TVB-Framework command initialisation

```
In [1]: import os
        from collections import OrderedDict
        import time
        import numpy as np
        from tvb.basic.profile import TvbProfile
        TvbProfile.set profile(TvbProfile.LIBRARY PROFILE)
        from tvb_multiscale.tvb_nest.config import *
        work path = os.getcwd()
        data path = work path
        tvb_conn_filepath = os.path.join(data_path, "Connectivity_res100_596_regions
        outputs path = os.path.join(work path, "outputs/cereb")
        config = Config(output base=outputs path)
        config.figures.SHOW FLAG = True
        config.figures.SAVE FLAG = True
        config.figures.FIG FORMAT = 'png'
        config.figures.DEFAULT_SIZE= config.figures.NOTEBOOK_SIZE
        FIGSIZE = config.figures.DEFAULT SIZE
        from tvb_multiscale.core.plot.plotter import Plotter
        plotter = Plotter(config.figures)
        WEIGHTED AVERAGE CENTRES = True
        TRACT_LENGTHS_MODE = "weighted_average" # "weighted_average", "euclidean"
        # "average" is not correct at all for tract lenghts,
        # because we are merging recursively one major structure after the other,
        # which could work only if the merged regions have a larger weights in a wei
        # for subsequent mergings of other regions.
        FORCE_MERGED_ZERO_DIAGONAL = True
        # For interactive plotting:
        # %matplotlib notebook
        # Otherwise:
        %matplotlib inline
```

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2023-01-30 11:44:24,833 - INFO - tvb_multiscale.tvb_nest.config - Loading a
NEST instance...
2023-01-30 11:44:24,833 - INFO - tvb multiscale.tvb nest.config - Loading a
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2023-01-30 11:44:24,839 - INFO - tvb multiscale.tvb nest.config - NEST INST
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This loads the just-prepared data:

```
In [2]:
        import h5py
        from tvb.datatypes.connectivity import Connectivity
        # Load connnectivity from file:
        f = h5py.File(tvb conn filepath)
        # We have to transpose the square connectivity matrices for TVB, because of
        result connectivity = Connectivity(weights=np.array(f["weights"][()]).T,
                                            tract_lengths=np.array(f["tract_lengths"]
                                            centres=np.array(f["centres"][()]), # hen
                                            region labels=np.array(f["region labels"]
        f.close()
        # For the full connectome, self-connections are meaningless:
        np.fill_diagonal(result_connectivity.weights, 0.0)
        np.fill_diagonal(result_connectivity.tract_lengths, 0.0)
        result connectivity.tract lengths[result connectivity.weights == 0.0] = 0.0
        result connectivity.configure()
        print('number_of_regions:\n', result_connectivity.number_of_regions)
        print('\nweights.shape:\n', result_connectivity.weights.shape)
        print('\ntract_lengths.shape:\n', result_connectivity.tract_lengths.shape)
        print('\nregion_labels:\n', result_connectivity.region_labels)
        print('\nweights:\n', result_connectivity.weights)
```

```
number_of_regions:
 596
weights.shape:
 (596, 596)
tract lengths.shape:
 (596, 596)
region labels:
 ['Right Frontal pole, cerebral cortex' 'Right Primary motor area'
 'Right Secondary motor area' 'Right Primary somatosensory area, nose'
 'Right Primary somatosensory area, barrel field'
 'Right Primary somatosensory area, lower limb'
 'Right Primary somatosensory area, mouth'
 'Right Primary somatosensory area, upper limb'
 'Right Primary somatosensory area, trunk'
 'Right Primary somatosensory area, unassigned'
 'Right Supplemental somatosensory area' 'Right Gustatory areas'
 'Right Visceral area' 'Right Dorsal auditory area'
 'Right Primary auditory area' 'Right Posterior auditory area'
 'Right Ventral auditory area' 'Right Anterolateral visual area'
 'Right Anteromedial visual area' 'Right Lateral visual area'
 'Right Primary visual area' 'Right Posterolateral visual area'
 'Right posteromedial visual area' 'Right Laterointermediate area'
 'Right Postrhinal area' 'Right Anterior cinqulate area, dorsal part'
 'Right Anterior cingulate area, ventral part' 'Right Prelimbic area'
 'Right Infralimbic area' 'Right Orbital area, lateral part'
 'Right Orbital area, medial part'
 'Right Orbital area, ventrolateral part'
 'Right Agranular insular area, dorsal part'
 'Right Agranular insular area, posterior part'
 'Right Agranular insular area, ventral part'
 'Right Retrosplenial area, lateral agranular part'
 'Right Retrosplenial area, dorsal part'
 'Right Retrosplenial area, ventral part' 'Right Anterior area'
 'Right Rostrolateral visual area' 'Right Temporal association areas'
 'Right Perirhinal area' 'Right Ectorhinal area'
 'Right Main olfactory bulb' 'Right Accessory olfactory bulb'
 'Right Anterior olfactory nucleus' 'Right Taenia tecta'
 'Right Dorsal peduncular area' 'Right Piriform area'
 'Right Nucleus of the lateral olfactory tract'
 'Right Cortical amygdalar area, anterior part'
 'Right Cortical amygdalar area, posterior part'
 'Right Piriform-amygdalar area' 'Right Postpiriform transition area'
 'Right Field CA1' 'Right Field CA2' 'Right Field CA3'
 'Right Dentate gyrus' 'Right Induseum griseum'
 'Right Entorhinal area, lateral part'
 'Right Entorhinal area, medial part, dorsal zone' 'Right Parasubiculum'
 'Right Postsubiculum' 'Right Presubiculum' 'Right Subiculum'
 'Right Prosubiculum' 'Right Area prostriata' 'Right Claustrum'
 'Right Endopiriform nucleus, dorsal part'
 'Right Endopiriform nucleus, ventral part'
 'Right Lateral amygdalar nucleus' 'Right Basolateral amygdalar nucleus'
 'Right Basomedial amygdalar nucleus' 'Right Posterior amygdalar nucleus'
 'Right Caudoputamen' 'Right Nucleus accumbens' 'Right Fundus of striatum'
 'Right Olfactory tubercle'
 'Right Lateral septal nucleus, caudal (caudodorsal) part'
 'Right Lateral septal nucleus, rostral (rostroventral) part'
 'Right Lateral septal nucleus, ventral part'
 'Right Septofimbrial nucleus' 'Right Anterior amygdalar area'
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'Right Bed nucleus of the accessory olfactory tract' 'Right Central amygdalar nucleus' 'Right Intercalated amygdalar nucleus' 'Right Medial amygdalar nucleus' 'Right Globus pallidus, external segment' 'Right Globus pallidus, internal segment' 'Right Substantia innominata' 'Right Magnocellular nucleus' 'Right Medial septal nucleus' 'Right Diagonal band nucleus' 'Right Triangular nucleus of septum' 'Right Bed nuclei of the stria terminalis' 'Right Bed nucleus of the anterior commissure' 'Right Ventral anterior-lateral complex of the thalamus' 'Right Ventral medial nucleus of the thalamus' 'Right Ventral posterolateral nucleus of the thalamus' 'Right Ventral posterolateral nucleus of the thalamus, parvicellular part' 'Right Ventral posteromedial nucleus of the thalamus' 'Right Ventral posteromedial nucleus of the thalamus, parvicellular part' 'Right Posterior triangular thalamic nucleus' 'Right Subparafascicular nucleus, magnocellular part' 'Right Subparafascicular nucleus, parvicellular part' 'Right Subparafascicular area' 'Right Peripeduncular nucleus' 'Right Medial geniculate complex' 'Right Dorsal part of the lateral geniculate complex' 'Right Lateral posterior nucleus of the thalamus' 'Right Posterior complex of the thalamus' 'Right Posterior limiting nucleus of the thalamus' 'Right Suprageniculate nucleus' 'Right Anteroventral nucleus of thalamus' 'Right Anteromedial nucleus' 'Right Anterodorsal nucleus' 'Right Interanteromedial nucleus of the thalamus' 'Right Interanterodorsal nucleus of the thalamus' 'Right Lateral dorsal nucleus of thalamus' 'Right Intermediodorsal nucleus of the thalamus' 'Right Mediodorsal nucleus of thalamus' 'Right Submedial nucleus of the thalamus' 'Right Perireunensis nucleus' 'Right Paraventricular nucleus of the thalamus' 'Right Parataenial nucleus' 'Right Nucleus of reuniens' 'Right Xiphoid thalamic nucleus' 'Right Rhomboid nucleus' 'Right Central medial nucleus of the thalamus' 'Right Paracentral nucleus' 'Right Central lateral nucleus of the thalamus' 'Right Parafascicular nucleus' 'Right Posterior intralaminar thalamic nucleus' 'Right Reticular nucleus of the thalamus' 'Right Intergeniculate leaflet of the lateral geniculate complex' 'Right Intermediate geniculate nucleus' 'Right Ventral part of the lateral geniculate complex' 'Right Medial habenula' 'Right Lateral habenula' 'Right Accessory supraoptic group' 'Right Paraventricular hypothalamic nucleus' 'Right Periventricular hypothalamic nucleus, anterior part' 'Right Periventricular hypothalamic nucleus, intermediate part' 'Right Arcuate hypothalamic nucleus' 'Right Anterodorsal preoptic nucleus' 'Right Anteroventral preoptic nucleus' 'Right Anteroventral periventricular nucleus' 'Right Dorsomedial nucleus of the hypothalamus' 'Right Median preoptic nucleus' 'Right Medial preoptic area' 'Right Vascular organ of the lamina terminalis' 'Right Posterodorsal preoptic nucleus' 'Right Parastrial nucleus' 'Right Periventricular hypothalamic nucleus, posterior part' 'Right Periventricular hypothalamic nucleus, preoptic part' 'Right Subparaventricular zone' 'Right Suprachiasmatic nucleus' 'Right Ventromedial preoptic nucleus'

'Right Ventrolateral preoptic nucleus' 'Right Anterior hypothalamic nucleus' 'Right Lateral mammillary nucleus' 'Right Medial mammillary nucleus' 'Right Supramammillary nucleus' 'Right Tuberomammillary nucleus, dorsal part' 'Right Tuberomammillary nucleus, ventral part' 'Right Medial preoptic nucleus' 'Right Dorsal premammillary nucleus' 'Right Ventral premammillary nucleus' 'Right Paraventricular hypothalamic nucleus, descending division' 'Right Ventromedial hypothalamic nucleus' 'Right Posterior hypothalamic nucleus' 'Right Lateral hypothalamic area' 'Right Lateral preoptic area' 'Right Preparasubthalamic nucleus' 'Right Parasubthalamic nucleus' 'Right Perifornical nucleus' 'Right Retrochiasmatic area' 'Right Subthalamic nucleus' 'Right Tuberal nucleus' 'Right Zona incerta' 'Right Superior colliculus, sensory related' 'Right Inferior colliculus' 'Right Nucleus of the brachium of the inferior colliculus' 'Right Nucleus sagulum' 'Right Parabigeminal nucleus' 'Right Midbrain trigeminal nucleus' 'Right Subcommissural organ' 'Right Substantia nigra, reticular part' 'Right Ventral tegmental area' 'Right Paranigral nucleus' 'Right Midbrain reticular nucleus, retrorubral area' 'Right Midbrain reticular nucleus' 'Right Superior colliculus, motor related' 'Right Periaqueductal gray' 'Right Anterior pretectal nucleus' 'Right Medial pretectal area' 'Right Nucleus of the optic tract' 'Right Nucleus of the posterior commissure' 'Right Olivary pretectal nucleus' 'Right Posterior pretectal nucleus' 'Right Cuneiform nucleus' 'Right Red nucleus' 'Right Oculomotor nucleus' 'Right Medial accesory oculomotor nucleus' 'Right Edinger-Westphal nucleus' 'Right Trochlear nucleus' 'Right Paratrochlear nucleus' 'Right Ventral tegmental nucleus' 'Right Anterior tegmental nucleus' 'Right Lateral terminal nucleus of the accessory optic tract' 'Right Dorsal terminal nucleus of the accessory optic tract' 'Right Medial terminal nucleus of the accessory optic tract' 'Right Substantia nigra, compact part' 'Right Pedunculopontine nucleus' 'Right Interfascicular nucleus raphe' 'Right Interpeduncular nucleus' 'Right Rostral linear nucleus raphe' 'Right Central linear nucleus raphe' 'Right Dorsal nucleus raphe' 'Right Nucleus of the lateral lemniscus' 'Right Principal sensory nucleus of the trigeminal' 'Right Parabrachial nucleus' "Right Barrington's nucleus" 'Right Dorsal tegmental nucleus' 'Right Posterodorsal tegmental nucleus' 'Right Pontine central gray' 'Right Pontine gray' 'Right Pontine reticular nucleus, caudal part' 'Right Supratrigeminal nucleus' 'Right Tegmental reticular nucleus' 'Right Motor nucleus of trigeminal' 'Right Peritrigeminal zone' 'Right Intertrigeminal nucleus' 'Right Superior central nucleus raphe' 'Right Locus ceruleus' 'Right Laterodorsal tegmental nucleus' 'Right Nucleus incertus' 'Right Pontine reticular nucleus' 'Right Nucleus raphe pontis' 'Right Subceruleus nucleus' 'Right Sublaterodorsal nucleus' 'Right Dorsal cochlear nucleus' 'Right Ventral cochlear nucleus' 'Right Cuneate nucleus' 'Right Gracile nucleus' 'Right Nucleus of the trapezoid body' 'Right Nucleus of the solitary tract' 'Right Spinal nucleus of the trigeminal, caudal part' 'Right Spinal nucleus of the trigeminal, interpolar part' 'Right Spinal nucleus of the trigeminal, oral part' 'Right Paratrigeminal nucleus' 'Right Abducens nucleus' 'Right Facial motor nucleus' 'Right Nucleus ambiguus' 'Right Dorsal motor nucleus of the vagus nerve' 'Right Gigantocellular reticular nucleus'

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'Right Inferior olivary complex' 'Right Intermediate reticular nucleus'
'Right Inferior salivatory nucleus' 'Right Linear nucleus of the medulla'
'Right Lateral reticular nucleus' 'Right Magnocellular reticular nucleus'
'Right Medullary reticular nucleus'
'Right Medullary reticular nucleus, dorsal part'
'Right Medullary reticular nucleus, ventral part'
'Right Parvicellular reticular nucleus'
'Right Paragigantocellular reticular nucleus, dorsal part'
'Right Paragigantocellular reticular nucleus, lateral part'
'Right Nucleus of Roller' 'Right Nucleus prepositus'
'Right Parapyramidal nucleus' 'Right Lateral vestibular nucleus'
'Right Medial vestibular nucleus' 'Right Spinal vestibular nucleus'
'Right Superior vestibular nucleus' 'Right Nucleus x'
'Right Hypoglossal nucleus' 'Right Nucleus y'
'Right Nucleus raphe magnus' 'Right Nucleus raphe obscurus'
'Right Lingula (I)' 'Right Central lobule' 'Right Culmen'
'Right Declive (VI)' 'Right Folium-tuber vermis (VII)'
'Right Pyramus (VIII)' 'Right Uvula (IX)' 'Right Nodulus (X)'
'Right Simple lobule' 'Right Ansiform lobule' 'Right Paramedian lobule'
'Right Copula pyramidis' 'Right Paraflocculus' 'Right Flocculus'
'Right Fastigial nucleus' 'Right Interposed nucleus'
'Right Dentate nucleus' 'Right Vestibulocerebellar nucleus'
'Left Frontal pole, cerebral cortex' 'Left Primary motor area'
'Left Secondary motor area' 'Left Primary somatosensory area, nose'
'Left Primary somatosensory area, barrel field'
'Left Primary somatosensory area, lower limb'
'Left Primary somatosensory area, mouth'
'Left Primary somatosensory area, upper limb'
'Left Primary somatosensory area, trunk'
'Left Primary somatosensory area, unassigned'
'Left Supplemental somatosensory area' 'Left Gustatory areas'
'Left Visceral area' 'Left Dorsal auditory area'
'Left Primary auditory area' 'Left Posterior auditory area'
'Left Ventral auditory area' 'Left Anterolateral visual area'
'Left Anteromedial visual area' 'Left Lateral visual area'
'Left Primary visual area' 'Left Posterolateral visual area'
'Left posteromedial visual area' 'Left Laterointermediate area'
'Left Postrhinal area' 'Left Anterior cingulate area, dorsal part'
'Left Anterior cingulate area, ventral part' 'Left Prelimbic area'
'Left Infralimbic area' 'Left Orbital area, lateral part'
'Left Orbital area, medial part' 'Left Orbital area, ventrolateral part'
'Left Agranular insular area, dorsal part'
'Left Agranular insular area, posterior part'
'Left Agranular insular area, ventral part'
'Left Retrosplenial area, lateral agranular part'
'Left Retrosplenial area, dorsal part'
'Left Retrosplenial area, ventral part' 'Left Anterior area'
'Left Rostrolateral visual area' 'Left Temporal association areas'
'Left Perirhinal area' 'Left Ectorhinal area' 'Left Main olfactory bulb'
'Left Accessory olfactory bulb' 'Left Anterior olfactory nucleus'
'Left Taenia tecta' 'Left Dorsal peduncular area' 'Left Piriform area'
'Left Nucleus of the lateral olfactory tract'
'Left Cortical amygdalar area, anterior part'
'Left Cortical amygdalar area, posterior part'
'Left Piriform-amygdalar area' 'Left Postpiriform transition area'
'Left Field CA1' 'Left Field CA2' 'Left Field CA3' 'Left Dentate gyrus'
'Left Induseum griseum' 'Left Entorhinal area, lateral part'
'Left Entorhinal area, medial part, dorsal zone' 'Left Parasubiculum'
'Left Postsubiculum' 'Left Presubiculum' 'Left Subiculum'
'Left Prosubiculum' 'Left Area prostriata' 'Left Claustrum'
'Left Endopiriform nucleus, dorsal part'
```

interacting_with_Allen 'Left Endopiriform nucleus, ventral part' 'Left Lateral amygdalar nucleus' 'Left Basolateral amygdalar nucleus' 'Left Basomedial amygdalar nucleus' 'Left Posterior amygdalar nucleus' 'Left Caudoputamen' 'Left Nucleus accumbens' 'Left Fundus of striatum' 'Left Olfactory tubercle' 'Left Lateral septal nucleus, caudal (caudodorsal) part' 'Left Lateral septal nucleus, rostral (rostroventral) part' 'Left Lateral septal nucleus, ventral part' 'Left Septofimbrial nucleus' 'Left Anterior amygdalar area' 'Left Bed nucleus of the accessory olfactory tract' 'Left Central amygdalar nucleus' 'Left Intercalated amygdalar nucleus' 'Left Medial amygdalar nucleus' 'Left Globus pallidus, external segment' 'Left Globus pallidus, internal segment' 'Left Substantia innominata' 'Left Magnocellular nucleus' 'Left Medial septal nucleus' 'Left Diagonal band nucleus' 'Left Triangular nucleus of septum' 'Left Bed nuclei of the stria terminalis' 'Left Bed nucleus of the anterior commissure' 'Left Ventral anterior-lateral complex of the thalamus' 'Left Ventral medial nucleus of the thalamus' 'Left Ventral posterolateral nucleus of the thalamus' 'Left Ventral posterolateral nucleus of the thalamus, parvicellular part' 'Left Ventral posteromedial nucleus of the thalamus' 'Left Ventral posteromedial nucleus of the thalamus, parvicellular part' 'Left Posterior triangular thalamic nucleus' 'Left Subparafascicular nucleus, magnocellular part' 'Left Subparafascicular nucleus, parvicellular part' 'Left Subparafascicular area' 'Left Peripeduncular nucleus' 'Left Medial geniculate complex' 'Left Dorsal part of the lateral geniculate complex' 'Left Lateral posterior nucleus of the thalamus' 'Left Posterior complex of the thalamus' 'Left Posterior limiting nucleus of the thalamus' 'Left Suprageniculate nucleus' 'Left Anteroventral nucleus of thalamus' 'Left Anteromedial nucleus' 'Left Anterodorsal nucleus' 'Left Interanteromedial nucleus of the thalamus' 'Left Interanterodorsal nucleus of the thalamus' 'Left Lateral dorsal nucleus of thalamus' 'Left Intermediodorsal nucleus of the thalamus' 'Left Mediodorsal nucleus of thalamus' 'Left Submedial nucleus of the thalamus' 'Left Perireunensis nucleus' 'Left Paraventricular nucleus of the thalamus' 'Left Parataenial nucleus' 'Left Nucleus of reuniens' 'Left Xiphoid thalamic nucleus' 'Left Rhomboid nucleus' 'Left Central medial nucleus of the thalamus' 'Left Paracentral nucleus' 'Left Central lateral nucleus of the thalamus' 'Left Parafascicular nucleus' 'Left Posterior intralaminar thalamic nucleus' 'Left Reticular nucleus of the thalamus' 'Left Intergeniculate leaflet of the lateral geniculate complex' 'Left Intermediate geniculate nucleus' 'Left Ventral part of the lateral geniculate complex' 'Left Medial habenula' 'Left Lateral habenula' 'Left Accessory supraoptic group' 'Left Paraventricular hypothalamic nucleus' 'Left Periventricular hypothalamic nucleus, anterior part' 'Left Periventricular hypothalamic nucleus, intermediate part' 'Left Arcuate hypothalamic nucleus' 'Left Anterodorsal preoptic nucleus' 'Left Anteroventral preoptic nucleus' 'Left Anteroventral periventricular nucleus' 'Left Dorsomedial nucleus of the hypothalamus' 'Left Median preoptic nucleus' 'Left Medial preoptic area'

'Left Vascular organ of the lamina terminalis'

- interacting_with_Allen 'Left Posterodorsal preoptic nucleus' 'Left Parastrial nucleus' 'Left Periventricular hypothalamic nucleus, posterior part' 'Left Periventricular hypothalamic nucleus, preoptic part' 'Left Subparaventricular zone' 'Left Suprachiasmatic nucleus' 'Left Ventromedial preoptic nucleus' 'Left Ventrolateral preoptic nucleus' 'Left Anterior hypothalamic nucleus' 'Left Lateral mammillary nucleus' 'Left Medial mammillary nucleus' 'Left Supramammillary nucleus' 'Left Tuberomammillary nucleus, dorsal part' 'Left Tuberomammillary nucleus, ventral part' 'Left Medial preoptic nucleus' 'Left Dorsal premammillary nucleus' 'Left Ventral premammillary nucleus' 'Left Paraventricular hypothalamic nucleus, descending division' 'Left Ventromedial hypothalamic nucleus' 'Left Posterior hypothalamic nucleus' 'Left Lateral hypothalamic area' 'Left Lateral preoptic area' 'Left Preparasubthalamic nucleus' 'Left Parasubthalamic nucleus' 'Left Perifornical nucleus' 'Left Retrochiasmatic area' 'Left Subthalamic nucleus' 'Left Tuberal nucleus' 'Left Zona incerta' 'Left Superior colliculus, sensory related' 'Left Inferior colliculus' 'Left Nucleus of the brachium of the inferior colliculus' 'Left Nucleus sagulum' 'Left Parabigeminal nucleus' 'Left Midbrain trigeminal nucleus' 'Left Subcommissural organ' 'Left Substantia nigra, reticular part' 'Left Ventral tegmental area' 'Left Paranigral nucleus' 'Left Midbrain reticular nucleus, retrorubral area' 'Left Midbrain reticular nucleus' 'Left Superior colliculus, motor related' 'Left Periaqueductal gray' 'Left Anterior pretectal nucleus' 'Left Medial pretectal area' 'Left Nucleus of the optic tract' 'Left Nucleus of the posterior commissure' 'Left Olivary pretectal nucleus' 'Left Posterior pretectal nucleus' 'Left Cuneiform nucleus' 'Left Red nucleus' 'Left Oculomotor nucleus' 'Left Medial accesory oculomotor nucleus' 'Left Edinger-Westphal nucleus' 'Left Trochlear nucleus' 'Left Paratrochlear nucleus' 'Left Ventral tegmental nucleus' 'Left Anterior tegmental nucleus' 'Left Lateral terminal nucleus of the accessory optic tract' 'Left Dorsal terminal nucleus of the accessory optic tract' 'Left Medial terminal nucleus of the accessory optic tract' 'Left Substantia nigra, compact part' 'Left Pedunculopontine nucleus' 'Left Interfascicular nucleus raphe' 'Left Interpeduncular nucleus' 'Left Rostral linear nucleus raphe' 'Left Central linear nucleus raphe' 'Left Dorsal nucleus raphe' 'Left Nucleus of the lateral lemniscus' 'Left Principal sensory nucleus of the trigeminal' 'Left Parabrachial nucleus' "Left Barrington's nucleus" 'Left Dorsal tegmental nucleus' 'Left Posterodorsal tegmental nucleus' 'Left Pontine central gray' 'Left Pontine gray' 'Left Pontine reticular nucleus, caudal part' 'Left Supratrigeminal nucleus' 'Left Tegmental reticular nucleus' 'Left Motor nucleus of trigeminal' 'Left Peritrigeminal zone' 'Left Intertrigeminal nucleus' 'Left Superior central nucleus raphe' 'Left Locus ceruleus' 'Left Laterodorsal tegmental nucleus' 'Left Nucleus incertus' 'Left Pontine reticular nucleus' 'Left Nucleus raphe pontis' 'Left Subceruleus nucleus' 'Left Sublaterodorsal nucleus' 'Left Dorsal cochlear nucleus' 'Left Ventral cochlear nucleus' 'Left Cuneate nucleus' 'Left Gracile nucleus' 'Left Nucleus of the trapezoid body' 'Left Nucleus of the solitary tract' 'Left Spinal nucleus of the trigeminal, caudal part' 'Left Spinal nucleus of the trigeminal, interpolar part' 'Left Spinal nucleus of the trigeminal, oral part'
- localhost:8888/nbconvert/html/packages/tvb-multiscale/examples/data/cerebellum/interacting_with_Allen.ipynb?download=false

```
'Left Paratrigeminal nucleus' 'Left Abducens nucleus'
 'Left Facial motor nucleus' 'Left Nucleus ambiguus'
 'Left Dorsal motor nucleus of the vagus nerve'
 'Left Gigantocellular reticular nucleus' 'Left Inferior olivary complex'
 'Left Intermediate reticular nucleus' 'Left Inferior salivatory nucleus'
 'Left Linear nucleus of the medulla' 'Left Lateral reticular nucleus'
 'Left Magnocellular reticular nucleus' 'Left Medullary reticular nucleus'
 'Left Medullary reticular nucleus, dorsal part'
 'Left Medullary reticular nucleus, ventral part'
 'Left Parvicellular reticular nucleus'
 'Left Paragigantocellular reticular nucleus, dorsal part'
 'Left Paragigantocellular reticular nucleus, lateral part'
 'Left Nucleus of Roller' 'Left Nucleus prepositus'
 'Left Parapyramidal nucleus' 'Left Lateral vestibular nucleus'
 'Left Medial vestibular nucleus' 'Left Spinal vestibular nucleus'
 'Left Superior vestibular nucleus' 'Left Nucleus x'
 'Left Hypoglossal nucleus' 'Left Nucleus y' 'Left Nucleus raphe magnus'
 'Left Nucleus raphe obscurus' 'Left Lingula (I)' 'Left Central lobule'
 'Left Culmen' 'Left Declive (VI)' 'Left Folium-tuber vermis (VII)'
 'Left Pyramus (VIII)' 'Left Uvula (IX)' 'Left Nodulus (X)'
 'Left Simple lobule' 'Left Ansiform lobule' 'Left Paramedian lobule'
 'Left Copula pyramidis' 'Left Paraflocculus' 'Left Flocculus'
 'Left Fastigial nucleus' 'Left Interposed nucleus' 'Left Dentate nucleus'
 'Left Vestibulocerebellar nucleus']
weights:
 [[0.00000000e+00 3.63441680e-02 8.47970561e-02 ... 8.07329172e-07
 8.07329172e-07 1.02162597e-041
 [3.32728901e-02 0.00000000e+00 5.49682272e-02 ... 2.73112949e-05
 2.85348052e-05 2.77993498e-051
 [9.89313689e-02 6.43270451e-02 0.00000000e+00 ... 3.36029153e-05
 3.54584335e-05 9.15227954e-061
 [1.36803986e-06 6.32084139e-05 2.99611362e-05 ... 0.00000000e+00
 1.00408312e-01 1.37087573e-02]
 [1.95515882e-07 5.04100941e-05 4.17897432e-05 ... 5.17439950e-02
 0.00000000e+00 2.65998675e-031
 [2.71852623e-05 4.45484959e-04 2.03744779e-04 ... 3.86534058e-02
 3.67525396e-02 0.00000000e+0011
```

Summarize the major structures, except for the cerebellum

```
In [3]: # Open the xls file and create a mapping between all regions and the major s

#pip install xlrd
try:
    import xlrd

except:
    import sys
    !{sys.executable} -m pip install xlrd
    import xlrd

sheet_data = []
wb = xlrd.open_workbook(os.path.join(data_path, 'oh_table1.xls'))
p = wb.sheet_names()
#p[1]
#for y in p:
sh = wb.sheet_by_name(p[1])
```

```
print(sh)
for rownum in range(sh.nrows):
    sheet_data.append((sh.row_values(rownum)))
found list = []
voxel count = []
rows_to_be_saved = []
regions missing = []
cc=0
for j in result connectivity.region labels: #go over the region labels
    j=j.split(" ",1)[1]
    #let us get rid of the left or right
    a=len(found list)
    for i in sheet data:
        if i[3] == j: #or i[2] == "string2" or i[2] == "string3" or i[2] ==
            found list.append(i[4])
            if i[5]:
                voxel_count.append(int(i[5]))
            else:
                voxel count.append(-1)
        else:
            rows_to_be_saved.append(i)
    b=len(found list)
    if a == b:
        regions missing.append(j)
        found list.append('X')
        voxel_count.append(-1)
        print(a,b)
    b=len(found list)
    cc+=1
    if b != cc:
        print(b,cc)
print("Regions missing:\n%s" % str(regions_missing))
n_regs = len(found_list)
print("Number of regions: %d" % n_regs)
n regs2 = n regs/2
major_structures_labels = ["Right " + msl if iL <n_regs2 else "Left " + msl</pre>
                           for iL, msl in enumerate(found_list)]
voxel count = np.array(voxel count).astype('i')
major structures = np.unique(major structures labels)
print("\nmajor_structures:\n", major_structures)
```

```
Sheet 1:<Voxel Count 295 Structures>
Regions missing:
[]
Number of regions: 596
maior structures:
 ['Left Cerebellar Cortex' 'Left Cerebellar Nuclei'
 'Left Cortical Subplate' 'Left Hippocampal Formation' 'Left Hypothalamus'
 'Left Isocortex' 'Left Medulla' 'Left Midbrain'
 'Left Nonspecific Thalamus' 'Left Olfactory Areas' 'Left Pallidum'
 'Left Pons Behavioral' 'Left Pons Motor' 'Left Pons Sensory'
 'Left Spinal nucleus of the trigeminal' 'Left Striatum' 'Left Thalamus'
 'Right Cerebellar Cortex' 'Right Cerebellar Nuclei'
 'Right Cortical Subplate' 'Right Hippocampal Formation'
 'Right Hypothalamus' 'Right Isocortex' 'Right Medulla' 'Right Midbrain'
 'Right Nonspecific Thalamus' 'Right Olfactory Areas' 'Right Pallidum'
 'Right Pons Behavioral' 'Right Pons Motor' 'Right Pons Sensory'
 'Right Spinal nucleus of the trigeminal' 'Right Striatum'
 'Right Thalamus']
```

Find the connections FROM Isocortex nodes TO the Reticular Thalamic Nucleus and store them:

```
In [4]: crtx_inds = np.where(["Isocortex" in lbl for lbl in major_structures_labels]
    rtn_inds = np.where(["Reticular nucleus of the thalamus" in reg for reg in r
    crtx_to_rtn_conns = result_connectivity.weights[rtn_inds][:, crtx_inds]
    print("crtx_to_rtn_conns.shape: ", crtx_to_rtn_conns.shape)
    # print("\ncrtx_to_rtn_conns:\n","crtx_to_rtn_conns)
    crtx_to_rtn_conns.shape: (2, 86)
```

Some functions to use for merging regions to larger structures

```
def voxel_count_sum(arr, axis=0, **kwargs):
    voxel_count_sum = np.sum(arr[arr>0], axis=axis)
    print("Voxel count sum: %s" % str(voxel_count_sum))
    return voxel_count_sum

def weighted_average(arr, axis=0, **kwargs):
    weights = kwargs.pop('weights', 1.0)
    if weights.ndim < arr.ndim:
        weights = np.expand_dims(weights, 1-axis)
    assert np.nansum(weights) > 0.0
    wav = np.nansum(arr * weights, axis=axis, **kwargs) / np.nansum(weights, return wav)

def repeat(arr, sub, axis=0, **kwargs):
```

```
"""This function will tile a subarray
      to create an array of shape similar to the input array's arr,
       except for the axis given in the input, where size will be 1.
       It is used to substitute many labels by a single one."""
    shape = list(arr.shape)
    shape[axis] = 1
    return np.tile(sub, tuple(shape))
def delete(arr, axis=0, **kwargs):
    """This function returns an empty array
       of shape similar to the input array's arr,
       except for the axis given in the input, where size will be 0.
       It is used to delete a subarray."""
    shape = list(arr.shape)
    shape[axis] = 0
    return np.empty(tuple(shape))
def insert axis(arr, arr to insert=None, inds=None, axis=0, def value=0.0):
    if inds is None:
        inds = [arr.shape[axis]]
    if arr_to_insert is None:
        arr_to_insert = np.tile(def_value, np.take(arr, [-1], axis=axis).sha
    return np.insert(arr, inds, arr to insert, axis=axis)
def merge_axis(inds, arr, axis=0, fun=np.nansum, **funkwargs):
    """This function will merge a subarray of the input array arr,
       as defined by the input indices inds, along the input axis,
       applying the function fun, in order to summarize the values."""
    new_arr = np.delete(arr, inds, axis)
    array_to_be_merged = np.take(arr, inds, axis)
    if funkwargs.get('weights', None) is not None:
        # we need to reduce weights just like arr
        funkwarqs['weights'] = np.take(funkwarqs['weights'], inds, axis=axis
    merged_arr = fun(array_to_be_merged, axis, keepdims=True, **funkwargs)
    return insert_axis(new_arr, merged_arr, [np.minimum(inds[0], new_arr.sha
    # return np.insert(new arr, [np.minimum(inds[0], new arr.shape[axis])],
def merge nD(inds, arr, fun=np.nansum, weights=None, **funkwargs):
    """This function will merge a subarray of the input array arr,
       as defined by the input indices inds,
       along all the axes of arr (assuming same dimensionality along all axe
       applying the function fun, in order to summarize the values."""
    new_arr = arr.copy()
    for ax in range(arr.ndim):
        if weights is not None:
            new_arr = merge_axis(inds, new_arr, axis=ax, fun=fun, weights=we
            # we need to reduce weights just like arr
            weights = merge_axis(inds, weights, axis=ax, fun=np.nansum)
        else:
            new_arr = merge_axis(inds, new_arr, axis=ax, fun=fun, **funkwarg
    return new arr
def euclidean distance(p1, p2, mask=None, axis=1):
    return np.sqrt(np.sum(np.square(p1 - p2), axis=axis))
```

```
def compute_euclidean_tract_lengths(centres, weights):
    N = centres.shape[0]
    tl = np.zeros((N, N))
    for iR1 in range(N-1):
        for iR2 in range(iR1+1, N):
            if weights[iR1, iR2] > 0.0:
                tl[iR1, iR2] = euclidean_distance(centres[iR1][np.newaxis],
            else:
                tl[iR1, iR2] = 0.0
            tl[iR2, iR1] = tl[iR1, iR2]
    return tl
def merge conn(conn, regions, new label, voxel count,
               weight_fun=np.nansum, configure=False):
    """This function will merge an input TVB connectivity conn,
       for the input regions (indices or labels),
       substituting them with a summarized region of label new_label,
       applying the summary function for the connectivity weights weight_fun
       If configure is True, the new connectivity will also be configured.""
    if np.issubdtype(regions.dtype, np.integer):
        inds = regions
    else:
        inds = []
        for iR, region in conn.region labels:
            if region in regions:
                inds.append(iR)
    new conn = deepcopy(conn)
    repeat_fun = lambda arr, axis=0, **kwargs: repeat(arr, new_label, axis,
    new_conn.region_labels = merge_axis(inds, conn.region_labels, axis=0, fu
    new conn.weights = merge nD(inds, conn.weights, fun=weight fun)
    if FORCE_MERGED_ZERO_DIAGONAL:
        np.fill diagonal(new conn.weights, 0.0)
    # If WEIGHTED_AVERAGE_CENTRES...
    if WEIGHTED AVERAGE CENTRES:
        # ...compute merged regions' centres as weighted averages of the con
        # weighted by the relative volume (i.e., voxel count) of each region
        new_conn.centres = merge_axis(inds, conn.centres, axis=0, fun=weight
    else: # ...otherwise...
        # ...approximate merged regions' centres as simple averages of the d
        new_conn.centres = merge_axis(inds, conn.centres, axis=0, fun=np.nar
    if TRACT LENGTHS MODE == "euclidean":
        # Compute tract lengths as euclidean distances from the newly formed
        new_conn tract_lengths = compute_euclidean_tract_lengths(new_conn.ce
    else:
        # Compute tract lengths as weighted averages of the tract lengths
        # of all connections referring to/from a merged region,
        # weighted by corresponding tracts' weights:
        new_conn.tract_lengths = merge_nD(inds, conn.tract_lengths, fun=weig
    if FORCE MERGED ZERO DIAGONAL:
        np.fill_diagonal(new_conn.tract_lengths, 0.0)
    new_conn.tract_lengths[new_conn.weights == 0.0] = 0.0
    if configure:
        new_conn.configure()
    return new conn
def merge_major_structure(conn, major_struct_to_merge, major_structs_labels,
                          exclude_regions=[], weight_fun=np.nansum, configur
    """This function will merge an input TVB connectivity conn,
       for the input major structure label major_struct_to_merge,
```

substituting merged regions with a summarized region of the major str

```
and applying the summary function for the connectivity weights weight
               If configure is True, the new connectivity will also be configured.""
            regions_inds = np.where([major_structs_label == major_struct_to_merge ar
                                      region not in exclude regions
                                      for region, major_structs_label in zip(conn.reg
            print("...%d regions' indices of %s:\n%s" % (len(regions_inds), major_st
            repeat fun = lambda arr, axis=0, **kwargs: repeat(arr, major struct to m
            return merge_conn(conn, regions_inds, major_struct_to_merge, voxel_count
                              weight_fun=weight_fun, configure=configure), \
                   merge_axis(regions_inds, major_structs_labels, axis=0, fun=repeat
                   merge axis(regions inds, voxel count, axis=0, fun=voxel count sum
        def merge_major_structures(conn, major_structs_to_merge, major_structs_label
                                   exclude_regions={}, weight_fun=np.nansum):
            """This function will merge an input TVB connectivity conn,
               for the input major structures labels major_structs_to_merge,
               assuming an input vector major structs labels, mapping all regions to
               substituting merged regions with a summarized region of the respectiv
               and applying the summary function for the connectivity weights weight
               If configure is True, the new connectivity will also be configured.""
            new_conn = deepcopy(conn)
            new_major_structs_labels = major_structs_labels.copy()
            new voxel count = voxel count.copy()
            for major_struct_to_merge in major_structs_to_merge:
                print("Merging %s ..." % major_struct_to_merge)
                new_conn, new_major_structs_labels, new_voxel_count = \
                    merge_major_structure(new_conn, major_struct_to_merge,
                                           new major structs labels, new voxel count,
                                           exclude_regions=exclude_regions.get(major_
                                           weight fun=weight fun)
            new conn.configure()
            return new_conn, new_major_structs_labels, new_voxel_count
In [6]: def unilateral_to_bilateral_exclude_regions(major_structure, hemi, exclude_r
            new_exclude_regions = []
            if len(exclude regions):
                if hemi == "Left":
                    hemistr = "Left "
                    hemidel = "Right "
                else:
                    hemistr = "Right "
                    hemidel = "Left "
                for reg in exclude regions:
                    if hemistr in reg:
                        new_exclude_regions.append(reg)
                    elif hemidel in reg:
                        pass
                    else:
                        new_exclude_regions.append(hemistr + reg)
            return new_exclude_regions
        def unilateral_to_bilateral(major_structures_to_merge, exclude_regions):
            # For bilateral merge:
            new_major_structures_to_merge = []
            new_exclude_regions = {}
```

```
for mstm in major_structures_to_merge:
    if "Left " not in mstm and "Right " not in mstm:
        left = "Left " + mstm
        new_major_structures_to_merge.append("Left " + mstm)
        new major structures to merge.append("Right " + mstm)
    else:
        new_major_structures_to_merge.append(mstm)
new_major_structures_to_merge = np.unique(new_major_structures_to_merge)
for hemimstm in new major structures to merge:
    if "Left " in hemimstm:
        hemi = "Left"
        mstm = hemimstm.split("Left ")[-1]
    else:
        hemi = "Right"
        mstm = hemimstm.split("Right ")[-1]
    new exclude regions[hemimstm] = \
        exclude regions.get(hemimstm,
                            unilateral_to_bilateral_exclude_regions(mstm
return new_major_structures_to_merge, new_exclude_regions
```

Determine regions and structures to be merged, uni- and/or bi-laterally

```
In [7]: # Summarize all subcortical major structures except for the Cerebellar Corti
        # for _SummedSubcortical connectomes.
        # Specific Thalami have to be merged and then distributed to the Isocortex n
        major structures to merge = ['Thalamus',
                                        'Cortical Subplate', 'Hippocampal Formation',
                                        'Medulla', 'Spinal nucleus of the trigeminal',
                                        'Midbrain', 'Olfactory Areas', 'Pallidum',
                                        'Pons Motor', 'Pons Sensory', 'Pons Behavioral'
'Striatum', 'Cerebellar Cortex', 'Cerebellar Nu
        exclude_regions = {'Cerebellar Cortex': ["Ansiform lobule"],
                             # !!! Final decision to merge all Cerebellar Nuclei toge
                             # 'Cerebellar Nuclei': ["Interposed nucleus"]
                             'Medulla': ["Inferior olivary complex", "Facial motor nuc
                             'Pons Sensory': ["Principal sensory nucleus of the trigen
                             'Midbrain': ["Superior colliculus, motor related"]}
        # For bilateral merging:
        major structures to merge, exclude regions = unilateral to bilateral(major s
        print('major_structures_to_merge:\n', major_structures_to_merge)
        print('\nexclude_regions from mergin:\n', exclude_regions)
```

```
major_structures_to_merge:
['Left Cerebellar Cortex' 'Left Cerebellar Nuclei'
'Left Cortical Subplate' 'Left Hippocampal Formation' 'Left Hypothalamus'
'Left Medulla' 'Left Midbrain' 'Left Nonspecific Thalamus'
'Left Olfactory Areas' 'Left Pallidum' 'Left Pons Behavioral'
'Left Pons Motor' 'Left Pons Sensory'
'Left Spinal nucleus of the trigeminal' 'Left Striatum' 'Left Thalamus'
'Right Cerebellar Cortex' 'Right Cerebellar Nuclei'
'Right Cortical Subplate' 'Right Hippocampal Formation'
'Right Hypothalamus' 'Right Medulla' 'Right Midbrain'
'Right Nonspecific Thalamus' 'Right Olfactory Areas' 'Right Pallidum'
'Right Pons Behavioral' 'Right Pons Motor' 'Right Pons Sensory'
'Right Spinal nucleus of the trigeminal' 'Right Striatum'
'Right Thalamus']
```

exclude regions from mergin:

{'Left Cerebellar Cortex': ['Left Ansiform lobule'], 'Left Cerebellar Nucl ei': [], 'Left Cortical Subplate': [], 'Left Hippocampal Formation': [], 'L eft Hypothalamus': [], 'Left Medulla': ['Left Inferior olivary complex', 'L eft Facial motor nucleus'], 'Left Midbrain': ['Left Superior colliculus, mo tor related'], 'Left Nonspecific Thalamus': [], 'Left Olfactory Areas': [], 'Left Pallidum': [], 'Left Pons Behavioral': [], 'Left Pons Motor': [], 'Left Pons M ft Pons Sensory': ['Left Principal sensory nucleus of the trigeminal'], 'Le ft Spinal nucleus of the trigeminal': [], 'Left Striatum': [], 'Left Thalam us': [], 'Right Cerebellar Cortex': ['Right Ansiform lobule'], 'Right Cereb ellar Nuclei': [], 'Right Cortical Subplate': [], 'Right Hippocampal Format ion': [], 'Right Hypothalamus': [], 'Right Medulla': ['Right Inferior oliva ry complex', 'Right Facial motor nucleus'], 'Right Midbrain': ['Right Super ior colliculus, motor related'], 'Right Nonspecific Thalamus': [], 'Right O lfactory Areas': [], 'Right Pallidum': [], 'Right Pons Behavioral': [], 'Ri ght Pons Motor': [], 'Right Pons Sensory': ['Right Principal sensory nucleu s of the trigeminal'], 'Right Spinal nucleus of the trigeminal': [], 'Right Striatum': [], 'Right Thalamus': []}

Execute merging

```
Merging Left Cerebellar Cortex ...
...13 regions' indices of Left Cerebellar Cortex:
[578 579 580 581 582 583 584 585 586 588 589 590 591]
Voxel count sum: 33696
Merging Left Cerebellar Nuclei ...
...4 regions' indices of Left Cerebellar Nuclei:
[580 581 582 583]
Voxel count sum: 1116
Merging Left Cortical Subplate ...
...7 regions' indices of Left Cortical Subplate:
[365 366 367 368 369 370 371]
Voxel count sum: 6669
Merging Left Hippocampal Formation ...
...13 regions' indices of Left Hippocampal Formation:
[352 353 354 355 356 357 358 359 360 361 362 363 364]
Voxel count sum: 34819
Merging Left Hypothalamus ...
...41 regions' indices of Left Hypothalamus:
[419 420 421 422 423 424 425 426 427 428 429 430 431 432 433 434 435 436
 437 438 439 440 441 442 443 444 445 446 447 448 449 450 451 452 453 454
 455 456 457 458 4591
Voxel count sum: 13871
Merging Left Medulla ...
...34 regions' indices of Left Medulla:
[481 482 483 484 485 486 490 491 493 494 495 497 498 499 500 501 502 503
 504 505 506 507 508 509 510 511 512 513 514 515 516 517 518 5191
Voxel count sum: 17303
Merging Left Midbrain ...
...38 regions' indices of Left Midbrain:
[420 421 422 423 424 425 426 427 428 429 430 431 433 434 435 436 437 438
 439 440 441 442 443 444 445 446 447 448 449 450 451 452 453 454 455 456
 457 4581
Voxel count sum: 19804
Merging Left Nonspecific Thalamus ...
...9 regions' indices of Left Nonspecific Thalamus:
[403 404 406 407 408 409 410 411 412]
Voxel count sum: 2324
Merging Left Olfactory Areas ...
...11 regions' indices of Left Olfactory Areas:
[341 342 343 344 345 346 347 348 349 350 351]
Voxel count sum: 41043
Merging Left Pallidum ...
...9 regions' indices of Left Pallidum:
[357 358 359 360 361 362 363 364 365]
Voxel count sum: 9308
Merging Left Pons Behavioral ...
...8 regions' indices of Left Pons Behavioral:
[410 411 412 413 414 415 416 417]
Voxel count sum: 3890
Merging Left Pons Motor ...
...11 regions' indices of Left Pons Motor:
[399 400 401 402 403 404 405 406 407 408 409]
Voxel count sum: 5971
Merging Left Pons Sensory ...
... 2 regions' indices of Left Pons Sensory:
[396 398]
Voxel count sum: 1585
Merging Left Spinal nucleus of the trigeminal ...
...3 regions' indices of Left Spinal nucleus of the trigeminal:
[401 402 403]
Voxel count sum: 3107
```

```
Merging Left Striatum ...
...13 regions' indices of Left Striatum:
[344 345 346 347 348 349 350 351 352 353 354 355 356]
Voxel count sum: 40219
Merging Left Thalamus ...
...34 regions' indices of Left Thalamus:
[346 347 348 349 350 351 352 353 354 355 356 357 358 359 360 361 362 363
 364 365 366 367 368 369 370 371 372 374 375 376 377 378 379 380]
Voxel count sum: 13346
Merging Right Cerebellar Cortex ...
...13 regions' indices of Right Cerebellar Cortex:
[280 281 282 283 284 285 286 287 288 290 291 292 293]
Voxel count sum: 33696
Merging Right Cerebellar Nuclei ...
...4 regions' indices of Right Cerebellar Nuclei:
[282 283 284 285]
Voxel count sum: 1116
Merging Right Cortical Subplate ...
...7 regions' indices of Right Cortical Subplate:
[67 68 69 70 71 72 73]
Voxel count sum: 6669
Merging Right Hippocampal Formation ...
...13 regions' indices of Right Hippocampal Formation:
[54 55 56 57 58 59 60 61 62 63 64 65 66]
Voxel count sum: 34819
Merging Right Hypothalamus ...
...41 regions' indices of Right Hypothalamus:
[121 122 123 124 125 126 127 128 129 130 131 132 133 134 135 136 137 138
 139 140 141 142 143 144 145 146 147 148 149 150 151 152 153 154 155 156
 157 158 159 160 161]
Voxel count sum: 13871
Merging Right Medulla ...
...34 regions' indices of Right Medulla:
[183 184 185 186 187 188 192 193 195 196 197 199 200 201 202 203 204 205
 206 207 208 209 210 211 212 213 214 215 216 217 218 219 220 221]
Voxel count sum: 17303
Merging Right Midbrain ...
...38 regions' indices of Right Midbrain:
[122 123 124 125 126 127 128 129 130 131 132 133 135 136 137 138 139 140
 141 142 143 144 145 146 147 148 149 150 151 152 153 154 155 156 157 158
 159 1601
Voxel count sum: 19804
Merging Right Nonspecific Thalamus ...
...9 regions' indices of Right Nonspecific Thalamus:
[105 106 108 109 110 111 112 113 114]
Voxel count sum: 2324
Merging Right Olfactory Areas ...
...11 regions' indices of Right Olfactory Areas:
[43 44 45 46 47 48 49 50 51 52 53]
/tmp/ipykernel_222/2251895505.py:15: RuntimeWarning: invalid value encounte
red in true divide
  wav = np.nansum(arr * weights, axis=axis, **kwargs) / np.nansum(weights,
axis=axis, **kwarqs)
```

```
Voxel count sum: 41043
Merging Right Pallidum ...
...9 regions' indices of Right Pallidum:
[59 60 61 62 63 64 65 66 67]
Voxel count sum: 9308
Merging Right Pons Behavioral ...
...8 regions' indices of Right Pons Behavioral:
[112 113 114 115 116 117 118 119]
Voxel count sum: 3890
Meraina Right Pons Motor ...
...11 regions' indices of Right Pons Motor:
[101 102 103 104 105 106 107 108 109 110 111]
Voxel count sum: 5971
Merging Right Pons Sensory ...
...2 regions' indices of Right Pons Sensory:
[ 98 100]
Voxel count sum: 1585
Merging Right Spinal nucleus of the trigeminal ...
...3 regions' indices of Right Spinal nucleus of the trigeminal:
[103 104 105]
Voxel count sum: 3107
Merging Right Striatum ...
...13 regions' indices of Right Striatum:
[46 47 48 49 50 51 52 53 54 55 56 57 58]
Voxel count sum: 40219
Merging Right Thalamus ...
...34 regions' indices of Right Thalamus:
[48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71
72 73 74 76 77 78 79 80 81 82]
Voxel count sum: 13346
Merged major structures and their voxel counts:
 [['Right Frontal pole, cerebral cortex' 'Right Isocortex' '514']
 ['Right Primary motor area' 'Right Isocortex' '11760']
 ['Right Secondary motor area' 'Right Isocortex' '10098']
 ['Right Primary somatosensory area, nose' 'Right Isocortex' '1358']
 ['Right Primary somatosensory area, barrel field' 'Right Isocortex'
  '10306']
 ['Right Primary somatosensory area, lower limb' 'Right Isocortex' '3254']
 ['Right Primary somatosensory area, mouth' 'Right Isocortex' '2924']
 ['Right Primary somatosensory area, upper limb' 'Right Isocortex' '5406']
 ['Right Primary somatosensory area, trunk' 'Right Isocortex' '4799']
 ['Right Primary somatosensory area, unassigned' 'Right Isocortex' '2958']
 ['Right Supplemental somatosensory area' 'Right Isocortex' '5729']
 ['Right Gustatory areas' 'Right Isocortex' '2104']
 ['Right Visceral area' 'Right Isocortex' '1793']
 ['Right Dorsal auditory area' 'Right Isocortex' '2556']
 ['Right Primary auditory area' 'Right Isocortex' '2689']
 ['Right Posterior auditory area' 'Right Isocortex' '191']
 ['Right Ventral auditory area' 'Right Isocortex' '2554']
 ['Right Anterolateral visual area' 'Right Isocortex' '1981']
 ['Right Anteromedial visual area' 'Right Isocortex' '1795']
 ['Right Lateral visual area' 'Right Isocortex' '880']
 ['Right Primary visual area' 'Right Isocortex' '6227']
 ['Right Posterolateral visual area' 'Right Isocortex' '679']
 ['Right posteromedial visual area' 'Right Isocortex' '1265']
 ['Right Laterointermediate area' 'Right Isocortex' '-1']
 ['Right Postrhinal area' 'Right Isocortex' '-1']
 ['Right Anterior cingulate area, dorsal part' 'Right Isocortex' '2789']
 ['Right Anterior cingulate area, ventral part' 'Right Isocortex' '2418']
 ['Right Prelimbic area' 'Right Isocortex' '1790']
```

```
['Right Infralimbic area' 'Right Isocortex' '1764']
['Right Orbital area, lateral part' 'Right Isocortex' '1853']
['Right Orbital area, medial part' 'Right Isocortex' '1024']
['Right Orbital area, ventrolateral part' 'Right Isocortex' '1556']
['Right Agranular insular area, dorsal part' 'Right Isocortex' '2364']
['Right Agranular insular area, posterior part' 'Right Isocortex' '1348']
['Right Agranular insular area, ventral part' 'Right Isocortex' '1120']
['Right Retrosplenial area, lateral agranular part' 'Right Isocortex'
'937'1
['Right Retrosplenial area, dorsal part' 'Right Isocortex' '2738']
['Right Retrosplenial area, ventral part' 'Right Isocortex' '3997']
['Right Anterior area' 'Right Isocortex' '-1']
['Right Rostrolateral visual area' 'Right Isocortex' '-1']
['Right Temporal association areas' 'Right Isocortex' '3773']
['Right Perirhinal area' 'Right Isocortex' '1412']
['Right Ectorhinal area' 'Right Isocortex' '3390']
['Right Olfactory Areas' 'Right Olfactory Areas' '41043']
['Right Hippocampal Formation' 'Right Hippocampal Formation' '34819']
['Right Cortical Subplate' 'Right Cortical Subplate' '6669']
['Right Striatum' 'Right Striatum' '40219']
['Right Pallidum' 'Right Pallidum' '9308']
['Right Thalamus' 'Right Thalamus' '13346']
['Right Nonspecific Thalamus' 'Right Nonspecific Thalamus' '2324']
['Right Hypothalamus' 'Right Hypothalamus' '13871']
['Right Midbrain' 'Right Midbrain' '19804']
['Right Superior colliculus, motor related' 'Right Midbrain' '6779']
['Right Pons Sensory' 'Right Pons Sensory' '1585']
['Right Principal sensory nucleus of the trigeminal'
'Right Pons Sensory' '1093']
['Right Pons Motor' 'Right Pons Motor' '5971']
['Right Pons Behavioral' 'Right Pons Behavioral' '3890']
['Right Medulla' 'Right Medulla' '17303']
['Right Spinal nucleus of the trigeminal'
'Right Spinal nucleus of the trigeminal' '3107']
['Right Facial motor nucleus' 'Right Medulla' '660']
['Right Inferior olivary complex' 'Right Medulla' '567']
['Right Cerebellar Cortex' 'Right Cerebellar Cortex' '33696']
['Right Ansiform lobule' 'Right Cerebellar Cortex' '9044']
['Right Cerebellar Nuclei' 'Right Cerebellar Nuclei' '1116']
['Left Frontal pole, cerebral cortex' 'Left Isocortex' '514']
['Left Primary motor area' 'Left Isocortex' '11760']
['Left Secondary motor area' 'Left Isocortex' '10098']
['Left Primary somatosensory area, nose' 'Left Isocortex' '1358']
['Left Primary somatosensory area, barrel field' 'Left Isocortex'
'10306']
['Left Primary somatosensory area, lower limb' 'Left Isocortex' '3254']
['Left Primary somatosensory area, mouth' 'Left Isocortex' '2924']
['Left Primary somatosensory area, upper limb' 'Left Isocortex' '5406']
['Left Primary somatosensory area, trunk' 'Left Isocortex' '4799']
['Left Primary somatosensory area, unassigned' 'Left Isocortex' '2958']
['Left Supplemental somatosensory area' 'Left Isocortex' '5729']
['Left Gustatory areas' 'Left Isocortex' '2104']
['Left Visceral area' 'Left Isocortex' '1793']
['Left Dorsal auditory area' 'Left Isocortex' '2556']
['Left Primary auditory area' 'Left Isocortex' '2689']
['Left Posterior auditory area' 'Left Isocortex' '191']
['Left Ventral auditory area' 'Left Isocortex' '2554']
['Left Anterolateral visual area' 'Left Isocortex' '1981']
['Left Anteromedial visual area' 'Left Isocortex' '1795']
['Left Lateral visual area' 'Left Isocortex' '880']
['Left Primary visual area' 'Left Isocortex' '6227']
```

```
['Left Posterolateral visual area' 'Left Isocortex' '679']
['Left posteromedial visual area' 'Left Isocortex' '1265']
['Left Laterointermediate area' 'Left Isocortex' '-1']
['Left Postrhinal area' 'Left Isocortex' '-1']
['Left Anterior cingulate area, dorsal part' 'Left Isocortex' '2789']
['Left Anterior cingulate area, ventral part' 'Left Isocortex' '2418']
['Left Prelimbic area' 'Left Isocortex' '1790']
['Left Infralimbic area' 'Left Isocortex' '1764']
['Left Orbital area, lateral part' 'Left Isocortex' '1853']
['Left Orbital area, medial part' 'Left Isocortex' '1024']
['Left Orbital area, ventrolateral part' 'Left Isocortex' '1556']
['Left Agranular insular area, dorsal part' 'Left Isocortex' '2364']
['Left Agranular insular area, posterior part' 'Left Isocortex' '1348']
['Left Agranular insular area, ventral part' 'Left Isocortex' '1120']
['Left Retrosplenial area, lateral agranular part' 'Left Isocortex'
'937'1
['Left Retrosplenial area, dorsal part' 'Left Isocortex' '2738']
['Left Retrosplenial area, ventral part' 'Left Isocortex' '3997']
['Left Anterior area' 'Left Isocortex' '-1']
['Left Rostrolateral visual area' 'Left Isocortex' '-1']
['Left Temporal association areas' 'Left Isocortex' '3773']
['Left Perirhinal area' 'Left Isocortex' '1412']
['Left Ectorhinal area' 'Left Isocortex' '3390']
['Left Olfactory Areas' 'Left Olfactory Areas' '41043']
['Left Hippocampal Formation' 'Left Hippocampal Formation' '34819']
['Left Cortical Subplate' 'Left Cortical Subplate' '6669']
['Left Striatum' 'Left Striatum' '40219']
['Left Pallidum' 'Left Pallidum' '9308']
['Left Thalamus' 'Left Thalamus' '13346']
['Left Nonspecific Thalamus' 'Left Nonspecific Thalamus' '2324']
['Left Hypothalamus' 'Left Hypothalamus' '13871']
['Left Midbrain' 'Left Midbrain' '19804']
['Left Superior colliculus, motor related' 'Left Midbrain' '6779']
['Left Pons Sensory' 'Left Pons Sensory' '1585']
['Left Principal sensory nucleus of the trigeminal' 'Left Pons Sensory'
'1093'l
['Left Pons Motor' 'Left Pons Motor' '5971']
['Left Pons Behavioral' 'Left Pons Behavioral' '3890']
['Left Medulla' 'Left Medulla' '17303']
['Left Spinal nucleus of the trigeminal'
 'Left Spinal nucleus of the trigeminal' '3107']
['Left Facial motor nucleus' 'Left Medulla' '660']
['Left Inferior olivary complex' 'Left Medulla' '567']
['Left Cerebellar Cortex' 'Left Cerebellar Cortex' '33696']
['Left Ansiform lobule' 'Left Cerebellar Cortex' '9044']
['Left Cerebellar Nuclei' 'Left Cerebellar Nuclei' '1116']]
```

Testing merging

```
In [9]: # Testing that everything went well:
    print("Testing merging, voxel counts, sum of (self/from/to) weights, and sel
    for iR, (reg, msl, vc) in enumerate(zip(new_conn.region_labels, new_major_st
        print("\n%s - %s:" % (reg, msl))
        if reg in result_connectivity.region_labels:
            iR2 = np.where([reg == reg2 for reg2 in result_connectivity.region_l
            print("...%d = %d voxels" % (vc, voxel_count[iR2]))
        assert voxel_count[iR2] - vc == 0
        print("...centres: %s = %s" % (new_conn.centres[iR], result_connectivity.centres
```

```
ws = np.nansum(new conn.weights[iR])
    ws2 = np.nansum(result connectivity.weights[iR2])
    print("...weights to: %g = %g" % (ws, ws2))
    assert np.abs(ws - ws2) < 1e-6
    ws = np.nansum(new conn.weights[:, iR])
    ws2 = np.nansum(result connectivity.weights[:, iR2])
    print("...weights from: %g = %g" % (ws, ws2))
    assert np.abs(ws - ws2) < 1e-6
    print("...self tract lengths: %g = %g" % (new conn.tract lengths[iR,
    assert np.abs(new conn.tract lengths[iR, iR] - result connectivity.t
else:
    excl_regs = exclude_regions.get(reg, [])
    iR2s = np.where([msl == msl2 and reg2 not in excl regs
                     for reg2, msl2 in zip(result connectivity.region la
    old vc = voxel_count[iR2s]
    old_vc = old_vc[old_vc >= 0]
    old vc sum = old vc.sum()
    print("...%d voxels" % old_vc_sum)
    assert old_vc_sum - vc == 0
    if WEIGHTED AVERAGE CENTRES:
        assert np.allclose(new_conn.centres[iR],
                           np.average(result connectivity.centres[iR2s,
                                      axis=0, weights=voxel_count[iR2s])
    else:
        oldcntrs = np.nanmean(result connectivity.centres[iR2s], axis=0)
        print("...centres: %s = %s" % (new_conn.centres[iR], oldcntrs))
        assert np.allclose(new conn.centres[iR], oldcntrs)
    ws = np.nansum(new conn.weights[iR])
    ws2 = np.nansum(result_connectivity.weights[iR2s])
    print("...weights to: %g = %g" % (ws, ws2))
    if not FORCE MERGED ZERO DIAGONAL:
        assert np.abs(ws - ws2) < 1e-6
    ws = np.nansum(new conn.weights[:, iR])
    ws2 = np.nansum(result_connectivity.weights[:, iR2s])
    print("...weights from: %g = %g" % (ws, ws2))
    if not FORCE MERGED ZERO DIAGONAL:
        assert np.abs(ws - ws2) < 1e-6
    ws2 = np.nansum(result_connectivity.weights[iR2s][:, iR2s])
    print("...self weights: %g = %g" % (new_conn.weights[iR, iR], ws2))
    # assert np.abs(new_conn.weights[iR, iR] - ws2) < 1e-6
    if TRACT_LENGTHS_MODE == "euclidean":
        tl = new conn.tract lengths[iR, :]
        weights = new_conn.weights[iR, :] > 0.0
        tl2 = weights*euclidean distance(new conn.centres[iR, :], new cd
        print("...average tract_lengths to/from: %g = %g" % (tl.mean(),
        assert np.allclose(tl, tl2)
    # TODO: Think of a test for weighhed average!!!
```

Testing merging, voxel counts, sum of (self/from/to) weights, and self-trac t lengths: Right Frontal pole, cerebral cortex - Right Isocortex: ...514 = 514 voxels...centres: [45.5942029 22.72256729 51.2484472] = [[45.5942029 22.722567 29 51.2484472]] ...weights to: 3.14868 = 3.14868...weights from: 6.36079 = 6.36079...self tract lengths: 0 = 0 Right Primary motor area - Right Isocortex: ...11760 = 11760 voxels ...centres: [35.30080448 42.96502274 55.40433718] = [[35.30080448 42.965022 74 55,40433718]] ...weights to: 1.87382 = 1.87382...weights from: 6.37274 = 6.37274...self tract_lengths: 0 = 0 Right Secondary motor area - Right Isocortex: 1.10098 = 10098 voxels...centres: [43.65873016 36.05326618 57.91804029] = [[43.65873016 36.053266 18 57.91804029]] ...weights to: 2.33534 = 2.33534...weights from: 6.99391 = 6.99391...self tract lengths: 0 = 0 Right Primary somatosensory area, nose - Right Isocortex: ...1358 = 1358 voxels ...centres: [21.02302632 56.74407895 54.42039474] = [[21.02302632 56.744078 95 54.4203947411 ...weights to: 1.20136 = 1.20136...weights from: 5.39897 = 5.39897...self tract_lengths: 0 = 0 Right Primary somatosensory area, barrel field - Right Isocortex: 1.10306 = 10306 voxels...centres: [23.73700988 67.39846988 61.48453937] = [[23.73700988 67.398469]]88 61,4845393711 ...weights to: 1.37367 = 1.37367...weights from: 6.26508 = 6.26508...self tract lengths: 0 = 0 Right Primary somatosensory area, lower limb - Right Isocortex: ...3254 = 3254 voxels...centres: [39.3573854 59.67402377 66.73344652] = [[39.3573854 59.674023 77 66.73344652]] ...weights to: 1.83371 = 1.83371...weights from: 5.63341 = 5.63341...self tract_lengths: 0 = 0 Right Primary somatosensory area, mouth - Right Isocortex: ...2924 = 2924 voxels...centres: [22.99775281 46.18266453 48.86420546] = [[22.99775281 46.182664 53 48.86420546]] ...weights to: 1.3537 = 1.3537...weights from: 5.09649 = 5.09649...self tract_lengths: 0 = 0 Right Primary somatosensory area, upper limb - Right Isocortex:

...5406 = 5406 voxels

```
...centres: [32.6544196 54.61182109 61.69755059] = [[32.6544196 54.611821
09 61.69755059]]
...weights to: 1.39079 = 1.39079
...weights from: 2.70357 = 2.70357
...self tract lengths: 0 = 0
Right Primary somatosensory area, trunk - Right Isocortex:
...4799 = 4799 \text{ voxels}
...centres: [38.95590327 66.81507824 69.51920341] = [[38.95590327 66.815078]
24 69.51920341]]
...weights to: 2.32431 = 2.32431
...weights from: 4.8304 = 4.8304
...self tract lengths: 0 = 0
Right Primary somatosensory area, unassigned - Right Isocortex:
...2958 = 2958 \text{ voxels}
...centres: [29.06161137 58.57661927 61.72827804] = [[29.06161137 58.576619
27 61.72827804]]
...weights to: 1.71146 = 1.71146
...weights from: 3.56411 = 3.56411
...self tract_lengths: 0 = 0
Right Supplemental somatosensory area - Right Isocortex:
...5729 = 5729 \text{ voxels}
...centres: [15.35346359 61.09413854 45.52775311] = [[15.35346359 61.094138
54 45.52775311]]
...weights to: 2.45012 = 2.45012
...weights from: 7.48341 = 7.48341
...self tract_lengths: 0 = 0
Right Gustatory areas - Right Isocortex:
...2104 = 2104 \text{ voxels}
...centres: [19.56851642 46.67610419 34.30124575] = [[19.56851642 46.676104
19 34.30124575]]
...weights to: 3.59145 = 3.59145
...weights from: 8.43749 = 8.43749
...self tract lengths: 0 = 0
Right Visceral area - Right Isocortex:
...1793 = 1793 voxels
...centres: [12.67567568 62.97043919 34.46706081] = [[12.67567568 62.970439
19 34,46706081]]
...weights to: 4.12891 = 4.12891
...weights from: 9.98387 = 9.98387
...self tract_lengths: 0 = 0
Right Dorsal auditory area - Right Isocortex:
...2556 = 2556 \text{ voxels}
...centres: [14.68144499 75.45812808 54.78981938] = [[14.68144499 75.458128
08 54.78981938]]
...weights to: 3.36174 = 3.36174
...weights from: 4.9361 = 4.9361
...self tract lengths: 0 = 0
Right Primary auditory area - Right Isocortex:
...2689 = 2689 voxels
...centres: [11.802595
                         79.33920297 50.82205746] = [[11.802595
97 50.82205746]]
...weights to: 3.70985 = 3.70985
...weights from: 3.86101 = 3.86101
...self tract_lengths: 0 = 0
```

```
Right Posterior auditory area - Right Isocortex:
...191 = 191 \text{ voxels}
...centres: [14.58333333 84.17 56.06666667] = [[14.58333333 84.17
56.0666666711
...weights to: 4.81068 = 4.81068
...weights from: 4.55941 = 4.55941
...self tract_lengths: 0 = 0
Right Ventral auditory area - Right Isocortex:
...2554 = 2554 \text{ voxels}
...centres: [10.50774336 78.00110619 45.04535398] = [[10.50774336 78.001106
19 45.04535398]]
...weights to: 3.64962 = 3.64962
...weights from: 7.59361 = 7.59361
...self tract lengths: 0 = 0
Right Anterolateral visual area - Right Isocortex:
...1981 = 1981 \text{ voxels}
...centres: [19.92327366 82.52173913 62.23529412] = [[19.92327366 82.521739
13 62.23529412]]
...weights to: 3.36884 = 3.36884
...weights from: 6.11907 = 6.11907
...self tract_lengths: 0 = 0
Right Anteromedial visual area - Right Isocortex:
...1795 = 1795 \text{ voxels}
...centres: [40.48704663 76.65025907 71.68911917] = [[40.48704663 76.650259
07 71.68911917]]
...weights to: 3.6538 = 3.6538
...weights from: 7.05138 = 7.05138
...self tract_lengths: 0 = 0
Right Lateral visual area - Right Isocortex:
...880 = 880 voxels
...centres: [21.07281553 91.04530744 60.5566343 ] = [[21.07281553 91.045307
44 60.5566343 11
...weights to: 3.31325 = 3.31325
...weights from: 5.33257 = 5.33257
...self tract_lengths: 0 = 0
Right Primary visual area - Right Isocortex:
...6227 = 6227 \text{ voxels}
...centres: [31.24740533 90.72734923 66.94922861] = [[31.24740533 90.727349
23 66.94922861]]
...weights to: 2.40566 = 2.40566
...weights from: 4.65915 = 4.65915
...self tract_lengths: 0 = 0
Right Posterolateral visual area - Right Isocortex:
...679 = 679 \text{ voxels}
...centres: [ 24.43859649 100.0726817 56.95989975] = [[ 24.43859649 100.0
         56.9598997511
726817
...weights to: 2.98092 = 2.98092
...weights from: 5.24074 = 5.24074
...self tract_lengths: 0 = 0
Right posteromedial visual area - Right Isocortex:
...1265 = 1265 \text{ voxels}
...centres: [40.06730769 83.88076923 71.89615385] = [[40.06730769 83.880769
23 71.89615385]]
```

```
...weights to: 3.1971 = 3.1971
...weights from: 9.94985 = 9.94985
...self tract lengths: 0 = 0
Right Laterointermediate area - Right Isocortex:
\dots -1 = -1 voxels
...centres: [17.85185185 \ 90.53497942 \ 57.65432099] = [[17.85185185 \ 90.534979]
42 57.65432099]]
...weights to: 3.73671 = 3.73671
...weights from: 6.57665 = 6.57665
...self tract_lengths: 0 = 0
Right Postrhinal area - Right Isocortex:
\dots -1 = -1 voxels
...centres: [15.60658307 95.63793103 51.69435737] = [[15.60658307 95.637931
03 51.69435737]]
...weights to: 4.35521 = 4.35521
...weights from: 6.61262 = 6.61262
...self tract_lengths: 0 = 0
Right Anterior cingulate area, dorsal part - Right Isocortex:
...2789 = 2789 \text{ voxels}
...centres: [52.53279243 44.29952671 59.43002028] = [[52.53279243 44.299526
71 59.43002028]]
...weights to: 3.40323 = 3.40323
...weights from: 7.78532 = 7.78532
...self tract_lengths: 0 = 0
Right Anterior cingulate area, ventral part - Right Isocortex:
...2418 = 2418 \text{ voxels}
...centres: [53.40868794 48.55319149 54.30141844] = [[53.40868794 48.553191
49 54.3014184411
...weights to: 4.55297 = 4.55297
...weights from: 10.0158 = 10.0158
...self tract_lengths: 0 = 0
Right Prelimbic area - Right Isocortex:
...1790 = 1790 \text{ voxels}
...centres: [52.33045729 31.06039689 51.18205349] = [[52.33045729 31.060396
89 51.18205349]]
...weights to: 4.54218 = 4.54218
...weights from: 10.5647 = 10.5647
...self tract_lengths: 0 = 0
Right Infralimbic area - Right Isocortex:
...1764 = 1764 \text{ voxels}
...centres: [53.15632754 34.98759305 41.86848635] = [[53.15632754 34.987593
05 41.8684863511
...weights to: 7.0478 = 7.0478
...weights from: 7.95842 = 7.95842
...self tract_lengths: 0 = 0
Right Orbital area, lateral part - Right Isocortex:
...1853 = 1853 voxels
...centres: [41.75661765 29.64191176 40.67426471] = [[41.75661765 29.641911
76 40.67426471]]
...weights to: 4.16078 = 4.16078
...weights from: 6.72122 = 6.72122
...self tract_lengths: 0 = 0
```

Right Orbital area, medial part - Right Isocortex:

```
...1024 = 1024 \text{ voxels}
...centres: [53.2739726 28.0456621 43.21613394] = [[53.2739726 28.045662
1 43.21613394]]
...weights to: 5.11325 = 5.11325
...weights from: 7.6126 = 7.6126
...self tract lengths: 0 = 0
Right Orbital area, ventrolateral part - Right Isocortex:
...1556 = 1556 voxels
...centres: [49.09438202 27.48539326 42.88426966] = [[49.09438202 27.485393
26 42.88426966]]
...weights to: 4.6912 = 4.6912
...weights from: 7.00555 = 7.00555
...self tract lengths: 0 = 0
Right Agranular insular area, dorsal part - Right Isocortex:
...2364 = 2364 \text{ voxels}
...centres: [29.77396893 34.61810391 36.22656668] = [[29.77396893 34.618103
91 36.2265666811
...weights to: 4.4157 = 4.4157
...weights from: 11.3421 = 11.3421
...self tract lengths: 0 = 0
Right Agranular insular area, posterior part - Right Isocortex:
...1348 = 1348 \text{ voxels}
...centres: [15.16350124 59.63996697 27.96944674] = [[15.16350124 59.639966
97 27.96944674]]
...weights to: 5.1191 = 5.1191
...weights from: 8.72044 = 8.72044
...self tract_lengths: 0 = 0
Right Agranular insular area, ventral part - Right Isocortex:
...1120 = 1120 voxels
...centres: [29.87514318 37.21076747 30. ] = [[29.87514318 37.210767
47 30.
              ]]
...weights to: 5.56127 = 5.56127
...weights from: 5.79887 = 5.79887
...self tract_lengths: 0 = 0
Right Retrosplenial area, lateral agranular part - Right Isocortex:
...937 = 937 voxels
...centres: [44.0636833 81.13253012 70.83390706] = [[44.0636833 81.132530
12 70.8339070611
...weights to: 2.48287 = 2.48287
...weights from: 4.5897 = 4.5897
...self tract_lengths: 0 = 0
Right Retrosplenial area, dorsal part - Right Isocortex:
...2738 = 2738 \text{ voxels}
...centres: [48.19167983 80.32490785 70.98683518] = [[48.19167983 80.324907
85 70.98683518]]
...weights to: 1.95983 = 1.95983
...weights from: 4.54976 = 4.54976
...self tract_lengths: 0 = 0
Right Retrosplenial area, ventral part - Right Isocortex:
...3997 = 3997 voxels
...centres: [51.74832536 78.26985646 66.47607656] = [[51.74832536 78.269856
46 66,4760765611
...weights to: 2.52084 = 2.52084
...weights from: 7.47747 = 7.47747
```

```
...self tract_lengths: 0 = 0
Right Anterior area - Right Isocortex:
\dots -1 = -1 voxels
...centres: [36.72853186 72.52493075 70.37257618] = [[36.72853186 72.524930
75 70.37257618]]
...weights to: 2.75957 = 2.75957
...weights from: 5.86238 = 5.86238
...self tract lengths: 0 = 0
Right Rostrolateral visual area - Right Isocortex:
\dots -1 = -1 voxels
...centres: [26.21960784 77.9745098 66.57254902] = [[26.21960784 77.974509
8 66.57254902]]
...weights to: 2.27166 = 2.27166
...weights from: 6.42253 = 6.42253
...self tract_lengths: 0 = 0
Right Temporal association areas - Right Isocortex:
...3773 = 3773 voxels
...centres: [11.30600387 82.74112331 44.43124597] = [[11.30600387 82.741123
31 44.43124597]]
...weights to: 4.7417 = 4.7417
...weights from: 6.0477 = 6.0477
...self tract_lengths: 0 = 0
Right Perirhinal area - Right Isocortex:
...1412 = 1412 \text{ voxels}
...centres: [10.31155779 82.23366834 33.93467337] = [[10.31155779 82.233668
34 33.93467337]]
...weights to: 4.78609 = 4.78609
...weights from: 8.2565 = 8.2565
...self tract_lengths: 0 = 0
Right Ectorhinal area - Right Isocortex:
...3390 = 3390 \text{ voxels}
...centres: [11.0908046 80.43908046 36.10689655] = [[11.0908046 80.439080
46 36.10689655]]
...weights to: 5.28341 = 5.28341
...weights from: 7.69002 = 7.69002
...self tract_lengths: 0 = 0
Right Olfactory Areas - Right Olfactory Areas:
...41043 voxels
...weights to: 39.4701 = 48.8479
...weights from: 53.8398 = 63.2176
...self weights: 0 = 9.37775
Right Hippocampal Formation - Right Hippocampal Formation:
...34819 voxels
...weights to: 34.0178 = 43.4767
...weights from: 48.5659 = 58.0249
...self weights: 0 = 9.45896
Right Cortical Subplate - Right Cortical Subplate:
...6669 voxels
...weights to: 44.7184 = 50.8921
...weights from: 44.4496 = 50.6234
...self weights: 0 = 6.17378
Right Striatum - Right Striatum:
```

```
...40219 voxels
...weights to: 98.9072 = 109.003
...weights from: 88.2897 = 98.3856
...self weights: 0 = 10.0959
Right Pallidum - Right Pallidum:
...9308 voxels
...weights to: 84.9946 = 89.3808
...weights from: 75.5767 = 79.9628
...self weights: 0 = 4.38613
Right Thalamus - Right Thalamus:
...13346 voxels
...weights to: 280.134 = 326.04
...weights from: 186.091 = 231.998
...self weights: 0 = 45.906
Right Nonspecific Thalamus - Right Nonspecific Thalamus:
...2324 voxels
...weights to: 115.961 = 118.485
...weights from: 61.0322 = 63.5561
...self weights: 0 = 2.5239
Right Hypothalamus - Right Hypothalamus:
...13871 voxels
...weights to: 354.926 = 630.732
...weights from: 366.33 = 642.136
...self weights: 0 = 275.806
Right Midbrain - Right Midbrain:
...19804 voxels
...weights to: 338.668 = 429.195
...weights from: 286.757 = 377.284
...self weights: 0 = 90.5271
Right Superior colliculus, motor related - Right Midbrain:
...6779 = 6779 \text{ voxels}
...centres: [47.53567587 91.42129796 55.60666906] = [[47.53567587 91.421297
96 55,6066690611
...weights to: 9.62674 = 9.62674
...weights from: 7.67702 = 7.67702
...self tract lengths: 0 = 0
Right Pons Sensory - Right Pons Sensory:
...1585 voxels
...weights to: 13.202 = 13.3324
...weights from: 19.1642 = 19.2947
...self weights: 0 = 0.130488
Right Principal sensory nucleus of the trigeminal - Right Pons Sensory:
...1093 = 1093 voxels
...centres: [ 36.33636364 101.84
                                        22.50363636] = [[ 36.33636364 101.8
         22.5036363611
...weights to: 3.14049 = 3.14049
...weights from: 9.72171 = 9.72171
...self tract_lengths: 0 = 0
Right Pons Motor - Right Pons Motor:
...5971 voxels
...weights to: 102.516 = 111.275
...weights from: 99.5803 = 108.339
```

```
...self weights: 0 = 8.75894
Right Pons Behavioral - Right Pons Behavioral:
...3890 voxels
...weights to: 115.281 = 126.613
...weights from: 81.4818 = 92.8137
...self weights: 0 = 11.3319
Right Medulla - Right Medulla:
...17303 voxels
...weights to: 162.776 = 241.373
...weights from: 187.547 = 266.144
...self weights: 0 = 78.5974
Right Spinal nucleus of the trigeminal - Right Spinal nucleus of the trigem
inal:
...3107 voxels
...weights to: 7.41493 = 7.94454
...weights from: 18.6598 = 19.1894
...self weights: 0 = 0.529612
Right Facial motor nucleus - Right Medulla:
...660 = 660 \text{ voxels}
...centres: [ 43.41327623 108.59100642 11.23768737] = [[ 43.41327623 108.5
9100642 11.23768737]]
...weights to: 9.15874 = 9.15874
...weights from: 11.0785 = 11.0785
...self tract_lengths: 0 = 0
Right Inferior olivary complex - Right Medulla:
...567 = 567 \text{ voxels}
4279835
         8.0617284 11
...weights to: 4.99518 = 4.99518
...weights from: 10.4086 = 10.4086
...self tract lengths: 0 = 0
Right Cerebellar Cortex - Right Cerebellar Cortex:
...33696 voxels
...weights to: 14.3996 = 16.6808
...weights from: 23.8543 = 26.1355
...self weights: 0 = 2.28123
Right Ansiform lobule - Right Cerebellar Cortex:
...9044 = 9044 \text{ voxels}
49.59369202]]
67718
...weights to: 0.843584 = 0.843584
...weights from: 0.209096 = 0.209096
...self tract_lengths: 0 = 0
Right Cerebellar Nuclei - Right Cerebellar Nuclei:
...1116 voxels
...weights to: 6.58665 = 7.09595
...weights from: 22.0603 = 22.5696
...self weights: 0 = 0.5093
Left Frontal pole, cerebral cortex - Left Isocortex:
...514 = 514 \text{ voxels}
...centres: [68.4057971 22.72256729 51.2484472 ] = [[68.4057971 22.722567
29 51.2484472 ]]
```

```
...weights to: 3.14868 = 3.14868
...weights from: 6.36079 = 6.36079
...self tract lengths: 0 = 0
Left Primary motor area - Left Isocortex:
...11760 = 11760 voxels
...centres: [78.69919552 42.96502274 55.40433718] = [[78.69919552 42.965022
74 55,4043371811
...weights to: 1.87382 = 1.87382
...weights from: 6.37274 = 6.37274
...self tract lengths: 0 = 0
Left Secondary motor area - Left Isocortex:
...10098 = 10098 voxels
...centres: [70.34126984 36.05326618 57.91804029] = [[70.34126984 36.053266
18 57.91804029]]
...weights to: 2.33534 = 2.33534
...weights from: 6.99391 = 6.99391
...self tract_lengths: 0 = 0
Left Primary somatosensory area, nose - Left Isocortex:
...1358 = 1358 voxels
...centres: [92.97697368\ 56.74407895\ 54.42039474] = [[92.97697368\ 56.744078]
95 54.42039474]]
...weights to: 1.20136 = 1.20136
...weights from: 5.39897 = 5.39897
...self tract_lengths: 0 = 0
Left Primary somatosensory area, barrel field - Left Isocortex:
...10306 = 10306 voxels
...centres: [90.26299012 67.39846988 61.48453937] = [[90.26299012 67.398469
88 61.4845393711
...weights to: 1.37367 = 1.37367
...weights from: 6.26508 = 6.26508
...self tract_lengths: 0 = 0
Left Primary somatosensory area, lower limb - Left Isocortex:
...3254 = 3254 \text{ voxels}
...centres: [74.6426146 59.67402377 66.73344652] = [[74.6426146 59.674023
77 66.73344652]]
...weights to: 1.83371 = 1.83371
...weights from: 5.63341 = 5.63341
...self tract_lengths: 0 = 0
Left Primary somatosensory area, mouth - Left Isocortex:
...2924 = 2924 \text{ voxels}
...centres: [91.00224719 46.18266453 48.86420546] = [[91.00224719 46.182664
53 48.8642054611
...weights to: 1.3537 = 1.3537
...weights from: 5.09649 = 5.09649
...self tract_lengths: 0 = 0
Left Primary somatosensory area, upper limb - Left Isocortex:
...5406 = 5406 \text{ voxels}
...centres: [81.3455804 54.61182109 61.69755059] = [[81.3455804 54.611821
09 61.69755059]]
...weights to: 1.39079 = 1.39079
...weights from: 2.70357 = 2.70357
...self tract_lengths: 0 = 0
```

Left Primary somatosensory area, trunk - Left Isocortex:

```
...4799 = 4799 \text{ voxels}
...centres: [75.04409673 66.81507824 69.51920341] = [[75.04409673 66.815078
24 69.51920341]]
...weights to: 2.32431 = 2.32431
...weights from: 4.8304 = 4.8304
...self tract lengths: 0 = 0
Left Primary somatosensory area, unassigned - Left Isocortex:
...2958 = 2958 voxels
...centres: [84.93838863 58.57661927 61.72827804] = [[84.93838863 58.576619
27 61.72827804]]
...weights to: 1.71146 = 1.71146
...weights from: 3.56411 = 3.56411
...self tract lengths: 0 = 0
Left Supplemental somatosensory area - Left Isocortex:
...5729 = 5729 voxels
...centres: [98.64653641 61.09413854 45.52775311] = [[98.64653641 61.094138
54 45.5277531111
...weights to: 2.45012 = 2.45012
...weights from: 7.48341 = 7.48341
...self tract lengths: 0 = 0
Left Gustatory areas - Left Isocortex:
...2104 = 2104 \text{ voxels}
...centres: [94.43148358 46.67610419 34.30124575] = [[94.43148358 46.676104
19 34.30124575]]
...weights to: 3.59145 = 3.59145
...weights from: 8.43749 = 8.43749
...self tract_lengths: 0 = 0
Left Visceral area - Left Isocortex:
...1793 = 1793 \text{ voxels}
...centres: [101.32432432 62.97043919 34.46706081] = [[101.32432432 62.9
7043919 34.46706081]]
...weights to: 4.12891 = 4.12891
...weights from: 9.98387 = 9.98387
...self tract_lengths: 0 = 0
Left Dorsal auditory area - Left Isocortex:
...2556 = 2556 voxels
...centres: [99.31855501 75.45812808 54.78981938] = [[99.31855501 75.458128
08 54.78981938]]
...weights to: 3.36174 = 3.36174
...weights from: 4.9361 = 4.9361
...self tract_lengths: 0 = 0
Left Primary auditory area - Left Isocortex:
...2689 = 2689 voxels
...centres: [102.197405
                          79.33920297 50.82205746] = [[102.197405
                                                                        79.3
3920297 50.82205746]]
...weights to: 3.70985 = 3.70985
...weights from: 3.86101 = 3.86101
...self tract_lengths: 0 = 0
Left Posterior auditory area - Left Isocortex:
...191 = 191 \text{ voxels}
...centres: [99.41666667 84.17
                                    56.06666667] = [[99.41666667 84.17
56.0666666711
...weights to: 4.81068 = 4.81068
...weights from: 4.55941 = 4.55941
```

```
...self tract_lengths: 0 = 0
Left Ventral auditory area - Left Isocortex:
...2554 = 2554 \text{ voxels}
...centres: [103.49225664 78.00110619 45.04535398] = [[103.49225664 78.0]]
0110619 45.0453539811
...weights to: 3.64962 = 3.64962
...weights from: 7.59361 = 7.59361
...self tract lengths: 0 = 0
Left Anterolateral visual area - Left Isocortex:
...1981 = 1981 voxels
...centres: [94.07672634 82.52173913 62.23529412] = [[94.07672634 82.521739
13 62.23529412]]
...weights to: 3.36884 = 3.36884
...weights from: 6.11907 = 6.11907
...self tract_lengths: 0 = 0
Left Anteromedial visual area - Left Isocortex:
...1795 = 1795 \text{ voxels}
...centres: [73.51295337 76.65025907 71.68911917] = [[73.51295337 76.650259]
07 71.68911917]]
...weights to: 3.6538 = 3.6538
...weights from: 7.05138 = 7.05138
...self tract lengths: 0 = 0
Left Lateral visual area - Left Isocortex:
...880 = 880 \text{ voxels}
...centres: [92.92718447 91.04530744 60.5566343 ] = [[92.92718447 91.045307
44 60.5566343 ]]
...weights to: 3.31325 = 3.31325
...weights from: 5.33257 = 5.33257
...self tract_lengths: 0 = 0
Left Primary visual area - Left Isocortex:
...6227 = 6227 voxels
...centres: [82.75259467 90.72734923 66.94922861] = [[82.75259467 90.727349
23 66.94922861]]
...weights to: 2.40566 = 2.40566
...weights from: 4.65915 = 4.65915
...self tract_lengths: 0 = 0
Left Posterolateral visual area - Left Isocortex:
...679 = 679 \text{ voxels}
...centres: [ 89.56140351 100.0726817 56.95989975] = [[ 89.56140351 100.0
726817
        56.95989975]]
...weights to: 2.98092 = 2.98092
...weights from: 5.24074 = 5.24074
...self tract_lengths: 0 = 0
Left posteromedial visual area - Left Isocortex:
...1265 = 1265 \text{ voxels}
...centres: [73.93269231 83.88076923 71.89615385] = [[73.93269231 83.880769
23 71.89615385]]
...weights to: 3.1971 = 3.1971
...weights from: 9.94985 = 9.94985
...self tract_lengths: 0 = 0
Left Laterointermediate area - Left Isocortex:
\dots -1 = -1 voxels
...centres: [96.14814815 90.53497942 57.65432099] = [[96.14814815 90.534979
```

```
42 57.65432099]]
...weights to: 3.73671 = 3.73671
...weights from: 6.57665 = 6.57665
...self tract lengths: 0 = 0
Left Postrhinal area - Left Isocortex:
\dots -1 = -1 voxels
...centres: [98.39341693 95.63793103 51.69435737] = [[98.39341693 95.637931
03 51,6943573711
...weights to: 4.35521 = 4.35521
...weights from: 6.61262 = 6.61262
...self tract_lengths: 0 = 0
Left Anterior cinqulate area, dorsal part - Left Isocortex:
...2789 = 2789 \text{ voxels}
...centres: [61.46720757 44.29952671 59.43002028] = [[61.46720757 44.299526
71 59.4300202811
...weights to: 3.40323 = 3.40323
...weights from: 7.78532 = 7.78532
...self tract lengths: 0 = 0
Left Anterior cingulate area, ventral part - Left Isocortex:
...2418 = 2418 \text{ voxels}
...centres: [60.59131206 48.55319149 54.30141844] = [[60.59131206 48.553191
49 54.30141844]]
...weights to: 4.55297 = 4.55297
...weights from: 10.0158 = 10.0158
...self tract_lengths: 0 = 0
Left Prelimbic area - Left Isocortex:
...1790 = 1790 \text{ voxels}
...centres: [61.66954271 31.06039689 51.18205349] = [[61.66954271 31.060396
89 51.18205349]]
...weights to: 4.54218 = 4.54218
...weights from: 10.5647 = 10.5647
...self tract lengths: 0 = 0
Left Infralimbic area - Left Isocortex:
...1764 = 1764 \text{ voxels}
...centres: [60.84367246 34.98759305 41.86848635] = [[60.84367246 34.987593
05 41.86848635]]
...weights to: 7.0478 = 7.0478
...weights from: 7.95842 = 7.95842
...self tract_lengths: 0 = 0
Left Orbital area, lateral part - Left Isocortex:
...1853 = 1853 voxels
...centres: [72.24338235 29.64191176 40.67426471] = [[72.24338235 29.641911
76 40.67426471]]
...weights to: 4.16078 = 4.16078
...weights from: 6.72122 = 6.72122
...self tract_lengths: 0 = 0
Left Orbital area, medial part - Left Isocortex:
...1024 = 1024 \text{ voxels}
...centres: [60.7260274 28.0456621 43.21613394] = [[60.7260274 28.045662
1 43.21613394]]
...weights to: 5.11325 = 5.11325
...weights from: 7.6126 = 7.6126
...self tract_lengths: 0 = 0
```

```
Left Orbital area, ventrolateral part - Left Isocortex:
...1556 = 1556 voxels
...centres: [64.90561798 27.48539326 42.88426966] = [[64.90561798 27.485393
26 42.8842696611
...weights to: 4.6912 = 4.6912
...weights from: 7.00555 = 7.00555
...self tract_lengths: 0 = 0
Left Agranular insular area, dorsal part - Left Isocortex:
...2364 = 2364 \text{ voxels}
...centres: [84.22603107 34.61810391 36.22656668] = [[84.22603107 34.618103
91 36.2265666811
...weights to: 4.4157 = 4.4157
...weights from: 11.3421 = 11.3421
...self tract lengths: 0 = 0
Left Agranular insular area, posterior part - Left Isocortex:
...1348 = 1348 \text{ voxels}
...centres: [98.83649876 59.63996697 27.96944674] = [[98.83649876 59.639966
97 27.9694467411
...weights to: 5.1191 = 5.1191
...weights from: 8.72044 = 8.72044
...self tract_lengths: 0 = 0
Left Agranular insular area, ventral part - Left Isocortex:
...1120 = 1120 \text{ voxels}
...centres: [84.12485682 37.21076747 30. ] = [[84.12485682 37.210767
                           11
...weights to: 5.56127 = 5.56127
...weights from: 5.79887 = 5.79887
...self tract lengths: 0 = 0
Left Retrosplenial area, lateral agranular part - Left Isocortex:
...937 = 937 \text{ voxels}
...centres: [69.9363167 81.13253012 70.83390706] = [[69.9363167 81.132530
12 70.83390706]]
...weights to: 2.48287 = 2.48287
...weights from: 4.5897 = 4.5897
...self tract_lengths: 0 = 0
Left Retrosplenial area, dorsal part - Left Isocortex:
...2738 = 2738 \text{ voxels}
...centres: [65.80832017 \ 80.32490785 \ 70.98683518] = [[65.80832017 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.324907 \ 80.3
85 70.98683518]]
...weights to: 1.95983 = 1.95983
...weights from: 4.54976 = 4.54976
...self tract lengths: 0 = 0
Left Retrosplenial area, ventral part - Left Isocortex:
...3997 = 3997 \text{ voxels}
...centres: [62.25167464 78.26985646 66.47607656] = [[62.25167464 78.269856
46 66.47607656]]
...weights to: 2.52084 = 2.52084
...weights from: 7.47747 = 7.47747
...self tract_lengths: 0 = 0
Left Anterior area - Left Isocortex:
\dots -1 = -1 voxels
...centres: [77.27146814 72.52493075 70.37257618] = [[77.27146814 72.524930
75 70.37257618]]
...weights to: 2.75957 = 2.75957
```

```
...weights from: 5.86238 = 5.86238
...self tract lengths: 0 = 0
Left Rostrolateral visual area - Left Isocortex:
\dots -1 = -1 voxels
...centres: [87.78039216 77.9745098 66.57254902] = [[87.78039216 77.974509
8 66.57254902]]
...weights to: 2.27166 = 2.27166
...weights from: 6.42253 = 6.42253
...self tract lengths: 0 = 0
Left Temporal association areas - Left Isocortex:
...3773 = 3773 voxels
...centres: [102.69399613 82.74112331 44.43124597] = [[102.69399613 82.74112331 44.43124597] = [102.69399613 82.74112331 44.43124597] = [102.69399613 82.74112331 44.43124597] = [102.69399613 82.74112331 44.43124597] = [102.69399613 82.74112331 44.43124597] = [102.69399613 82.74112331 44.43124597] = [102.69399613 82.74112331 44.43124597] = [102.69399613 82.74112331 44.43124597] = [102.69399613 82.74112331 44.43124597] = [102.69399613 82.74112331 44.43124597] = [102.69399613 82.74112331 44.43124597] = [102.69399613 82.74112331 44.43124597] = [102.69399613 82.74112331 44.43124597] = [102.69399613 82.74112331 44.43124597] = [102.69399613 82.74112331] = [102.69399613 82.74112331] = [102.69399613 82.74112331] = [102.69399613 82.74112331] = [102.69399613 82.74112331] = [102.69399613 82.74112331] = [102.69399613 82.74112331] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.69399613] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.6939961] = [102.693999] = [102.693999] = [102.693999] = [102.69399] = [102.69399] = [102.69399] = [102.69399] = [102.69399] = [102.69399] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6939] = [102.6
4112331 44.43124597]]
...weights to: 4.7417 = 4.7417
...weights from: 6.0477 = 6.0477
...self tract_lengths: 0 = 0
Left Perirhinal area - Left Isocortex:
...1412 = 1412 \text{ voxels}
...centres: [103.68844221 82.23366834 33.93467337] = [[103.68844221 82.2]]
3366834 33.93467337]]
...weights to: 4.78609 = 4.78609
...weights from: 8.2565 = 8.2565
...self tract lengths: 0 = 0
Left Ectorhinal area - Left Isocortex:
...3390 = 3390 voxels
...centres: [102.9091954
                                                         80.43908046 36.10689655] = [[102.9091954
                                                                                                                                                        80.4
3908046 36,1068965511
...weights to: 5.28341 = 5.28341
...weights from: 7.69002 = 7.69002
...self tract_lengths: 0 = 0
Left Olfactory Areas - Left Olfactory Areas:
...41043 voxels
...weights to: 39.4701 = 48.8479
...weights from: 53.8398 = 63.2176
...self weights: 0 = 9.37775
Left Hippocampal Formation - Left Hippocampal Formation:
...34819 voxels
...weights to: 34.0178 = 43.4767
...weights from: 48.5659 = 58.0249
...self weights: 0 = 9.45896
Left Cortical Subplate - Left Cortical Subplate:
...6669 voxels
...weights to: 44.7184 = 50.8921
...weights from: 44.4496 = 50.6234
...self weights: 0 = 6.17378
Left Striatum - Left Striatum:
...40219 voxels
...weights to: 98.9072 = 109.003
...weights from: 88.2897 = 98.3856
...self weights: 0 = 10.0959
Left Pallidum - Left Pallidum:
...9308 voxels
```

```
...weights to: 84.9946 = 89.3808
...weights from: 75.5767 = 79.9628
...self weights: 0 = 4.38613
Left Thalamus - Left Thalamus:
...13346 voxels
...weights to: 280.134 = 326.04
...weights from: 186.091 = 231.998
...self weights: 0 = 45.906
Left Nonspecific Thalamus - Left Nonspecific Thalamus:
...2324 voxels
...weights to: 115.961 = 118.485
...weights from: 61.0322 = 63.5561
...self weights: 0 = 2.5239
Left Hypothalamus - Left Hypothalamus:
...13871 voxels
...weights to: 354.926 = 630.732
...weights from: 366.33 = 642.136
...self weights: 0 = 275.806
Left Midbrain - Left Midbrain:
...19804 voxels
...weights to: 338.668 = 429.195
...weights from: 286.757 = 377.284
...self weights: 0 = 90.5271
Left Superior colliculus, motor related - Left Midbrain:
...6779 = 6779 \text{ voxels}
...centres: [66.46432413 91.42129796 55.60666906] = [[66.46432413 91.421297
96 55.6066690611
...weights to: 9.62674 = 9.62674
...weights from: 7.67702 = 7.67702
...self tract_lengths: 0 = 0
Left Pons Sensory - Left Pons Sensory:
...1585 voxels
...weights to: 13.202 = 13.3324
...weights from: 19.1642 = 19.2947
...self weights: 0 = 0.130488
Left Principal sensory nucleus of the trigeminal - Left Pons Sensory:
...1093 = 1093 \text{ voxels}
                                         22.50363636] = [[ 77.66363636 101.8
...centres: [ 77.66363636 101.84
         22.50363636]]
...weights to: 3.14049 = 3.14049
...weights from: 9.72171 = 9.72171
...self tract_lengths: 0 = 0
Left Pons Motor - Left Pons Motor:
...5971 voxels
...weights to: 102.516 = 111.275
...weights from: 99.5803 = 108.339
...self weights: 0 = 8.75894
Left Pons Behavioral - Left Pons Behavioral:
...3890 voxels
...weights to: 115.281 = 126.613
...weights from: 81.4818 = 92.8137
...self weights: 0 = 11.3319
```

```
Left Medulla - Left Medulla:
...17303 voxels
...weights to: 162.776 = 241.373
...weights from: 187.547 = 266.144
...self weights: 0 = 78.5974
Left Spinal nucleus of the trigeminal - Left Spinal nucleus of the trigemin
al:
...3107 voxels
...weights to: 7.41493 = 7.94454
...weights from: 18.6598 = 19.1894
...self weights: 0 = 0.529612
Left Facial motor nucleus - Left Medulla:
...660 = 660 \text{ voxels}
...centres: [ 70.58672377 108.59100642 11.23768737] = [[ 70.58672377 108.5
9100642 11.23768737]]
...weights to: 9.15874 = 9.15874
...weights from: 11.0785 = 11.0785
...self tract_lengths: 0 = 0
Left Inferior olivary complex - Left Medulla:
...567 = 567 \text{ voxels}
4279835 8,0617284 11
...weights to: 4.99518 = 4.99518
...weights from: 10.4086 = 10.4086
...self tract_lengths: 0 = 0
Left Cerebellar Cortex - Left Cerebellar Cortex:
...33696 voxels
...weights to: 14.3996 = 16.6808
...weights from: 23.8543 = 26.1355
...self weights: 0 = 2.28123
Left Ansiform lobule - Left Cerebellar Cortex:
...9044 = 9044 \text{ voxels}
...centres: [ 86.66716141 118.267718
                                      49.59369202] = [[ 86.66716141 118.2
        49.59369202]]
...weights to: 0.843584 = 0.843584
...weights from: 0.209096 = 0.209096
...self tract lengths: 0 = 0
Left Cerebellar Nuclei - Left Cerebellar Nuclei:
...1116 voxels
...weights to: 6.58665 = 7.09595
...weights from: 22.0603 = 22.5696
...self weights: 0 = 0.5093
```

Plot merged connectome

```
In [10]: # Plot the resulting connectivity weights at linear and log scale
import matplotlib.pyplot as plt

fig, axes = plt.subplots(2, 2, figsize=(15, 15))
cs_lin = axes[0, 0].imshow(new_conn.weights, cmap='jet', aspect='equal', int axes[0, 0].set_title('Structural connectivity matrix', fontsize=12)
axcb_lin = plt.colorbar(cs_lin, ax=axes[0, 0])
```

```
interacting_with_Allen
axcb_lin.set_label('Weights', fontsize=12)
cs log = axes[0, 1].imshow(np.log10(new conn.weights), cmap='jet', aspect='<math>\epsilon
axes[0, 1].set title('Structural connectivity matrix (log scale)', fontsize=
axcb_log = plt.colorbar(cs_log, ax=axes[0, 1])
axcb log.set label('Log10(Weights)', fontsize=12)
ls lin = axes[1, 0].imshow(new conn.tract lengths, cmap='jet', aspect='equal
axcb_lin = plt.colorbar(ls_lin, ax=axes[1, 0])
axcb_lin.set_label('Lengths', fontsize=12)
ls log = axes[1, 1].imshow(np.log10(new conn.tract lengths), cmap='jet', asp
axcb log = plt.colorbar(ls log, ax=axes[1, 1])
axcb log.set label('Log10(Lengths)', fontsize=12)
/tmp/ipykernel 222/2993400780.py:10: RuntimeWarning: divide by zero encount
ered in log10
  cs_log = axes[0, 1].imshow(np.log10(new_conn.weights), cmap='jet', aspect
='equal', interpolation='none')
/tmp/ipykernel 222/2993400780.py:20: RuntimeWarning: divide by zero encount
ered in log10
  ls_log = axes[1, 1].imshow(np.log10(new_conn.tract_lengths), cmap='jet',
aspect='equal', interpolation='none')
        Structural connectivity matrix
                                                Structural connectivity matrix (log scale)
                                   0.8
                                           80
100
                                           100
60
100
                                           100
                                                                              0.8
```

Construct useful indices

120

```
In [11]: inds = {}
         inds["crtx"] = []
         inds["m1"] = []
         inds["s1brl"] = []
         inds["supcol"] = []
         inds["facial"] = []
         inds["trigeminal"] = []
         inds["ponssens trigeminal"] = []
         inds["ponssens"] = []
         inds["ponsmotor"] = []
         inds["thal"] = []
         inds["thalspec"] = []
         inds["m1thal"] = []
         inds["s1brlthal"] = []
         inds["ansilob"] = []
         inds["dentate"] = []
         inds["interposed"] = []
         inds["oliv"] = []
         inds['cereb nuclei'] = []
         inds['cereb crtx'] = []
         for iR, (reg, lbl) in enumerate(zip(new_conn.region_labels, new_major_struct
             if "Isocortex" in lbl:
                 inds["crtx"].append(iR)
             if "Primary motor area" in reg and "Specific" not in reg:
                 inds["m1"].append(iR)
             if "barrel" in reg and "Specific" not in reg:
                 inds["s1brl"].append(iR)
             if "Superior colliculus, motor related" in reg:
                 inds["supcol"].append(iR)
             if "Primary motor area" in reg and "Specific" in reg:
                 inds["m1thal"].append(iR)
             if "barrel" in reg and "Specific" in reg:
                 inds["s1brlthal"].append(iR)
             if "Superior colliculus, motor related" in req:
                 inds["supcol"].append(iR)
             if "Facial" in req:
                 inds["facial"] .append(iR)
             if "Spinal nucleus of the trigeminal" in reg:
                 inds["trigeminal"].append(iR)
             if "Principal sensory nucleus of the trigeminal" in reg:
                 inds["ponssens_trigeminal"] .append(iR)
             if "Pons Sensory" in lbl:
                 inds["ponssens"] .append(iR)
             if "Pons Motor" in lbl:
                 inds["ponsmotor"].append(iR)
             if "Nonspecific Thalamus" in lbl:
                 inds["thal"].append(iR)
             if "Thalamus" in lbl:
                 inds["thalspec"].append(iR)
             if "Ansiform lobule" in rea:
                 inds["ansilob"].append(iR)
             elif "Cerebellar Cortex" in lbl:
                 inds['cereb_crtx'].append(iR)
             if "Dentate nucleus" in reg:
                 inds["dentate"].append(iR)
             if "Interposed nucleus" in reg:
                 inds["interposed"].append(iR)
             elif "Cerebellar Nuclei" in lbl:
                 inds['cereb nuclei'].append(iR)
             if "Inferior olivary complex" in reg:
```

```
inds["oliv"].append(iR)
         inds["m1s1brl"] = inds["m1"] + inds["s1brl"]
         inds["sens"] = inds["trigeminal"] + inds["ponssens trigeminal"] + inds["s1b
         inds["motor"] = inds["m1thal"] + inds["facial"] # inds["supcol"] +
         inds["cereb"] = inds["ansilob"] + inds['cereb nuclei'] + inds["dentate"] + i
         inds['cereb merged'] = inds['cereb nuclei'] + inds['cereb crtx']
         allspecial = inds["m1s1brl"] + inds["motor"] + inds["sens"] + inds["thal"] +
         for key, val in inds.items():
             inds[key] = np.array(val)
         print("ROIs:\n", new conn.region labels[allspecial])
         ROTs:
          ['Right Primary motor area' 'Left Primary motor area'
          'Right Primary somatosensory area, barrel field'
          'Left Primary somatosensory area, barrel field'
          'Right Facial motor nucleus' 'Left Facial motor nucleus'
          'Right Spinal nucleus of the trigeminal'
          'Left Spinal nucleus of the trigeminal'
          'Right Principal sensory nucleus of the trigeminal'
          'Left Principal sensory nucleus of the trigeminal'
          'Right Nonspecific Thalamus' 'Left Nonspecific Thalamus' 'Right Thalamus'
          'Right Nonspecific Thalamus' 'Left Thalamus' 'Left Nonspecific Thalamus'
          'Right Ansiform lobule' 'Left Ansiform lobule' 'Right Cerebellar Nuclei'
          'Left Cerebellar Nuclei' 'Right Inferior olivary complex'
          'Left Inferior olivary complex']
In [12]: import h5py
         # the method returns the tract lengths between the brain areas in the select
         def construct tract lengths(centres):
             n_regions = len(centres)
             tracts = np.zeros((n regions, n regions), dtype=float)
             for i_ind, inj in enumerate(centres):
                 for i targ, targ in enumerate(centres):
                     tracts[i_ind, i_targ] = np.sqrt(
                         (inj[0] - targ[0]) ** 2 + (inj[1] - targ[1]) ** 2 + (inj[2])
             return tracts
         def write connectivity to h5 file(connectivity, filename):
             #save the structural conn matrix
             f = h5py.File(os.path.join(data path, filename), "w")
             for attr in ["weights", "tract_lengths", "centres", "region_labels"]:
                 if attr == "region labels":
                     f.create dataset(attr, data=getattr(connectivity, attr).astype(h
                 else:
                     f.create_dataset(attr, data=getattr(connectivity, attr))
             f.close()
         def write all files(connectivity, major structs labels, voxel count, inds,
                             conn_name="Connectivity", conn_type=""):
             if WEIGHTED AVERAGE CENTRES:
```

if TRACT_LENGTHS_MODE == "euclidean":

conn_name += "_wavCntrs"

conn_name += "_avCntrs"

conn_name += "_TLed"

conn name += " TLwav"

else:

else:

```
# Write the resulting connectivity to .h5 TVB compatible file...
print("Writing %s_%s.h5" % (conn_name, conn_type))
write_connectivity_to_h5_file(connectivity, "%s_%s.h5" % (conn_name, con
# ... along with the corresponding mapping from regions to major structu
print("Writing major_structs_labels_%s.npy..." % conn_type)
np.save("major_structs_labels_%s.npy" % conn_type, major_structs_labels)
print("Writing voxel_count_%s.npy..." % conn_type)
np.save("voxel_count_%s.npy" % conn_type, voxel_count)
print("Writing inds_%s.npy..." % conn_type)
np.save("inds_%s.npy" % conn_type, inds)
print("Major structures:\n", np.unique(new_major_structs_labels))
```

```
In [13]: print("Major structures:\n", np.unique(new_major_structs_labels))

Major structures:
    ['Left Cerebellar Cortex' 'Left Cerebellar Nuclei'
    'Left Cortical Subplate' 'Left Hippocampal Formation' 'Left Hypothalamus'
    'Left Isocortex' 'Left Medulla' 'Left Midbrain'
    'Left Nonspecific Thalamus' 'Left Olfactory Areas' 'Left Pallidum'
    'Left Pons Behavioral' 'Left Pons Motor' 'Left Pons Sensory'
    'Left Spinal nucleus of the trigeminal' 'Left Striatum' 'Left Thalamus'
    'Right Cerebellar Cortex' 'Right Cerebellar Nuclei'
    'Right Cortical Subplate' 'Right Hippocampal Formation'
    'Right Hypothalamus' 'Right Isocortex' 'Right Medulla' 'Right Midbrain'
    'Right Nonspecific Thalamus' 'Right Olfactory Areas' 'Right Pallidum'
    'Right Pons Behavioral' 'Right Pons Motor' 'Right Pons Sensory'
    'Right Spinal nucleus of the trigeminal' 'Right Striatum'
    'Right Thalamus']
```

Writing files for Connectivity with summarized subcortical structures

Splitting Thalamus to create specific thalamic nuclei

```
In [15]: # Now we need to create the specific Thalamic nuclei, one for each Isocortex

# Thalamus indices:
iThR = np.where(["Right Thalamus" in lbl for lbl in new_major_structs_labels
iThL = np.where(["Left Thalamus" in lbl for lbl in new_major_structs_labels]
assert np.all(new_conn.region_labels[[iThR, iThL]] == ['Right Thalamus', 'Le

# Isocortex indices
crtx_inds = np.where(["Isocortex" in lbl for lbl in new_major_structs_labels
n_crtx = int(len(crtx_inds))
n_crtx2 = int(n_crtx/2)
crtx_inds_R = crtx_inds[:n_crtx2]
```

```
crtx_inds_L = crtx_inds[n_crtx2:]
# All the rest indices, subcortical but not specific thalamic, but including
subcrtx inds = np.arange(len(new major structs labels)).astype('i')
subcrtx inds = np.delete(subcrtx inds,
                         crtx inds.tolist() + [iThR, iThL])
# Create new major_structs_labels
insert inds R = crtx inds[n crtx2-1]+1
final major structs labels = insert axis(new major structs labels,
                                         arr to insert=np.tile('Right Specif
                                         inds=[insert_inds_R])
insert inds L = 1+ np.where(["Left Isocortex" in lbl for lbl in final major
final major structs labels = insert axis(final major structs labels,
                                         arr to insert=np.tile('Left Specifi
                                         inds=[insert inds L])
# print(final major structs labels)
# Get the new indices of major structures:
final crtx inds = np.where(["Isocortex" in lbl for lbl in final major struct
final_crtx_inds_R = final_crtx_inds[:n_crtx2]
final crtx inds L = final crtx inds[n crtx2:]
final_spec_thal_inds_R = final_crtx_inds[:n_crtx2] + n_crtx2
final spec thal inds L = final crtx inds[n crtx2:] + n crtx2
final spec thal inds = final spec thal inds R.tolist() + final spec thal ind
final iThR = np.where(["Right Nonspecific Thalamus" in lbl for lbl in final
final iThL = np.where(["Left Nonspecific Thalamus" in lbl for lbl in final m
final_n_regions = len(final_major_structs_labels)
# All the rest indices, subcortical but not thalamic:
final_subcrtx_inds = np.arange(final_n_regions).astype('i')
final subcrtx inds = np.delete(final subcrtx inds,
                               final_crtx_inds.tolist() +
                               final_spec_thal_inds_R.tolist() +
                               final_spec_thal_inds_L.tolist() +
                               [final_iThR, final_iThL])
assert final_n_regions == new_conn.number_of_regions + n_crtx
assert np.all(final major structs labels[final crtx inds R] == "Right Isocor"
assert np.all(final_major_structs_labels[final_crtx_inds_L] == "Left Isocort"
assert np.all(final_major_structs_labels[final_spec_thal_inds_R] == "Right S
assert np.all(final_major_structs_labels[final_spec_thal_inds_L] == "Left Sp
assert np.all(final_major_structs_labels[[final_iThR, final_iThL]] == ['Righ
                                                                        'Left
# Create new voxel counts:
# Thalamus voxel counts
voxel_count_th = new_voxel_count[iThL]
assert new voxel count[iThL] == new voxel count[iThR] == voxel count th
voxel_count_spec_th = int(voxel_count_th / n_crtx2)
final_voxel_count = insert_axis(new_voxel_count,
                                arr_to_insert=np.tile(voxel_count_spec_th, r
                                inds=[insert_inds_R])
final_voxel_count = insert_axis(final_voxel_count,
                                arr to insert=np.tile(voxel count spec th, r
                                inds=[insert_inds_L])
assert np.all(final_voxel_count[final_spec_thal_inds_R] == voxel_count_spec_
assert np.all(final_voxel_count[final_spec_thal_inds_L] == voxel_count_spec_
# Create new region labels
final_region_labels = insert_axis(new_conn.region_labels,
                                  arr_to_insert=np.array(['Right Specific Th
                                                          for reg in new_cor
```

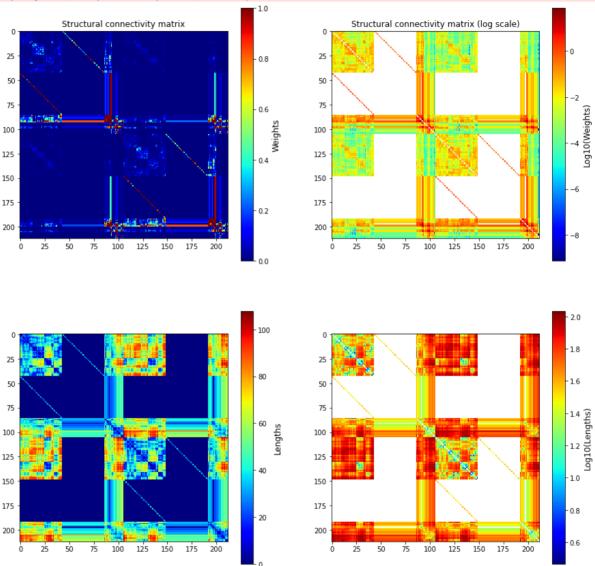
```
inds=[insert_inds_R])
final region labels = insert axis(final region labels,
                                  arr to insert=np.array(['Left Specific Tha
                                                          for reg in new cor
                                  inds=[insert inds L])
assert np.all(np.array([req.split("Right")[-1] for req in final region labe
              np.array([reg.split("Right Specific Thalamus to ")[-1] for reg
assert np.all(np.array([reg.split("Left ")[-1] for reg in final_region_label")
              np.array([reg.split("Left Specific Thalamus to ")[-1] for reg
# Create new centers:
final_centres = insert_axis(new_conn.centres,
                            arr to insert=np.repeat([new conn.centres[iThR]]
                            inds=[insert inds R])
final centres = insert axis(final centres,
                            arr to insert=np.repeat([new conn.centres[iThL]]
                            inds=[insert inds L])
def expand 2D with zeros(arr, inds, nregs):
    new arr = insert axis(arr,
                          arr to insert=np.tile(0.0, (nregs, arr.shape[1])),
                          inds=[inds[0]], axis=0)
    new_arr = insert_axis(new_arr,
                          arr to insert=np.tile(0.0, (nregs, new arr.shape[1
                          inds=[inds[1]], axis=0)
    new_arr = insert_axis(new_arr,
                          arr_to_insert=np.tile(0.0, (new_arr.shape[0], nreg
                          inds=[inds[0]], axis=1)
    new_arr = insert_axis(new_arr,
                          arr to insert=np.tile(0.0, (new arr.shape[0], nred
                          inds=[inds[1]], axis=1)
    return new arr
# Create new weights:
final weights = expand 2D with zeros(new conn.weights, [insert inds R, insert
# Connections Specific Thalami -> Cortex = Specific Thalamus -> Cortex:
final_weights[final_crtx_inds_R, final_spec_thal_inds_R] = new_conn.weights[
final_weights[final_crtx_inds_L, final_spec_thal_inds_L] = new_conn.weights[
# Connections Cortex -> Specific Thalami = Cortex -> Specific Thalamus:
final_weights[final_spec_thal_inds_R, final_crtx_inds_R] = new_conn.weights[
final weights[final spec thal inds L, final crtx inds L] = new conn.weights[
# Connections Subcortex -> Specific Thalami = Subcortex -> Specific Thalamus
final_weights[final_spec_thal_inds_R[:, None], final_subcrtx_inds[None, :]]
final_weights[final_spec_thal_inds_L[:, None], final_subcrtx_inds[None, :]]
# Connections Specific Thalami -> Subcortex = Specific Thalamus -> Subcort
final_weights[final_subcrtx_inds[:, None], final_spec_thal_inds_R[None, :]]
final_weights[final_subcrtx_inds[:, None], final_spec_thal_inds_L[None, :]]
# No connections Specific Thalami <-> Specific Thalami!!!
# Create new tract_lengths:
final_tract_lengths = expand_2D_with_zeros(new_conn.tract_lengths, [insert_i
# Connections Specific Thalami -> Cortex = Specific Thalamus -> Cortex:
final_tract_lengths[final_crtx_inds_R, final_spec_thal_inds_R] = new_conn.tr
final_tract_lengths[final_crtx_inds_L, final_spec_thal_inds_L] = new_conn.tr
# Connections Cortex -> Specific Thalami = Cortex -> Specific Thalamus:
final_tract_lengths[final_spec_thal_inds_R, final_crtx_inds_R] = new_conn.tr
final tract lengths[final spec thal inds L, final crtx inds L] = new conn.tr
# Connections Subcortex -> Specific Thalami = Subcortex -> Specific Thalamus
# Connections Subcortex -> Specific Thalami = Subcortex -> Specific Thalamus
```

```
final_tract_lengths[final_spec_thal_inds_R[:, None], final_subcrtx_inds[None
final tract lengths[final spec thal inds L[:, None], final subcrtx inds[None
# Connections Specific Thalami -> Subcortex = Specific Thalamus -> Subcort
final_tract_lengths[final_subcrtx_inds[:, None], final_spec_thal_inds_R[None
final_tract_lengths[final_subcrtx_inds[:, None], final_spec_thal_inds_L[None
# No connections Specific Thalami <-> Specific Thalami!!!
# Finally delete the merged Right/Left Thalamus:
iTh = []
iTh.append(np.where(["Right Thalamus" in lbl for lbl in final region labels]
iTh.append(np.where(["Left Thalamus" in lbl for lbl in final region labels])
iTh = np.unique(iTh)
final major structs labels = np.delete(final major structs labels, iTh)
final voxel count = np.delete(final voxel count, iTh)
final region labels = np.delete(final region labels, iTh)
final_centres = np.delete(final_centres, iTh)
final weights = np.delete(final weights, iTh, axis=0)
final weights = np.delete(final weights, iTh, axis=1)
final_tract_lengths = np.delete(final_tract_lengths, iTh, axis=0)
final tract lengths = np.delete(final tract lengths, iTh, axis=1)
final connectivity = Connectivity(weights=final weights,
                                  tract_lengths=final_tract_lengths,
                                  centres=final centres,
                                  region labels=final region labels)
```

Plot the connectivity with the specific thalami

```
In [16]: # Plot the resulting connectivity weights at linear and log scale
         import matplotlib.pyplot as plt
         fig, axes = plt.subplots(2, 2, figsize=(15, 15))
         cs_lin = axes[0, 0].imshow(final_connectivity.weights, cmap='jet', aspect='e
         axes[0, 0].set_title('Structural connectivity matrix', fontsize=12)
         axcb_lin = plt.colorbar(cs_lin, ax=axes[0, 0])
         axcb_lin.set_label('Weights', fontsize=12)
         cs log = axes[0, 1].imshow(np.log10(final connectivity.weights), cmap='jet',
         axes[0, 1].set title('Structural connectivity matrix (log scale)', fontsize=
         axcb log = plt.colorbar(cs log, ax=axes[0, 1])
         axcb_log.set_label('Log10(Weights)', fontsize=12)
         ls lin = axes[1, 0].imshow(final connectivity.tract lengths, cmap='jet', asp
         axcb_lin = plt.colorbar(ls_lin, ax=axes[1, 0])
         axcb_lin.set_label('Lengths', fontsize=12)
         ls_log = axes[1, 1].imshow(np.log10(final_connectivity.tract_lengths), cmap=
         axcb log = plt.colorbar(ls log, ax=axes[1, 1])
         axcb_log.set_label('Log10(Lengths)', fontsize=12)
```

/tmp/ipykernel_222/2136995184.py:10: RuntimeWarning: divide by zero encount
ered in log10
 cs_log = axes[0, 1].imshow(np.log10(final_connectivity.weights), cmap='je
t', aspect='equal', interpolation='none')
/tmp/ipykernel_222/2136995184.py:20: RuntimeWarning: divide by zero encount
ered in log10
 ls_log = axes[1, 1].imshow(np.log10(final_connectivity.tract_lengths), cm
ap='jet', aspect='equal', interpolation='none')



Plot the resulting connectivity weights and tract lengths beween Isocortex and Specific Thalamic Nuclei

```
In [17]: fig, axes = plt.subplots(2, 2, figsize=(15, 15))
    cs_spec_R = axes[0, 0].plot(final_connectivity.weights[final_crtx_inds_R, fi
    cs_spec_L = axes[0, 0].plot(final_connectivity.weights[final_crtx_inds_L, fi
    axes[0, 0].set_title('Isocortex -> Specific Thalamic Nuclei', fontsize=12)
    axes[0, 0].set_ylabel('Weights', fontsize=12)
    axes[0, 0].legend()

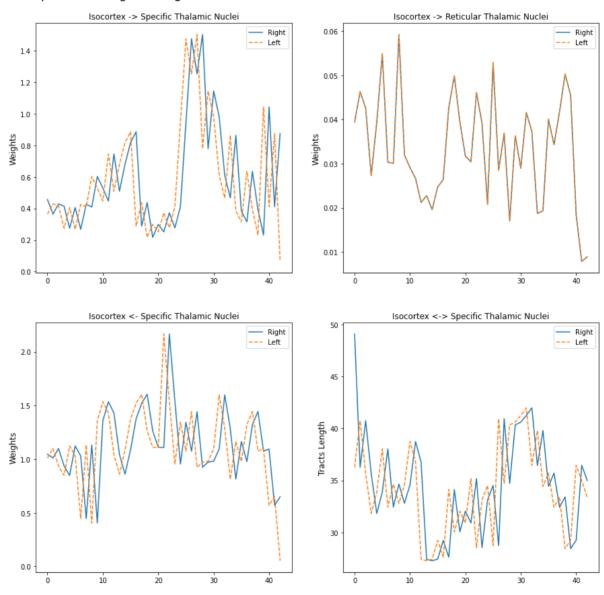
cs_rtn_R = axes[0, 1].plot(crtx_to_rtn_conns[0, :43], "-", label="Right")
    cs_rtn_L = axes[0, 1].plot(crtx_to_rtn_conns[1, 43:], "--", label="Left")
    axes[0, 1].set_title('Isocortex -> Reticular Thalamic Nuclei', fontsize=12)
    axes[0, 1].set_ylabel('Weights', fontsize=12)
```

```
axes[0, 1].legend()

cs_lin_R = axes[1, 0].plot(final_connectivity.weights[final_spec_thal_inds_F
cs_lin_L = axes[1, 0].plot(final_connectivity.weights[final_spec_thal_inds_L
axes[1, 0].set_title('Isocortex <- Specific Thalamic Nuclei', fontsize=12)
axes[1, 0].set_ylabel('Weights', fontsize=12)
axes[1, 0].legend()

tl_lin_R = axes[1, 1].plot(final_connectivity.tract_lengths[final_spec_thal_tl_lin_L = axes[1, 1].plot(final_connectivity.tract_lengths[final_spec_thal_axes[1, 1].set_title('Isocortex <-> Specific Thalamic Nuclei', fontsize=12)
axes[1, 1].set_ylabel('Tracts Length', fontsize=12)
axes[1, 1].legend()
```

Out[17]: <matplotlib.legend.Legend at 0x7ff189060e50>



Construct useful indices

```
In [18]: inds = {}
  inds["crtx"] = []
  inds["m1"] = []
  inds["s1brl"] = []
  inds["supcol"] = []
  inds["facial"] = []
```

```
inds["trigeminal"] = []
inds["ponssens trigeminal"] = []
inds["ponssens"] = []
inds["ponsmotor"] = []
inds["thal"] = []
inds["thalspec"] = []
inds["m1thal"] = []
inds["s1brlthal"] = []
inds["ansilob"] = []
inds["dentate"] = []
inds["interposed"] = []
inds["oliv"] = []
inds['cereb nuclei'] = []
inds['cereb crtx'] = []
for iR, (reg, lbl) in enumerate(zip(final_connectivity.region_labels, final_
    if "Isocortex" in lbl:
        inds["crtx"].append(iR)
    if "Primary motor area" in reg and "Specific" not in reg:
        inds["m1"].append(iR)
    if "barrel" in reg and "Specific" not in reg:
        inds["s1brl"].append(iR)
    if "Superior colliculus, motor related" in reg:
        inds["supcol"].append(iR)
    if "Primary motor area" in reg and "Specific" in reg:
        inds["m1thal"].append(iR)
    if "barrel" in reg and "Specific" in reg:
        inds["s1brlthal"].append(iR)
    if "Superior colliculus, motor related" in reg:
        inds["supcol"].append(iR)
    if "Facial" in reg:
        inds["facial"] .append(iR)
    if "Spinal nucleus of the trigeminal" in reg:
        inds["trigeminal"].append(iR)
    if "Principal sensory nucleus of the trigeminal" in reg:
        inds["ponssens_trigeminal"] .append(iR)
    if "Pons Sensory" in lbl:
        inds["ponssens"] .append(iR)
    if "Pons Motor" in lbl:
        inds["ponsmotor"].append(iR)
    if "Nonspecific Thalamus" in lbl:
        inds["thal"].append(iR)
    if "Specific Thalamus" in lbl:
        inds["thalspec"].append(iR)
    if "Ansiform lobule" in reg:
        inds["ansilob"].append(iR)
    elif "Cerebellar Cortex" in lbl:
        inds['cereb crtx'].append(iR)
    if "Dentate nucleus" in reg:
        inds["dentate"].append(iR)
    if "Interposed nucleus" in req:
        inds["interposed"].append(iR)
    elif "Cerebellar Nuclei" in lbl:
        inds['cereb nuclei'].append(iR)
    if "Inferior olivary complex" in reg:
        inds["oliv"].append(iR)
inds["m1s1brl"] = inds["m1"] + inds["s1brl"]
inds["sens"] = inds["trigeminal"] + inds["ponssens_trigeminal"] + inds["s1t
inds["motor"] = inds["m1thal"] + inds["facial"] # inds["supcol"] +
inds["cereb"] = inds["ansilob"] + inds['cereb_nuclei'] + inds["dentate"] + i
inds['cereb_merged'] = inds['cereb_nuclei'] + inds['cereb_crtx']
```

```
allspecial = inds["m1s1brl"] + inds["motor"] + inds["sens"] + inds["thal"] +
         for key, val in inds.items():
             inds[key] = np.array(val)
         print("ROIs:\n", final connectivity.region labels[allspecial])
          ['Right Primary motor area' 'Left Primary motor area'
          'Right Primary somatosensory area, barrel field'
          'Left Primary somatosensory area, barrel field'
          'Right Specific Thalamus to Primary motor area'
          'Left Specific Thalamus to Primary motor area'
          'Right Facial motor nucleus' 'Left Facial motor nucleus'
          'Right Spinal nucleus of the trigeminal'
          'Left Spinal nucleus of the trigeminal'
          'Right Principal sensory nucleus of the trigeminal'
          'Left Principal sensory nucleus of the trigeminal'
          'Right Specific Thalamus to Primary somatosensory area, barrel field'
          'Left Specific Thalamus to Primary somatosensory area, barrel field'
          'Right Nonspecific Thalamus' 'Left Nonspecific Thalamus'
          'Right Ansiform lobule' 'Left Ansiform lobule' 'Right Cerebellar Nuclei'
          'Left Cerebellar Nuclei' 'Right Inferior olivary complex'
          'Left Inferior olivary complex']
In [19]: print("Major structures:\n", np.unique(final_major_structs_labels))
         Maior structures:
          ['Left Cerebellar Cortex' 'Left Cerebellar Nuclei'
          'Left Cortical Subplate' 'Left Hippocampal Formation' 'Left Hypothalamus'
          'Left Isocortex' 'Left Medulla' 'Left Midbrain'
          'Left Nonspecific Thalamus' 'Left Olfactory Areas' 'Left Pallidum'
          'Left Pons Behavioral' 'Left Pons Motor' 'Left Pons Sensory'
          'Left Specific Thalamus' 'Left Spinal nucleus of the trigeminal'
          'Left Striatum' 'Right Cerebellar Cortex' 'Right Cerebellar Nuclei'
          'Right Cortical Subplate' 'Right Hippocampal Formation'
          'Right Hypothalamus' 'Right Isocortex' 'Right Medulla' 'Right Midbrain'
          'Right Nonspecific Thalamus' 'Right Olfactory Areas' 'Right Pallidum'
          'Right Pons Behavioral' 'Right Pons Motor' 'Right Pons Sensory'
          'Right Specific Thalamus' 'Right Spinal nucleus of the trigeminal'
          'Right Striatum']
```

Writing files for Connectivity with summarized subcortical structures and Specific Thalami

Print out statistics of final regions' sizes and connectivity

```
In [21]: from six import string types
         from collections import OrderedDict
         from tvb.contrib.scripts.utils.data_structures_utils import is_integer, is_f
         from examples.tvb nest.notebooks.cerebellum.utils import print conn
         def maxrow(maxrow, weights):
             if maxrow < 1:</pre>
                 cum = np.maximum(maxrow, 0.1)
                 maxrow_label = "%g%%" % (100*cum)
                 weights sum = 0.0
                 weights total = weights.sum()
                 maxrow = 0
                 for weight in weights:
                     maxrow += 1
                     weights sum += weight
                     if weights sum / weights total >= cum:
             else:
                 maxrow = int(maxrow)
                 maxrow label = "%d" % maxrow
             return maxrow, maxrow label
         def print_weights_from(label, inds, region_labels, weights, maxrow=0.9):
             for iH, hemi in enumerate(["Right", "Left"]):
                 arginds = np.argsort(weights[inds[iH], :])[::-1]
                 out_maxrow, maxrow_label = _maxrow(maxrow, weights[inds[iH], arginds
                 d = OrderedDict()
                 d[("Region", 55)] = ["%d.%s" % (iR, region_labels[iR]) for iR in arg
                 d[("Weight", 30)] = weights[inds[iH], arginds]
                 d[("Total weights to", 30)] = np.nansum(weights[arginds], axis=1)
                 print conn(d, prnt="\nStronger %s connections from %s %s to:\n" % (m
                             maxrow=out_maxrow, printit=True);
         def print_weights_to(label, inds, region_labels, weights, maxrow=0.9):
             for iH, hemi in enumerate(["Right", "Left"]):
                 arginds = np.argsort(weights[:, inds[iH]])[::-1]
                 out_maxrow, maxrow_label = _maxrow(maxrow, weights[inds[iH], arginds
                 d = OrderedDict()
                 d[("Region", 55)] = ["%d.%s" % (iR, region_labels[iR]) for iR in arg
                 d[("Weight", 30)] = weights[arginds, inds[iH]]
                 d[("Total weights from", 30)] = np.nansum(weights[:, arginds], axis=
                 print_conn(d, prnt="\nStronger %s connections to %s %s from:\n" % (m
                            maxrow=out maxrow, printit=True);
In [22]: d = OrderedDict()
         d[("Region", 55)] = ["%d.%s" % (iR, reg[:50]) for iR, reg in enumerate(final)
         d[("Major Structure", 30)] = final_major_structs_labels
         d[("Voxels", 10)] = final voxel count
         d[("Total weights to/from", 21)] = np.array(["%0.3f/%0.3f" % ( wt, wf)
                                                       for wt, wf in zip(np.nansum(fir
                                                                         np.nansum(fir
         print_conn(d, prnt="", printit=True);
```

interacting_with_Allen	_
RegionMajor eVoxelsTotal weights to/from	Structur
<pre>0.Right Frontal pole, cerebral cortexRight x5143.107/5.948</pre>	Isocorte
1.Right Primary motor areaRight	Isocorte
x	Isocorte
x100982.315/6.759	
3.Right Primary somatosensory area, noseRight x13581.197/5.138	Isocorte
4.Right Primary somatosensory area, barrel fieldRight x103061.371/6.229	Isocorte
5.Right Primary somatosensory area, lower limbRight	Isocorte
x32541.829/5.456	Isocorte
x	
7.Right Primary somatosensory area, upper limbRight x54061.388/2.691	Isocorte
8.Right Primary somatosensory area, trunkRight x47992.315/4.736	Isocorte
9.Right Primary somatosensory area, unassignedRight	Isocorte
x29581.709/3.534	Isocorte
x57292.436/7.186 11.Right Gustatory areas	
x21043.481/7.911	
12.Right Visceral areaRight x17934.021/9.503	Isocorte
13.Right Dorsal auditory areaRight	Isocorte
x	Isocorte
x	Tsocorte
x1914.806/4.555	
16.Right Ventral auditory areaRight x25543.647/7.515	Isocorte
17.Right Anterolateral visual areaRight x19813.365/6.113	Isocorte
18.Right Anteromedial visual areaRight	Isocorte
x	Isocorte
x	
20.Right Primary visual areaRight x62272.396/4.574	Isocorte
21.Right Posterolateral visual areaRight x6792.945/5.238	Isocorte
22.Right posteromedial visual areaRight	Isocorte
x12653.178/9.601	Isocorte
x3.731/6.566	
x4.321/6.548	
25.Right Anterior cingulate area, dorsal partRight x27893.355/7.577	Isocorte
26.Right Anterior cingulate area, ventral partRight x24184.413/9.728	Isocorte
27.Right Prelimbic areaRight	Isocorte
x17904.390/9.983	Isocorte
5	

x17646.726/7.545	
29.Right Orbital area, lateral partRight	Isocorte
x18534.088/6.327	
30.Right Orbital area, medial part	Isocorte
31.Right Orbital area, ventrolateral partRight	Isocorte
x	
32.Right Agranular insular area, dorsal partRight	Isocorte
x	Tananta
33.Right Agranular insular area, posterior partRight x13485.005/8.246	Isocorte
34.Right Agranular insular area, ventral partRight	Isocorte
x11205.441/5.439	
35.Right Retrosplenial area, lateral agranular partRight	Isocorte
x	Tsocorte
x	13000100
37.Right Retrosplenial area, ventral partRight	Isocorte
x39972.476/7.241	_
38.Right Anterior area	Isocorte
39.Right Rostrolateral visual area	Isocorte
x2.267/6.416	
40.Right Temporal association areasRight	Isocorte
X	Tananta
41.Right Perirhinal areaRight x14124.662/8.128	Isocorte
42.Right Ectorhinal areaRight	Isocorte
x33905.197/7.544	
43.Right Specific Thalamus to Frontal pole, cerebralRight	Specific Thala
mus3105.630/3.463	Specific Thala
mus3105.595/3.371	Specific mata
45.Right Specific Thalamus to Secondary motor areaRight	Specific Thala
mus3105.683/3.436	Considia Thala
46.Right Specific Thalamus to Primary somatosensory aRight mus3105.521/3.421	Specific Inala
47.Right Specific Thalamus to Primary somatosensory aRight	Specific Thala
mus3105.432/3.282	
48.Right Specific Thalamus to Primary somatosensory aRight	Specific Thala
mus3105.707/3.413	Specific Thala
mus3105.615/3.275	Specific filata
50.Right Specific Thalamus to Primary somatosensory aRight	Specific Thala
mus3105.031/3.433	C 'C' TI 1
51.Right Specific Thalamus to Primary somatosensory aRight mus3105.716/3.416	Specific Thata
52.Right Specific Thalamus to Primary somatosensory aRight	Specific Thala
mus3104.989/3.610	•
53.Right Specific Thalamus to Supplemental somatosensRight	Specific Thala
mus3105.943/3.535 54.Right Specific Thalamus to Gustatory areasRight	Specific Thala
mus3106.119/3.454	Specific mata
55.Right Specific Thalamus to Visceral areaRight	Specific Thala
mus3106.013/3.752	a
56.Right Specific Thalamus to Dorsal auditory areaRight mus3105.627/3.516	Specific Thala
57.Right Specific Thalamus to Primary auditory areaRight	Specific Thala
mus3105.444/3.687	,
58.Right Specific Thalamus to Posterior auditory areaRight	Specific Thala
mus3105.671/3.824	

59.Right Specific Thalamus to Ventral auditory areaRight Specific Thala
mus3105.959/3.893 60.Right Specific Thalamus to Anterolateral visual arRight Specific Thala
mus3106.105/3.295
61.Right Specific Thalamus to Anteromedial visual areRight Specific Thala
mus3106.188/3.445 62.Right Specific Thalamus to Lateral visual areaRight Specific Thala
mus3105.847/3.224
63.Right Specific Thalamus to Primary visual areaRight Specific Thala
mus3105.694/3.306
64.Right Specific Thalamus to Posterolateral visual aRight Specific Thala mus3105.693/3.259
65.Right Specific Thalamus to posteromedial visual arRight Specific Thala
mus3106.751/3.380
66.Right Specific Thalamus to Laterointermediate areaRight Specific Thala mus3106.127/3.284
67.Right Specific Thalamus to Postrhinal areaRight Specific Thala
mus3105.538/3.418
68.Right Specific Thalamus to Anterior cingulate areaRight Specific Thala
mus3105.928/3.951 69.Right Specific Thalamus to Anterior cingulate areaRight Specific Thala
mus3105.658/4.482
70.Right Specific Thalamus to Prelimbic areaRight Specific Thala
mus3106.026/4.259
71.Right Specific Thalamus to Infralimbic areaRight Specific Thalamus3105.508/4.508
72.Right Specific Thalamus to Orbital area, lateral pRight Specific Thala
mus3105.556/3.785
73.Right Specific Thalamus to Orbital area, medial paRight Specific Thala mus3105.565/4.152
74.Right Specific Thalamus to Orbital area, ventrolatRight Specific Thala
mus3105.679/3.987
75.Right Specific Thalamus to Agranular insular area,Right Specific Thala mus3106.182/3.616
76.Right Specific Thalamus to Agranular insular area,Right Specific Thala
mus3105.870/3.474
77.Right Specific Thalamus to Agranular insular area,Right Specific Thala
mus3105.398/3.870 78.Right Specific Thalamus to Retrosplenial area, latRight Specific Thala
mus3105.748/3.391
79.Right Specific Thalamus to Retrosplenial area, dorRight Specific Thala
mus3105.562/3.322 80.Right Specific Thalamus to Retrosplenial area, venRight Specific Thala
mus3105.906/3.641
81.Right Specific Thalamus to Anterior areaRight Specific Thala
mus3106.028/3.399 82.Right Specific Thalamus to Rostrolateral visual arRight Specific Thala
mus3105.660/3.239
83.Right Specific Thalamus to Temporal association arRight Specific Thala
mus3105.679/4.050
84.Right Specific Thalamus to Perirhinal areaRight Specific Thala mus3105.154/3.418
85.Right Specific Thalamus to Ectorhinal areaRight Specific Thala
mus3105.233/3.882
86.Right Olfactory Areas
87.Right Hippocampal Formation
rmation3481934.018/48.566
88.Right Cortical SubplateRight Cortical Subplate666944.718/44.450
89.Right Striatum

40040 00 00 000	
m	
90.Right PallidumRigh	t Pallidu
m	. N
91.Right Nonspecific Thalamus	t Nonspecific in
alamus232497.939/46.910	+ 11
92.Right Hypothalamus	т нуротпатати
s	
93.Right MidbrainRigh	t Midbrai
n	
94.Right Superior colliculus, motor relatedRigh	t Midbrai
n	
95.Right Pons SensoryRigh	t Pons Sensor
y	
96.Right Principal sensory nucleus of the trigeminalRigh	t Pons Sensor
y	. D. M. I
97.Right Pons MotorRigh	t Pons Moto
r	t Dana Dahawiana
98.Right Pons Behavioral	t Pons Benaviora
l3890115.281/81.482	+ Mad
99.Right MedullaRigh	t Medull
a17303162.776/187.547	+ Cninal muslaus
100.Right Spinal nucleus of the trigeminalRigh	t Spinat nucleus
of the trigeminal31077.415/18.660	+ Madull
	t Medutt
a	+ Madull
a	t Medutt
103.Right Cerebellar CortexRigh	t Caraballar Cor
tex3369614.400/23.854	t cerebettar cor
104.Right Ansiform lobule	t Cerebellar Cor
tex90440.844/0.209	c cerebettar cor
105.Right Cerebellar Nuclei	
	t Cerebellar Nuc
	t Cerebellar Nuc
lei11166.587/22.060	
lei11166.587/22.060 106.Left Frontal pole, cerebral cortexLeft	
lei11166.587/22.060 106.Left Frontal pole, cerebral cortexLeft x5143.107/5.948	Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortexLeft x5143.107/5.948 107.Left Primary motor areaLeft	Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortexLeft x5143.107/5.948 107.Left Primary motor areaLeft x117601.855/6.174	Isocorte Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortexLeft x5143.107/5.948 107.Left Primary motor areaLeft x117601.855/6.174 108.Left Secondary motor areaLeft	Isocorte Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortexLeft x5143.107/5.948 107.Left Primary motor areaLeft x117601.855/6.174	Isocorte Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortexLeft x5143.107/5.948 107.Left Primary motor areaLeft x117601.855/6.174 108.Left Secondary motor areaLeft xLeft	Isocorte Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortex	Isocorte Isocorte Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortexLeft x5143.107/5.948 107.Left Primary motor areaLeft x117601.855/6.174 108.Left Secondary motor areaLeft xLeft xLeft xLeft x	Isocorte Isocorte Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortex	Isocorte Isocorte Isocorte Isocorte Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortex	Isocorte Isocorte Isocorte Isocorte Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortex	Isocorte Isocorte Isocorte Isocorte Isocorte Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortex	Isocorte Isocorte Isocorte Isocorte Isocorte Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortex	Isocorte Isocorte Isocorte Isocorte Isocorte Isocorte Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortex	Isocorte Isocorte Isocorte Isocorte Isocorte Isocorte Isocorte Isocorte Isocorte
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lei11166.587/22.060	Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortex	Isocorte
lei	Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortex	Isocorte
lei11166.587/22.060 106.Left Frontal pole, cerebral cortexLeft x	Isocorte
lei11166.587/22.060	Isocorte
lei11166.587/22.060	Isocorte
lei11166.587/22.060	Isocorte
lei11166.587/22.060	Isocorte
lei11166.587/22.060	Isocorte

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120.Left Primary auditory areaLeft Isocorte x26893.708/3.858
121.Left Posterior auditory areaLeft Isocorte x1914.806/4.555
122.Left Ventral auditory areaLeft Isocorte x25543.647/7.515
123.Left Anterolateral visual areaLeft Isocorte x19813.365/6.113
124.Left Anteromedial visual areaLeft Isocorte x17953.629/6.890
125.Left Lateral visual areaLeft Isocorte x8803.308/5.298
126.Left Primary visual areaLeft Isocorte x62272.396/4.574
127.Left Posterolateral visual areaLeft Isocorte x6792.945/5.238
128.Left posteromedial visual areaLeft Isocorte x12653.178/9.601
129.Left Laterointermediate areaLeft Isocorte x1
130.Left Postrhinal areaLeft Isocorte x
131.Left Anterior cingulate area, dorsal partLeft Isocorte x
132.Left Anterior cingulate area, ventral partLeft Isocorte x24184.413/9.728
133.Left Prelimbic area4.390/9.983Left Isocorte
134.Left Infralimbic area6.726/7.545Left Isocorte
135.Left Orbital area, lateral partLeft Isocorte x18534.088/6.327
136.Left Orbital area, medial partLeft Isocorte x
137.Left Orbital area, ventrolateral partLeft Isocorte x15564.611/6.571
138.Left Agranular insular area, dorsal partLeft Isocorte x23644.273/10.528
139.Left Agranular insular area, posterior partLeft Isocorte x13485.005/8.246
140.Left Agranular insular area, ventral partLeft Isocorte x
141.Left Retrosplenial area, lateral agranular partLeft Isocorte x9372.446/4.479
142.Left Retrosplenial area, dorsal partLeft Isocorte x27381.937/4.418
143.Left Retrosplenial area, ventral partLeft Isocorte x39972.476/7.241
144.Left Anterior area2.749/5.801Left Isocorte
145.Left Rostrolateral visual areaLeft Isocorte x
146.Left Temporal association areasLeft Isocorte x
147.Left Perirhinal area4.662/8.128Left Isocorte
148.Left Ectorhinal area5.197/7.544Left Isocorte
149.Left Specific Thalamus to Frontal pole, cerebral c.Left Specific Thalam us3105.630/3.463
150.Left Specific Thalamus to Primary motor areaLeft Specific Thalam

interacting_witn_Allen
us3105.595/3.371 151.Left Specific Thalamus to Secondary motor areaLeft Specific Thalam
us3105.683/3.436
152.Left Specific Thalamus to Primary somatosensory ar.Left Specific Thalam
us3105.521/3.421 153.Left Specific Thalamus to Primary somatosensory ar.Left Specific Thalam
us3105.432/3.282
154.Left Specific Thalamus to Primary somatosensory ar.Left Specific Thalam us3105.707/3.413
155.Left Specific Thalamus to Primary somatosensory ar.Left Specific Thalam
us3105.615/3.275 156.Left Specific Thalamus to Primary somatosensory ar.Left Specific Thalam
us3105.031/3.433 157.Left Specific Thalamus to Primary somatosensory ar.Left Specific Thalam
us3105.716/3.416
158.Left Specific Thalamus to Primary somatosensory ar.Left Specific Thalam us3104.989/3.610
159.Left Specific Thalamus to Supplemental somatosenso.Left Specific Thalam
us3105.943/3.535
us3106.119/3.454
161.Left Specific Thalamus to Visceral areaLeft Specific Thalam us3106.013/3.752
162.Left Specific Thalamus to Dorsal auditory areaLeft Specific Thalam
us3105.627/3.516 163.Left Specific Thalamus to Primary auditory areaLeft Specific Thalam
us3105.444/3.687 164.Left Specific Thalamus to Posterior auditory areaLeft Specific Thalam
us3105.671/3.824
165.Left Specific Thalamus to Ventral auditory areaLeft Specific Thalam us3105.959/3.893
166.Left Specific Thalamus to Anterolateral visual are.Left Specific Thalam us3106.105/3.295
167.Left Specific Thalamus to Anteromedial visual area.Left Specific Thalam
us3106.188/3.445 168.Left Specific Thalamus to Lateral visual areaLeft Specific Thalam
us3105.847/3.224
169.Left Specific Thalamus to Primary visual areaLeft Specific Thalam us3105.694/3.306
170.Left Specific Thalamus to Posterolateral visual ar.Left Specific Thalam us3105.693/3.259
171.Left Specific Thalamus to posteromedial visual are.Left Specific Thalam
us3106.751/3.380
172.Left Specific Thalamus to Laterointermediate areaLeft Specific Thalam us3106.127/3.284
173.Left Specific Thalamus to Postrhinal areaLeft Specific Thalam us3105.538/3.418
174.Left Specific Thalamus to Anterior cingulate area,.Left Specific Thalam
us3105.928/3.951 175.Left Specific Thalamus to Anterior cingulate area,.Left Specific Thalam
us3105.658/4.482
us3106.026/4.259
177.Left Specific Thalamus to Infralimbic areaLeft Specific Thalam us3105.508/4.508
178.Left Specific Thalamus to Orbital area, lateral pa.Left Specific Thalam us3105.556/3.785
179.Left Specific Thalamus to Orbital area, medial par.Left Specific Thalam
us3105.565/4.152 180.Left Specific Thalamus to Orbital area, ventrolate.Left Specific Thalam
us3105.679/3.987

181.Left Specific Thalamus to Agranular insular area, .Left Specific Thalam

```
us......310......6.182/3.616.....
       182.Left Specific Thalamus to Agranular insular area, .Left Specific Thalam
       us......310......5.870/3.474.....
       183.Left Specific Thalamus to Agranular insular area, .Left Specific Thalam
       us......310......5.398/3.870.....
       184.Left Specific Thalamus to Retrosplenial area, late.Left Specific Thalam
       us......310......5.748/3.391.....
       185.Left Specific Thalamus to Retrosplenial area, dors.Left Specific Thalam
       us......310......5.562/3.322.....
       186.Left Specific Thalamus to Retrosplenial area, vent.Left Specific Thalam
       us.......310......5.906/3.641.....
       187.Left Specific Thalamus to Anterior area.....Left Specific Thalam
       us......310......6.028/3.399.....
       188.Left Specific Thalamus to Rostrolateral visual are.Left Specific Thalam
       us......310......5.660/3.239.....
       189.Left Specific Thalamus to Temporal association are.Left Specific Thalam
       us......310......5.679/4.050.....
       190.Left Specific Thalamus to Perirhinal area.....Left Specific Thalam
       us......310......5.154/3.418.....
       191.Left Specific Thalamus to Ectorhinal area.....Left Specific Thalam
       us......310......5.233/3.882.....
       192.Left Olfactory Areas.....Left Olfactory Area
       s.....41043.....39.470/53.840.....
       193.Left Hippocampal Formation.....Left Hippocampal For
       mation....34819.....34.018/48.566......
       194.Left Cortical Subplate.....Left Cortical Subpla
       te.........6669......44.718/44.450......
       195.Left Striatum.....Left Striatu
       196.Left Pallidum.....Left Pallidu
       m......9308......84.995/75.577......
       197.Left Nonspecific Thalamus.....Left Nonspecific Tha
       lamus.....2324......97.939/46.910......
       198.Left Hypothalamus.....Left Hypothalamu
       199.Left Midbrain.....Left Midbrai
       print("SENSORY PATHWAY:\n")
In [23]:
       print_weights_from("Spinal nucleus of the trigeminal", inds["trigeminal"],
                        final_connectivity.region_labels, final_connectivity.weig
       print weights to ("Principal sensory nucleus of the trigeminal", inds ["ponsse
                      final_connectivity.region_labels, final_connectivity.weight
       print_weights_from("Principal sensory nucleus of the trigeminal", inds["pons
                       final_connectivity.region_labels, final_connectivity.weig
       print_weights_to("Ansiform lobule", inds["ansilob"],
                     final connectivity region labels, final connectivity weights
       print("\n\n\nMOTOR PATHWAY:\n")
       print_weights_from("Primary motor area", inds["m1"], final_connectivity.regi
       # print_weights_to("Superior colliculus, motor related", inds["supcol"], fin
       # print_weights_from("Superior colliculus, motor related", inds["supcol"], f
       print_weights_to("Facial motor nucleus", inds["facial"], final_connectivity.
```

SENSORY PATHWAY:

Stronger 90% connections from Right Spinal nucleus of the trigeminal to: RegionWeigh tTotal weights to	
99.Right Medulla	
63.481	
Stronger 90% connections from Left Spinal nucleus of the trigeminal to: RegionWeigh tTotal weights to	
205.Left Medulla	

interacting_with_Allen
49.15874
199.Left Midbrain0.10354
2
201.Left Pons Sensory0.095514
913.202
202.Left Principal sensory nucleus of the trigeminal0.094956
13.14049
98.Right Pons Behavioral0.075833
3115.281
204.Left Pons Behavioral0.070241
1
6.Right Primary somatosensory area, mouth0.066440
102.Right Inferior olivary complex0.047695
2
11.Right Gustatory areas
6
Stronger 90% connections to Right Principal sensory nucleus of the trigemin
al from:
RegionWeigh
t
99.Right Medulla1.9306
3187.547
93.Right Midbrain1.1311
5286.757
97.Right Pons Motor0.9280
4
98.Right Pons Behavioral
199.Left Midbrain
2
205.Left Medulla0.78760
8
204.Left Pons Behavioral
281.4818
203.Left Pons Motor
9
95.Right Pons Sensory0.20954
4
103.Right Cerebellar Cortex
223.8543
198.Left Hypothalamus0.15884
4
91.Right Nonspecific Thalamus0.13078
9
197.Left Nonspecific Thalamus
92.Right Hypothalamus
8
100.Right Spinal nucleus of the trigeminal0.094956
1
201.Left Pons Sensory
3
Stronger 90% connections to Left Principal sensory nucleus of the trigemina
l from:
RegionWeigh
t

203.Left Pons Motor......0.9280 204.Left Pons Behavioral.....0.82995 9.....81.4818..... 93.Right Midbrain.....0.81668 2......81.4818..... 201.Left Pons Sensory......0.20954 209.Left Cerebellar Cortex.....0.16389 197.Left Nonspecific Thalamus.....0.13078 91.Right Nonspecific Thalamus......0.10733 198.Left Hypothalamus......0.095162 8......366.33..... 206.Left Spinal nucleus of the trigeminal.....0.094956 Stronger 90% connections from Right Principal sensory nucleus of the trigem inal to: Region......Weigh t.....Total weights to..... _____ _____ 3......102.516..... 100.Right Spinal nucleus of the trigeminal......0.24210 7......7.41493..... 203.Left Pons Motor.....0.23626 95.Right Pons Sensory......0.12026 3......13.202...... 4......338.668...... 204.Left Pons Behavioral.......................0.053991 103.Right Cerebellar Cortex.....0.049105 8......14.3996.....

98.Right Pons Behavioral0.048010
9115.281
206.Left Spinal nucleus of the trigeminal0.038664
7
Stronger 90% connections from Left Principal sensory nucleus of the trigem:
nal to:
RegionWeigh
tTotal weights to
205.Left Medulla
9
203.Left Pons Motor
3
99.Right Medulla0.33570
8162.776
206.Left Spinal nucleus of the trigeminal0.24210
97.Right Pons Motor
3
201.Left Pons Sensory0.12026
313.202
199.Left Midbrain0.10825
4
7
98.Right Pons Behavioral0.053991
4
209.Left Cerebellar Cortex0.049105
8
204.Left Pons Behavioral
100.Right Spinal nucleus of the trigeminal0.038664
77.41493
Stronger 95% connections to Right Ansiform lobule from:
RegionWeigh tTotal weights from
103.Right Cerebellar Cortex0.072371
423.8543
105.Right Cerebellar Nuclei0.050667
1
7
97.Right Pons Motor
999.5803
99.Right Medulla0.0070936
4
1
93.Right Midbrain0.003581
7286.757
87.Right Hippocampal Formation0.0024682
148.5659
YX.Right Pons Rehavioral
•
8
98.Right Pons Behavioral0.0017793

Left Pons Behavioral0.001391	
Left Cerebellar Cortex	
23.8543	
ight Pons Sensory0.00098330	
Left Medulla0.00093713	
Left Hypothalamus0.00085860	
ight Principal concern ruleus of the tripopinal (2.0064501	
ight Principal sensory nucleus of the trigeminal0.00064591	
Right Spinal nucleus of the trigeminal0.00055884	
nger 95% connections to Left Ansiform lobule from:	
onWeigh	
Total weights from	
Left Cerebellar Cortex0.072371	
23.8543	
Left Cerebellar Nuclei	
ight Pons Motor0.034214	
Left Medulla0.0070936	
ight Midbrain0.0043502	
Left Midbrain0.003581	
Left Hippocampal Formation	
48.5659	
Left Pons Behavioral0.0017793	
ight Hippocampal Formation0.0017001	
48.5659	
ight Pons Behavioral0.001391	
81.4818	
Right Cerebellar Cortex0.0012268	
Left Pons Sensory	
ight Medulla0.00093713	
ight Hypothalamus0.00085860	
Left Principal sensory nucleus of the trigeminal0.00064591	
Left Principal sensory nucleus of the trigeminal0.00064591	

MOTOR PATHWAY:

Stronger 90% connections from Right Primary motor area to:

t	Danidari Madala
44.Right Specific Thalamus to Primary motor area	RegionWeigh
2	
2	44.Right Specific Thalamus to Primary motor area0.36461
4	2
32.Right Agranular insular area, dorsal part	6.Right Primary somatosensory area, mouth0.083217
8	
11.Right Gustatory areas	· · · · · · · · · · · · · · · · · · ·
92.Right Hypothalamus	11.Right Gustatory areas0.080676
6	53.481
88.Right Cortical Subplate	
8	
4	844.7184
91.Right Nonspecific Thalamus	•
3	
12.Right Visceral area	•
2.Right Secondary motor area	12.Right Visceral area0.058169
2	
3.Right Primary somatosensory area, nose	
33.Right Agranular insular area, posterior part	3.Right Primary somatosensory area, nose0.054398
4	51.19654
107.Left Primary motor area	
3	
4	31.85526
7.Right Primary somatosensory area, upper limb	
2	
3	2
0.Right Frontal pole, cerebral cortex. 0.033272 9	138.Left Agranular insular area, dorsal part0.036374
9	
108.Left Secondary motor area. 0.029602 6	
8.Right Primary somatosensory area, trunk	108.Left Secondary motor area0.029602
89.Right Striatum	
89.Right Striatum	
4.Right Primary somatosensory area, barrel field0.024229 5	89.Right Striatum0.02563
5	398.9072
86.Right Olfactory Areas	
8	
5	839.4701
93.Right Midbrain	
5	
4	5
25.Right Anterior cingulate area, dorsal part0.016782	87.Right Hippocampal Formation0.017704
6	
	6
	117.Left Gustatory areas0.016399
	93.481
	34.Right Agranular insular area, ventral part0.015923
	112.Left Primary somatosensory area, mouth0.015613

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6
2
29.Right Orbital area, lateral part0.013663
94.08821
21.82863
38.Right Anterior area
194.Left Cortical Subplate0.01046
7
94.02148
90.Right Pallidum0.010135
6
6
35.Right Retrosplenial area, lateral agranular part0.0097695
31.Right Orbital area, ventrolateral part0.0096894
24.6111
139.Left Agranular insular area, posterior part0.0095774
Stronger 90% connections from Left Primary motor area to: RegionWeigh
tTotal weights to
150.Left Specific Thalamus to Primary motor area0.36461
25.5953
112.Left Primary somatosensory area, mouth0.083217
112.Left Primary somatosensory area, mouth0.083217 4
112.Left Primary somatosensory area, mouth
112.Left Primary somatosensory area, mouth. 0.083217 4
112.Left Primary somatosensory area, mouth 0.083217 4
112.Left Primary somatosensory area, mouth 0.083217 4
112.Left Primary somatosensory area, mouth 0.083217 4
112.Left Primary somatosensory area, mouth. 0.083217 4
112.Left Primary somatosensory area, mouth 0.083217 4
112.Left Primary somatosensory area, mouth
112.Left Primary somatosensory area, mouth

106.Left Frontal pole, cerebral cortex0.033272
93.1065
2.Right Secondary motor area0.029602
6
114.Left Primary somatosensory area, trunk0.025778
82.31494
195.Left Striatum0.02563
398.9072
110.Left Primary somatosensory area, barrel field0.024229
5
192.Left Olfactory Areas
8
115.Left Primary somatosensory area, unassigned0.021958
5
199.Left Midbrain
5
193.Left Hippocampal Formation0.017704
434.0178
131.Left Anterior cingulate area, dorsal part0.016782
63.35503
11.Right Gustatory areas0.016399
93.481
140.Left Agranular insular area, ventral part0.015923
45.44112
6.Right Primary somatosensory area, mouth0.015613
6
O.Right Frontal pole, cerebral cortex0.013729
23.1065
135.Left Orbital area, lateral part0.013663
94.08821
5.Right Primary somatosensory area, lower limb0.013589
2
144.Left Anterior area0.012594
1
88.Right Cortical Subplate0.01046
7
12.Right Visceral area0.010141
94.02148
196.Left Pallidum
684.9946
143.Left Retrosplenial area, ventral part0.0099800
6
141.Left Retrosplenial area, lateral agranular part0.0097695
3
137.Left Orbital area, ventrolateral part0.0096894
24.6111
33.Right Agranular insular area, posterior part0.0095774
75.00549
Stronger 90% connections to Right Facial motor nucleus from:
RegionWeigh
tTotal weights from
99.Right Medulla3.3347
8187.547
205.Left Medulla
9
97.Right Pons Motor0.80366
799.5803
93.Right Midhrain

5
3
Stronger 90% connections to Left Facial motor nucleus from: RegionWeigh tTotal weights from
8
9
9
9
9
9
9
9
9

Brain network graph

```
In [24]: # import networkx as nx # https://networkx.org/
         # from mpl toolkits.mplot3d import Axes3D
         # def conn_to_graph(conn, major_structs_labels, voxel_count):
               G = nx.DiGraph()
               # G.add edge('A', 'B', weight=4)
         #
         #
               nodes = []
         #
               for iR, (reg, centre, msl, vc) in enumerate(zip(conn.region labels, co
         #
                                                                 major_structs_labels,
         #
                    nodes.append(reg)
         #
                   G.add node(nodes[-1], ind=iR, centre=centre, major structure label
         #
               for iR1, reg1 in enumerate(nodes):
         #
                    for iR2, reg2 in enumerate(nodes):
         #
                       G.add_edge(reg1, reg2,
                                   weight=conn.weights[iR2, iR1],
                                   distance=1.0/conn.weights[iR2, iR1] if conn.weights
         #
         #
                                   tract_length=conn.tract_lengths[iR2, iR1])
               return G
         # def plot G conn 3D(G, nodes kwgs={'s': 100, 'ec':"w"},
                               edges_kwgs={'color':"tab:gray", 'alpha': 0.01}, show=Tr
               node_centres = nx.get_node_attributes(G, "centre")
               # 3d layout
               pos = nx.rescale_layout_dict(node_centres)
               # Extract node and edge positions from the layout
         #
               node_xyz = np.array([pos[v] for v in sorted(G)])
               edge_xyz = np.array([(pos[u], pos[v]) for u, v in G.edges()])
               # Create the 3D figure
               fig = plt.figure(figsize=(15, 15))
         #
               ax = fig.add_subplot(111, projection="3d")
               # Plot the nodes - alpha is scaled by "depth" automatically
         #
               ax.scatter(*node_xyz.T, **nodes_kwgs)
               # Plot the edges
         #
               for vizedge in edge xyz:
                   ax.plot(*vizedge.T, **edges kwgs)
         #
               def _format_axes(ax):
                   """Visualization options for the 3D axes."""
         #
                   # Turn gridlines off
         #
                   ax.grid(False)
                   # Suppress tick labels
```

#

for dim in (ax.xaxis, ax.yaxis, ax.zaxis):

```
#
                        dim.set ticks([])
         #
                   # Set axes labels
         #
                   ax.set xlabel("x")
         #
                   ax.set ylabel("y")
                   ax.set zlabel("z")
               format axes(ax)
         #
               fig.tight layout()
         #
               if show:
         #
                   plt.show()
               return fig, ax
In [25]: # G = conn_to_graph(final_connectivity, final_major_structs_labels, final_vo
         # source = "Right Spinal nucleus of the trigeminal"
         # target = 'Right Ansiform lobule'
         # print(G.nodes())
         # sp = nx.shortest path(G, source, target, weight='distance') # distance =
         # print("Shortest Path:\n", sp)
         # spl = nx.shortest_path_length(G, source, target, weight='distance')
         # print("Shortest Path Length:\n", spl)
         # w = 1.0 / spl / (len(sp) - 1)
         # print("Shortest Path Weight = 1.0 / ShortestPathLength / (len(ShortestPath
         # # For interactive plotting:
         # %matplotlib notebook
         # plot G conn 3D(G)
In [26]: # # Summarize also the Cerebellar Cortices and Nuclei:
         # major structures to merge = ['Cerebellar Cortex', "Cerebellar Nuclei"]
         # exclude_regions={}
         # # For bilateral merging:
         # major structures to merge, exclude regions = unilateral to bilateral(major
         # merged_cereb_conn, merged_cereb_major_structs_labels, merged_cereb_voxel_d
               merge_major_structures(final_connectivity,
         #
                                       major_structures_to_merge,
         #
                                       final_major_structs_labels, final_voxel_count,
         #
                                       exclude regions=exclude regions, weight fun=np.
         \# inds = \{\}
         # inds["crtx"] = []
         # inds["m1"] = []
         # inds["s1brl"] = []
         # inds["supcol"] = []
         # inds["facial"] = []
         # inds["trigeminal"] = []
         # inds["ponssens_trigeminal"] = []
         # inds["ponssens"] = []
         # inds["ponsmotor"] = []
         # inds["thal"] = []
```

```
# inds["thalspec"] = []
         # inds["m1thal"] = []
         # inds["s1brlthal"] = []
         # inds["cerebcrtx"] = []
         # inds["cerebnucleus"] = []
         # inds["oliv"] = []
         # for iR, (reg, lbl) in enumerate(zip(merged_cereb_conn.region_labels, merge
               if "Isocortex" in lbl:
         #
                   inds["crtx"].append(iR)
               if "Primary motor area" in reg and "Specific" not in reg:
         #
         #
                   inds["m1"].append(iR)
               if "barrel" in reg and "Specific" not in reg:
         #
         #
                   inds["s1brl"].append(iR)
         #
               if "Superior colliculus, motor related" in reg:
                   inds["supcol"].append(iR)
         #
         #
               if "Primary motor area" in reg and "Specific" in reg:
         #
                   inds["m1thal"].append(iR)
         #
               if "barrel" in reg and "Specific" in reg:
         #
                   inds["s1brlthal"].append(iR)
         #
               if "Superior colliculus, motor related" in reg:
                   inds["supcol"].append(iR)
         #
               if "Facial" in reg:
         #
         #
                   inds["facial"] .append(iR)
         #
               if "Spinal nucleus of the trigeminal" in reg:
                   inds["trigeminal"].append(iR)
         #
               if "Principal sensory nucleus of the trigeminal" in reg:
         #
         #
                   inds["ponssens_trigeminal"] .append(iR)
         #
               if "Pons Sensory" in lbl:
                   inds["ponssens"] .append(iR)
         #
         #
               if "Pons Motor" in lbl:
         #
                   inds["ponsmotor"].append(iR)
               if "Nonspecific Thalamus" in lbl:
         #
         #
                   inds["thal"].append(iR)
               if "Specific Thalamus" in lbl:
         #
         #
                   inds["thalspec"].append(iR)
         #
               if "Cerebellar Cortex" in reg:
                   inds["cerebcrtx"].append(iR)
         #
         #
               if "Cerebellar Nuclei" in reg:
                   inds["cerebnucleus"].append(iR)
               if "Inferior olivary complex" in reg:
         #
                   inds["oliv"].append(iR)
         #
         # inds["mls1brl"] = inds["m1"] + inds["s1brl"]
         # inds["sens"] = inds["trigeminal"] + inds["ponssens_trigeminal"] + inds["s
         # inds["motor"] = inds["m1thal"] + inds["facial"] # inds["supcol"] +
         # inds["cereb"] = inds["cerebcrtx"] + inds["cerebnucleus"] + inds["oliv"]
         # allspecial = inds["m1s1brl"] + inds["motor"] + inds["sens"] + inds["thal"]
         # for key, val in inds.items():
               inds[key] = np.array(val)
         # print(merged_cereb_conn.region_labels[allspecial])
In [27]: # # Write the resulting connectivity to .h5 TVB compatible file...
         # write_connectivity_to_h5_file(merged_cereb_conn, "Connectivity_SummedSubco
         # # ... along with the corresponding mapping from regions to major structure
         # np.save("major_structs_labels_SummedSubcortical_Thals_SummedCereb.npy", me
         # np.save("voxel_count_SummedSubcortical_Thals_SummedCereb.npy", merged_cere
         # np.save("inds SummedSubcortical Thals SummedCereb.npy", inds)
In []:
```