# Space Relationship Modeler (SPAREMO)

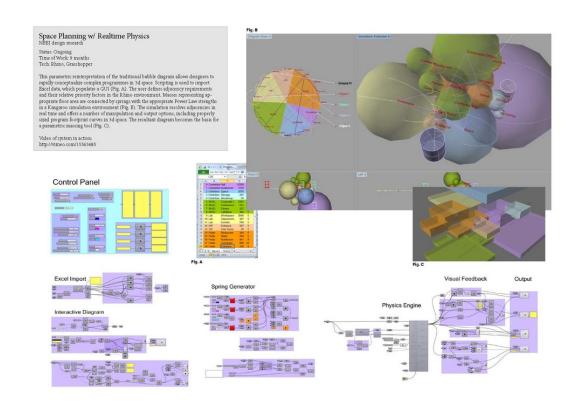
Marco Juliani
Morphogenetic Programming
ucbqjul@ucl.ac.uk

### Overall Objectives

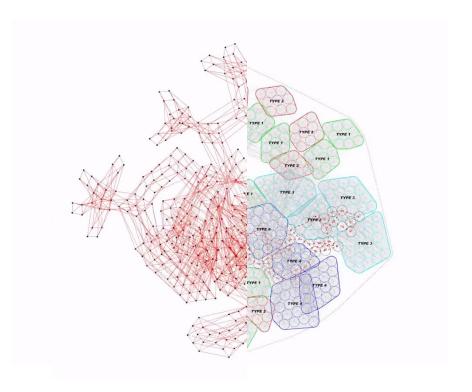
- 'Emulate physics' category
- Generative framework for deriving early space-planning solutions.
- A playful framework to visualize and rethink programmatic and spatial relationships.
- 'Relationship maps' produced by using relationship matrices combined with spring systems.
- Multiple ways of manipulating 'hyper-parameters' as representation options to guide the simulation to arrive at helpful solutions.

### Review

#### Space Planning w/ Real Time Physics



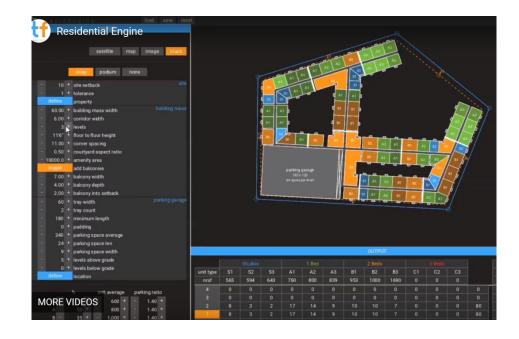
NBBJ - Physics-based space planning http://www.marcsyp.com/space-planning



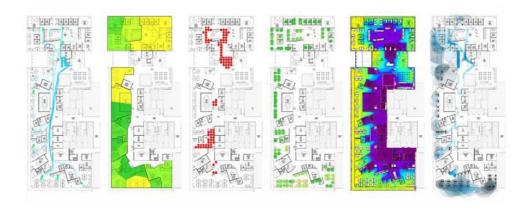
Yazdani Studio – Parametric Plan Tool https://vimeo.com/146462913

### Review

#### Plan Optimization



TestFitIO- Parametric Residential Planner https://testfit.io/



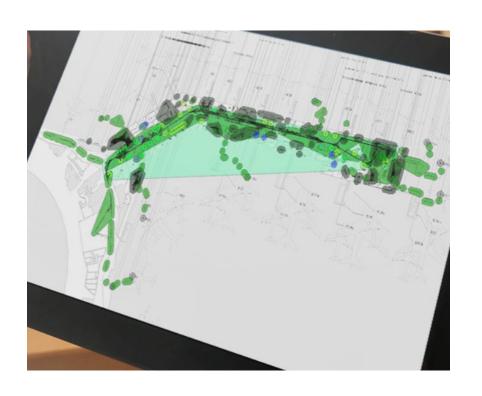
The Living – Generative Space Planning https://www.autodeskresearch.com/sites/default/files/Project -Discover-Generative-Design.pdf

### Review

#### Msc Algorithms



Stanislas Chaillou - GAN Generated Plans https://towardsdatascience.com/ai-architecturef9d78c6958e0



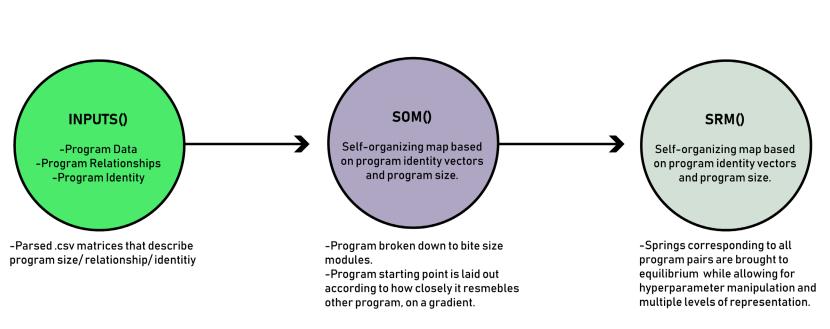
Woods Bagot- Program Relationship Tool

### Workflow Overall

Method 1: Global\_Program



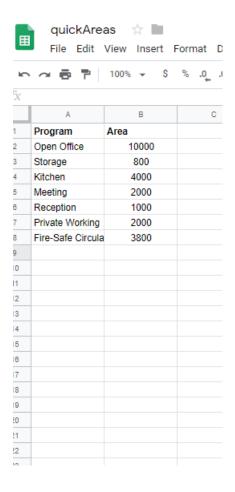
*Method 2: Kohonen\_Program* 



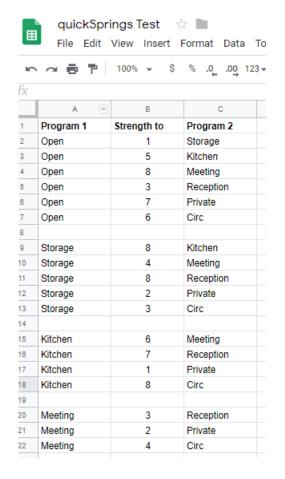
Marco Juliani

## Workflow Inputs

#### Program/ Areas



#### Program Relationships



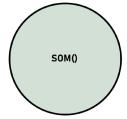
#### Program Identity Vectors

_	Kohonen Net Sample 🔯 🗎		
Ħ	File Edit	View Insert Form	at Data Tools Add-o
k	~ 5 7	100% 🕶 💲 %	.0 .00 123 ▼ Arial
ſχ			
	А	В	С
1	Open Office	null	0
2	Natural Light	0.4	1
3	Noise Allowed	0.2	2
4	Amenity Access	0.85	3
5	Openness	0.8	4
6	Pure function	0.7	5
7	Storage	null	6
8	Natural Light	0	7
9	Noise Allowed	1	8
10	Amenity Access	0.4	9
11	Openness	0	10
12	Pure function	1	11
13	Kitchen	null	12
14	Natural Light	0.7	13
15	Noise Allowed	1	14
16	Amenity Access	0.9	15
17	Openness	0.9	16
18	Pure function	0.5	17
19	Meeting	null	18
20	Natural Light	0.15	19
21	Noise Allowed	0.6	20
22	Amenity Access	0.14	21
23	Openness	0.12	22

## Pseudocode<sub>1</sub>

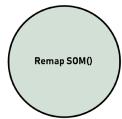


#### Data Processing()

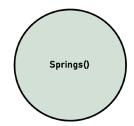


Feed Program Names and Areas();

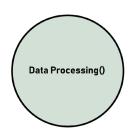
Feed Program Numerical Interrelationships();



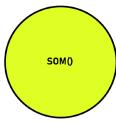
Feed Program Identity Vectors();



## Pseudocode<sub>2</sub>

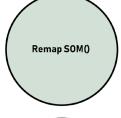


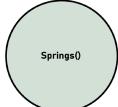
SOM()



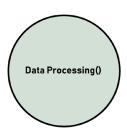
//Vanilla SOM adapted from Morphogenetic class

//Allows for a flexible matrix of variable subdivisions keeping with the total plan area required.





## Pseudocode<sub>3</sub>



#### SOM Remap()

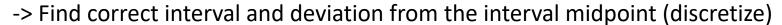
For each program



Find range of all summary vectors()

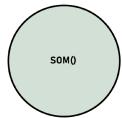
Divide range space into # intervals corresponding to number of program types();

Foreach particle

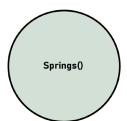


-> Store this multiplier value (instance) to be multiplied with a corresponding springConstant (type) to make up a weighted 'springConstant' for any given pair.

-> This performs a sort by how close each particle is from its corresponding BMU. This sort is visible in the SOM preview as a suffix of each particle's name . This gets carried forward to the spring system.







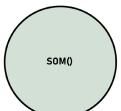
## Pseudocode<sub>4</sub>



#### Spring System()

Ref Dictionary - [Dictionary containing all program pair combinations (type) and corresponding springConstant for each]

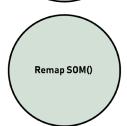
Construct all participating particles (instances) using their ranked name, their instance multiplier (how close they were to BMU), their starting location()



Generate a spring-chain of aforementioned particles (instances)

Iterate through 'ChainTension' which is a double for loop which iterates through all pairs:

-> Take particle-pair instance name and strip it of numbers/symbols to determine the 'type' pair'. ie. Office05\_ Storage11 would become Office\_Storage. If the type pair is accepted by the dictionary, then the type springConstant is multiplied by the instance springConstant. The radii of each instance pair is also taken into for restLength purposes.



//When 'attraction equilibrium' is reached, enable 'fitting' mode:

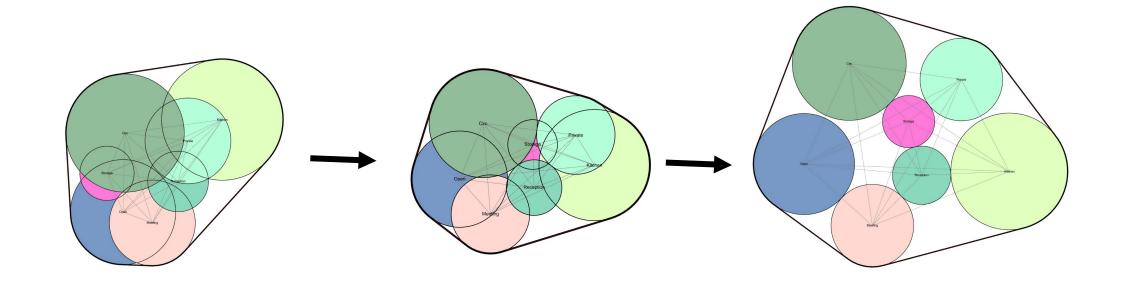
//Double for loop (similar to ChainReaction())to iterate through all pairs:

Find the restLength delta and vector between pairs to derive a new motion vector which in turn spurs a negative and equal motion vector on the other particle of the pair. Tally the number of interactions each particle has had with other particles.



Loop through the chain and add a new derived (average) position from above double loop, and divide by the number of interactions. This yields a motion that ends up separating all the particles while maintaining them at their desired restLength.

## Workflow Method 1 - Global\_Program

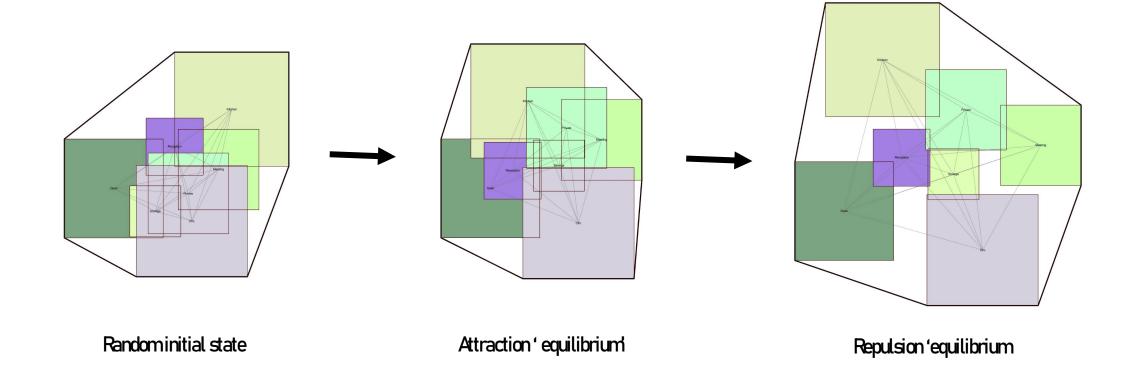


Randominitial state

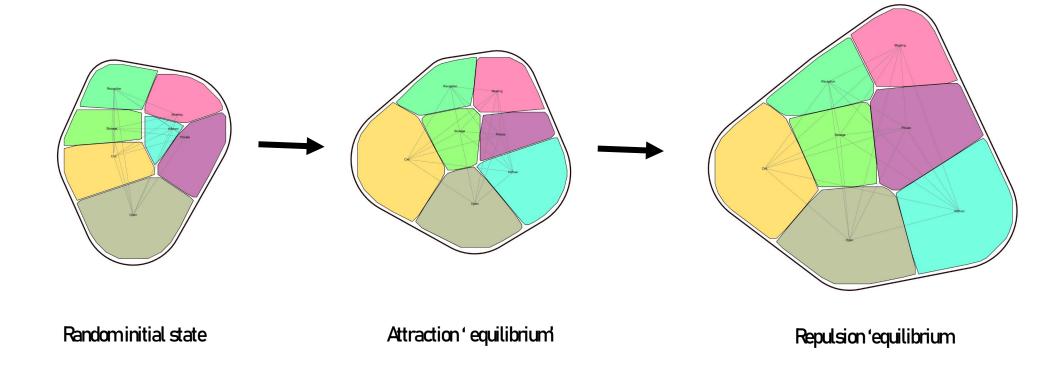
Attraction 'equilibrium'

Repulsion 'equilibrium

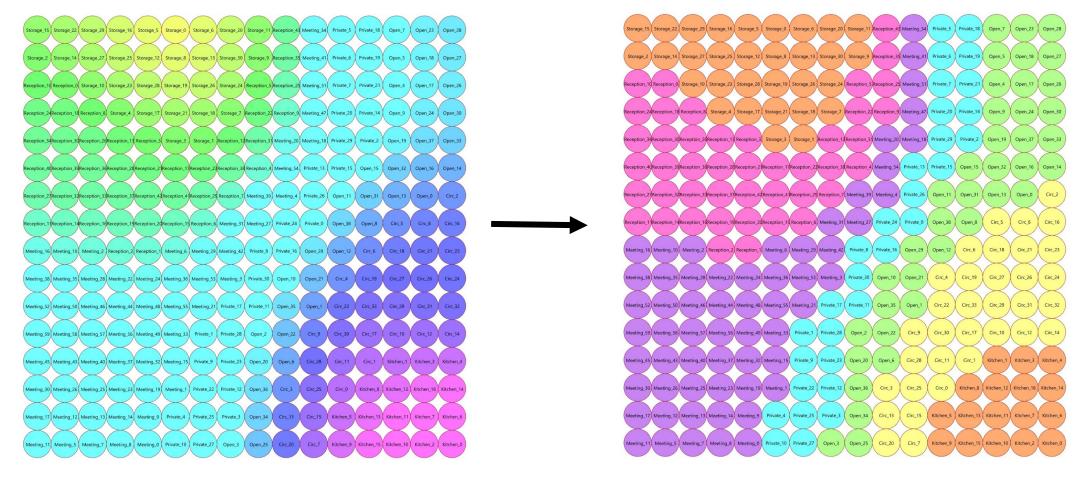
## Workflow Method 1 - Global\_Program



# Workflow Method 1 - Global\_Program



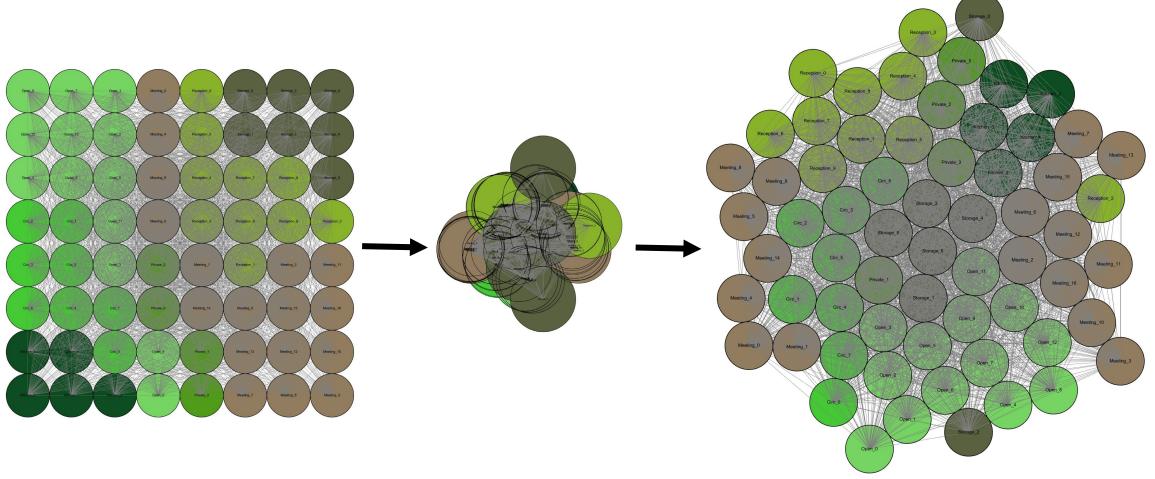
# Workflow Method 2 - Kohonen\_Program



Program Identity Gradient

Discretized Identity Gradient

# Workflow Method 2 - Kohonen\_Program



1) Initial State – all springs connected and weighted differently (similar to ANN)

2) Attraction 'equilibrium

3) Final 'fitted' separation state/ Repulsion equilibrium