

# BrainBuilder: Pseudocolumn

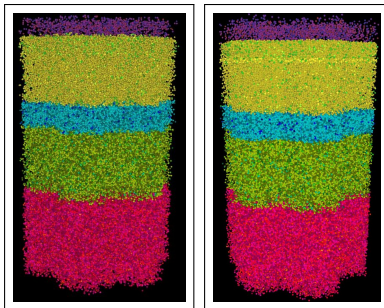
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Human Brain Project

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## Replication the column using BrainBuilder

Which one is which?



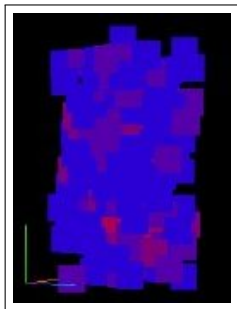
# Purpose

- ▶ See how well we can replicate the current column
- ▶ Have for check validation
- ▶ Have to do a 'vertical' column, since the later stages of the toolchain don't support arbitrary transformations

## Generate Densities

Take the original column, and 'voxel-ize' it to produce a cell density map

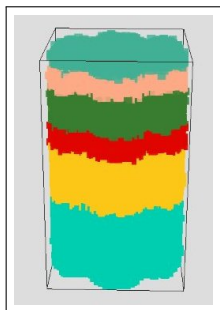
- ▶ Based only on soma positions
- ▶ Do not look at ME-Type or sClass



## Generate Annotation Map

Take the original column, and 'voxel-ize' it to produce a cell annotation map

- ▶ Per voxel, gather all the cells, and the most popular layer label 'wins'



## Generate Annotation Hierarchy

Take the original column, and 'voxel-ize' it to produce a cell density map

- ▶ Can't use the stock Allen Institute one: they use 1, 2/3, 4, 5, 6a, 6b, as SSCx layers
- ▶ Want to use our XML recipe, so have the same dimensions

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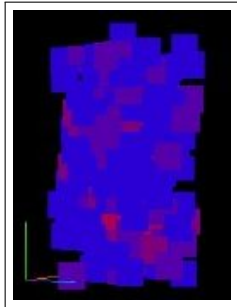
```
{  
  "id": 21,  
  "atlas_id": 977,  
  "ontology_id": 1,  
  "acronym": "SSp-111",  
  "name": "Primary somatosensory area, lower limb,  
    layer 1",  
  "color_hex_triplet": "188064",  
  "graph_order": 56,  
  "st_level": null,  
  "hemisphere_id": 3,
```

## Write Modules for BBP recipes

- ▶ Create 'Spatial Distributions' using the recipe
  - ▶ Need placement hints from NeuronDB.dat
  - ▶ Need cell percentages from builderRecipeAllPathways.xml

## Select the Region

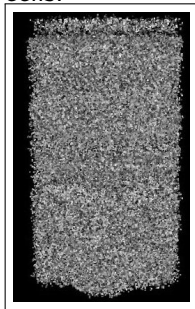
Select the whole column:





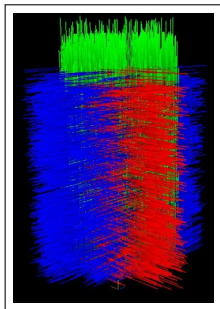
## Place the cell bodies

Colour doesn't mean anything, just needed contrast from different cells:



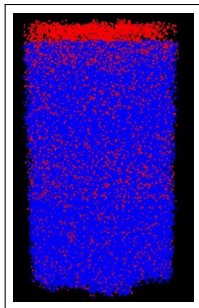
## Aside: Vector Fields

- ▶ For each voxel, need the nearest voxel that isn't of the same annotation
- ▶ Green is 'Up', which is all we need for the SSCx at the moment
- ▶ Other colours are for more complicated modules, if necessary



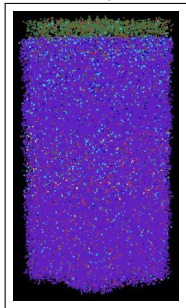
## Assign E-I-ness

Assign 'sclass' to cell bodies (Blue is Inhibitory):



## Assign ME-Types to cell bodies

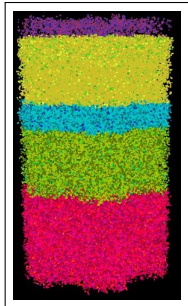
E-Type (Sorry about the colours):



Note: ME-Types are assigned in the same module

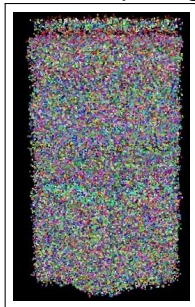
## Assign ME-Types to cell bodies

M-Type (Using same colour scheme as Platform column viewer):



## Assign morphologies to cell bodies

Each morphology is assigned a different colour:



## Future work

- ▶ Can synthesize the 'whole' brain, including 3D orientations, need new format to describe this freedom
- ▶ The scale of this pseudo-column matches the old column, so rat morphologies work, however the Allen Institute map is mouse. We can scale the voxel sizes appropriately
- ▶ Currently not worrying about, but that should be addressed:
  - ▶ Colliding soma positions
  - ▶ Morphologies protruding out of bounds
  - ▶ Coarseness of Allen Institute annotations: causes weird artifacts. Csaba has ways around this.