


Dataset for the Article "Predicting vessel diameter changes to up-regulate biphasic blood flow during activation in realistic microvascular networks"

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Dataset for the Article "Predicting vessel diameter changes to up-regulate biphasic blood flow during activation in realistic microvascular networks"

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Research article

Epp, R., Schmid, F., Weber, B., and Jenny, P. (2020). Predicting vessel diameter changes to up-regulate biphasic blood flow during activation in realistic microvascular networks. *Front. Physiol.* doi:10.3389/fphys.2020.566303

Accessing the result files

The result files of the realistic microvascular networks are saved as compressed **Python pickled formats** and can be easily accessed with the Python module `python-igraph` (<https://igraph.org/python/>, tested with Python 2.7.17 and `python-igraph` 0.7.1)

Example: Import the network including the results from file (here for Scenario I with the activated region being the most central barrel of the MVN):

```
g = igraph.Graph.Read_Picklez('Barrel_1/Scenario_I/full_graph.pkl')
```

Additionally, the same results are also given as **csv-files** to ensure long-term accessibility. For each scenario, the three files “edgedata.csv”, “nodedata.csv” and “graphdata.csv” contain the same information as the corresponding `igraph-pkl`-file (adjacency edge list is included in “graphdata.csv”). For the artificial hexagonal network, the results are only included in the csv-files.

The realistic microvascular networks are from the mouse parietal cortex and were first published in Blinder et al., 2013, *Nature Neuroscience* (<https://doi.org/10.1038/nn.3426>).

All results are time averaged values as described in the corresponding research article.

Attributes of realistic MVNs

Node attributes

Attribute name	Description	Unit	Remark
<code>p_tav_base_si</code>	Pressure (baseline)	Pa	
<code>p_tav_opt_si</code>	Pressure (activation)	Pa	
<code>r</code>	xyz coordinate of vertex	μm	
<code>pBC</code>	Prescribed fixed boundary pressure at boundary nodes	mmHg	A pressure value is given if the current node is a boundary node (None, if it is not a boundary node)

Examples:

- `g.vs['p_tav_base_si'][99]` returns the baseline pressure (in Pascal) in the node with index 99.

- *g.vs['r'][99]* returns the *x, y and z coordinates (in micro meters) of the node with index 99.*

Edge attributes

Attribute name	Description	Unit	Remark
activated_region	Indicates if current edge is in the activated region	-	1 if edge is in activated barrel; 0 otherwise
vessel_type	Vessel type	-	0: Pial arteriole; 1: Pial venule; 2: Descending arteriole; 3: Ascending venule; 4: Capillary
d_base_si	Edge diameter (baseline)	m	
d_opt_si	Edge diameter (activation)	m	
fplasma_tav_base_si	Blood flow rate (baseline)	m ³ /sec	
fplasma_tav_opt_si	Blood flow rate (activation)	m ³ /sec	
ht_tav_base_si	Tube haematocrit (baseline)	-	
ht_tav_opt_si	Tube haematocrit (activation)	-	
rbc_flux_base_si	RBC flux (baseline)	m ³ /sec	
rbc_flux_opt_si	RBC flux (activation)	m ³ /sec	
urbc_tav_base_si	RBC velocity (baseline)	m/sec	
urbc_tav_opt_si	RBC velocity (activation)	m/sec	
length	Edge length	μm	
httBC	Prescribed fixed tube haematocrit at boundary edges	-	A value is given if the current edge is a boundary edge (None, if it is not a boundary edge)

Examples:

- *g.es['d_base_si'][99]* returns the baseline diameter (in micro meters) of the edge with index 99.

- *g.es['rbc_flux_opt_si'][99]* returns the RBC flux (in m³/sec) during activation in the edge with index 99.

Graph attributes

Attribute name	Description	Unit	Remark
activation_center_coords_si	xyz coordinate of activation centre	m	
activation_radius_si	Radius of activated barrel	m	

Example: g['activation_center_coords_si'] returns the *x, y and z coordinates (in meters) of the activation centre.*