

REGINA H REYNOLDS



KEY PUBLICATIONS

2021

- **Cross-platform transcriptional profiling identifies common and distinct molecular pathologies in Lewy body disorders¹**

Acta Neuropathologica

- Feleke, R*, Reynolds, RH*, Smith, A*, Tilley, B, Gagliano Taliun, SA, Hardy, J, Matthews, PM, Gentleman, S, Owen , D, Johnson, MR, Srivastava, P, Ryten, M
- Role: Co-first author and analyst.
- Transcriptomic analysis of cell-type-specific changes in the Lewy body diseases. An overview of analyses conducted can be viewed here².

2020

- **Dystonia genes functionally converge in specific neurons and share neurobiology with psychiatric disorders³**

Brain

- Mencacci, NE*, Reynolds, RH*, Garcia Ruiz, S, Vandrovčova, J, Forabosco, P, Sánchez-Ferrer, A, Volpato, V, UK Brain Expression Consortium, International Parkinson's Disease Genomics Consortium, Weale, ME, Bhatia, KP, Webber, C, Hardy, J, Botía, JA, Ryten, M
- Role: Co-first author and analyst.
- Integrative omics analysis of monogenic dystonias, with the aim of improving our understanding of the pathways driving this clinically heterogeneous group of movement disorders.

2019

- **Informing disease modelling with brain-relevant functional genomic annotations⁴**

Brain

- Reynolds, RH, Hardy, J, Ryten, M, Gagliano Taliun, SA
- Role: First author.
- Review of conceptual advances in the generation of brain-relevant functional genomic annotations and among tools that allow integration of these annotations with genome-wide association summary statistics.

2019

- **Moving beyond neurons: the role of cell type-specific gene regulation in Parkinson's disease heritability⁵**

NPJ Parkinson's disease

- Reynolds, RH, Botía, JA, Nalls, MA, International Parkinson's Disease Genomic Consortium (IPDGC), System Genomics of Parkinson's Disease (SGPD), Hardy, J, Gagliano Taliun, SA, Ryten, M
- Role: First author and lead analyst.
- Analysis of Parkinson's disease common variation, with the aim of identifying cell types and pathways of importance to disease risk.

2018

- **Perturbations in the p53/miR-34a/SIRT1 pathway in the R6/2 Huntington's disease model⁶**

Molecular and Cellular Neuroscience

- Reynolds, RH, Petersen, MH, Willert, CW, Heinrich, M, Nyman, N, Dall, M, Treebak, JT, Björkqvist, M, Silahtaroglu, A, Hasholt, L and Nørremølle, A
- Role: First author, lead experimentalist and analyst.

View this publications list online with links at <https://rhreynolds.github.io/cv/>

CONTACT



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[GitHub](#)



[LinkedIn](#)



[ResearchGate](#)

Made with the R packages
[datadrivencv](#) and
[pagedown](#).

The source code is available
[GitHub](#).

Last updated on 2022-02-20.



PUBLICATIONS

2021

- **A genetic link between risk for Alzheimer's disease and severe COVID-19 outcomes via the OAS1 gene⁷**

Brain

- Magusali, N, Graham, AC, Piers, TM, Panichnantakul, P, Yaman, U, Shoai, M, Reynolds, RH et al.
- Role: Adviser. Reviewed and critiqued statistical analyses and manuscript.

2021

- **Mitochondrial-nuclear cross-talk in the human brain is modulated by cell type and perturbed in neurodegenerative disease⁸**

Communications Biology

- Fairbrother-Browne, A, Ali, A, Reynolds, RH et al.
- Role: Adviser. Advised on tools available for cell-type specificity and heritability analyses.

2021

- **A systems-level analysis highlights microglial activation as a modifying factor in common epilepsies⁹**

Neuropathology and Applied Neurobiology

- Altmann, A, Ryten, M, Di Nunzio, M, Ravizza, T, Tolomeo, D, Reynolds, RH et al.
- Role: Analyst. Collated gene lists designed to capture microglial states.

2021

- **An integrated genomic approach to dissect the genetic landscape regulating the cell-to-cell transfer of α-synuclein¹⁰**

Cell Reports

- Kara, E, Crimi, A, Wiedmer, A, Emmenegger, M, Manzoni, C, Bandres-Ciga, S, D'Sa, K, Reynolds, RH et al.
- Role: Analyst. Performed expression-weighted cell-type enrichment analyses.

2021

- **Heritability enrichment implicates microglia in Parkinson's disease pathogenesis¹¹**

Annals of Neurology

- Andersen, MS, Bandres-Ciga, S, Reynolds, RH et al.
- Role: Adviser. Reviewed and critiqued statistical analyses and manuscript.

2021

- **Human-lineage-specific genomic elements: relevance to neurodegenerative disease and APOE transcript usage¹²**

Nature Communications

- Chen, Z, Zhang, D, Reynolds, RH et al.
- Role: Analyst. Performed partitioned SNP-heritability analyses.

2021

- **Identification of Candidate Parkinson Disease Genes by Integrating Genome-Wide Association Study, Expression, and Epigenetic Data Sets¹³**

JAMA Neurology

- Kia, DA, Zhang, D, Guelfi, S, Manzoni, C, Hubbard, L, Reynolds, RH et al.
- Role: Analyst. Involved in cell-type annotation of prioritised genes.

2021

- **Modelling multifunctionality of genes with secondary gene co-expression networks in human brain provides novel disease insights¹⁴**

Bioinformatics

- Sánchez, JA, Gil-Martínez, AL, Cisterna, A, García-Ruiz, S, Gómez-Pascual, A, Reynolds, RH et al.
- Role: Adviser. Reviewed and critiqued statistical analyses and manuscript.

2021

- **Genome sequencing analysis identifies new loci associated with Lewy body dementia and provides insights into the complex genetic architecture¹⁵**

Nature Genetics

- Chia, R, Sabir, MS, Bandres-Ciga, S, Saez-Atienzar, S, Reynolds, RH et al.
- Role: Analyst. Performed colocalisation analyses and gene expression analyses.

2021

- **CoExp Web, a web tool for the exploitation of co-expression networks¹⁶**

Frontiers in Genetics

- García-Ruiz, S, Gil-Martínez, AL, Cisterna, A, Jurado, F, Reynolds, RH et al.
- Role: Analyst. Involved in cell-type annotation of gene co-expression networks.

2021

- **Genetic determinants of survival in progressive supranuclear palsy: a genome-wide association study¹⁷**

Lancet Neurology

- Jabbari, E, Koga, S, Valentino, R, Reynolds, RH et al.
- Role: Analyst. Performed colocalisation analyses.

2021

- **Genome-wide association studies of cognitive and motor progression in Parkinson's disease¹⁸**

Movement Disorders

- Tan, MMX, Lawton, MA, Jabbari, E, Reynolds, RH et al.
- Role: Adviser. Reviewed and critiqued statistical analyses and manuscript.

2021

- **Genetic analysis of amyotrophic lateral sclerosis identifies contributing pathways and cell types¹⁹**

Science Advances

- Saez-Atienzar, S, Bandres-Ciga, S, Langston, RG, Kim, JJ, Choi, SW, Reynolds, RH et al.
- Role: Adviser. Reviewed and critiqued statistical analyses and manuscript.

2020

- **Differences in network controllability and regional gene expression underlie hallucinations in Parkinson's disease²⁰**

Brain

- Zarkali, A, McColgan, P, Ryten, M, Reynolds, RH et al.
- Role: Adviser. Reviewed and critiqued statistical analyses and manuscript.

2020

- **Dementia risk in Parkinson's disease is associated with interhemispheric connectivity loss and determined by regional gene expression²¹**

NeuroImage: Clinical

- Zarkali, A, McColgan, P, Ryten, M, Reynolds, RH et al.
- Role: Adviser. Reviewed and critiqued statistical analyses and manuscript.

- 2020
- **Analysis of brain atrophy and local gene expression in genetic frontotemporal dementia²²**
Brain Communications
 - Altmann, A, Cash, DM, Bocchetta, M, Heller, C, Reynolds, RH et al.
 - Role: Analyst. Performed expression-weighted cell-type enrichment analyses.
- 2020
- **Incomplete annotation of disease-associated genes is limiting our understanding of Mendelian and complex neurogenetic disorders²³**
Science Advances
 - Zhang, D, Guelfi, S, Ruiz, SG, Costa, B, Reynolds, RH et al.
 - Role: Analyst. Performed expression-weighted cell-type enrichment analyses.
- 2020
- **Regulatory sites for known and novel splicing in human basal ganglia are enriched for disease-relevant information²⁴**
Nature Communications
 - Guelfi, S*, D'Sa, K*, Botía, JA*, Vandrovčová, J, Reynolds, RH et al.
 - Role: Experimentalist and analyst. Performed PCR validation of transcription and partitioned SNP-heritability analyses.
- 2019
- **Genetic variability in response to amyloid beta deposition influences Alzheimer's risk²⁵**
Brain Communications
 - Salih, DA, Bayram, S, Guelfi, S, Reynolds, RH et al.
 - Role: Analyst. Performed colocalisation analyses.
- 2019
- **Mitochondria function associated genes contribute to Parkinson's Disease risk and later age at onset²⁶**
NPJ Parkinson's disease
 - Billingsley, KJ, Barbosa, IA, Bandrés-Ciga S, Quinn, JP, Bubb, VJ, Deshpande, C, Botía, JA, Reynolds, RH et al.
 - Role: Analyst. Performed expression-weighted cell-type enrichment analyses.
- 2018
- **Variation at the TRIM11 locus modifies Progressive Supranuclear Palsy phenotype²⁷**
Annals of Neurology
 - Jabbari, E, John, W, Tan, MMX, Maryam, S, Pittman, A, Ferrari, R, Mok, KY, Zhang, D, Reynolds, RH et al.
 - Role: Analyst. Performed gene-based association testing and analysis of gene expression data.



PRE-PRINTS

2021

- **Detection of pathogenic splicing events from RNA-sequencing data using dasper²⁹**

bioRxiv

- Zhang, D, Reynolds, RH, et al.
- Role: Adviser. Helped guide and troubleshoot feature selection for outlier detection of splicing events.

2019

- **ERASE: Extended Randomization for assessment of annotation enrichment in ASE datasets³⁰**

bioRxiv

- D'Sa, K, Reynolds RH et al.
- Role: Analyst. Performed partitioned SNP-heritability analyses.

2018

- **G2P: Using machine learning to understand and predict genes causing rare neurological disorders³¹**

bioRxiv

- Botía, JA, Guelfi, S, Zhang, D, D'Sa, K, Reynolds, RH et al.
- Role: Adviser. Helped interpret the output of machine-learning-based models to distinguish disease genes from non-disease genes.