Daducci, A., Gerhard, S., Griffa, A., Lemkaddem, A., Cammoun, L., Gigandet, X., Meuli, R., Hagmann, P., Thiran, J.-P., 2012. The connectome mapper: an open-source processing pipeline to map connectomes with mri. PLoS ONE 7 (12), e48121.

Dall, J., Christensen, M., 2002. Random geometric graphs. Physical Review E 66 (1), 016121.

de Reus, M. A., van den Heuvel, M. P., 2014. Simulated rich club lesioning in brain networks: a scaffold for communication and integration? Frontiers in Human Neuroscience 8, 647.

Destrieux, C., Fischl, B., Dale, A., Halgren, E., 2010. Automatic parcellation of human cortical gyri and sulci using standard anatomical nomenclature. Neuroimage 53 (1), 1–15.

Ercsey-Ravasz, M., Markov, N. T., Lamy, C., Van Essen, D. C., Knoblauch, K., Toroczkai, Z., Kennedy, H., 2013. A predictive network model of cerebral cortical connectivity based on a distance rule. Neuron 80 (1), 184–197.

Fortunato, S., 2010. Community detection in graphs. Physics Reports 486 (3), 75–174.

Friedman, E. J., Landsberg, A. S., Owen, J., Hsieh, W., Kam, L., Mukherjee, P., 2015. Edge correlations in spatial networks. Journal of Complex Networks, cnv015.

Glasser, M. F., Sotiropoulos, S. N., Wilson, J. A., Coalson, T. S., Fischl, B., Andersson, J. L., Xu, J., Jbabdi, S., Webster, M., Polimeni, J. R., et al., 2013. The minimal preprocessing pipelines for the human connectome project. Neuroimage 80, 105–124.