

Accepted Manuscript

Spatial analysis and high resolution mapping of the human whole-brain transcriptome for integrative analysis in neuroimaging

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PII: S1053-8119(18)30388-4

DOI: [10.1016/j.neuroimage.2018.04.068](https://doi.org/10.1016/j.neuroimage.2018.04.068)

Reference: YNIMG 14915

To appear in: *NeuroImage*

Received Date: 26 January 2018

Revised Date: 30 March 2018

Accepted Date: 29 April 2018

Please cite this article as: Gryglewski, G., Seiger, René., James, G.M., Godbersen, G.M., Komorowski, A., Unterholzner, J., Michenthaler, P., Hahn, A., Wadsak, W., Mitterhauser, M., Kasper, S., Lanzenberger, R., Spatial analysis and high resolution mapping of the human whole-brain transcriptome for integrative analysis in neuroimaging, *NeuroImage* (2018), doi: 10.1016/j.neuroimage.2018.04.068.

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For submission to

NeuroImage

Original Article		
Word count	250	Abstract
	5161	Article Body
Tables: 2, figures: 4, references: 41		
Supplementary figures: 1; tables: 2		

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