridors

- Load a valid TIF raster with continuous values if skipping previous steps. No required to upload SR again
- Load a valid point shapefile if skipped previous steps
- 3. Select parameters
- Click on "Get corridors" or "heavy" for big rasters.
- Download the resulting raster in TIFF format



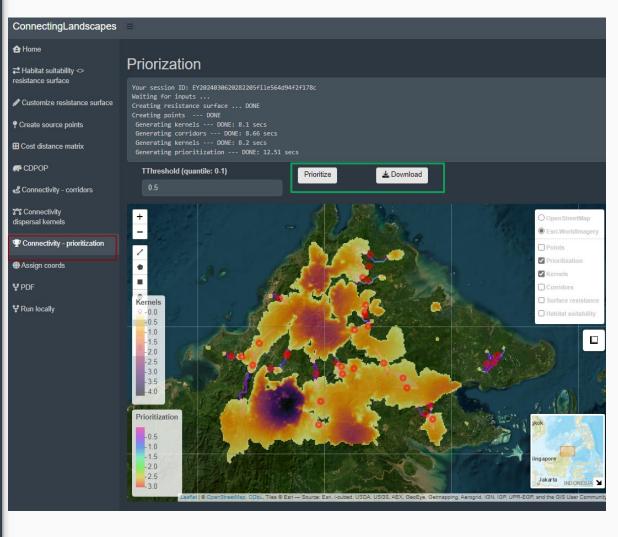
## 5. Kernels

- Load a valid TIF raster with continuous values if skipping previous steps. No required to upload SR again
- Load a valid point shapefile if skipped previous steps
- 3. Select parameters
- 4. Click on "Get kernels"
- Download the resulting raster in TIFF format



## 6. Prioritization

- Must have corridors and kernels developed on the app
- 2. Select parameters
- 3. Click on "Prioritize"
- Download the resulting raster in TIFF format





## Known issues

It will not stop the algorithm but is good to keep this in mind:

- 1. Use a valid NODATA value in your raster, and be aware of which it is
  - 2. Assign a coordinate reference system
    - 3. Your TIFs are big

## Thanks!

Contact us:

Patrick.Jantz@nau.edu

Iván González ig299@nau.edu

github.com/forestrev/

