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CellOrganizer basics

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The main functions (API)

- `img2slml`
- `slml2img`

LEARNING A MODEL – IMG2SLML

img2slml

- Main inputs
 - a collection of OME-TIFF files
 - an options block
 - an output path
- Main outputs
 - parameterizations of each cell (through file system)
 - model

Image requirements

- Images to be used to learn models are in OME-TIFF format
- All channels for each field are in the same file
- The resolution (microns per pixel or voxel) must be specified

Cell segmentation

- Each file must contain
 - A single cell (no need for segmentation)
 - or
 - A channel that contains a mask to isolate one cell
 - or
 - Regions of interest, one per cell

2D Model Classes and Types

Class	Type	Depends on	Produces
vesicle	gmm	nuclear_membrane & cell_membrane	vesicle
nuclear_membrane	medial_axis		nuclear_membrane
cell_membrane	ratio	nuclear_membrane	cell_membrane
framework	diffeomorphic		cell_membrane, nuclear_membrane

3D Model classes and types

Class	Type	Depends on	Produces
vesicle	gmm	nuclear_membrane & cell_membrane	vesicle
nuclear_membrane	cylindrical_surface		nuclear_membrane
cell_membrane	ratio	nuclear_membrane	cell_membrane
framework	diffeomorphic		cell_membrane, nuclear_membrane
network	microtubule_growth	nuclear_membrane & cell_membrane	microtubule
protein_distribution	morphing		protein spatial distribution
standardized_voxels	standardized_map_half-ellipsoid		
csgo	half-ellipsoid		

Options block

- Model class and type
- Model type specific settings
- Output settings (name, id, filename)
- Helper options (debug, verbose, etc.)

SYNTHESIZING AN IMAGE – SLML2IMG2

slml2img

- Main inputs
 - a model file
 - an options block
 - an output path
- Main outputs – one or more
 - Images
 - SBML spatial geometry
 - Blender object files