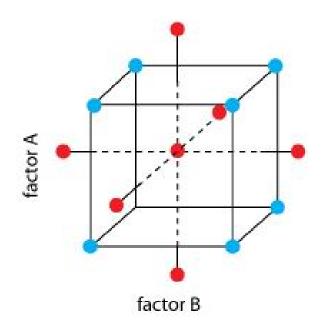
Supporting the Design of Experiments for Synthetic Biology with Clotho 3.0

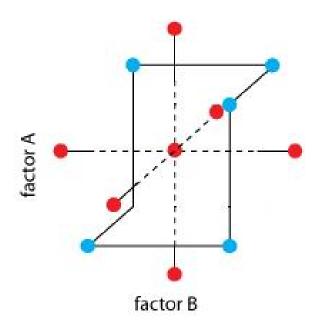


Nicholas Roehner
CIDAR (Densmore Lab)
Boston University

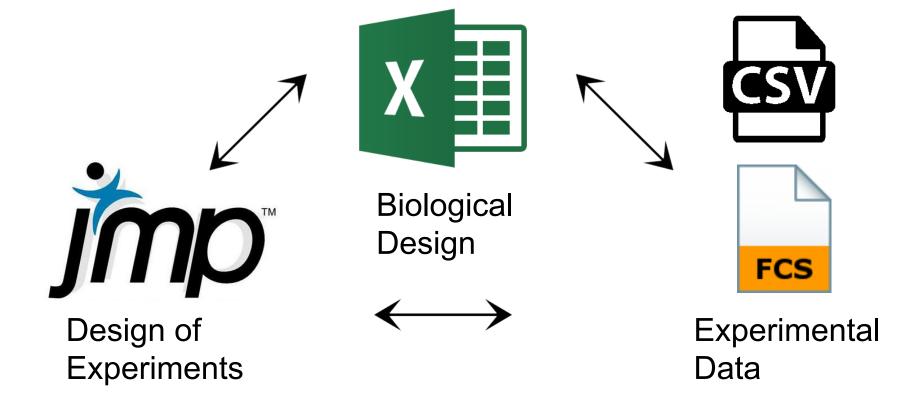
Design of Experiments (DOE)



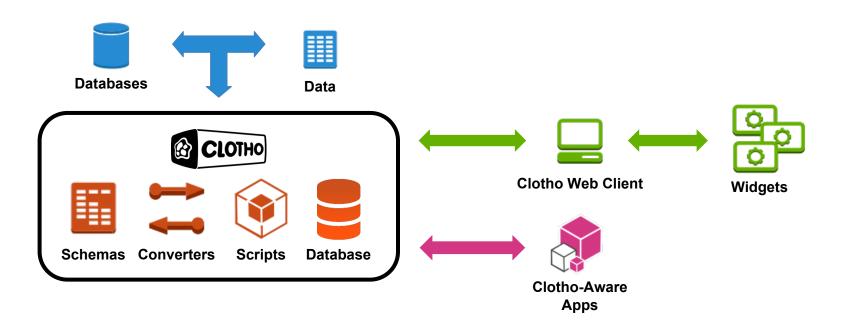
Design of Experiments (DOE)



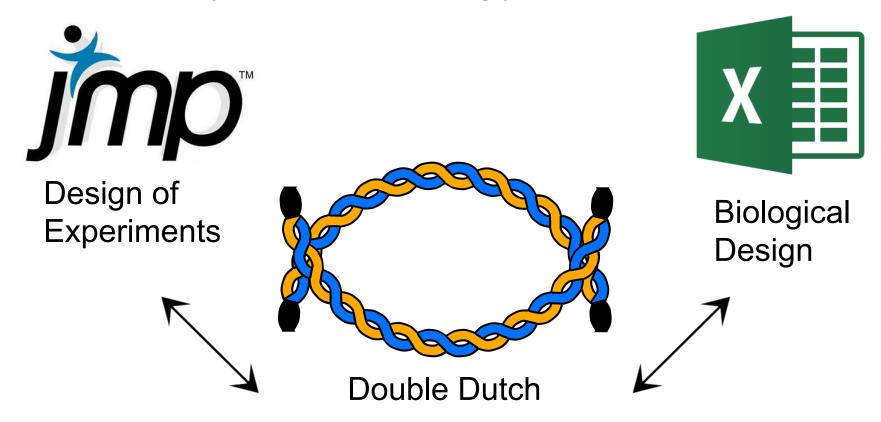
DOE for Synthetic Biology



Clotho 3.0



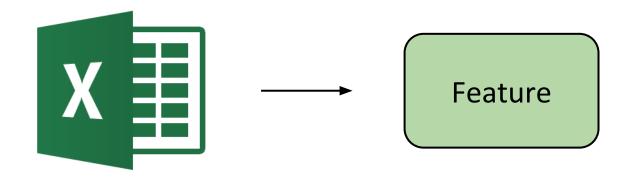
DOE for Synthetic Biology w/ Double Dutch



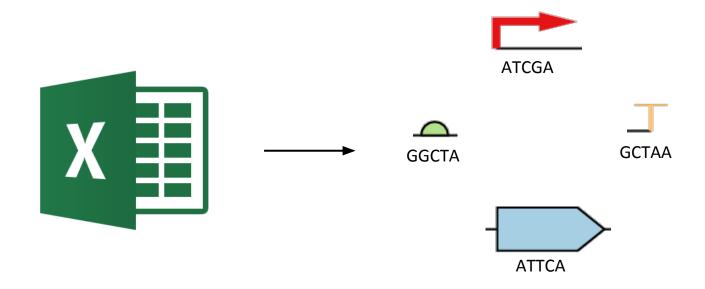
Talk Outline

- Overview of Double Dutch web app
- Double Dutch demo
- Discussion of app extensions

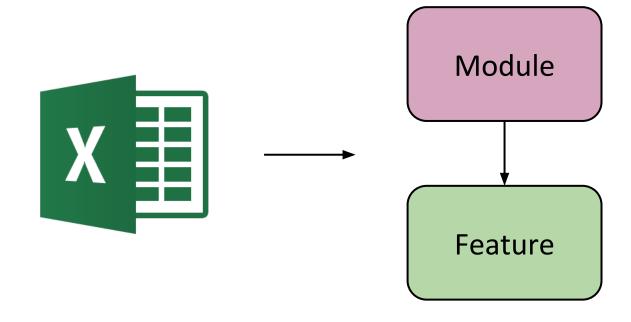
Step 1: Import Features



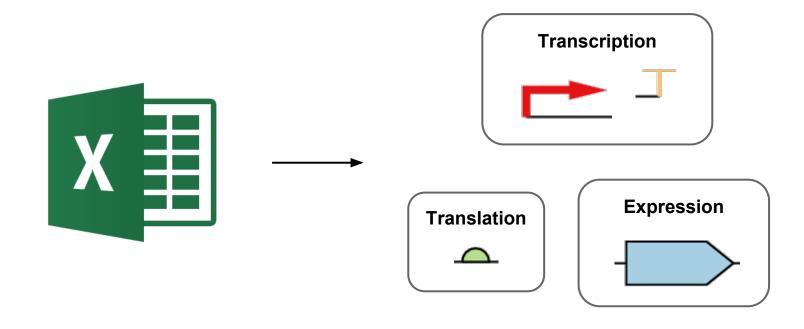
Step 1: Import Features



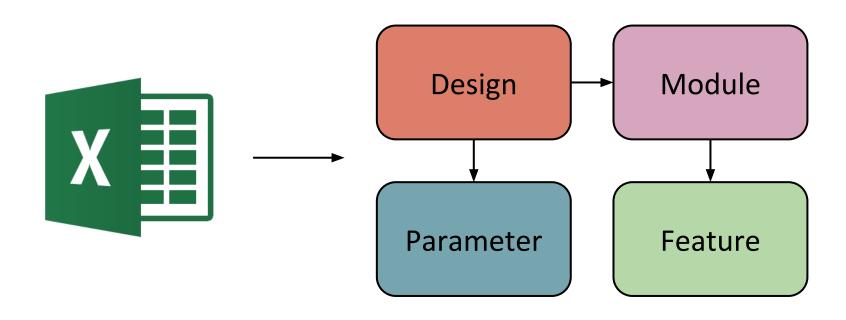
Step 2: Infer Modules



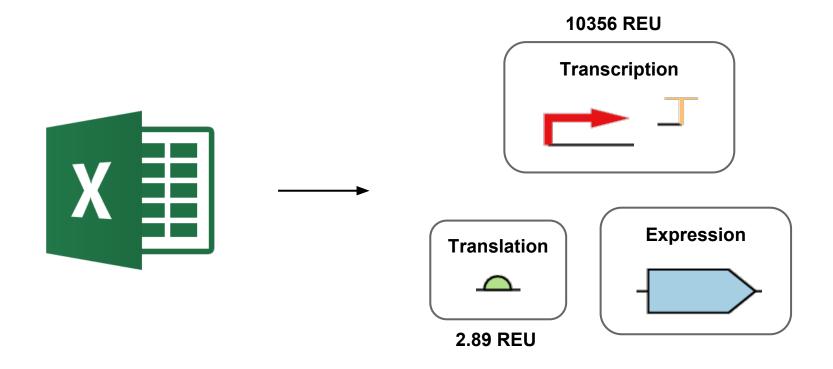
Step 2: Infer Modules



Step 3: Parameterize Modules

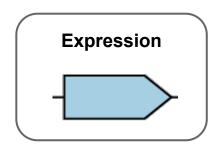


Step 3: Parameterize Module

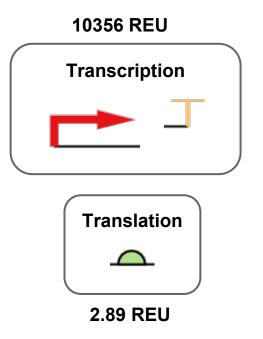


Step 4: Categorize as Factors or Levels

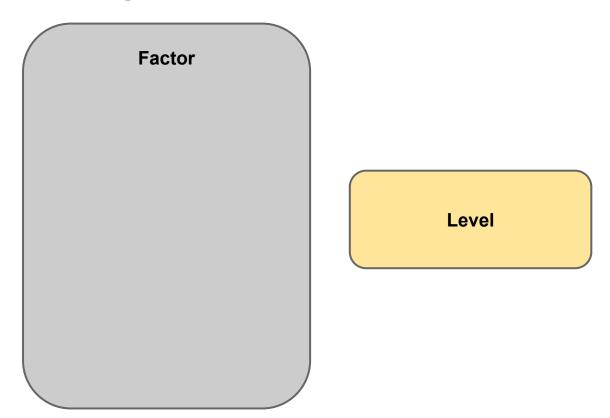
Available Factors



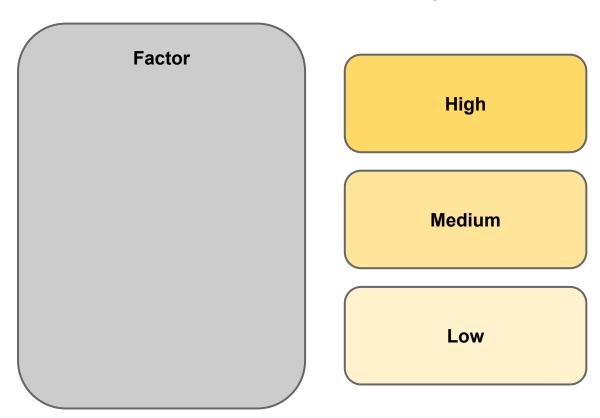
Available Levels



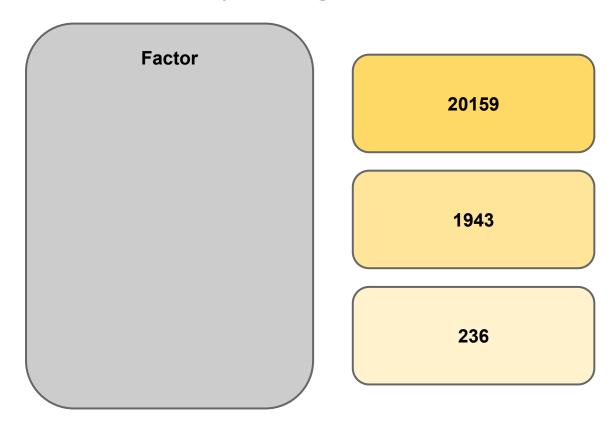
Step 5: Assign Levels to Factors



Step 5a: Choose # of Levels per Factor

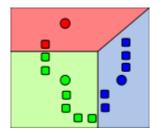


Step 5b: Quantify Target Levels

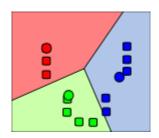


K-Means Clustering

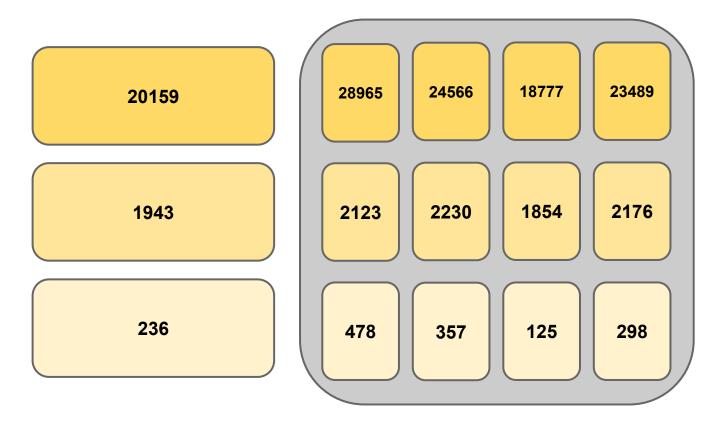




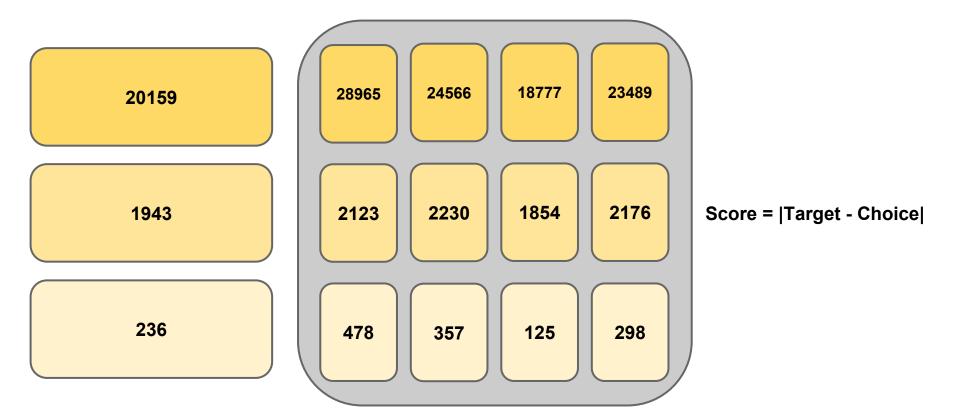




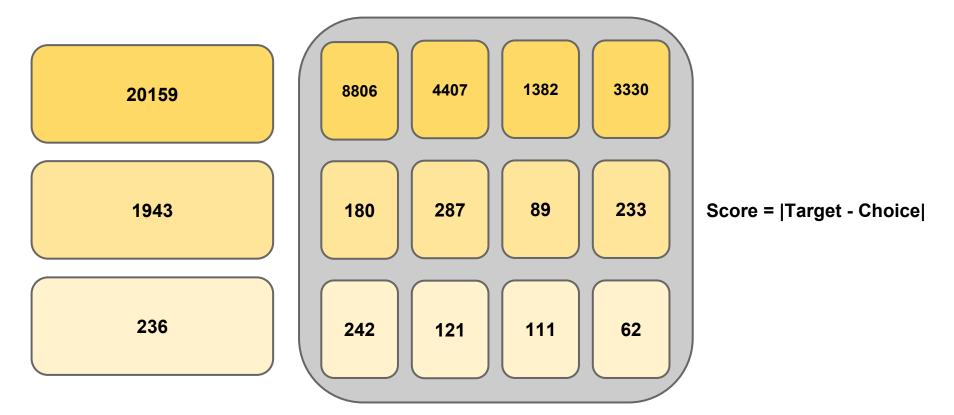
Step 5c: Partition Level Choices



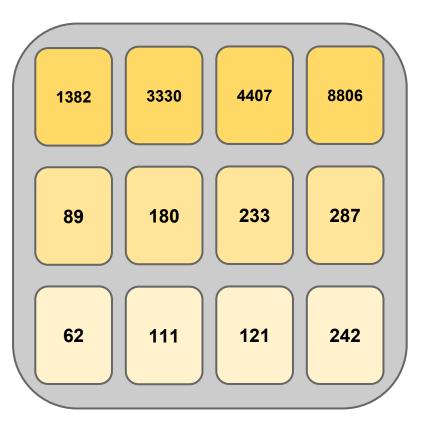
Step 5d: Score Level Choices

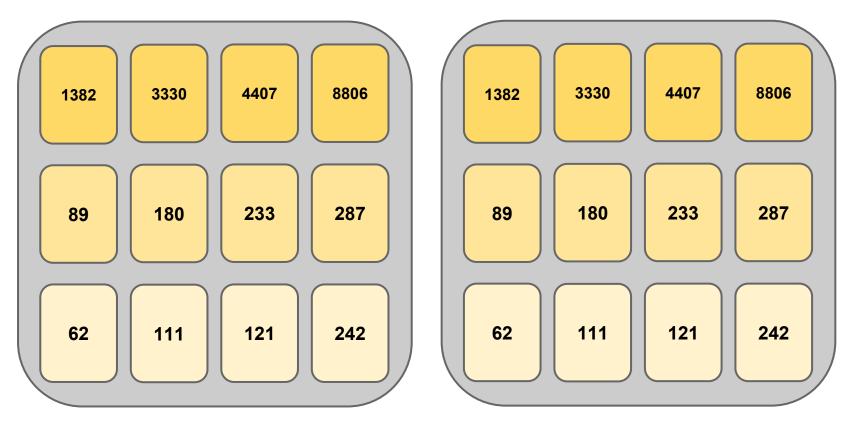


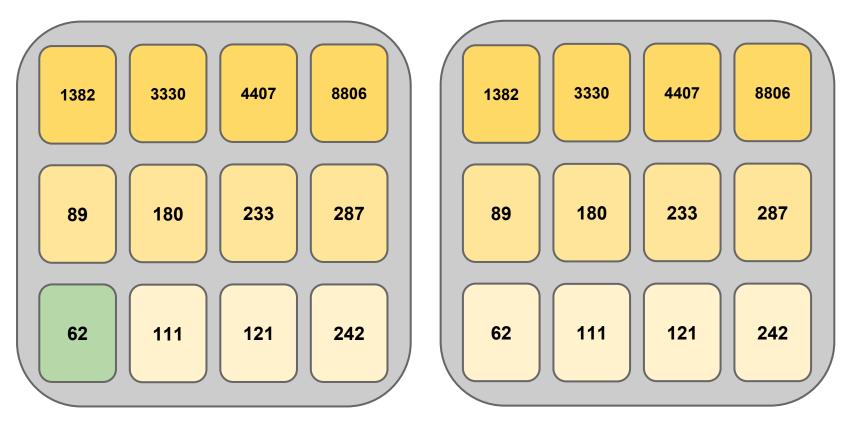
Step 5d: Score Level Choices

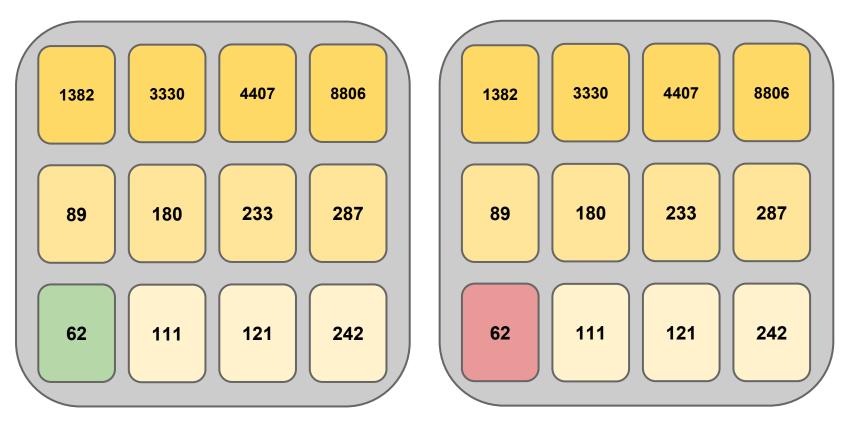


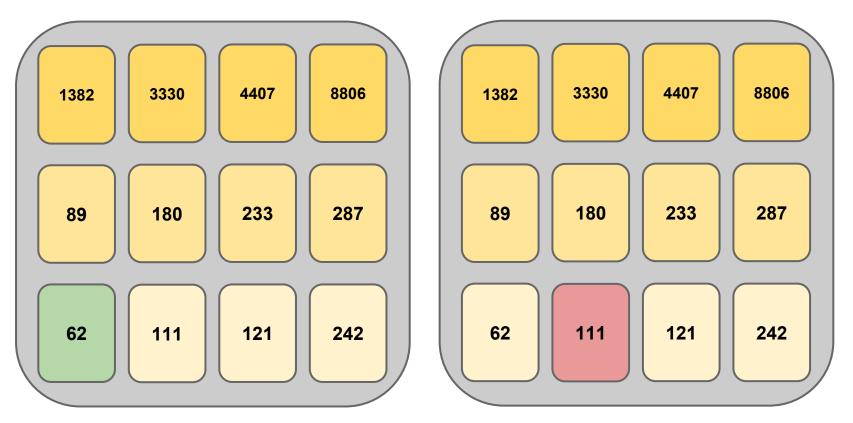
Step 5e: Sort Level Choices

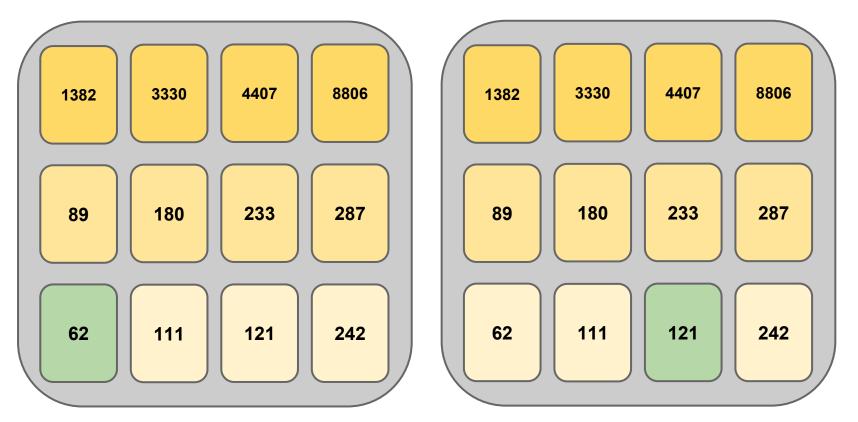


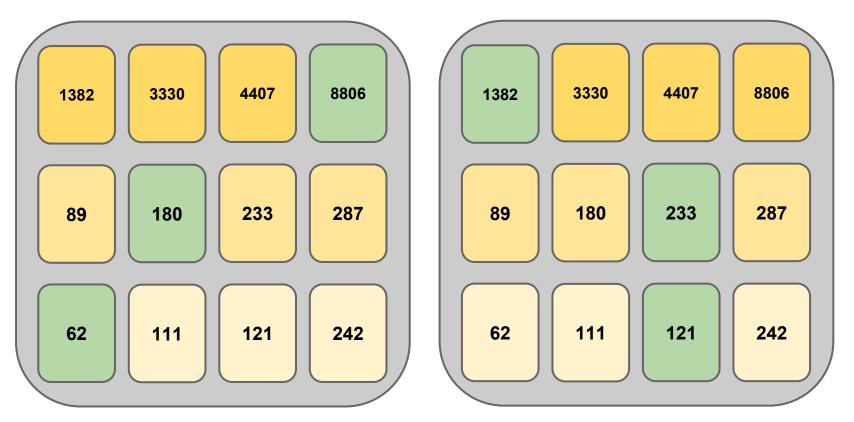




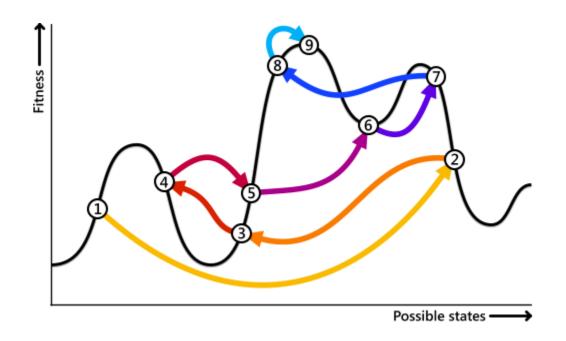


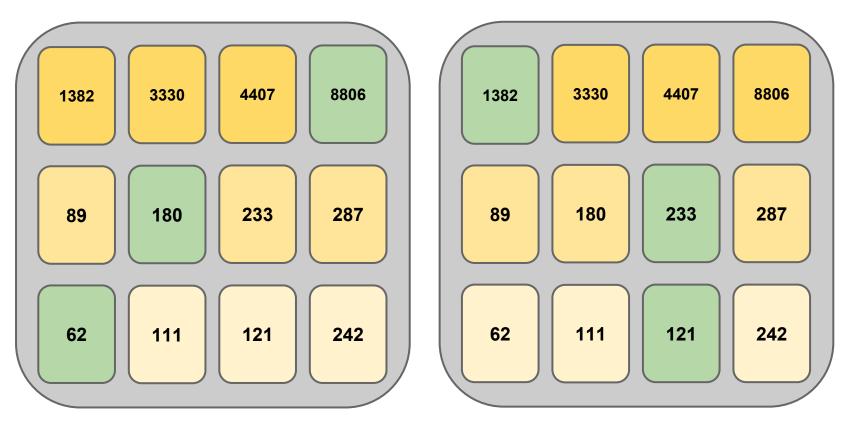


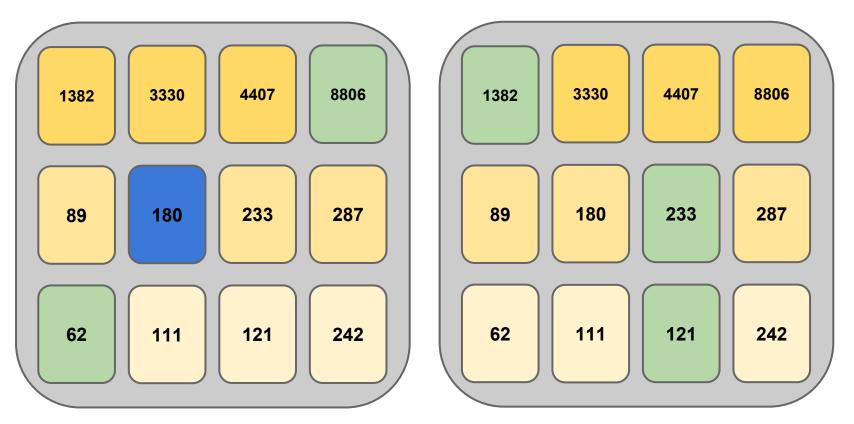


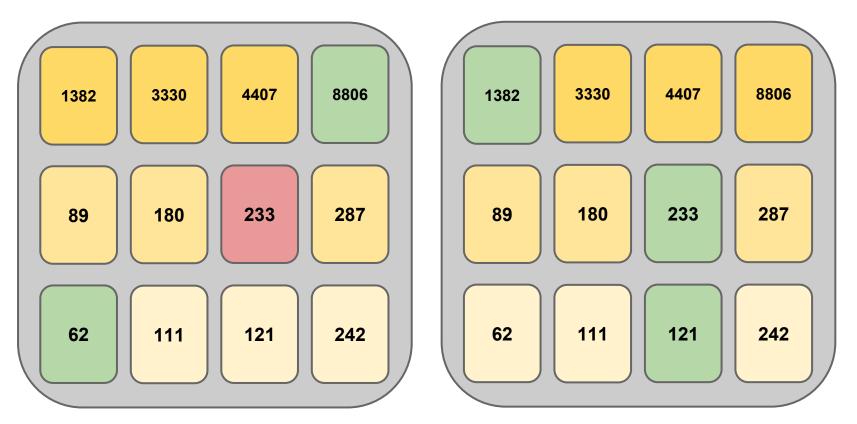


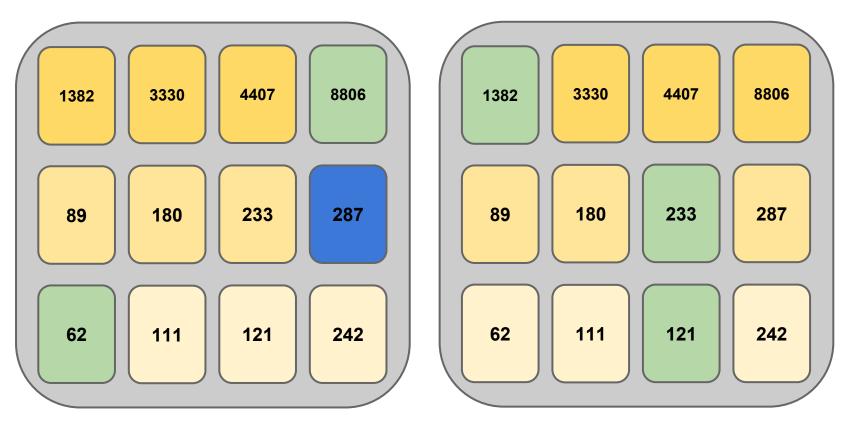
Local Optimum and Simulated Annealing

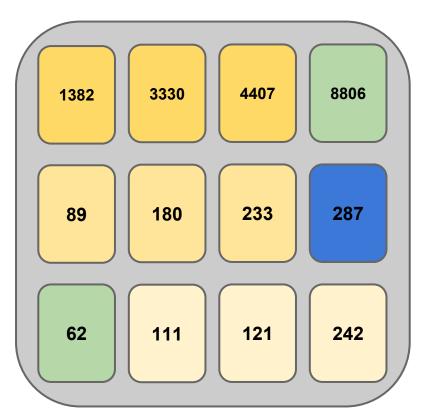




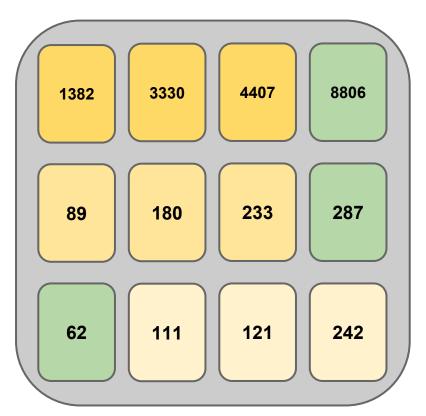






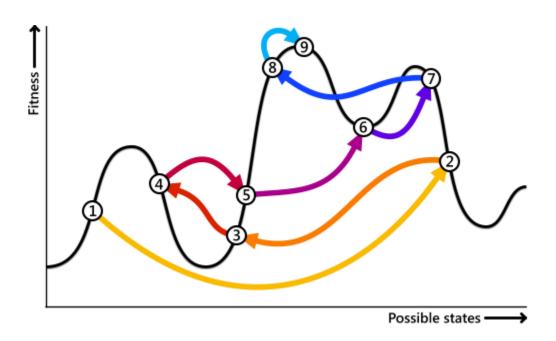


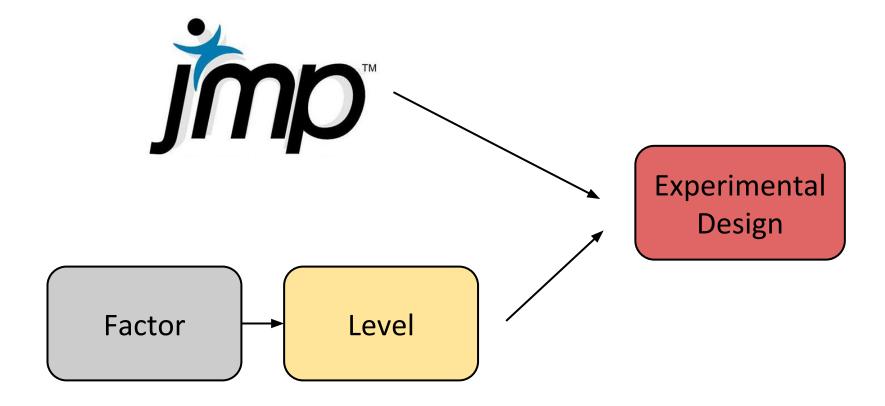
P(accept) = e^[(oldScore - newScore)/Temp]



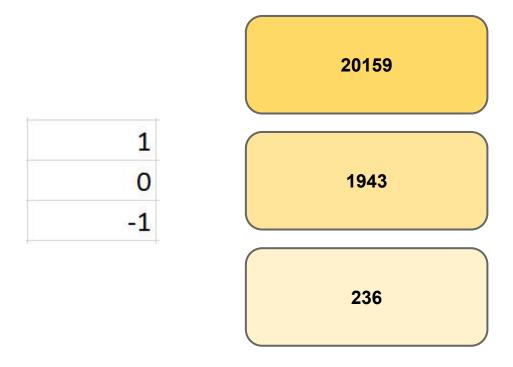
P(accept) = e^[(oldScore - newScore)/Temp]

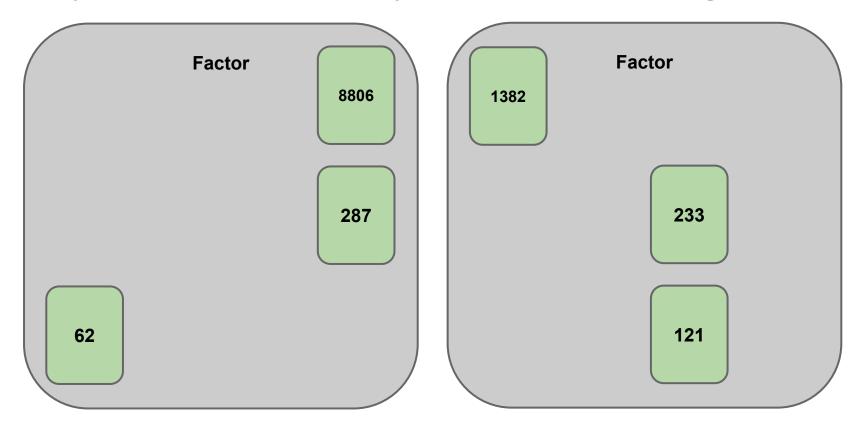
Simulated Annealing

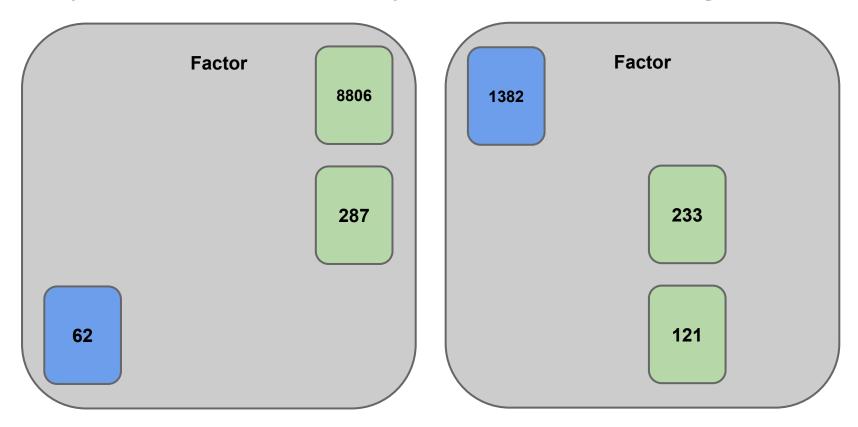




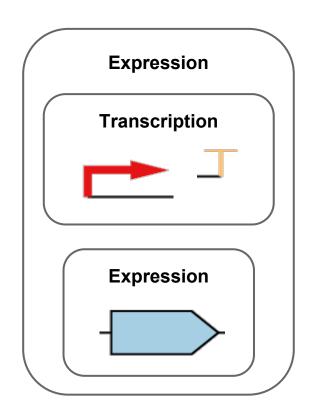
4th Order	A	В	C	D	E	F
1	0	1	1	1	-1	-1
2	0	1	0	0	0	0
3	1	1	1	1	-1	1
4	0	0	0	-1	1	0
5	-1	1	0	0	0	0
6	-1	-1	-1	1	1	0
7	0	-1	0	-1	0	0
8	1	1	-1	-1	1	-1
9	0	0	-1	0	1	-1
10	1	0	0	0	0	-1

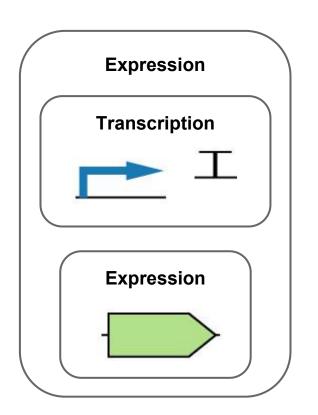






Step 7: Generate Biological Designs





Planned Extensions

- Validate composition of modules associated with factors and levels in experimental designs.
- Fit empirical models to experimental data.
- Incorporate Pigeon for design visualization.
- Export designs in additional file formats (Euegene, SBOL, JSON).
- Generate picklists for automated assembly.

Clotho Team



Douglas Densmore



BOSTON

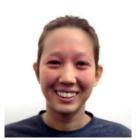
UNIVERSITY







J. Christopher Anderson



Stephanie **Paige**



Prashant Vaidyanatha



Bill Cao



Maxwell **Bates**





