

Making better conservation decisions



Jeffrey Hanson



jeffrey.hanson@uqconnect.edu.au



jeffrey-hanson.com

Acknowledgements

Amanda Martin

Angela Brennan

Brandon Edwards

Caitlyn Proctor

Emma Hudgins

Matthew Strimas-Mackey

Iadine Chadès

Hugh Possingham

Jaimie Vincent

Jenny McCune

Joseph Bennett

Josie Hughes

Lenore Fahrig

Nina Morell

Peter Arcese

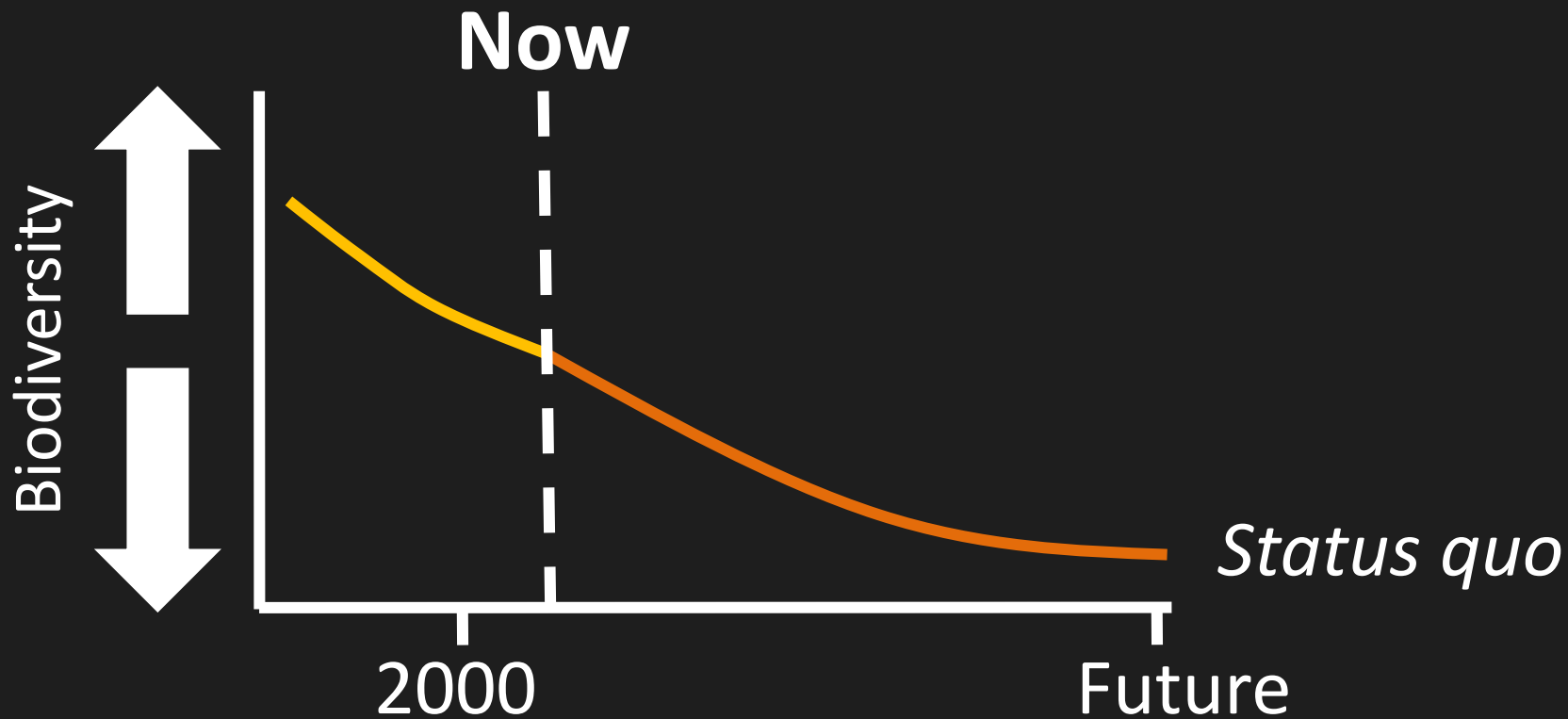
Richard Schuster

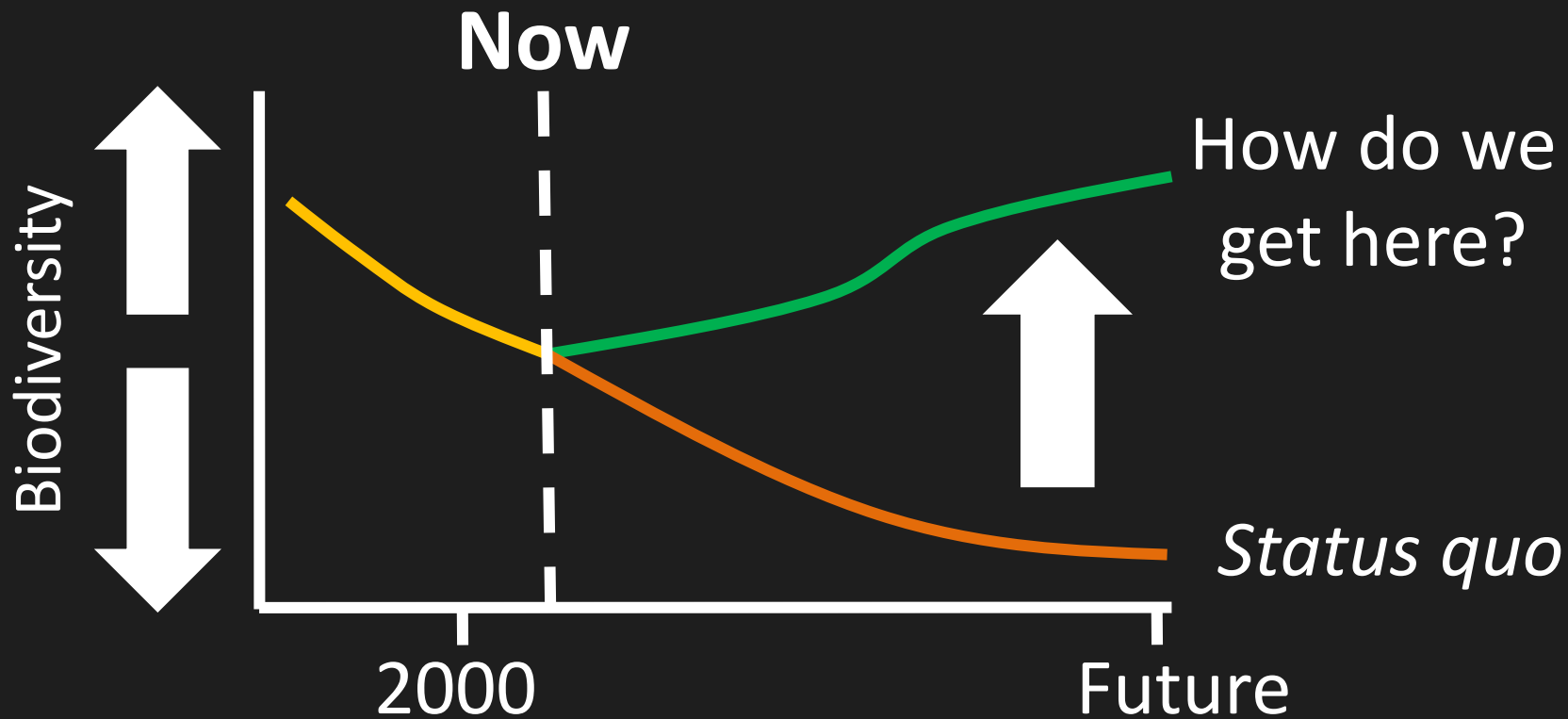
Richard Pither



Carleton
UNIVERSITY





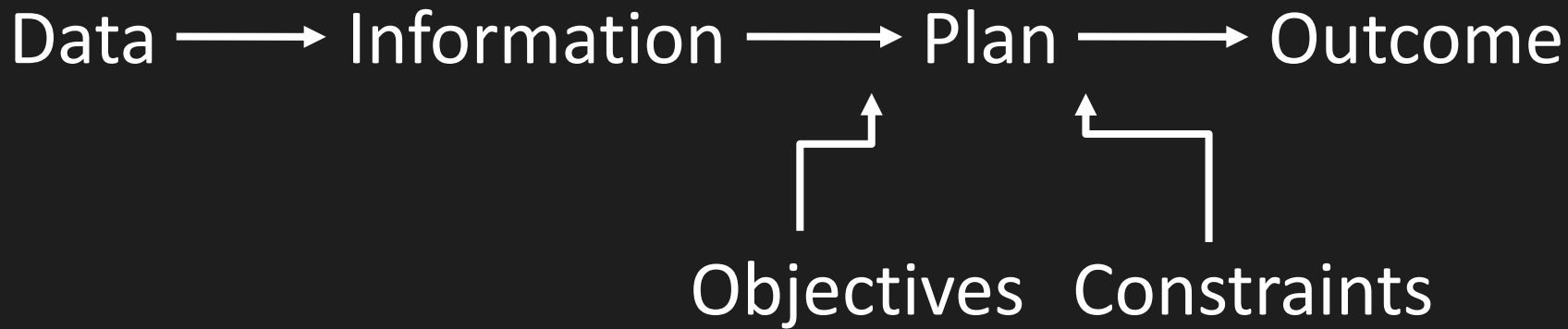
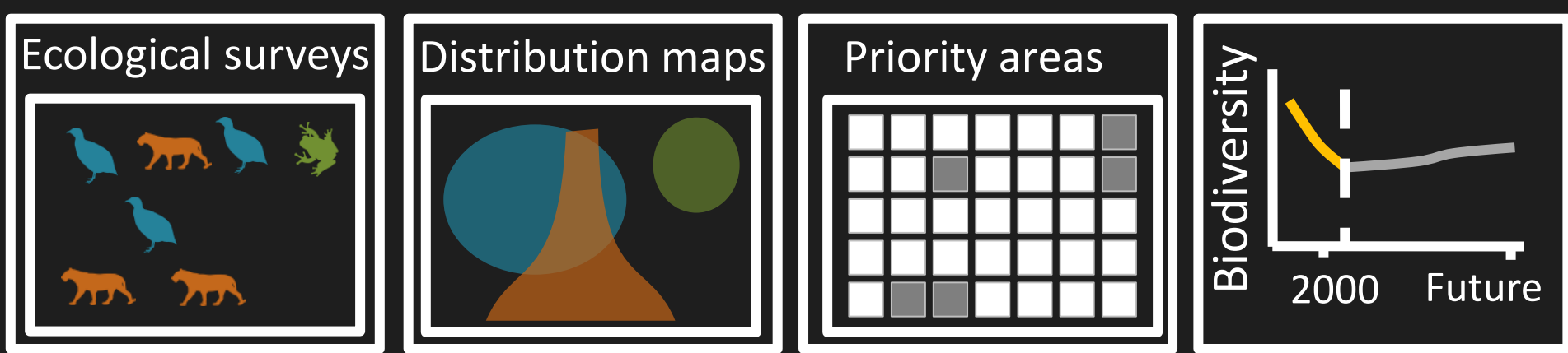


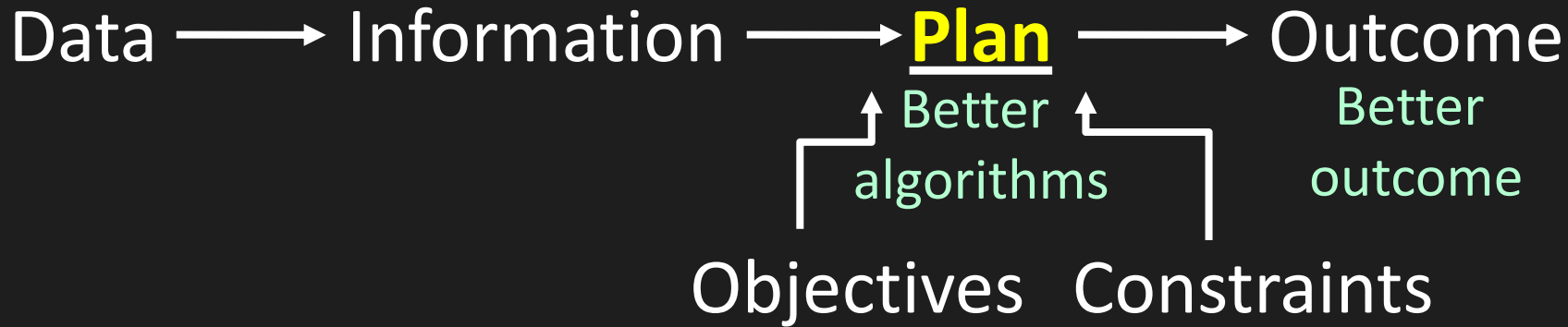
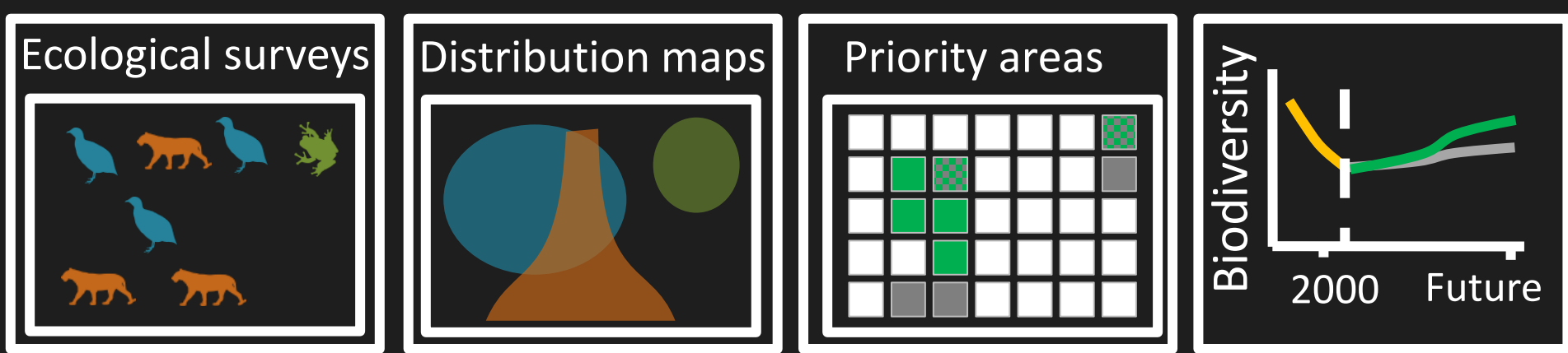
How can we get a better conservation decision?

(1) Better algorithms

(2) Better data

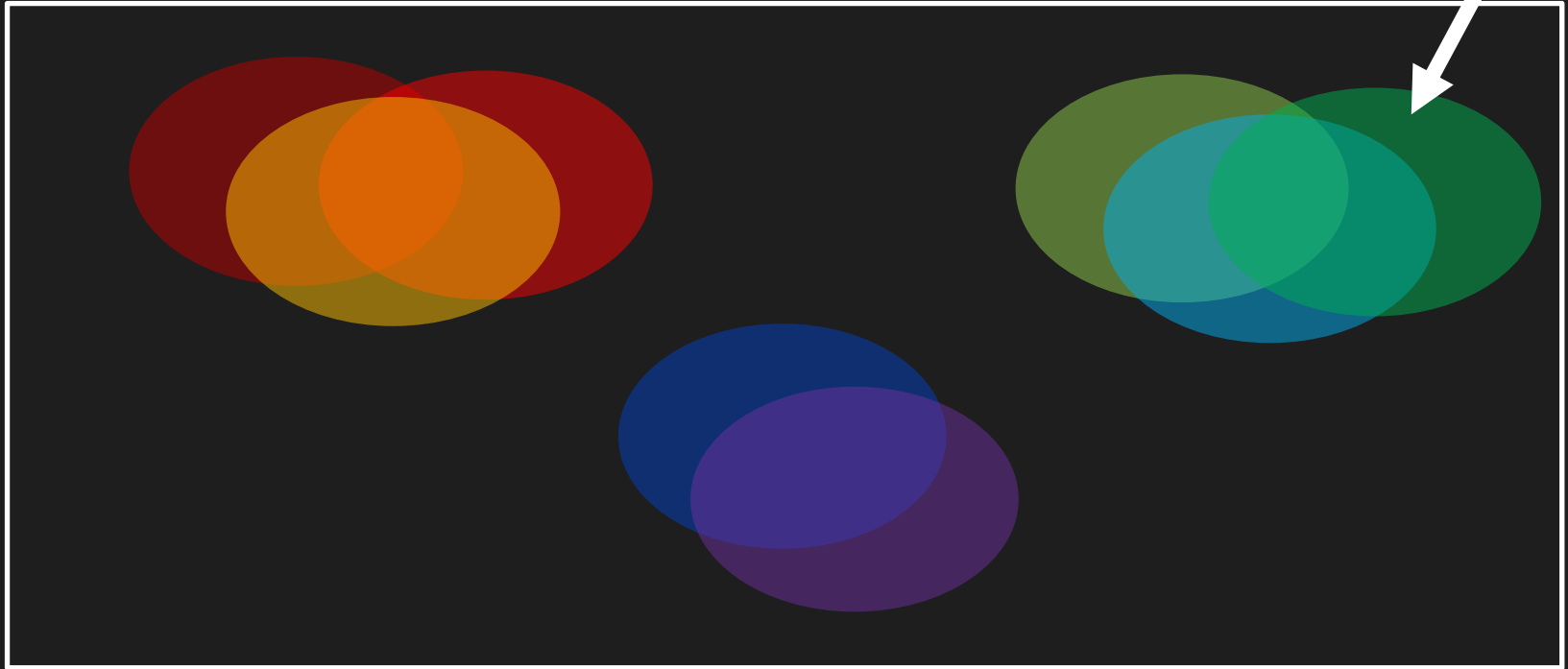
(3) Better surrogates





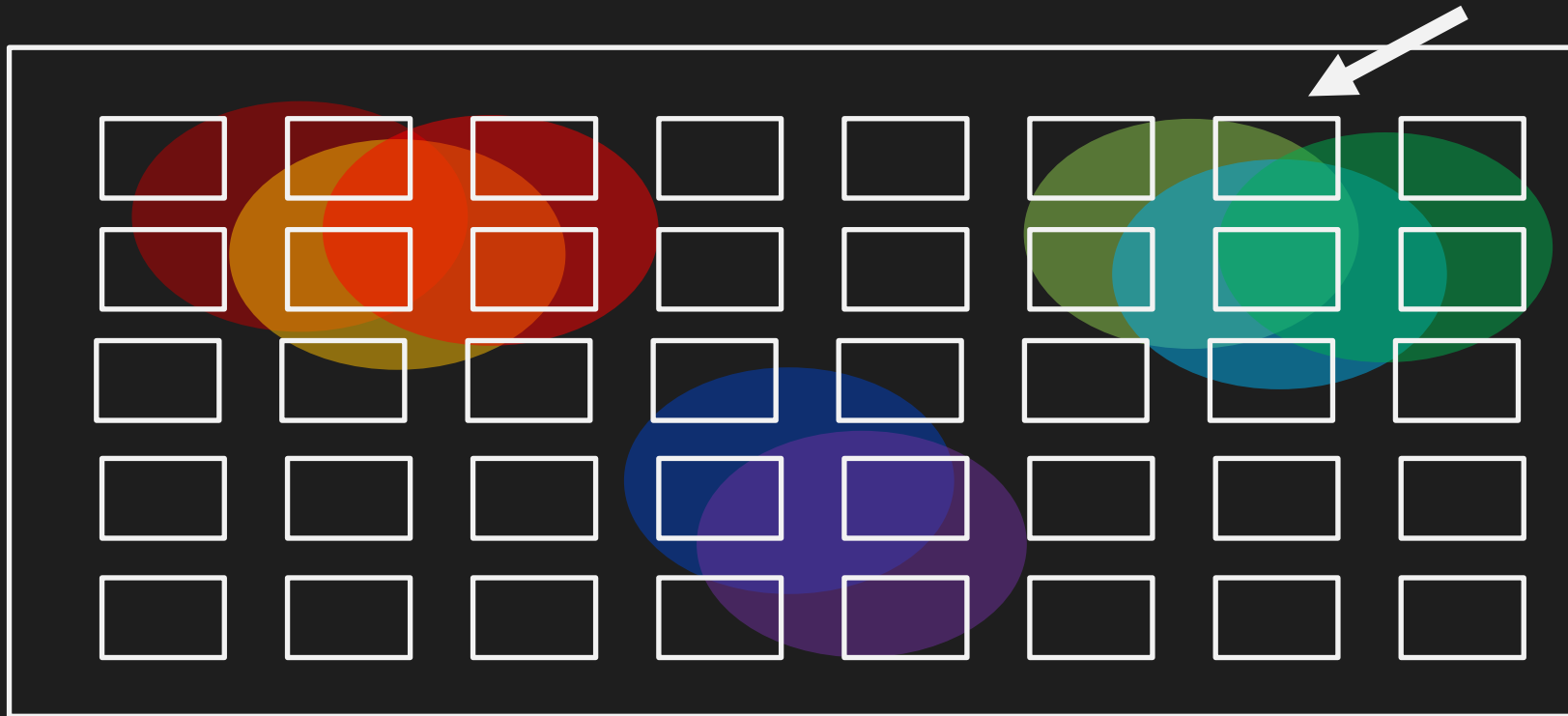
Reserve selection

Features

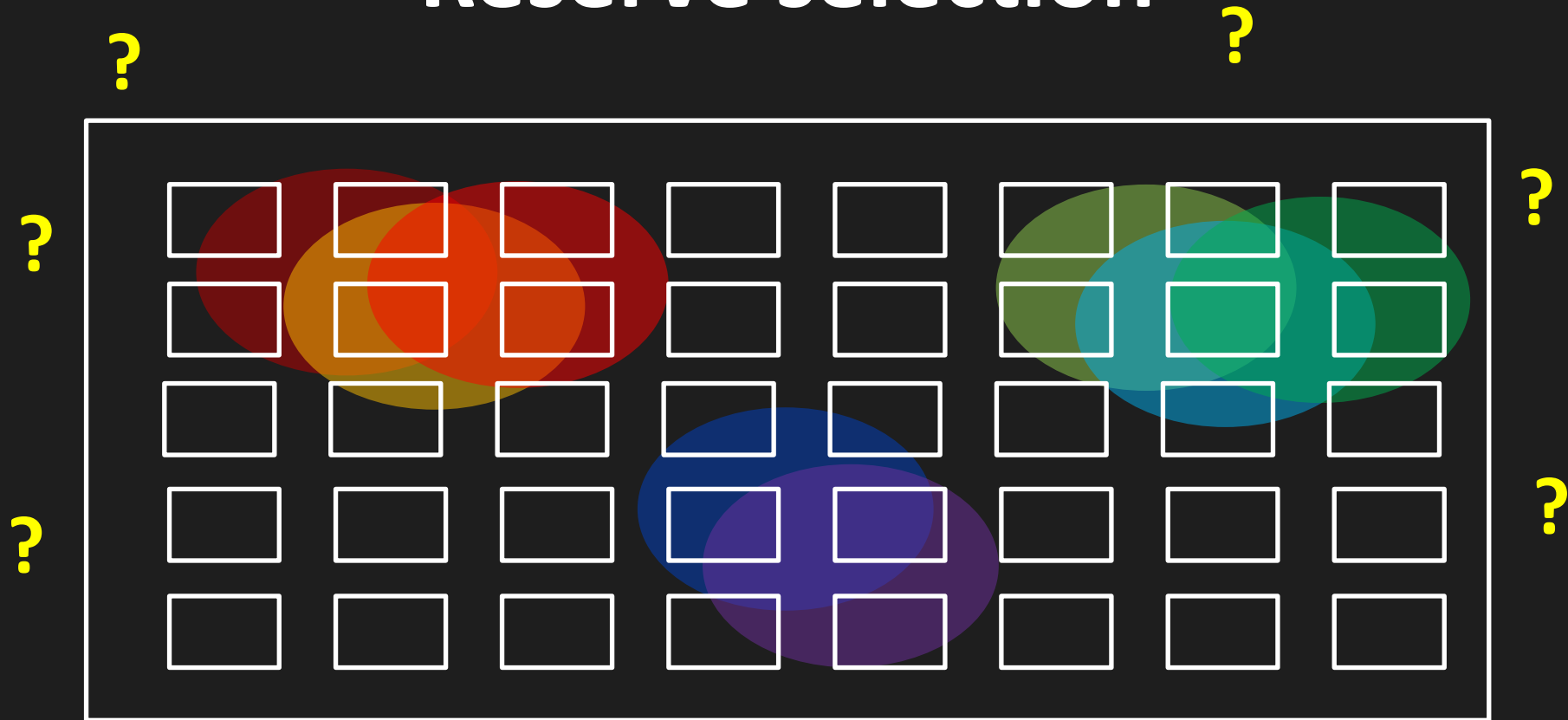


Reserve selection

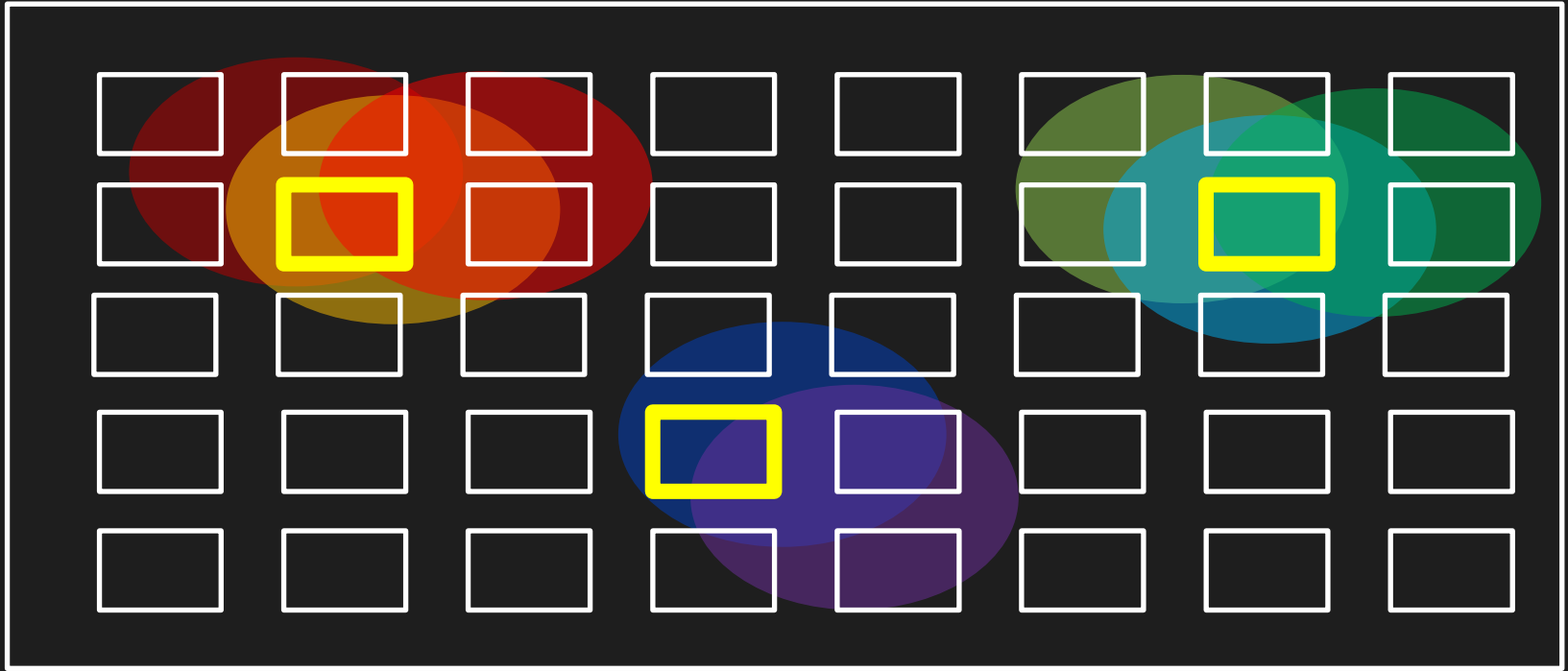
Planning units



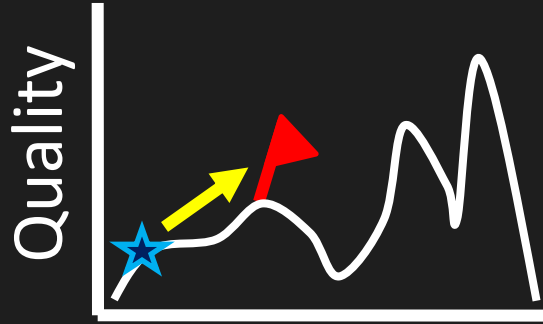
Reserve selection



Reserve selection



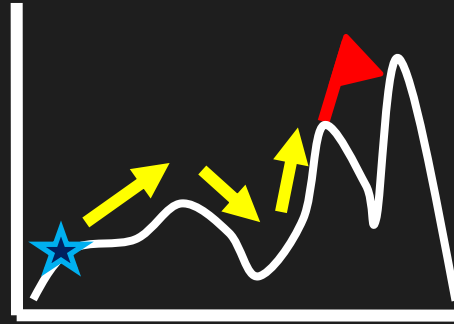
Heuristic algorithm



Different solutions



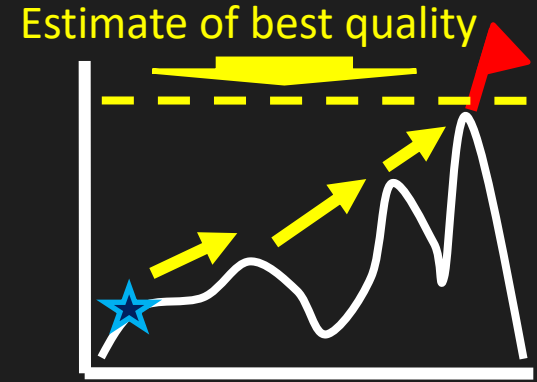
Meta-heuristic algorithms



Different solutions



Exact algorithms



Different solutions



prioritizr: Systematic conservation prioritization in R

- It's an R package (yes, this is good)
- First released on CRAN in 2016
- Highly flexible interface
- Supports tabular and GIS data formats
- Supports multiple zones/actions
- Powered by open source and commercial exact algorithm solvers
- Want to see a worked example? Ask me afterwards during question time

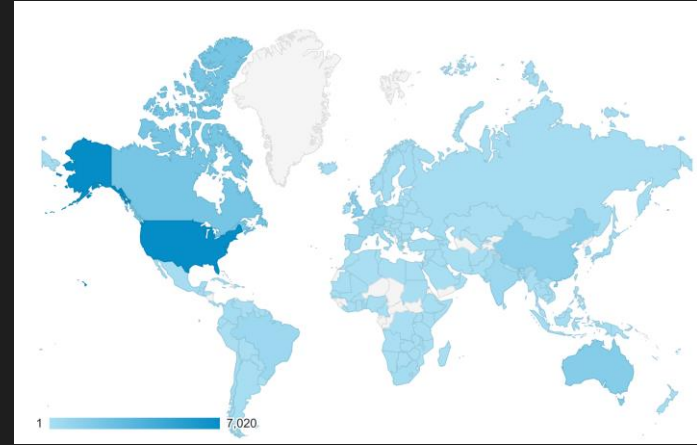


prioritizr.net

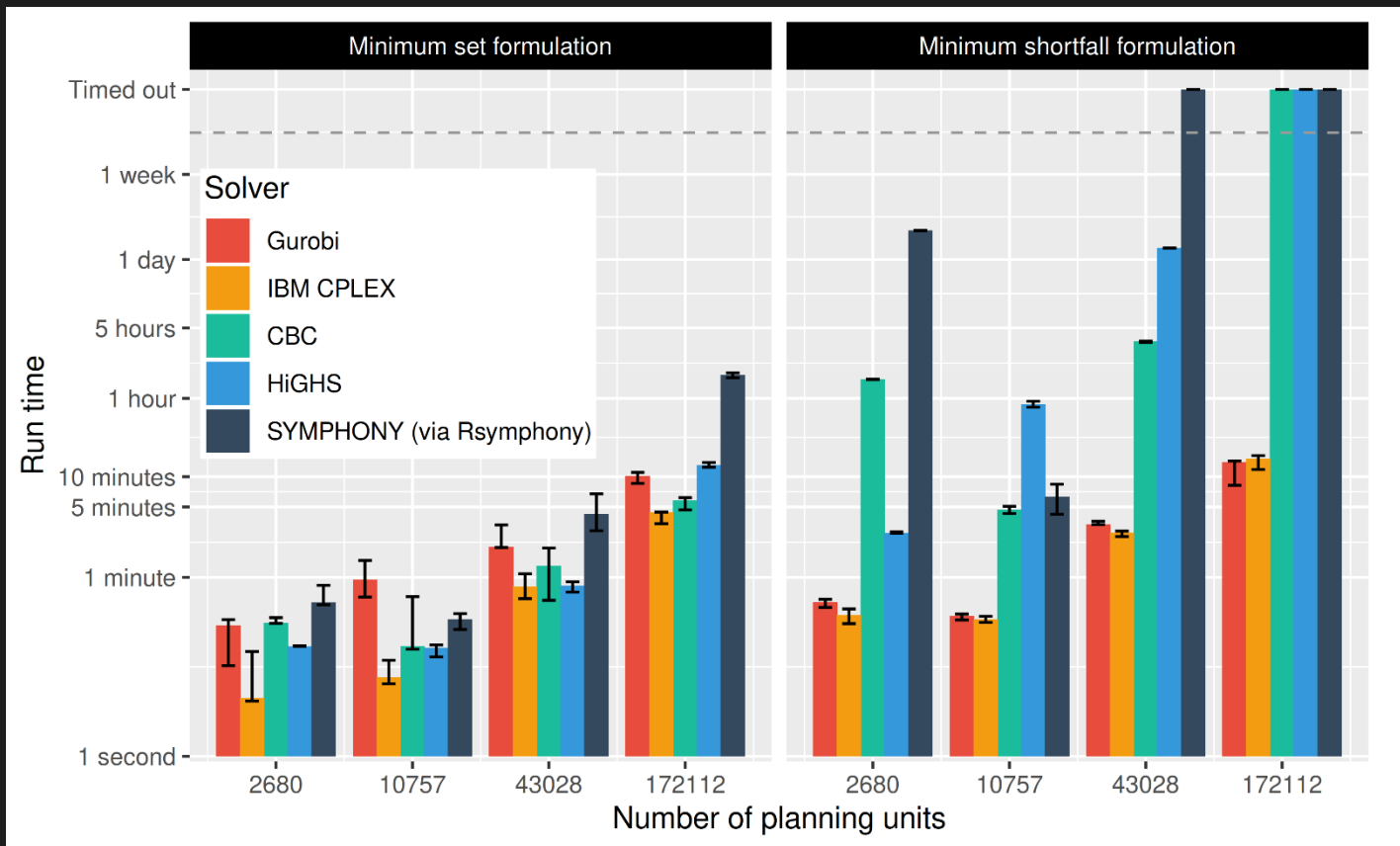
Who uses it?

- 85 scientific publications used it for generating prioritizations
- Conservation organizations: Nature conservancy Canada, UN Biodiversity Lab, Waitt Institute
- Helped inform planning for Government of Monsterrat, Scottish Government, US Geological Survey, Governments of the Maldives and the Federated States of Micronesia

Web site visitors



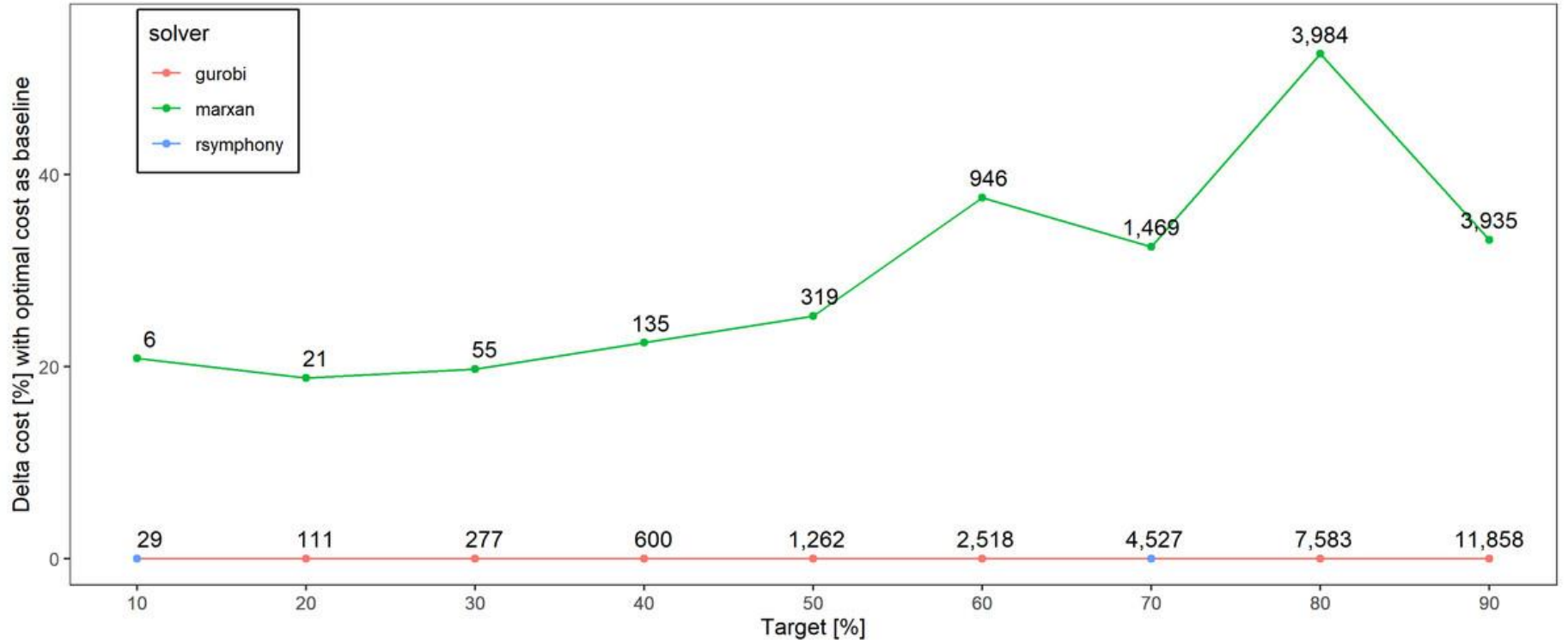
Solve problems pretty quickly!



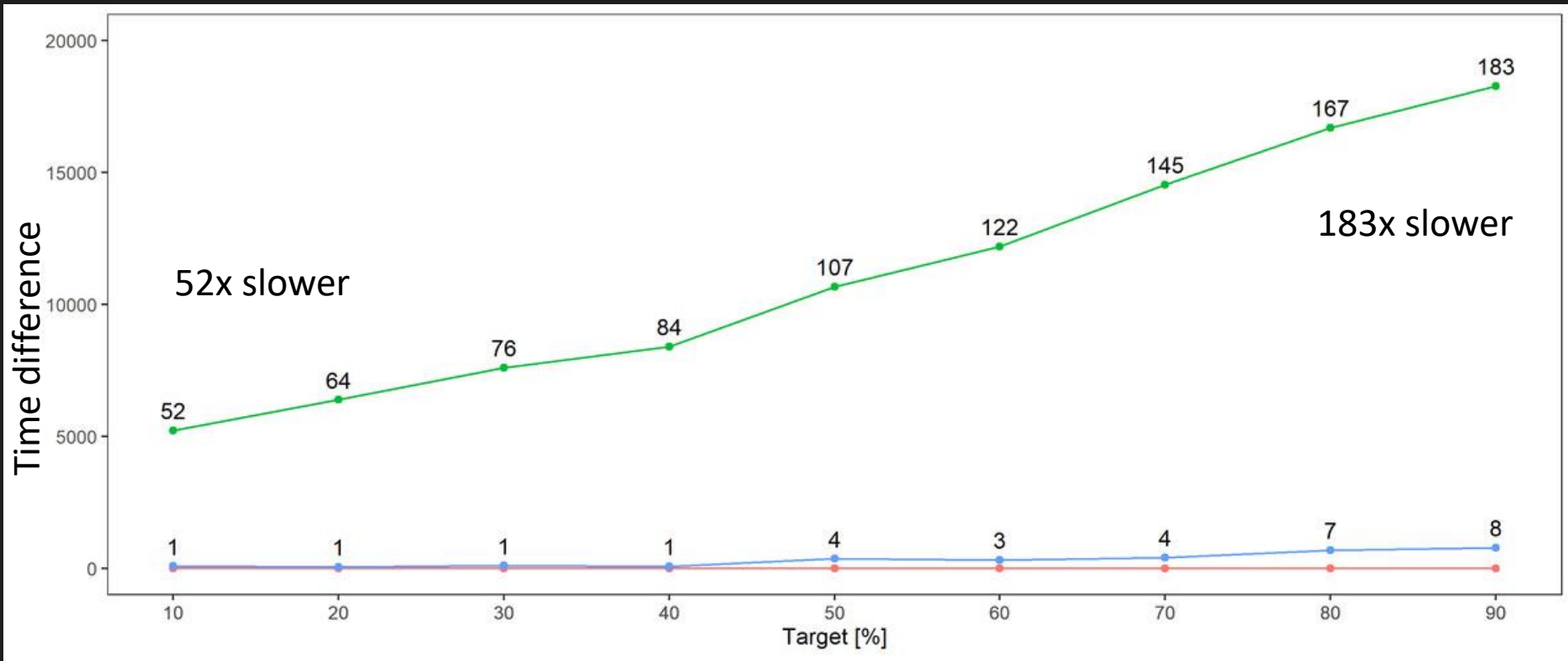
Notes: 396 features, no planning units locked in or out, prioritizations $\leq 10\%$ from optimality

Hanson *et al.* under review

Better solutions



Faster too

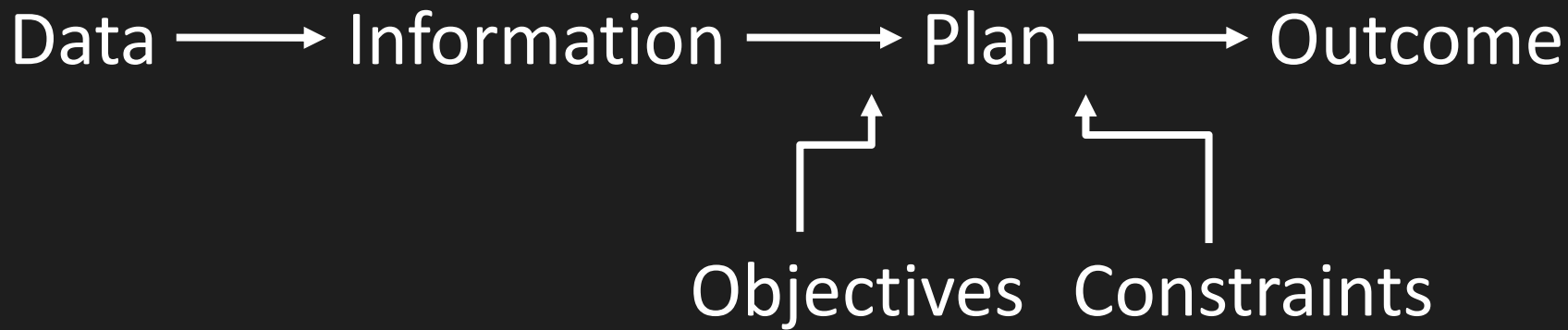
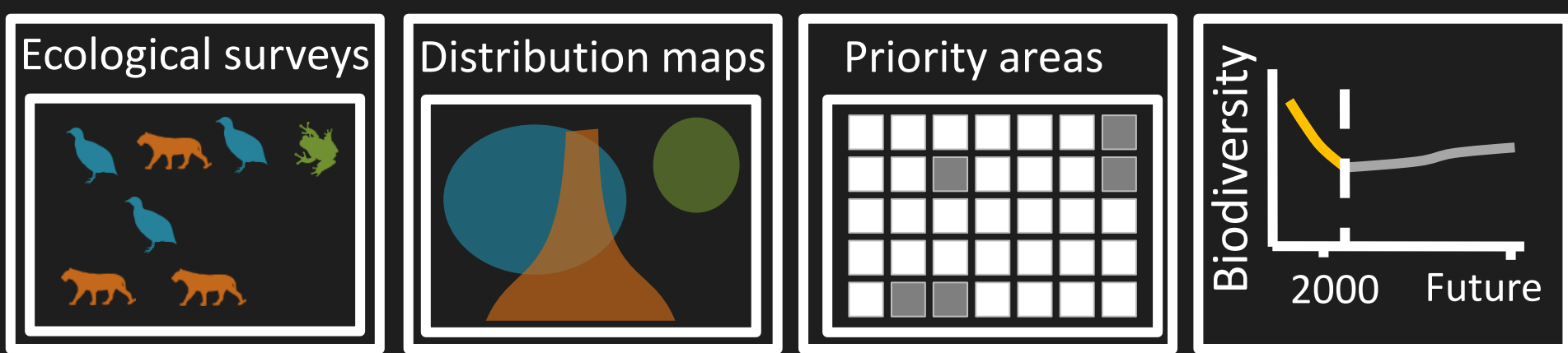


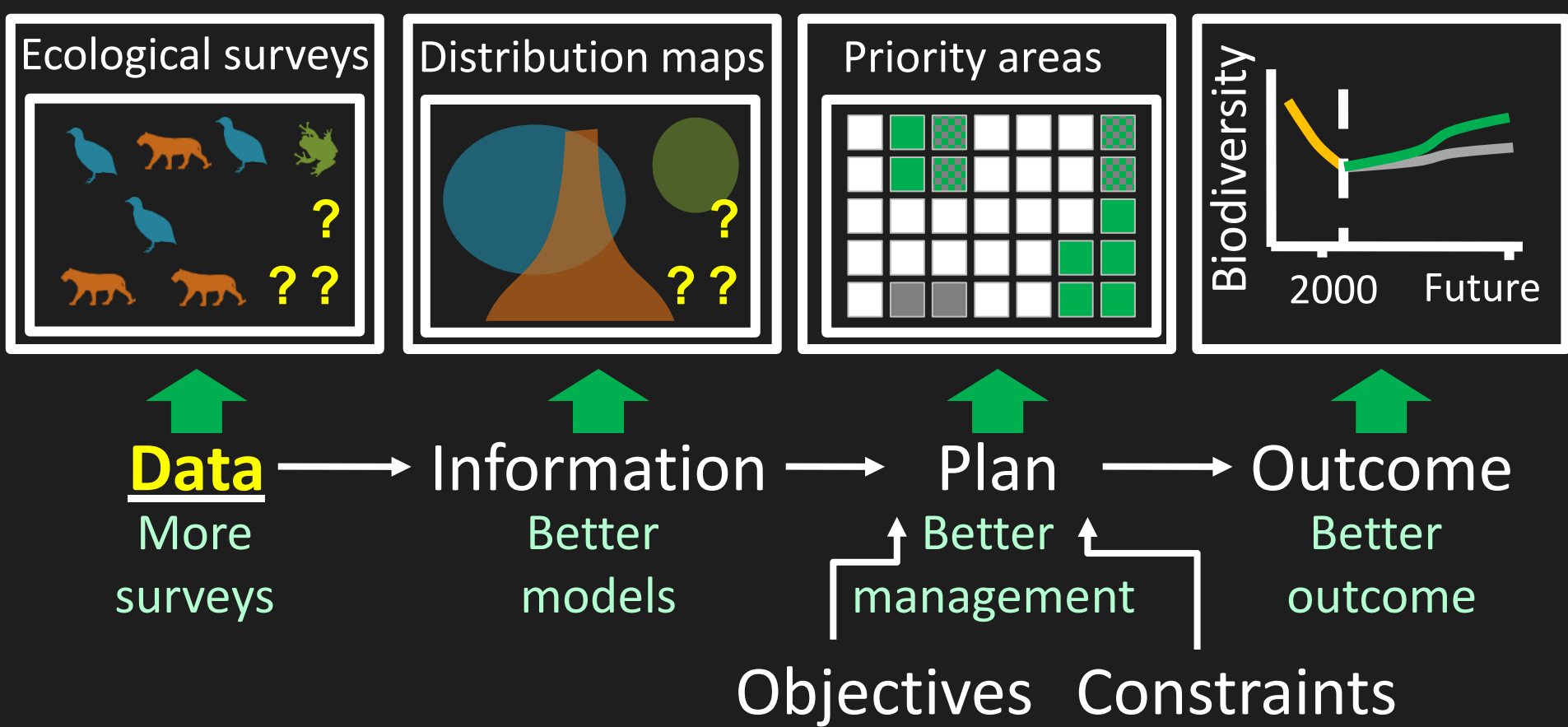
How can we get a better conservation decision?

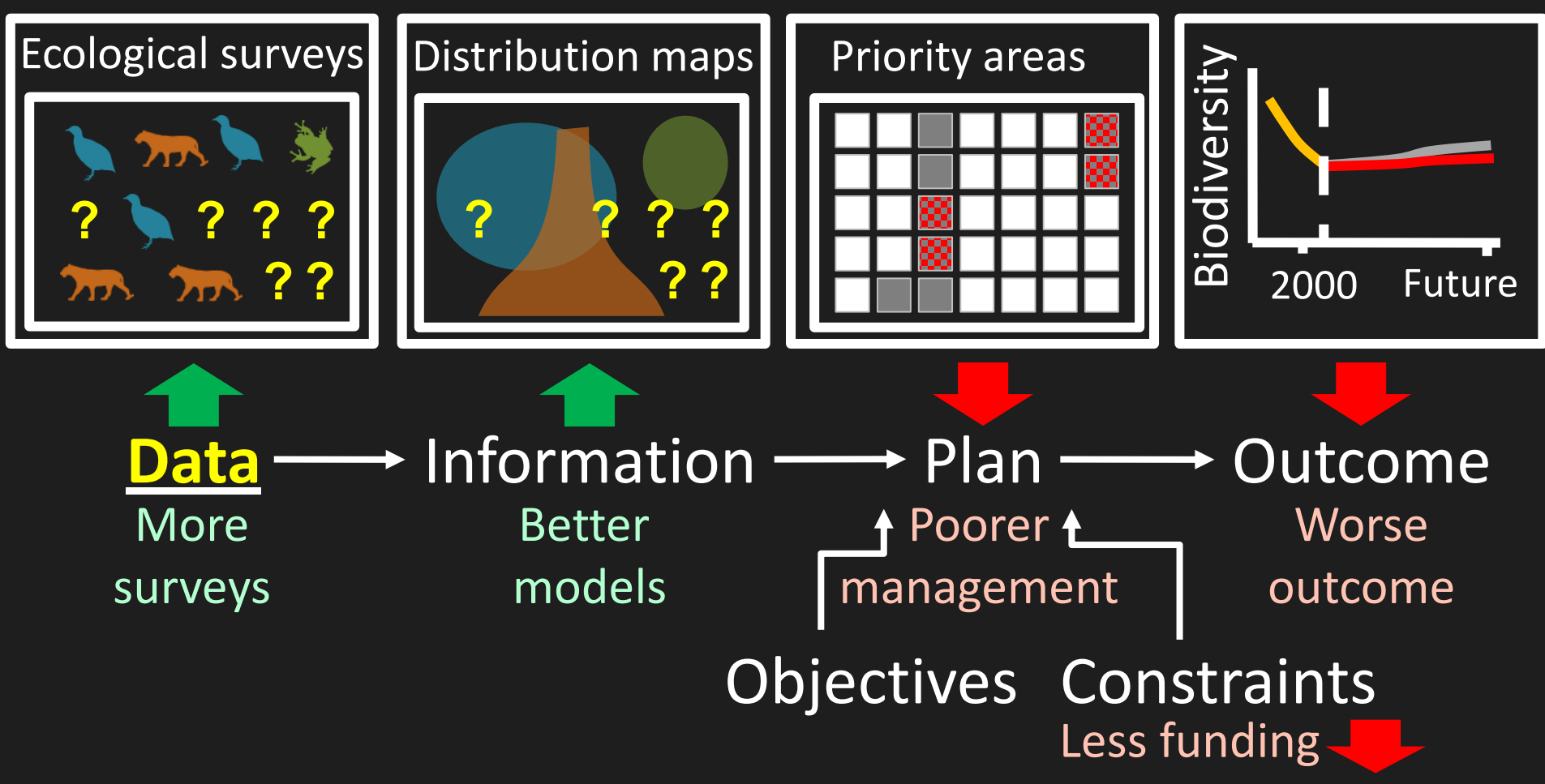
(1) Better algorithms

(2) Better data

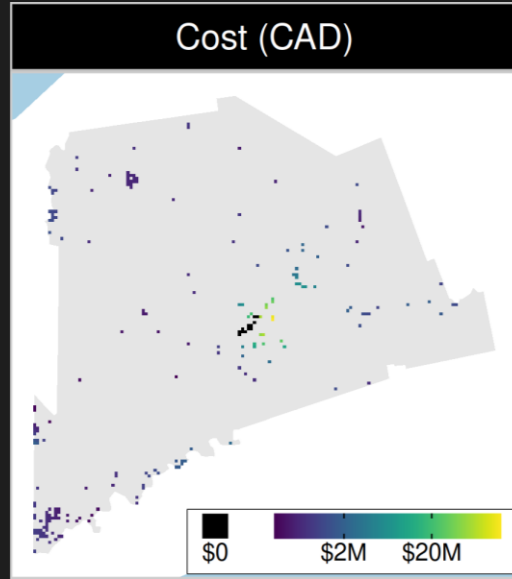
(3) Better surrogates



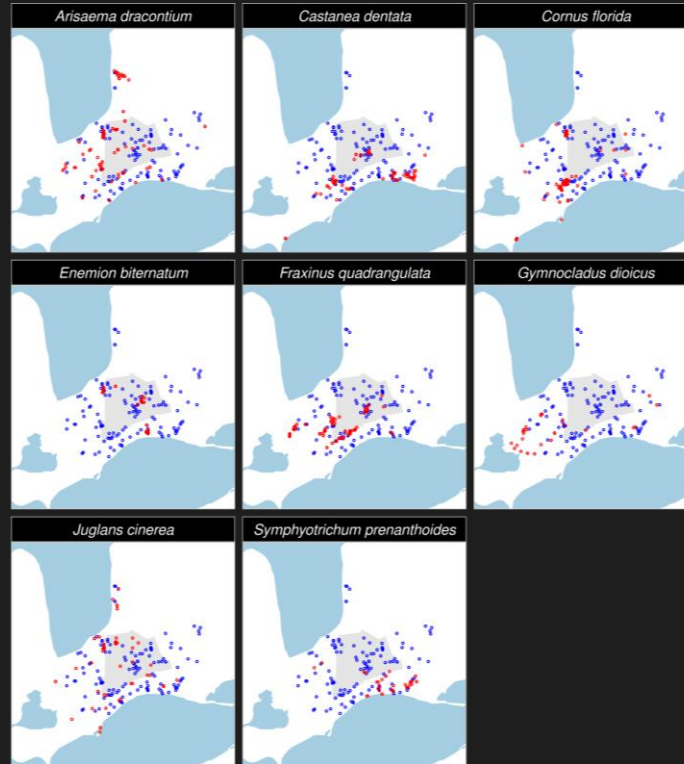




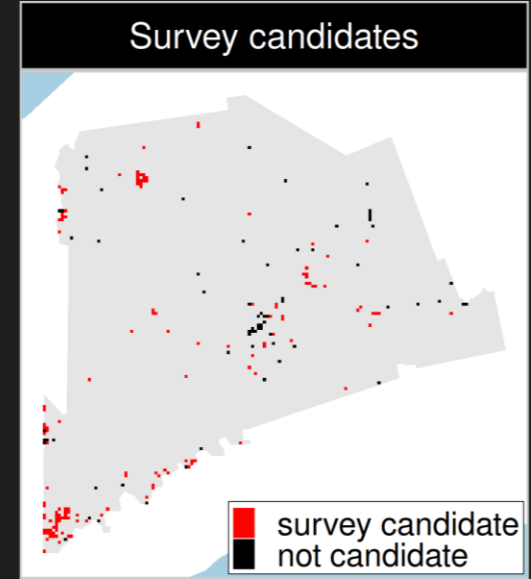
Study system: Middlesex county, Canada



199 places that could potentially be acquired for protected area establishment

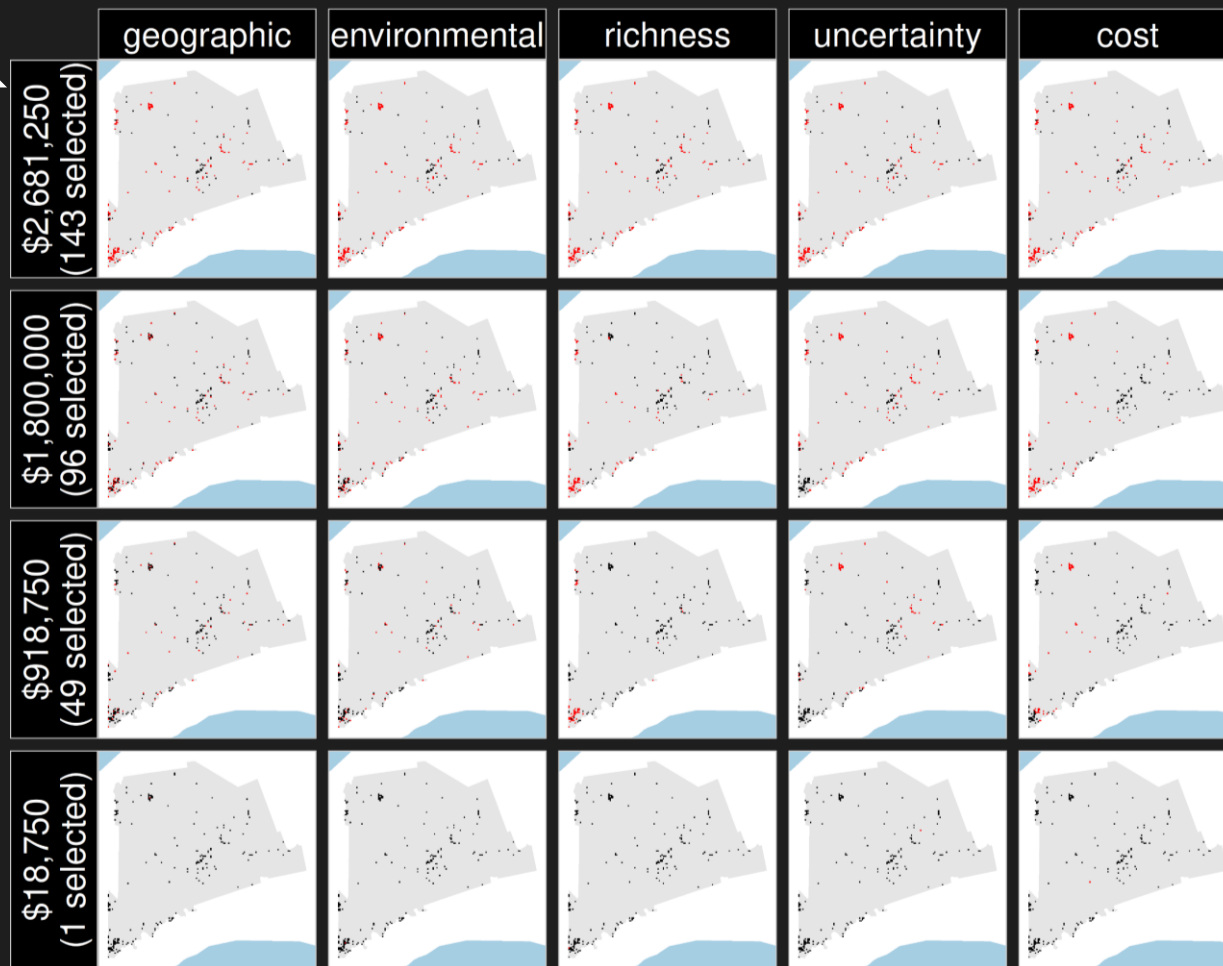


8 imperiled plant species



143 places that could potentially be surveyed to improve existing data

Number of places selected for surveys
(amount of funds allocated for collecting extra data)



Selected
for survey

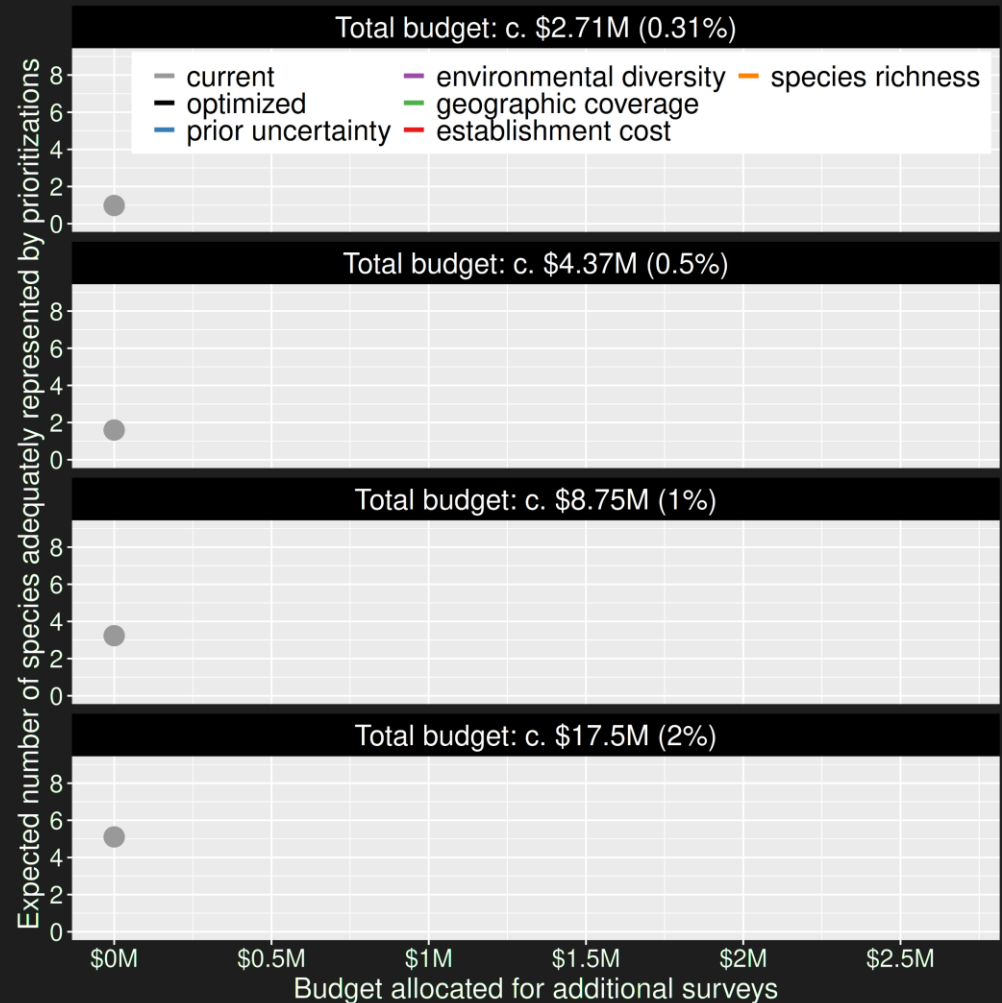


NOT selected
for survey

Different approaches for designing survey schemes

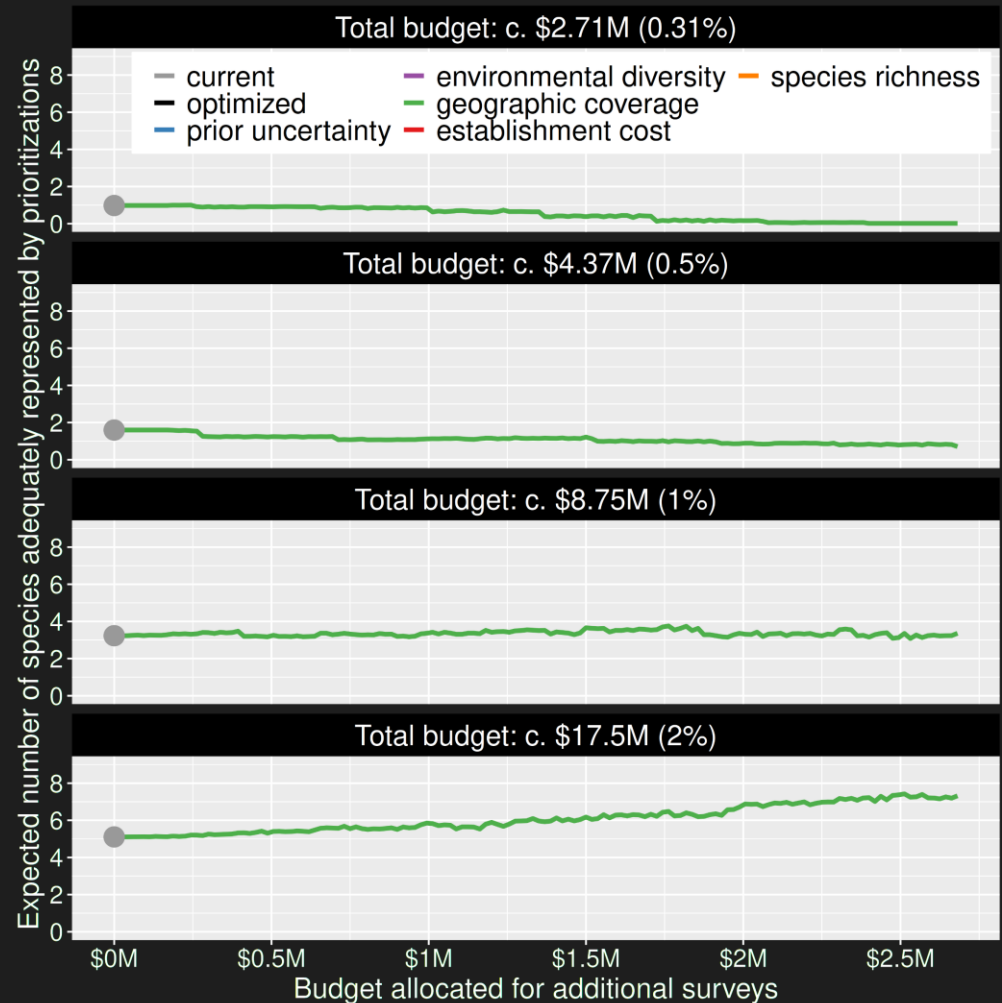
Value of information

- Existing data leads to positive outcomes
- More budget means better outcomes



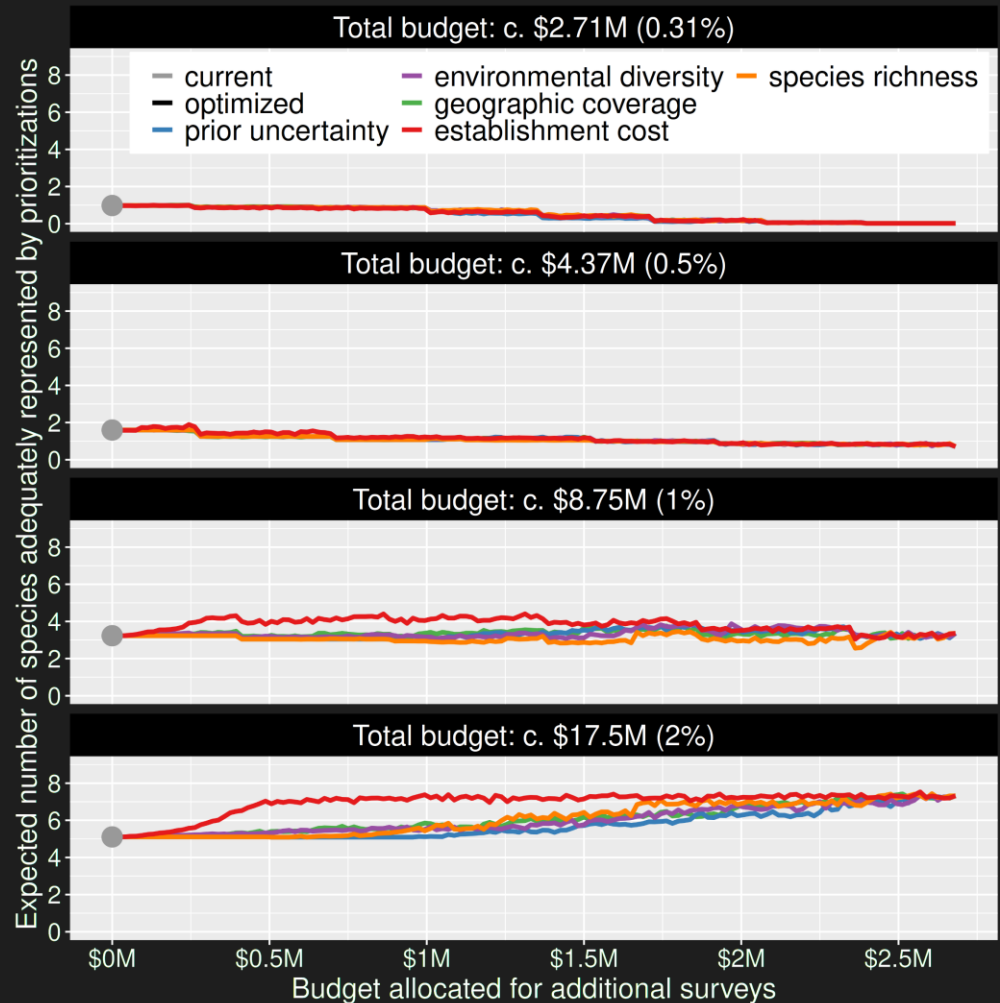
Value of information

- Allocating funds for gathering more data can mean worse outcomes
- Allocating funds for gathering more data can mean better outcomes too



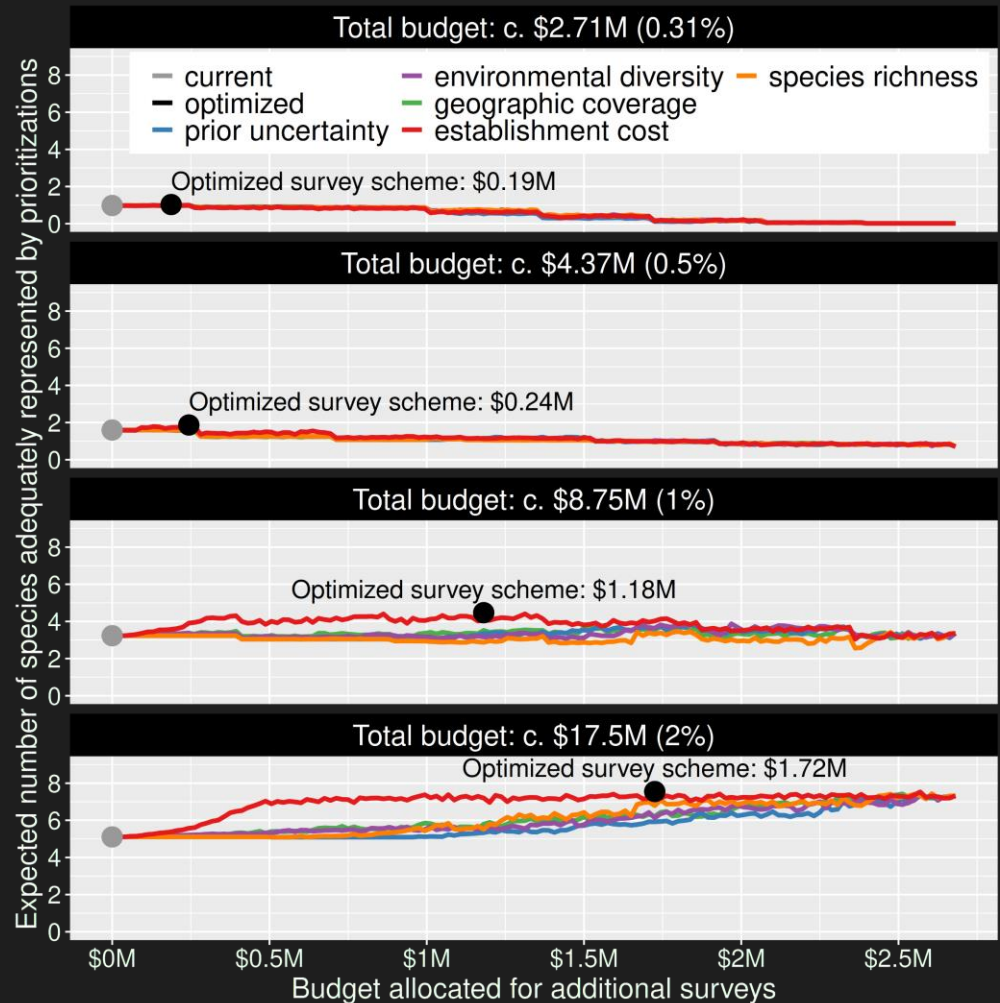
Value of information

- Conventional approaches for gathering additional evidence have different performance
- Performance of these approaches depends on available funds
- All of them could lead to worse outcomes



Value of information

- Directly maximizing return on investment is best method for additional data
- This considers objectives and constraints that underpin conservation plans and their success
- You can do this with: surveyvoi R package

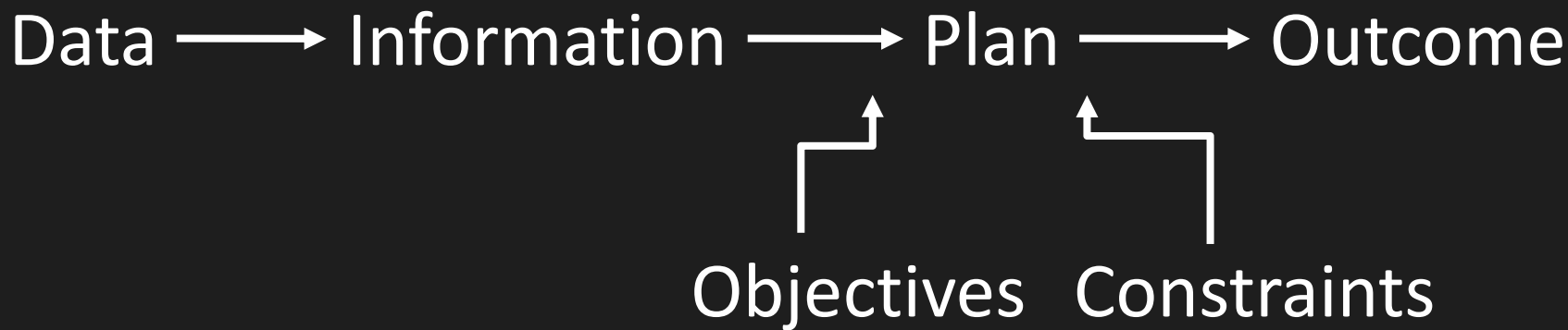
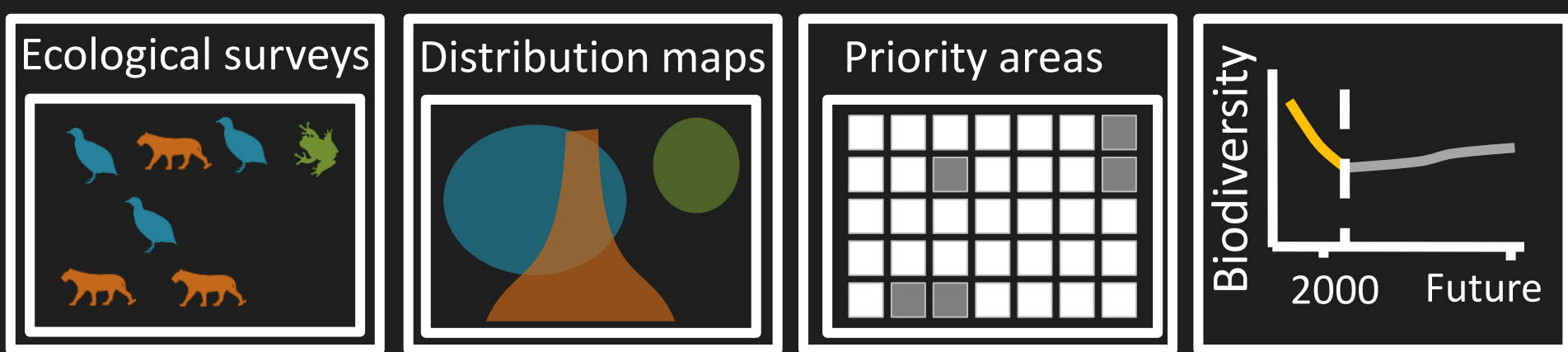


How can we get a better conservation decision?

(1) Better algorithms

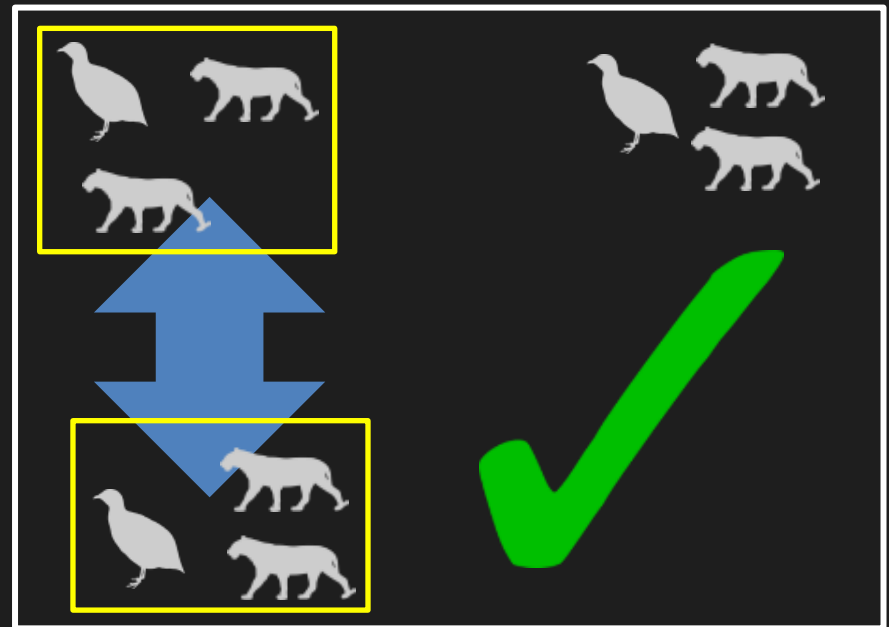
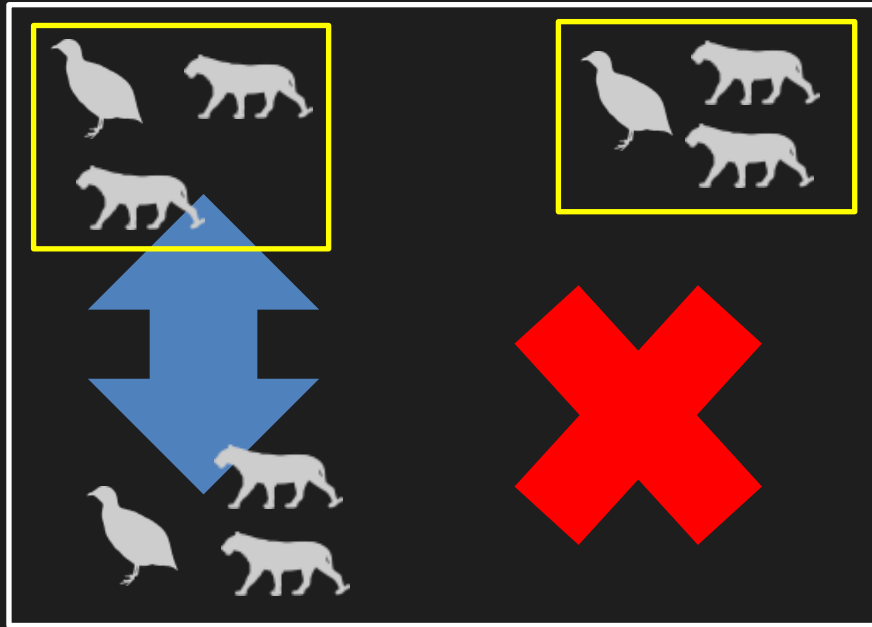
(2) Better data

(3) Better surrogates



Connectivity in reserve design

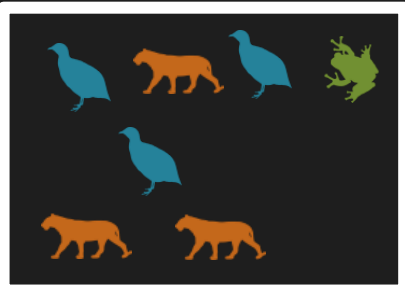
movement of individuals and genes between populations



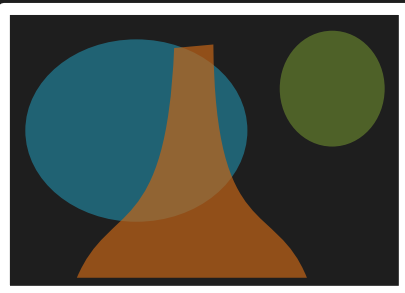
*all else being equal

Data → Information → Plan → Outcome

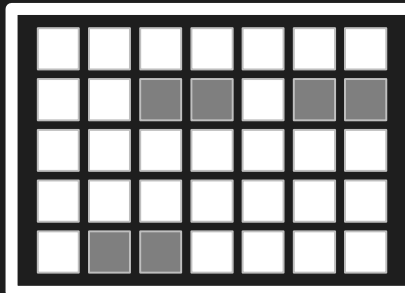
Ecological surveys



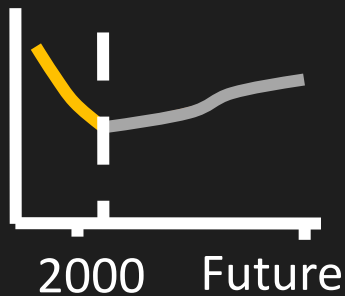
Distribution maps



Priority areas



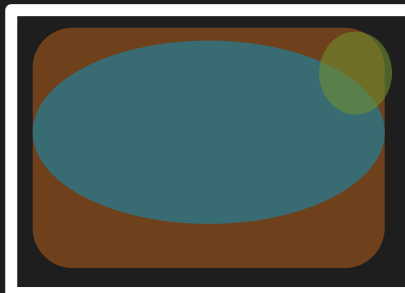
Biodiversity



Telemetry



Connectivity



But getting connectivity data is hard

Animal telemetry

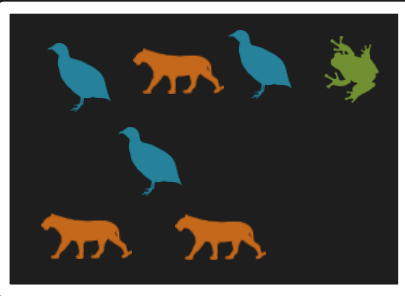


Molecular ecology

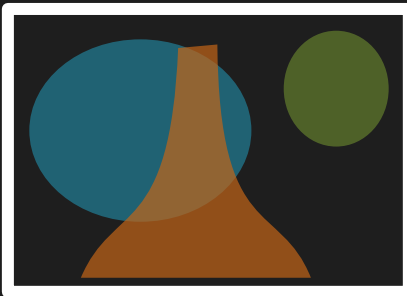


Data → Information → Plan → Outcome

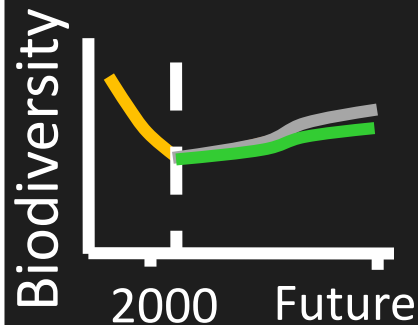
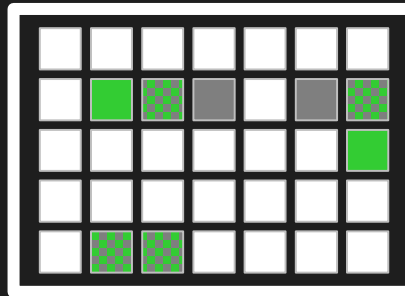
Ecological surveys



Distribution maps



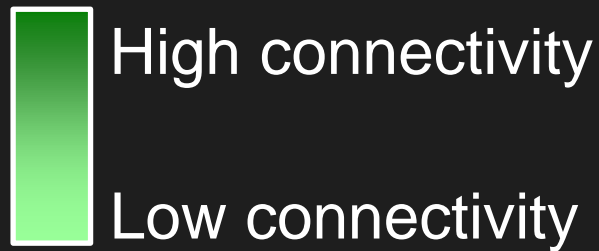
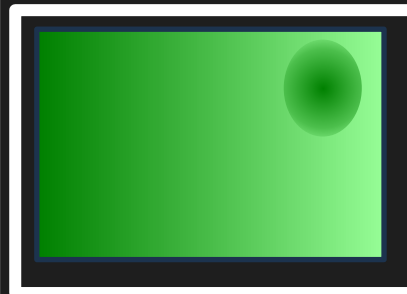
Priority areas



Surrogate



Surrogate



Many surrogates are
often available,
how do our choices
affect the results?

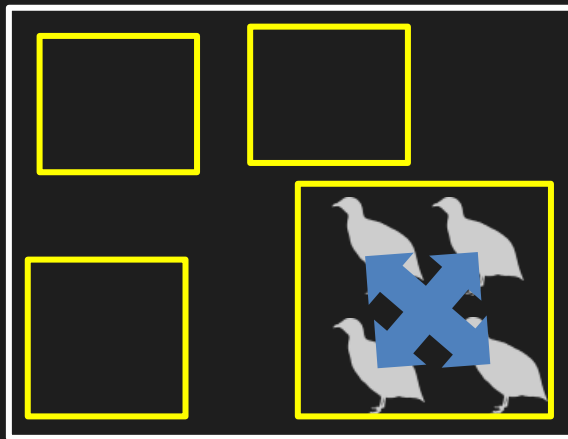
Rules of thumb for connectivity

More habitat = More connectivity

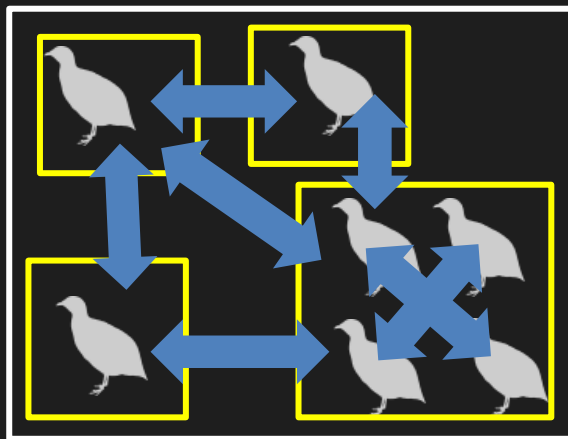
- Total area protected
- Species representation



Now



Soon

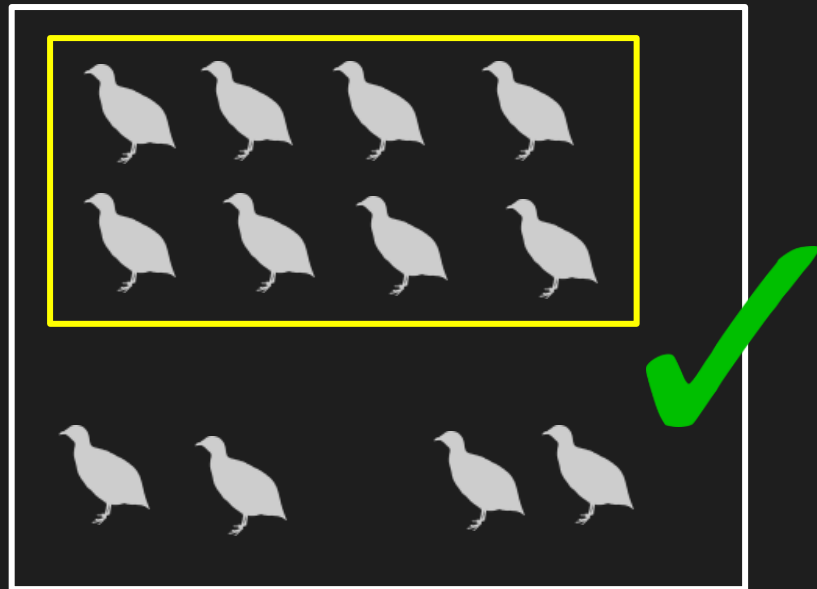
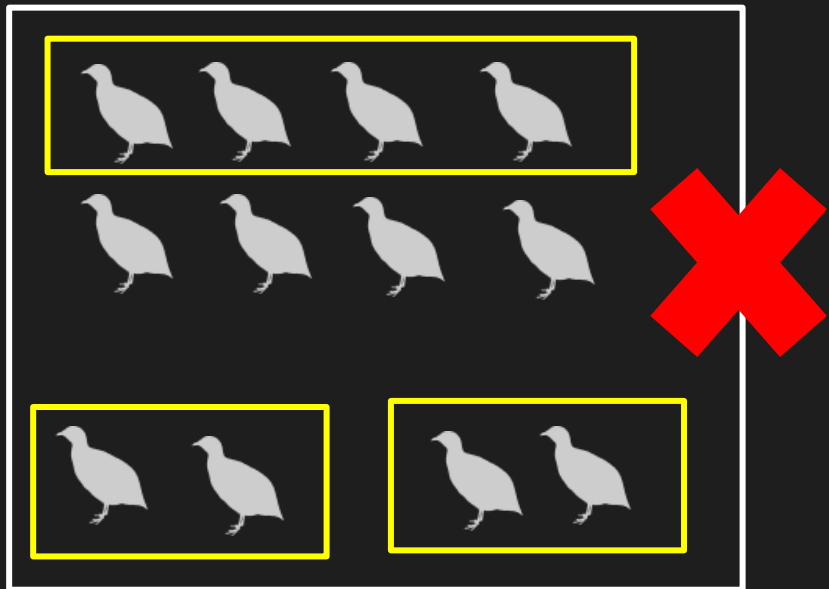


Later

Rules of thumb for connectivity

More spatial clustering = More connectivity

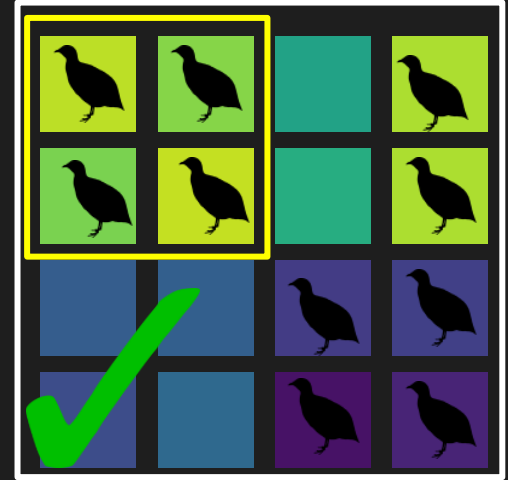
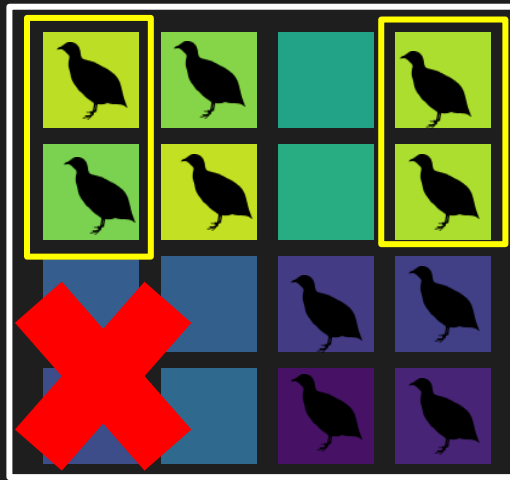
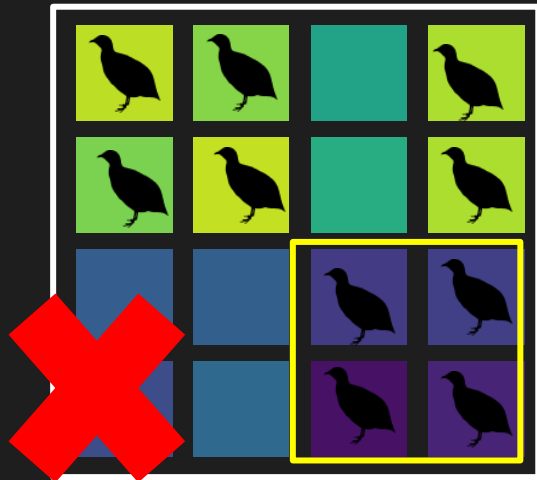
- Boundary length



Rules of thumb for connectivity

Protect clusters of low resistance = More connectivity

- Human pressure
- Naturalness based landscape resistance
- Focal species landscape resistance
- Habitat heterogeneity
- Environmental similarity



High resistance



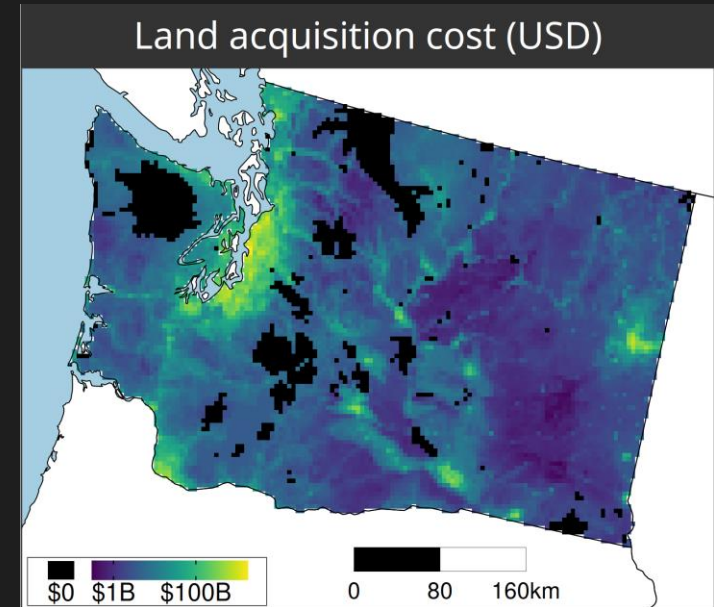
Low resistance

A comparison of approaches for including connectivity in systematic conservation planning

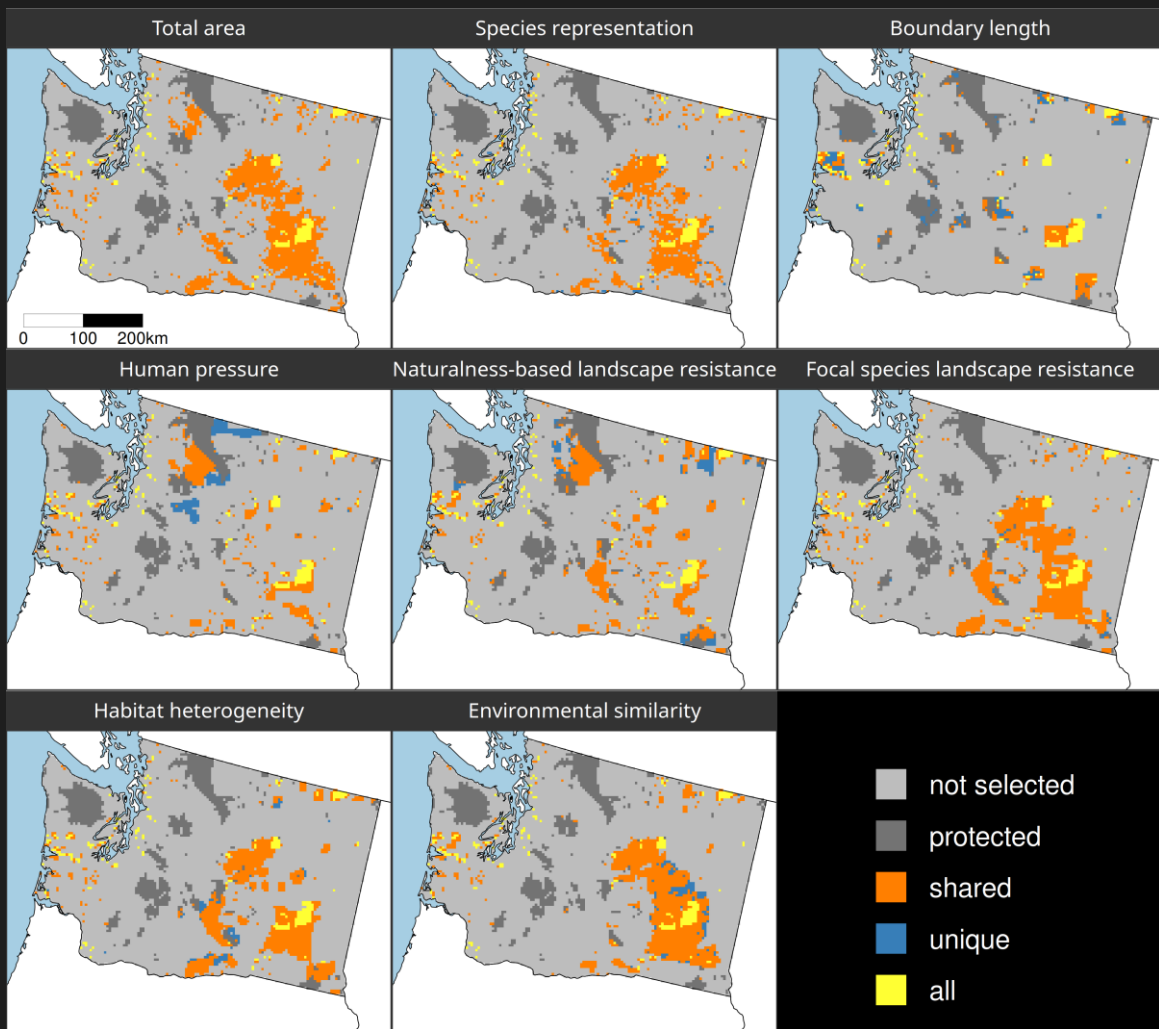
Washington State, USA

- 261 bird species
- Land acquisition costs
- Existing protected areas
- Multiple land-uses
- Multiple eco-systems

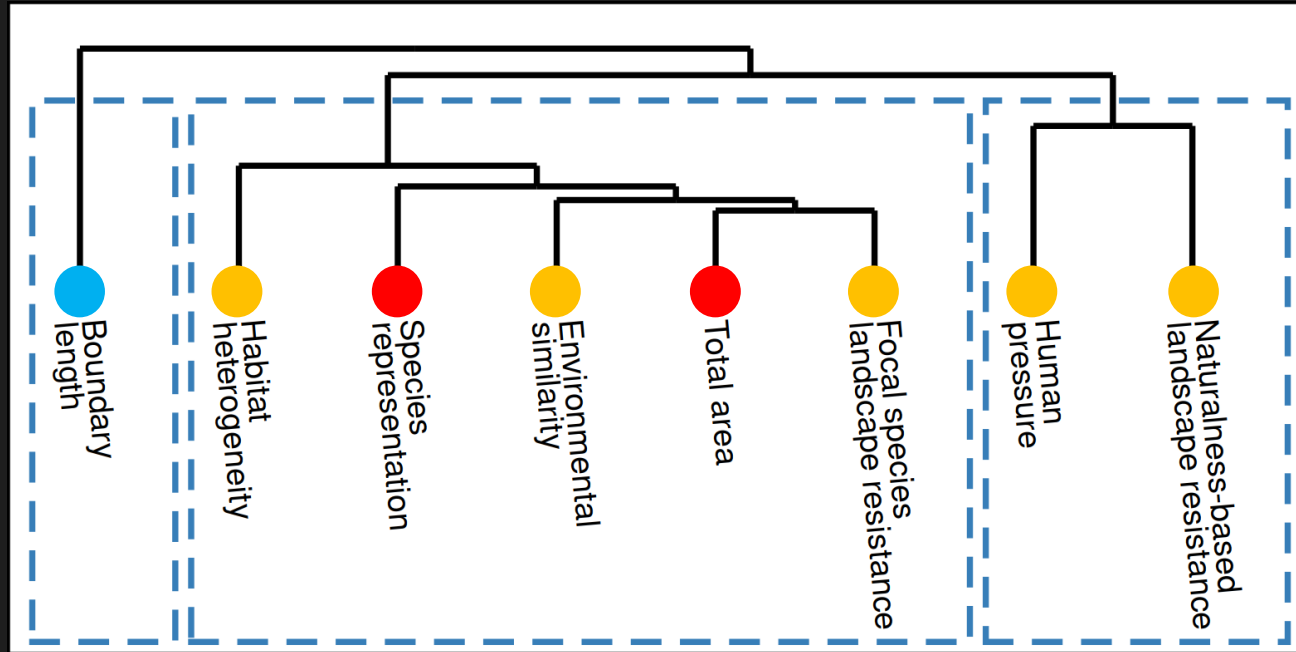
Hanson *et al.* 2022 J. Appl. Ecol



- Different connectivity approaches produce different prioritizations
- Different connectivity approaches can yield similar prioritizations

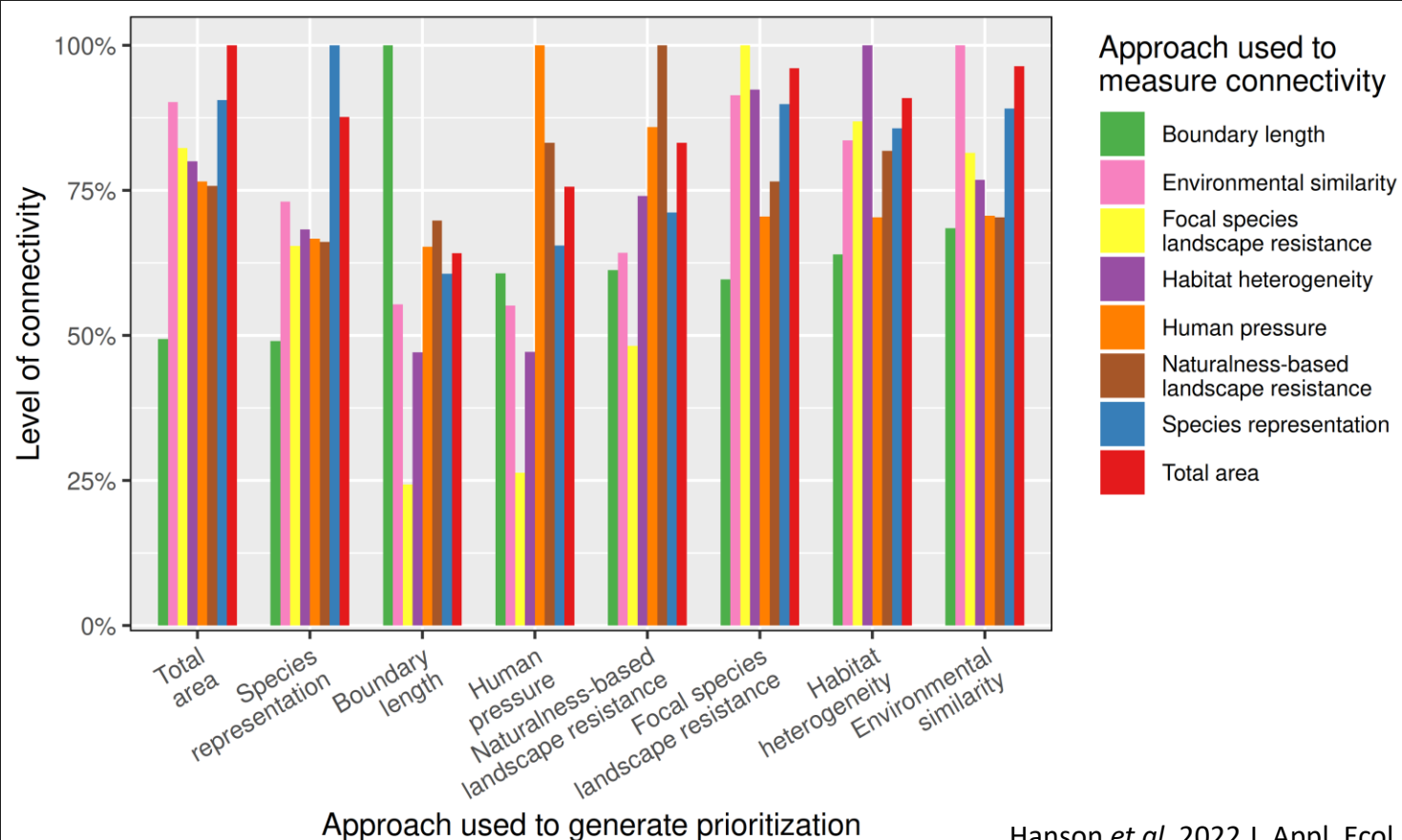


How do the prioritizations compare?



- Approach
- Habitat-based
 - Spatial clustering
 - Resistance-based

No obvious winner



Make better conservation decisions by using...

1. Better algorithms
2. Cost-effective data
3. Reliable surrogates



jeffrey.hanson@uqconnect.edu.au



github.com/jeffreyhanson



jeffrey-hanson.com



Worked example

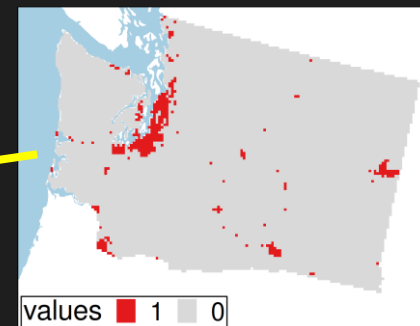
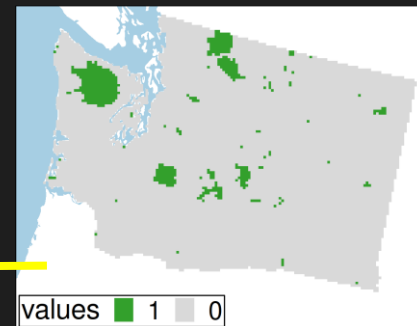
```
# load packages
library(prioritizr) # package for conservation planning
library(terra)      # package for raster data

# input data
## raster with continuous values indicating costs
planning_unit_data <- rast("pu.tif")
```

```
## multi-layer raster with relative abundance data
### feature_data[[1]] is the first feature,
### feature_data[[2]] is the second feature,
### and so on, with 396 features in total
feature_data <- rast("features.tif")
```

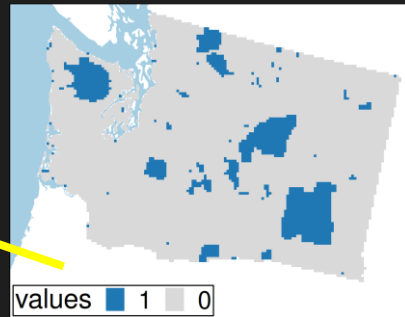
```
## raster with binary values indicating if each planning
## unit is covered by (1) protected areas or (0) not
protected_area_data <- rast("protected-areas.tif")
```

```
## raster with binary values indicating if each planning
## unit covered by (1) urban areas or (0) not
urban_area_data <- rast("urban-areas.tif")
```



```
# build problem
## specify the data, formulation, and solver
conservation_planning_problem <-
  ### initialize with planning unit and feature data
  problem(planning_unit_data, feature_data) %>%
  ### add minimum shortfall function with budget
  add_min_shortfall_objective(3917.631) %>%
  ### add representation targets for 20% coverage of each feature
  add_relative_targets(targets = 0.2) %>%
  ### add penalties to reduce spatial fragmentation
  add_boundary_penalties(penalty = 0.00001) %>%
  ### add constraints to ensure existing protected areas are selected
  add_locked_in_constraints(protected_area_data) %>%
  ### add constraints to ensure urban areas are not selected
  add_locked_out_constraints(urban_area_data) %>%
  ### specify that decision variables are binary (0 or 1 values)
  add_binary_decisions() %>%
  ### specify software to perform optimization,
  ### and set gap parameter for near-optimal solution
  add_gurobi_solver(gap = 0.1)
```

```
# solve problem
## output raster has binary values indicating if
## each planning unit is (1) selected or (0) not
prioritization <- solve(conservation_planning_problem)
```



```
# evaluate prioritization
## calculate overall cost of prioritization
eval_cost_summary(
  conservation_planning_problem, prioritization)
#> # A tibble: 1 × 2
#>   summary      cost
#>   <chr>      <dbl>
#> 1 overall 3911.832
```

```
## calculate relative importance of selected planning units
### output raster has continuous importance values
relative_importance <-
  eval_ferrier_importance(
    conservation_planning_problem, prioritization)
```

