



Application of neuroimaging to understand rare genetic disorders

Dr Kate Baker, Programme Leader and Honorary Consultant

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Outline

- 1. Introduction to an applied neuroimaging challenge
- 2. Discussion of opportunities and constraints
- 3. 1 project, 12 years (other examples exist)
- 4. Ideas for possible future directions





1. Introduction





Intellectual disability



DSM-V (APA, 2013)

- 1. Cognitive impairments
- 2. Adaptive functioning
- 3. Onset during development

Typically IQ < 70 = 2.5% (1-3%)



Intellectual disability



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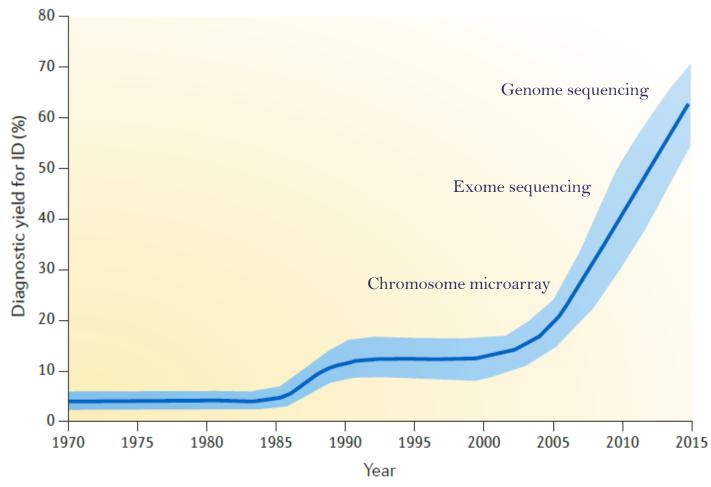


- Developmental trajectory
- Associated characteristics
 - Neurological heath
 - Physical health
 - Mental health
- Neurobiology
- Social and cultural context
- Aetiology





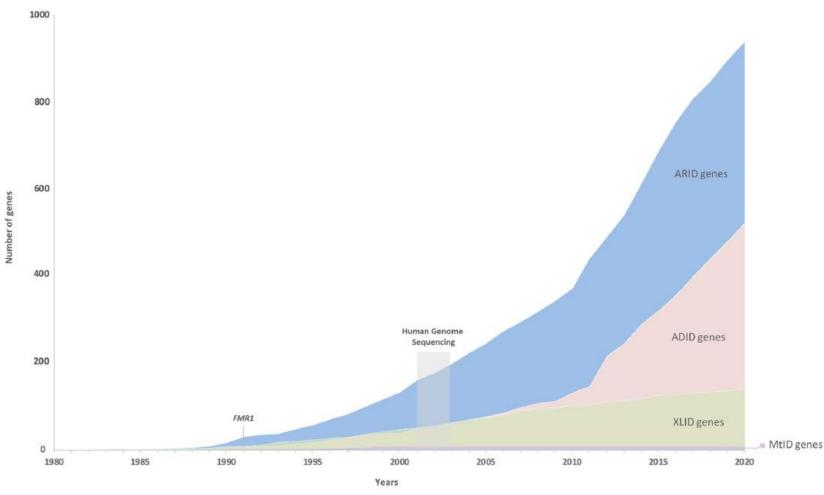
Genetic Diagnosis in ID – past, present...







Genetic Diagnosis in ID – past, present...







Opportunity



DSM-V (APA, 2013)

- 1. Cognitive impairments
- 2. Adaptive functioning
- 3. Onset during development Typically IQ < 70 = 2.5% (1-3%)



- Cognitive profile
- Adaptive impact
- Developmental trajectory
- Associated characteristics
 - Neurological heath
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- Neurobiology
- Social and cultural context
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2. Discussion

- A. What are the cognitive neuroscience questions arising from genetic diagnosis in ID?
- B. How can neuroimaging address these questions?
- C. What are the constraints?





3. One example

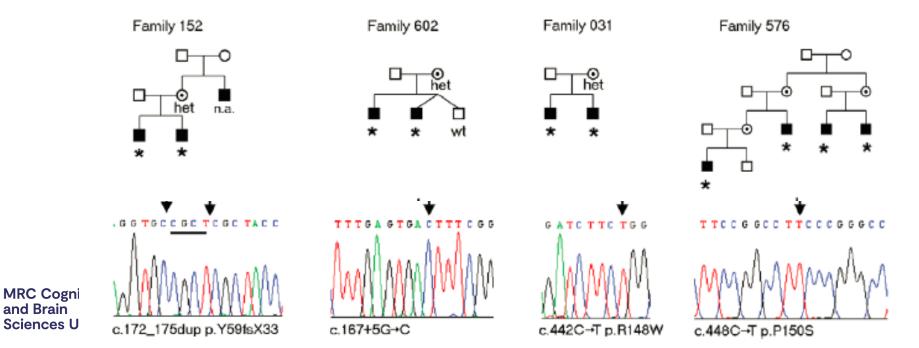




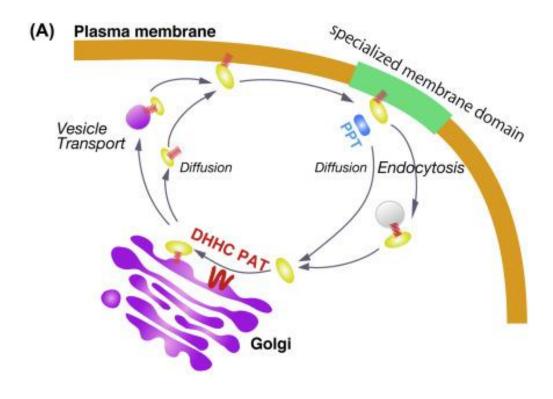
REPORT

Mutations in *ZDHHC9*, Which Encodes a Palmitoyltransferase of NRAS and HRAS, Cause X-Linked Mental Retardation Associated with a Marfanoid Habitus

F. Lucy Raymond,* Patrick S. Tarpey,* Sarah Edkins, Calli Tofts, Sarah O'Meara, Jon Teague, Adam Butler, Claire Stevens, Syd Barthorpe, Gemma Buck, Jennifer Cole, Ed Dicks, Kristian Gray, Kelly Halliday, Katy Hills, Jonathon Hinton, David Jones, Andrew Menzies, Janet Perry, Keiran Raine, Rebecca Shepherd, Alexandra Small, Jennifer Varian, Sara Widaa, Uma Mallya, Jenny Moon, Ying Luo, Marie Shaw, Jackie Boyle, Bronwyn Kerr, Gillian Turner, Oliver Quarrell, Trevor Cole, Douglas F. Easton, Richard Wooster, Martin Bobrow, Charles E. Schwartz, Jozef Gecz, Michael R. Stratton, and P. Andrew Futreal



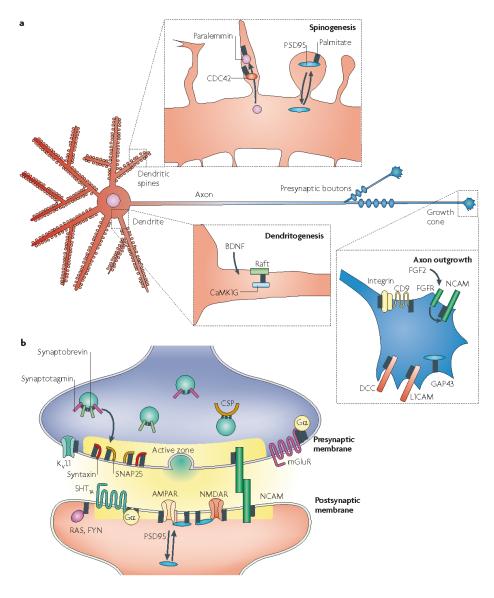
ZDHHC9 functions



Fukata, Yuko, Murakami, Tatsuro, Yokoi, Norihiko, & Fukata, Masaki. (2016). Current Topics in Membranes (Vol. 77, Dynamic Plasma Membranes - Portals Between Cells and Physiology). Elsevier.

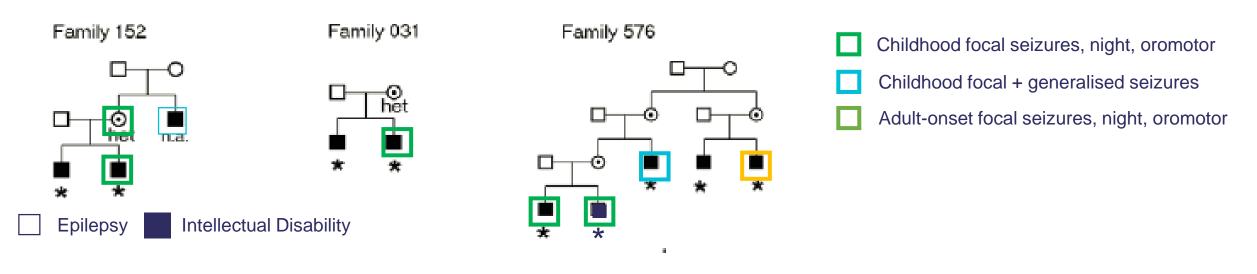


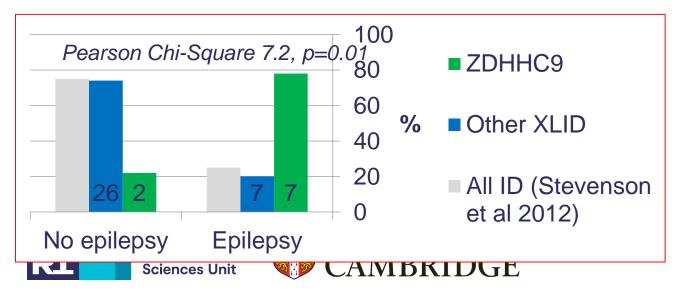


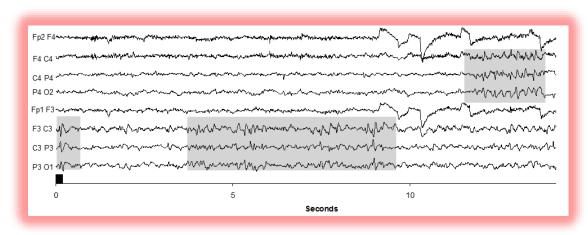


Fukata Y1, Fukata M. (2010) Nat Rev Neurosci. Protein palmitoylation in neuronal development and synaptic plasticity.

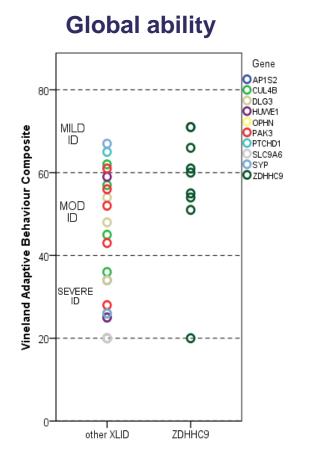
ZDHHC9-associated XLID: Neurology

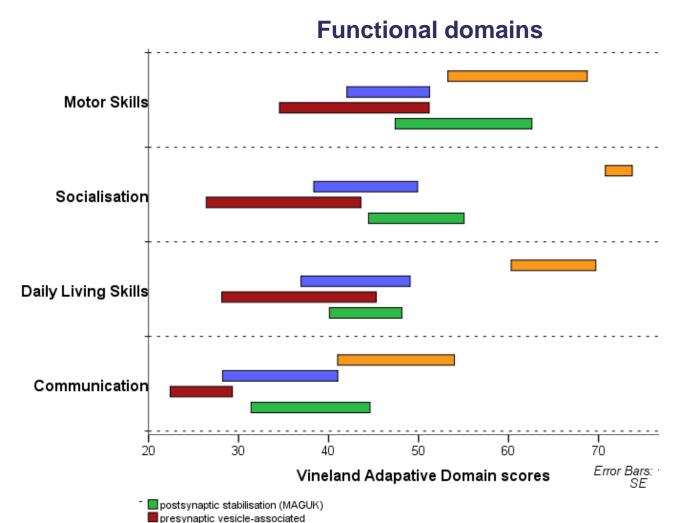






ZDHHC9-associated XLID: Adaptive function





☐ protein degradation (ubiquitin E3 ligase) ☐ postranslational modification (palmitoylation)

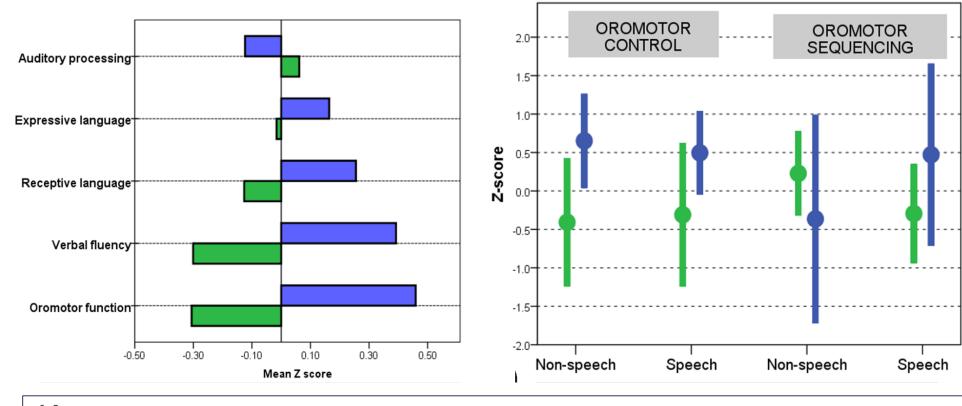




ZDHHC9-associated XLID: Language tests

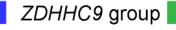
Standardised language test battery

Verbal Motor Production Assessment









Error bars: 95% confidence interval



ZDHHC9-associated XLID RE and DLD

What neuroimaging questions would you ask?





ZDHHC9-associated XLID RE and DLD

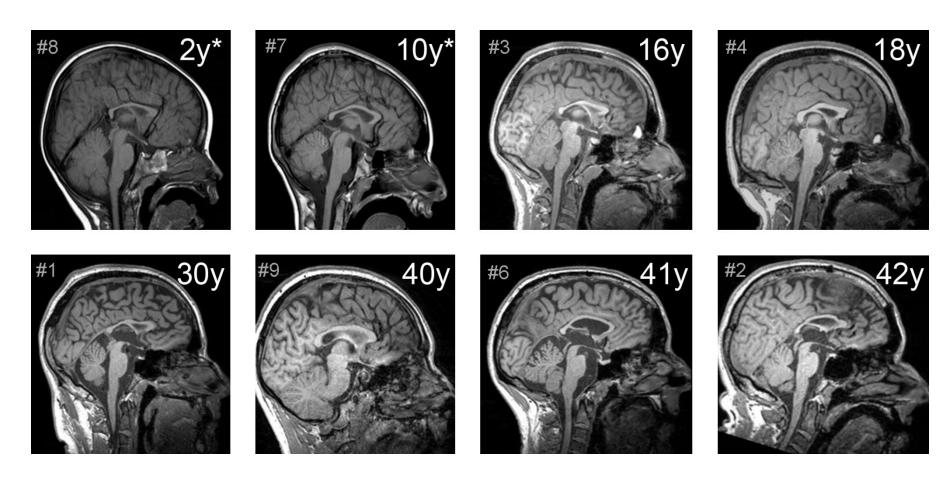
How is brain development structure and function altered?
How does this relate to communication difficulties?
How does this relate to ZDHHC9 expression?

Prediction from RE literature = we won't find anything much





ZDHHC9-associated XLID: Neuroradiology

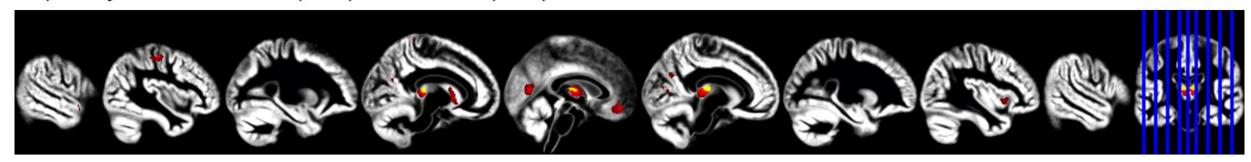


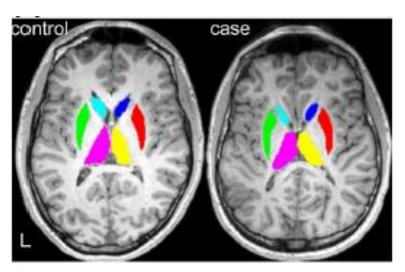


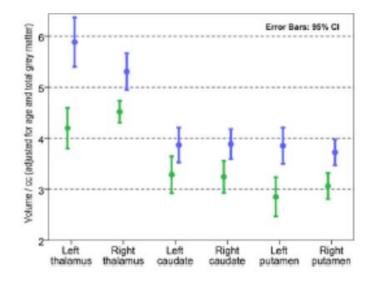


ZDHHC9-associated XLID: VBM

A) Grey matter: cases (n=7) < controls (n=7)











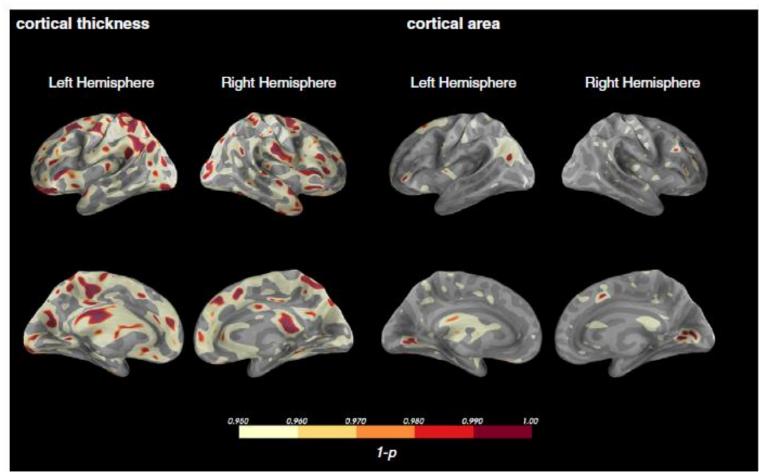


RESEARCH ARTICLE

Epilepsy, cognitive deficits and neuroanatomy in males with *ZDHHC*9 mutations

Kate Baker^{1,2}, Duncan E. Astle², Gaia Scerif³, Jessica Barnes², Jennie Smith⁴, Georgina Moffat⁴, Jonathan Gillard⁵, Torsten Baldeweg⁶ & F. Lucy Raymond¹

ZDHHC9-associated XLID: Cortical morphometry







Neurolmage: Clinical

journal homepage: www.elsevier.com/locate/ynicl



Structural brain abnormalities in a single gene disorder associated with epilepsy, language impairment and intellectual disability



Joe Bathelt^{a,*}, Duncan Astle^a, Jessica Barnes^a, F. Lucy Raymond^b, Kate Baker^{a,b}

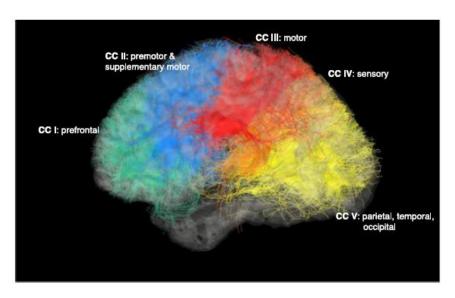
*MRC Cognition & Brain Sciences Unit, Cambridge, United Kingdom

^bDepartment of Medical Genetics, Cambridge Institute for Medical Research, University of Cambridge, Cambridge, United Kingdom





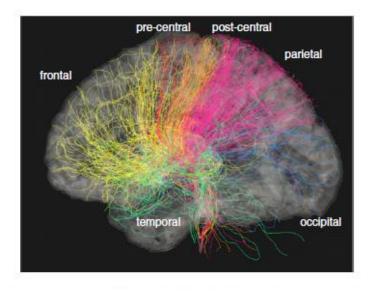
ZDHHC9-associated XLID: tractography



Cortico-spinal
Tract

A

Uncinate
Fasciculus



Reduced FA, increased MD / RD: CC I, CC II, **CC III**

Reduced FA, increased MD / RD: Right and left arcuate fasciculus Right and left uncinate fasciculus No differences in CST

Reduced FA, increased MD / RD: Right and left precentral and temporal thalamic projections only









Structural brain abnormalities in a single gene disorder associated with epilepsy, language impairment and intellectual disability



Joe Bathelt^{a,*}, Duncan Astle^a, Jessica Barnes^a, F. Lucy Raymond^b, Kate Baker^{a,b}

^aMRC Cognition & Brain Sciences Unit, Cambridge, United Kingdom ^bDepartment of Medical Genetics, Cambridge Institute for Medical Research, University of Cambridge, Cambridge, United Kingdom

ZDHHC9-associated XLID: Structural connectome

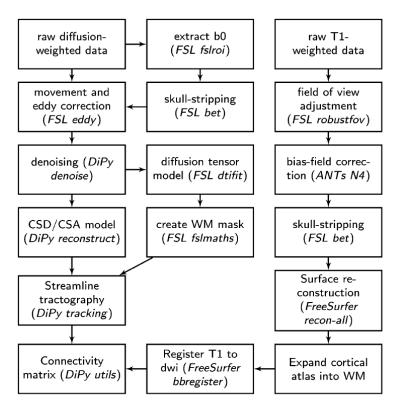
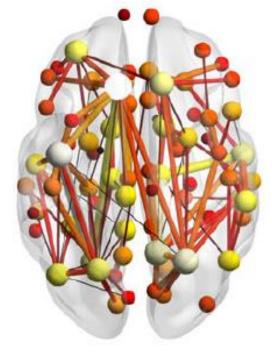
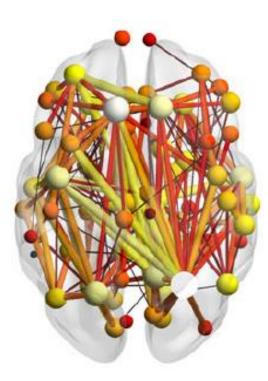


Figure 1. Overview of the processing steps to derive the diffusion-weighted structural connectome.





control



Cerebral Cortex, July 2017;27: 3806-3817

doi: 10.1093/cercor/bhx027 Advance Access Publication Date: 7 February 201 Original Article





ORIGINAL ARTICLE

Global and Local Connectivity Differences Converge With Gene Expression in a Neurodevelopmental Disorder of Known Genetic Origin

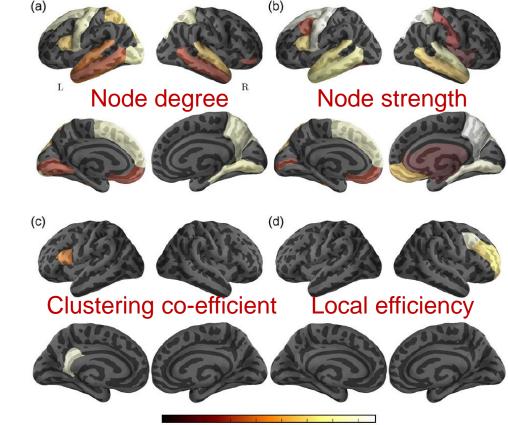
Joe Bathelt¹, Jessica Barnes¹, F Lucy Raymond², Kate Baker^{1,2,†} and Duncan Astle^{1,†}

ZDHHC9-associated XLID: Structural connectome

3814 | Cerebral Cortex, 2017, Vol. 27, No. 7

Global reductions in all FA-based graph metrics

- Node degree
- Node strength
- Clustering co-efficient
- Local efficiency





- p





ZDHHC9-associated XLID RE and DLD

How is brain development structure and function altered?
How does this relate to communication difficulties?
How does this relate to ZDHHC9 expression?

Prediction from RE literature = we won't find anything much

- Subcortical and CC volume reductions
- Cortical thickness reductions
- Extensive reductions in WM integrity
- Connectomic differences converging with typical ZDHHC9 expression





ZDHHC9-associated XLID RE and DLD

How is brain development structure altered?

- Subcortical and CC volume reductions
- Cortical thickness and WM integrity reductions
- Connectomic differences converging with ZDHHC9 expression
- Small case control study = multiple observations on small n with wide age range
- Rarity = impossible to replicate? Adding more subjects would be non-independent, biased.
- Multiple specificity problems
 - Comparing to high IQ group, not low IQ, language disordered, other genes, RE general, RE other causes...
 - Genotype vs phenotype
- Structure-function relationships are opaque
 - Cannot separate cause from consequence (language acquisition vs skill; language vs other abilities)
- Cross-disorder case control designs?
- Within-sample dimensional and brain-cognition analyses?
- Integrate with developmental cohort data?





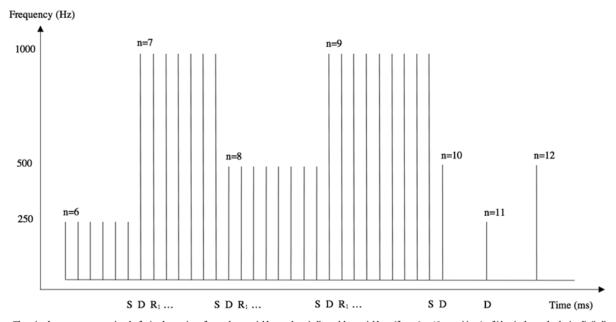
ZDHHC9-associated XLID: MEG







- 1. Resting state acquisition (2 x 6 minutes)
- 2. Passive auditory roving oddball acquisition (2 x 6 minutes)



The stimulus sequence consisted of stimulus trains of a random variable number, indicated by variable n (from 6 to 12 repetitions) of identical standard stimuli (indicated by vertical lines) within trains. The frequency varied randomly (between 250Hz, 500 and 1000Hz) from train to train, as indicated by the different height of the vertical lines. Tone lengths were 50ms, with inter-tone intervals of 500ms. S = preceding stimulus to deviant. D = Deviant. R1= Repeat 1.





ZDHHC9-associated XLID: MEG networks and HMM

Spatial maps of each Temporal statistics of Raw MEG data each state state acquisition Partial correlation of state State timecourses of when · 9 minute resting state timecourses onto parcel each state is 'active' used to 12 minute auditory oddball amplitude envelopes quantify state dynamics SPM12 and Workbench v1.2.3 GLEAN toolbox (OSL) in SPM12 Elekta VectorView MEG Gene expression **Preprocessing** Hidden Markov Model Gene expression level of External noise removal. Group level exploratory ZDHHC9 obtained for each downsampling, and temporal analysis of networks to infer 8 parcel, and weighted by the ICA to remove oculomotor states, from unique patterns of normalized activity level within and cardiac artefacts group-level covariance over each parcel the 38 amplitude envelopes Allen Brain Atlas, MATLAB Maxfilter v2.1, SPM12 GLEAN toolbox (OSL) in SPM12 Concatenation of Group comparison of Co-registration state temporal statistics participants Co-registration of MEG data to individual TI-weighted Imm Amplitude envelopes Permutation testing to temporally concatenated compare temporal statistics MRI scans for each state between groups OSL toolbox in SPM12 MATLAB OSL toolbox in SPM12 Test ZDHHC9 gene **Beamformer** Amplitude envelope expression in each state Lowpass filtering at 4-30Hz estimation Permutation testing for high Source reconstruction usine Estimate amplitude envelopes

of 38 parcel-wise oscillatory

timecourses using a Hilbert

transform

RESEARCH ARTICLE

Functional network dynamics in a neurodevelopmental disorder of known genetic origin

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Erin Hawkins<sup>1</sup> | Danyal Akarca<sup>1</sup> | Mengya Zhang<sup>1</sup> | Diandra Brkić<sup>1</sup> | Mark Woolrich<sup>2</sup> | Kate Baker<sup>1,3</sup> | Duncan Astle<sup>1</sup>
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8mm grid

OSL toolbox in SPM12



levels of ZDHHC9 gene

expression across each state

MATLAB

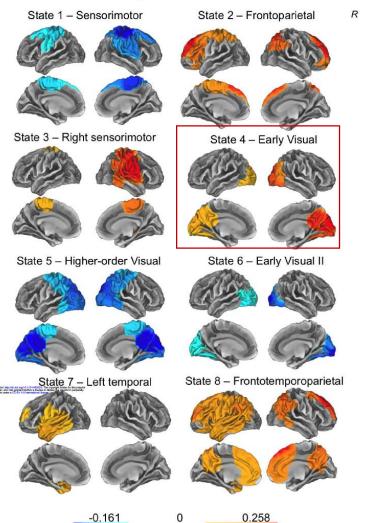
ZDHHC9-associated XLID: MEG networks

Resting state networks

Higher fractional occupancy and mean lifetime in ZDHHC9 subjects

MRC (and Br

Scienc



State 2 - Frontoparietal State 1 - Parietal State 3 - Fronto-occipital State 4 - Frontotemporal State 5 - Right temporoparietal State 6 - Bilateral temporal State 7 - Frontoparietal II State 8 - Fronto-occipital II -0.111

Partial correlation values

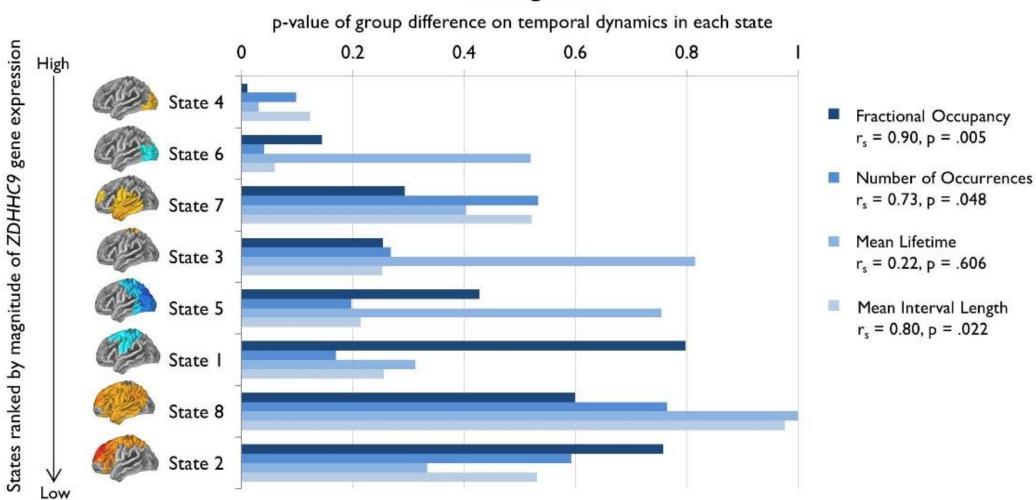
Oddball task networks

Reduced fractional occupancy, number of occurrences and mean interval length



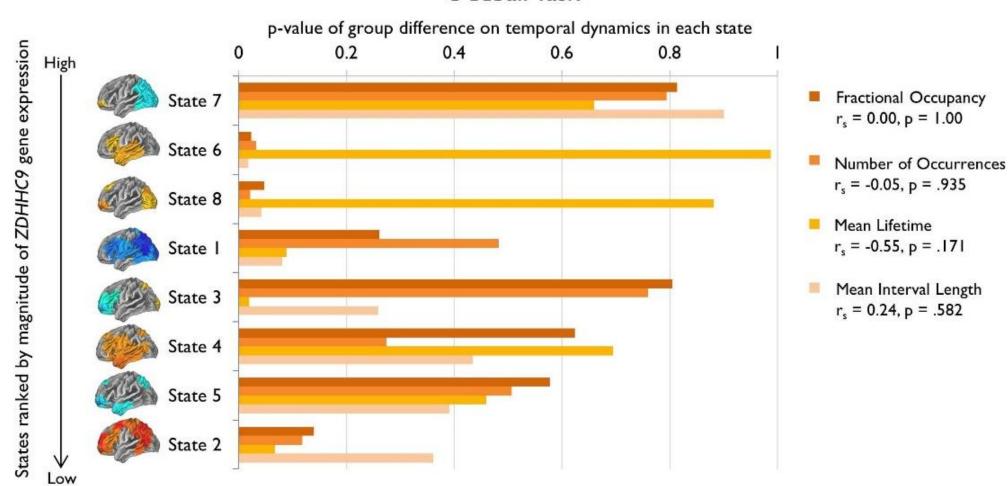
ZDHHC9-associated XLID: MEG networks

Resting State



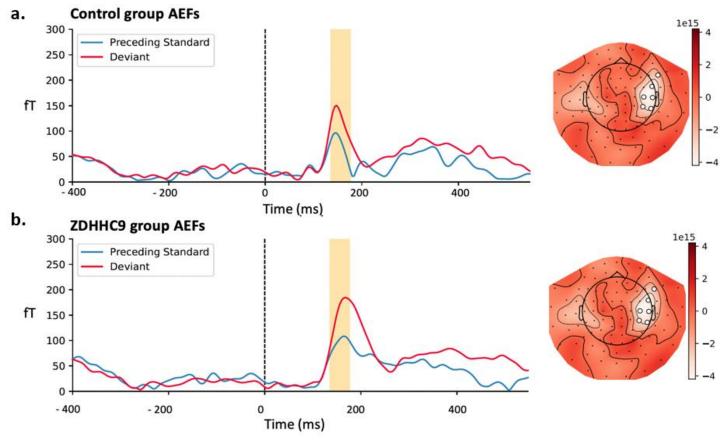
ZDHHC9-associated XLID: MEG networks

Oddball Task





ZDHHC9-associated XLID: MEG AEFs and MMN

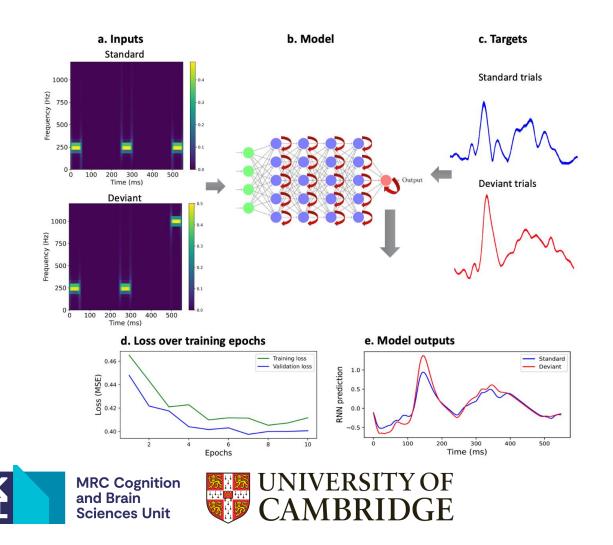


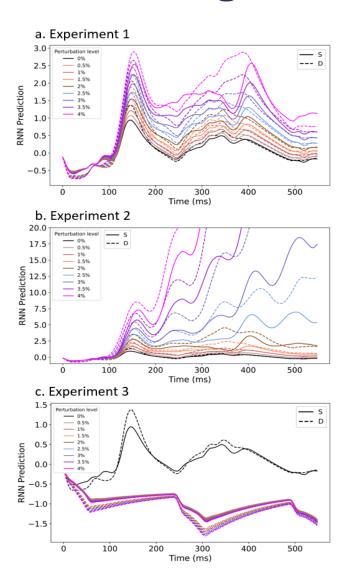






ZDHHC9-associated XLID: E-I RNN modelling





ZDHHC9-associated XLID

How is brain development function altered?

- Same functional networks present in cases / controls
- Resting state limited dynamic differences, correlating with expression topography
- Oddball networks dynamic differences appear phenotype-relevant, RNN model can recapitulate group differences and is sensitive to loss of inhibition
- All the same general limitations as MRI re N and comparison groups
- Dynamic network and RNN analysis is new we don't know what to expect across different developmental conditions





ZDHHC9-associated XLID: progress

How is brain development structure and function altered?

How does this relate to communication difficulties?

How does this relate to ZDHHC9 expression?



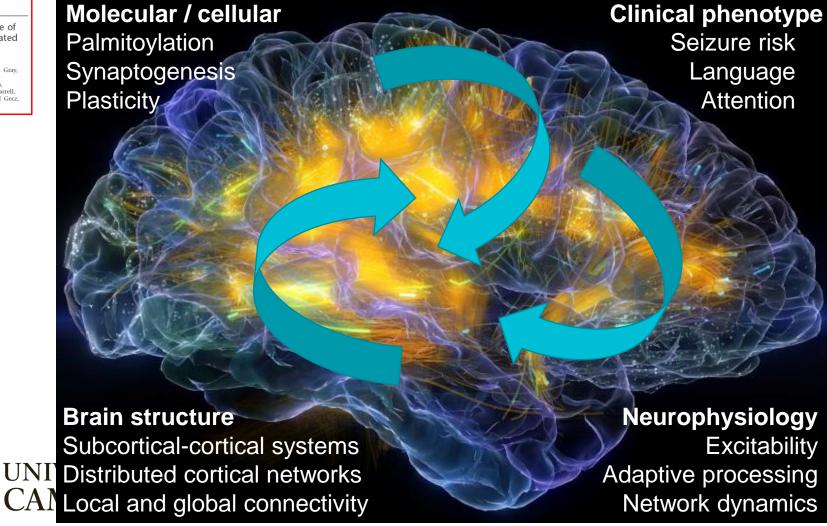


ZDHHC9-associated XLID: progress

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4. Ideas for future directions

How is brain development structure and function altered?

How does this relate to communication difficulties?

How does this relate to ZDHHC9 expression?

How to integrate?

Developmental and cognitive mechanisms?

Clinically useful?







MRC Cognition and Brain **Sciences Unit**









Gaia Scerif, Oxford



Lucy Raymond, UofCam



Torsten Baldeweg, UCL ICH



Elise Ng-Cordell, CBU



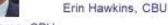


Diandra Brkic, CBU











Joe Bathelt, CBU



Rebeca Ianov, CBU















Would you like to join our team?



- Research assistant position currently advertised on the MRC CBU website
- 2. Open to PhD applications this term for starting in October 2025

