



Connectome based classification of BDNF Met allele carriers

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WHAT IS BDNF?

Brain-derived neurotrophic factor

Protein essential for brain development &
 long term potentiation
 Patterson SL, 1996, Neuron

Associated gene, **BDNF**

- Regulates the activity-dependent release of the
 BDNF protein
 Egan MF, 2003, Cell
- Alleles: **G** encodes *Valine*, **A** encodes *Methionine*
- Common single-nucleotide polymorphism (SNP): Val66Met = swap Val for Met at codon 6,6

SO WHAT?

- Carrying MET allele
 - reduced BDNF activity-dependent secretion in the brain

Egan MF, 2003, Cell

• BDNF role

pruning of silent axonal branches during brain development

> Singh KK, 2008, Nat Neurosci Cao L, 2007, Curr Biol



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STUDY DATA

- Population (n=36)
 - 15 Met carriers, 21 Val homozygotes
 - Highly regulated young healthy subjects (18-25y)
- MR acquisition
 - High-resolution T1 MPRAGE (1mm³ isotropic voxels)
 - Diffusion-weighted MRI
 - 7 unweighted (b=100) images
 - 61 directional (b=1000) images
 - 2.3 x 2.3 x 2.3 mm³ isotropic voxels

- Probabilistic Tractography
 - Spherical deconvolution \rightarrow ODFs
 - 300,000 fibers, randomly seeded in WM

Tournier JD, 2004 & 2007, Neuroimage



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- Anatomical segmentation & parcellation
 - Desikan-Killiany atlas (83 regions)
 - Lausanne 1015-region atlas

Desikan RS, 2006, *Neuroimage* Hagmann P, 2008, *PLoS Biol* Daducci A, 2012, *PLoS One*

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Connectome = 1015 x 1015 matrix
 − 1 fiber crossing 2 regions → +1 connection

STATISTICAL ANALYSIS

Compare connectivity matrices with

Network-based statistics (NBS)

Non-parametric test using NBS

Zalesky A, 2010, Neuroimage

NBS RESULTS

Increased fibers in Met carriers

NBS RESULTS

Fibers not present in Val homozygotes

NBS RESULTS

- Some connections only in Met carriers
- Increased connection strength in Met carriers:
 - between bilateral thalami and brainstem
 - sensorimotor areas of parietal and frontal cortex
 - ventromedial prefrontal cortex (anterior forceps)
 - occipital, posterior parietal, and temporal areas also differ to a lesser extent.

STATISTICAL ANALYSIS

Compare connectivity matrices with

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Non-parametric test using NBS

Zalesky A, 2010, Neuroimage

- Multivariate statistics
 - Gaussian Processes classification
 - Leave-one-out cross-validation procedure
 - Permutation testing

Rasmussen CE, 2006 Schrouff J, 2013, *Neuroinformatics*

CLASSIFICATION RESULTS

- Balanced accuracy: 87.1% (p<.001)
- Class predictive values (ValVal & Met-carrier): 94.4% & 77.8%

CLASSIFICATION RESULTS

Grouping according to

 adenosine deaminase (ADA) genotype: Classification of G/A (n=16) vs. G/G (n=20)

- accuracy 58.3% (n.s.)
- gender:

Classification of male (n=18) vs. female (n=18)

→ accuracy 63.9% (n.s.)

No significant classification for ADA nor gender!

DISCUSSION

- BDNF Met carriers prune less axons during brain development
- Resulting tracts stay in the brain

Seem to provide little or no benefit at young age

May protect against age-related deficits

Harris SE, 2006, Mol Psychiatry Gajewski PD, 2011, Neurobiol Aging

• Extraneous tracts are found as increases in fractional anisotropy
Chiang MC, 2011a and 2011b, Neuroimage

- BDNF genotype (allelic group) can be predicted based on brain connectivity
- Longitudinal BDNF studies are needed to cover youth-adult brain development
- Differences between Val66Met and Met66Met polymorphisms should be examined

ACKNOWLEDGMENTS

Open-source software:

- Nipype/Nibabel/Dipy http://nipy.org
- Mrtrix http://www.brain.org.au/software/mrtrix/
- ConnectomeViewer http://cmtk.org
- PRoNTo http://www.mlnl.cs.ucl.ac.uk/pronto/

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REFERENCES

- Cao L, et al. (2007) Genetic modulation of BDNF signaling affects the outcome of axonal competition in vivo. Curr Biol 17: 911–921.
- Chiang MC, et al. (2011) BDNF gene effects on brain circuitry replicated in 455 twins. Neuroimage 55: 448–454.
- Chiang MC, et al. Genetics of white matter development: A DTI study of 705 twins and their siblings aged 12 to 29. *NeuroImage* **54**, 2308–2317 (2011).
- Daducci A, et al, .(2012) The connectome mapper: an open-source processing pipeline to map connectomes with MRI. PLoS One 7: e48121.
- Egan MF, et al. (2003) The BDNF val66met polymorphism affects activity-dependent secretion of BDNF and human memory and hippocampal function. Cell 112: 257–269.
- Gorgolewski K, et al. (2011) Nipype: a flexible, lightweight and extensible neuroimaging data processing framework in python. Front Neuroinform 5: 13.
- Hagmann P, Cammoun L, Gigandet X, Meuli R, Honey CJ, et al. (2008) Mapping the structural core of human cerebral cortex. PLoS Biol 6: e159.
- Patterson SL, et al. (1996) Recombinant BDNF rescues deficits in basal synaptic transmission and hippocampal LTP in BDNF knockout mice. Neuron 16: 1137–1145.
- Rasmussen CE & Williams C (2006), Gaussian Processes for Machine Learning, the MIT Press.
- Schrouff J, et al. (2013) Pronto: Pattern recognition for neuroimaging toolbox. Neuroinformatics .
- Singh KK, et al. (2008) Developmental axon pruning mediated by BDNF-p75NTR-dependent axon degeneration. Nat Neurosci 11: 649–658.
- Zalesky A, Fornito A, Bullmore ET (2010) Network-based statistic: identifying differences in brain networks. Neuroimage 53: 1197–1207.

Poster #3840, Wednesday 13:30-15:30. Paper recently accepted in PLOS ONE.

Thank you for your attention!

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