

Dr. Ziarh Hawi
Psychology
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Biography

I am the lead of the functional genomics lab within the Turner Institute for Brain and Mental Health (School of Psychological Sciences) and a molecular geneticist with long experience in psychiatry/mental health research. The focus of my research has been the identification of susceptibility loci for attention deficit hyperactivity disorder (ADHD). Throughout my career in the field of psychiatric genetics, I have successfully identified risk loci (genes) for ADHD using the candidate approach (Hawi et al, 2002; Mol Psychiatry; Hawi et al, 2003; Mol Psychiatry; Sheehan et al 2005; Mol Psychiatry). More recently, I have mapped a novel locus for ADHD using the method of genome wide association (GWA) (Hawi et al, 2018; Translational Psychiatry). I have also published the first paper to apply whole exome sequencing to discover risk loci for ADHD (Hawi et al, 2017; Mol Psychiatry). As member of the ADHD subgroup of the Psychiatric Genetic Consortium (PGC) I have contributed to the identification of 12 novel risk loci for ADHD using GWAS, recently published in Nature Genetics (2019).

Since my relocation to the School of Psychological Sciences at Monash University (July 2012), I have been contributing to the development of genetic research projects by training staff from different groups including Huntington's disease, Sleep, Aging and Neurodegeneration, and cognitive control of action research, and have worked closely with my peers within the Turner Institute for Braun and mental Health. I have a strong record of successful HD students (Honours, MSc and Ph.D.) who I have supervised throughout my career. I have dedicated my life to research, discovery, teaching and will continue to do so.

Related Links:

Psych: Research Themes

Qualifications

Molecular Medical Genetics, PhD, Newcastle University (United Kingdom)

Award Date: 22 Dec 1994

Biochemical Genetics, Masters, University of Strathclyde

Award Date: 18 Dec 1989

Biology, Honours, University of Baghdad

Award Date: 2 Jun 1982

Employment

Senior Research Fellow

Psychology

MONASH UNIVERSITY

3 Jul 2012 → present

Research outputs

Potential role for immune-related genes in autism spectrum disorders: Evidence from genome-wide association meta-analysis of autistic traits

Arenella, M., Cadby, G., De Witte, W., Jones, R. M., Whitehouse, A. J. O., Moses, E. K., Fornito, A., Bellgrove, M. A., Hawi, Z., Johnson, B., Tiegó, J., Buitelaar, J. K., Kiemeny, L. A., Poelmans, G. & Bralten, J., Feb 2022, In: Autism. 26, 2, p. 361-372 12 p.

The Monash Autism-ADHD genetics and neurodevelopment (MAGNET) project design and methodologies: a dimensional approach to understanding neurobiological and genetic aetiology

Knott, R., Johnson, B. P., Tiegó, J., Mellahn, O., Finlay, A., Kallady, K., Kouspos, M., Mohanakumar Sindhu, V. P., Hawi, Z., Arnatkeviciute, A., Chau, T., Maron, D., Mercieca, E. C., Furley, K., Harris, K., Williams, K., Ure, A., Fornito, A., Gray, K., Coghil, D. & 7 others, Nicholson, A., Phung, D., Loth, E., Mason, L., Murphy, D., Buitelaar, J. & Bellgrove, M. A., 5 Aug 2021, In: Molecular Autism. 12, 1, 24 p., 55.

Genetic influences on hub connectivity of the human connectome

Arnatkeviciute, A., Fulcher, B. D., Oldham, S., Tiego, J., Paquola, C., Gerring, Z., Aquino, K., Hawi, Z., Johnson, B., Ball, G., Klein, M., Deco, G., Franke, B., Bellgrove, M. A. & Fornito, A., 9 Jul 2021, In: Nature Communications. 12, 1, 14 p., 4237.

A rare missense variant in the ATP2C2 gene is associated with language impairment and related measures

Martinelli, A., Rice, M. L., Talcott, J. B., Diaz, R., Smith, S., Raza, M. H., Snowling, M. J., Hulme, C., Stein, J., Hayiou-Thomas, M. E., Hawi, Z., Kent, L., Pitt, S. J., Newbury, D. F. & Paracchini, S., 15 Jun 2021, In: Human Molecular Genetics. 30, 12, p. 1160-1171 12 p.

Risk variants and polygenic architecture of disruptive behavior disorders in the context of attention-deficit/hyperactivity disorder

Demontis, D., Walters, R. K., Rajagopal, V. M., Waldman, I. D., Grove, J., Als, T. D., Dalsgaard, S., Ribasas, M., Bybjerg-Grauholm, J., Bækvad-Hansen, M., Werge, T., Nordentoft, M., Mors, O., Mortensen, P. B., ADHD Working Group of the Psychiatric Genomics Consortium (PGC), Cormand, B., Hougaard, D. M., Neale, B. M., Franke, B., Faraone, S. V. & 1 others, Børglum, A. D., 25 Jan 2021, In: Nature Communications. 12, 1, 12 p., 576.

Environmental exposures associated with elevated risk for autism spectrum disorder may augment the burden of deleterious de novo mutations among probands

Pugsley, K., Scherer, S. W., Bellgrove, M. A. & Hawi, Z., Jan 2021, In: Molecular Psychiatry. 27, p. 710–730 21 p.

Evidence against benefits from cognitive training and transcranial direct current stimulation in healthy older adults

Horne, K. S., Filmer, H. L., Nott, Z. E., Hawi, Z., Pugsley, K., Mattingley, J. B. & Dux, P. E., Jan 2021, In: Nature Human Behaviour. 5, 1, p. 146-158 13 p.

Functional validation of CHMP7 as an ADHD risk gene

Dark, C., Williams, C., Bellgrove, M. A., Hawi, Z. & Bryson-Richardson, R. J., 6 Nov 2020, In: Translational Psychiatry. 10, 1, 8 p., 385.

Shared genetic background between children and adults with attention deficit/hyperactivity disorder

Rovira, P., Demontis, D., Sanchez-Mora, C., Zayats, T., Klein, M., Roth Mota, N., Weber, H., Garcia-Martinez, I., Pagerols, M., Vilar-Ribo, L., Arribas, L., Richarte, V., Corrales, M., Fadeuilhe, C., Bosch, R., Espanol Martin, G., Almos, P., Doyle, A. E., Horacio Grevet, E., Grimm, O. & 36 others, Halmoy, A., Hoogman, M., Hutz, M., Jacob, C., Kittel-Schneider, S., Knappskog, P. M., Lundervold, A. J., Rivero, O., Luiz Rovaris, D., Salatino-Oliveira, A., Santos da Silva, B., Svirin, E., Sprooten, E., Strekalova, T., ADHD Working Group of the Psychiatric Genomics Consortium (PGC), 23andMe Research Team, Arias-Vasquez, A., Sonuga-Barke, E., Asherson, P. J., Bau, C. H. D., Buitelaar, J. K., Cormand, B., Faraone, S. V., Haavik, J., Johansson, S. E., Kuntsi, J., Larsson, H., Lesch, K. P., Reif, A., Rohde, L. A., Casas, M., Børglum, A. D., Franke, B., Ramos-Quiroga, J. A., Soler Artigas, M. & Ribases, M., Sep 2020, In: Neuropsychopharmacology. 45, p. 1617-1626 10 p.

A genetic profile of refractory individuals with major depressive disorder and their responsiveness to transcranial magnetic stimulation

Souza-Silva, N. G., Nicolau, E. S., Hoy, K., Hawi, Z., Bellgrove, M. A., Miranda, D. M., Romano-Silva, M. A. & Fitzgerald, P. B., Jul 2020, In: Brain Stimulation. 13, 4, p. 1091-1093 3 p.

Impact of CYP2C19 genotype-predicted enzyme activity on hippocampal volume, anxiety, and depression

Savadlou, A., Arnatkeviciute, A., Tiego, J., Hawi, Z., Bellgrove, M. A., Fornito, A. & Bousman, C., Jun 2020, In: Psychiatry Research. 288, 4 p., 112984.

The application of human pluripotent stem cells to model the neuronal and glial components of neurodevelopmental disorders

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Intensity matters: high-intensity interval exercise enhances motor cortex plasticity more than moderate exercise

Andrews, S. C., Curtin, D., Hawi, Z., Wongtrakun, J., Stout, J. C. & Coxon, J. P., 10 Jan 2020, In: *Cerebral Cortex*. 30, 1, p. 101-112 12 p.

Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders

Cross-Disorder Group of the Psychiatric Genomics Consortium, 12 Dec 2019, In: *Cell*. 179, 7, p. 1469-1482.e11 25 p.

Generation of four iPSC lines from peripheral blood mononuclear cells (PBMCs) of an attention deficit hyperactivity disorder (ADHD) individual and a healthy sibling in an Australia-Caucasian family

Tong, J., Lee, K. M., Liu, X., Nefzger, C. M., Vijayakumar, P., Hawi, Z., Pang, K. C., Parish, C. L., Polo, J. M. & Bellgrove, M. A., 1 Jan 2019, In: *Stem Cell Research*. 34, 5 p., 101353.

Discovery of the first genome-wide significant risk loci for attention deficit/hyperactivity disorder

Demontis, D., Walters, R. K., Martin, J., Manuel, M., Als, T., Agerbo, E., Baldursson, G., Belliveau, R., Bybjerg-Grauholm, J., Bækvad-Hansen, M., Cerrato, F., Chambert, K., Churchhouse, C., Dumont, A., Eriksson, N., Gandal, M., Goldstein, J. I., Grasby, K. L., Grove, J., Gudmundsson, O. O. & 3 others, ADHD Working Group of the Psychiatric Genomics Consortium (PGC), Early Lifecourse & Genetic Epidemiology (EAGLE) Consortium & 23andMe Research Team, Jan 2019, In: *Nature Genetics*. 51, 1, 16 p.

A case-control genome wide association study of childhood attention deficit hyperactivity disorder (ADHD)

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A case-control genome-wide association study of ADHD discovers a novel association with the tenascin R (TNR) gene

Hawi, Z., Yates, H., Pinar, A., Arnatkeviciuteb, A., Johnson, B., Tong, J., Pugsley, K., Dark, C., Pauper, M., Klein, M., Heussler, H. S., Hiscock, H., Fornito, A., Tiegó, J., Finlay, A., Vance, A., Gill, M., Kent, L. & Bellgrove, M. A., 18 Dec 2018, In: *Translational Psychiatry*. 8, 1, 8 p., 284.

Genome-wide association study reveals novel genetic locus associated with intra-individual variability in response time

Pinar, A., Hawi, Z., Cummins, T., Johnson, B., Pauper, M., Tong, J., Tiegó, J., Finlay, A., Klein, M., Franke, B., Fornito, A. & Bellgrove, M. A., 4 Oct 2018, In: *Translational Psychiatry*. 8, 1, 8 p., 207.

A Genetic Investigation of Sex Bias in the Prevalence of Attention-Deficit/Hyperactivity Disorder

Martin, J., Walters, R. K., Demontis, D., Mattheisen, M., Lee, S. H., Robinson, E., Brikell, I., Ghirardi, L., Larsson, H., Lichtenstein, P., Eriksson, N., 23andMe Research Team, Psychiatric Genomics Consortium: ADHD Subgroup, iPSYCH–Broad ADHD Workgroup, Werge, T., Mortensen, P. B., Pedersen, M. G., Mors, O., Nordentoft, M., Hougaard, D. M. & 8 others, Bybjerg-Grauholm, J., Wray, N. R., Franke, B., Faraone, S. V., O'Donovan, M. C., Thapar, A., Børglum, A. D. & Neale, B. M., 15 Jun 2018, In: *Biological Psychiatry*. 83, 12, p. 1044-1053 10 p.

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Trends in the Overlap of Autism Spectrum Disorder and Attention Deficit Hyperactivity Disorder: Prevalence, Clinical Management, Language and Genetics

May, T., Brignell, A., Hawi, Z., Brereton, A., Tonge, B., Bellgrove, M. A. & Rinehart, N., 1 Mar 2018, In: *Current Developmental Disorders Reports*. 5, 1, p. 49-57 9 p.

Rare DNA variants in the brain-derived neurotrophic factor gene increase risk for attention-deficit hyperactivity disorder: a next-generation sequencing study

Hawi, Z., Cummins, T. D. R., Tong, J., Arcos-Burgos, M., Zhao, Q., Matthews, N., Newman, D. P., Johnson, B., Vance, A., Heussler, H. S., Levy, F., Eastal, S., Wray, N., Kenny, E., Morris, D., Kent, L., Gill, M. & Bellgrove, M. A., Apr 2017, In: *Molecular Psychiatry*. 22, p. 580-584 5 p.

Allelic variation in dopamine D2 receptor gene is associated with attentional impulsiveness on the Barratt Impulsiveness Scale (BIS-11)

Taylor, J. B., Cummins, T. D. R., Fox, A. M., Johnson, B. P., Tong, J. H., Visser, T. A. W., Hawi, Z. & Bellgrove, M., 20 Jan 2017, In: *The World Journal of Biological Psychiatry*. 19, S2, p. S75-S83 9 p.

Attention Deficit Hyperactivity Disorder

Tong, J., Hawi, Z. & Bellgrove, M. A., 2017, *Developmental Disorders of the Brain*. Rinehart, N. J., Bradshaw, J. L. & Enticott, P. G. (eds.). 2nd ed. Routledge, p. 119-138 20 p. (Brain, Behaviour and Cognition Series).

Separating the wheat from the chaff: systematic identification of functionally relevant noncoding variants in ADHD

Tong, J.H.S., Hawi, Z., Dark, C., Cummins, T.D.R., Johnson, B.P., Newman, D.P., Lau, R., Vance, A., Heussler, H.S., Matthews, N., Bellgrove, M. A. & Pang, K.C., 1 Nov 2016, In: *Molecular Psychiatry*. 21, 11, p. 1589-1598 10 p.

Psychiatric gene discoveries shape evidence on ADHD's biology

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Genetically mediated resistance to distraction: Influence of dopamine transporter genotype on attentional selection

Bellgrove, M. A., Newman, D. P., Cummins, T., Tong, J., Johnson, B. P., Wagner, J., Goodrich, J., Hawi, Z. & Chambers, C., 24 Apr 2015.

The molecular genetic architecture of attention deficit hyperactivity disorder

Hawi, Z., Cummins, T.D.R., Tong, J., Johnson, B., Lau, R., Samarrai, W. & Bellgrove, M.A., Mar 2015, In: *Molecular Psychiatry*. 20, 3, p. 289-297 9 p.

An association between a dopamine transporter gene (SLC6A3) haplotype and ADHD symptom measures in nonclinical adults

Tong, J. H. S., Cummins, T., Johnson, B. P., McKinley, L-A., Pickering, H., Fanning, P., Stefanac, N., Newman, D. P., Hawi, Z. & Bellgrove, M. A., 2015, In: *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*. 168, 2, p. 89 - 96 8 p.

Identification and functional characterisation of a novel dopamine beta hydroxylase gene variant associated with attention deficit hyperactivity disorder

Tong, J. H. S., McKinley, L-A., Cummins, T., Johnson, B. P., Matthews, N. L., Vance, A. L. A., Heussler, H. S., Gill, M., Kent, L. S. W., Bellgrove, M. A. & Hawi, Z., 2015, In: *The World Journal of Biological Psychiatry*. 16, 8, p. 610 - 618 9 p.

Alpha-2A adrenergic receptor gene variants are associated with increased intra-individual variability in response time

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Dopamine transporter genotype is associated with a lateralized resistance to distraction during attention selection

Newman, D. P., Cummins, T., Tong, J. H. S., Johnson, B. P., Pickering, H., Fanning, P., Wagner, J., Goodrich, J. T. T., Hawi, Z., Chambers, C. D. & Bellgrove, M. A., 2014, In: *Journal of Neuroscience*. 34, 47, p. 15743 - 15750 8 p.

Neurodevelopmental and neuropsychiatric disorders represent an interconnected molecular system

Cristino, A. S., Williams, S. M., Hawi, Z., An, J. Y., Bellgrove, M. A., Schwartz, C. E., Costa, L. D. F. C. & Clodianos, C., 2014, In: *Molecular Psychiatry*. 19, 3, p. 294 - 301 8 p.

A high density linkage disequilibrium mapping in 14 noradrenergic genes: evidence of association between SLC6A2, ADRA1B and ADHD

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DNA variation in the SNAP25 gene confers risk to ADHD and is associated with reduced expression in prefrontal cortex

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Human amygdala volume is predicted by common DNA variation in the stathmin and serotonin transporter genes
Stjepanovic, D., Lorenzetti, V., Yucel, M., Hawi, Z. & Bellgrove, M. A., 2013, In: *Translational Psychiatry*. 3, 3 (Art. No.: e283), p. 1 - 7 7 p.

Methylphenidate side effect profile is influenced by genetic variation in the attention-deficit/hyperactivity disorder-associated CES1 gene

Johnson, K. A., Barry, E., Lambert, D., Fitzgerald, M. F., McNicholas, F., Kirley, A., Gill, M., Bellgrove, M. A. & Hawi, Z., 2013, In: *Journal of Child and Adolescent Psychopharmacology*. 23, 10, p. 655 - 664 10 p.

Norepinephrine genes predict response time variability and methylphenidate-induced changes in neuropsychological function in attention deficit hyperactivity disorder

Kim, B-N., Kim, J-W., Cummins, T. D., Bellgrove, M. A., Hawi, Z., Hong, S. B., Yang, Y. H., Kim, H. J., Shin, M. S., Cho, S-C., Kim, J. H., Son, J. W., Shin, Y. M., Chung, U. S. & Han, D. H., 2013, In: *Journal of Clinical Psychopharmacology*. 33, 3, p. 356 - 362 7 p.

Dopamine transporter genotype predicts behavioural and neural measures of response inhibition

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Investigating the contribution of common genetic variants to the risk and pathogenesis of ADHD

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Epistasis between neurochemical gene polymorphisms and risk for ADHD

Segurado, R., Bellgrove, M., Manconi, F., Gill, M. & Hawi, Z., 2011, In: *European Journal of Human Genetics*. 19, 5, p. 577 - 582 6 p.

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Case-control genome-wide association study of attention-deficit/ hyperactivity disorder

Neale, B. M., Medland, S. E., Ripke, S., Anney, R. J., Asherson, P. J., Buitelaar, J. K., Franke, B., Gill, M., Kent, L. S. W., Holmans, P. A., Middleton, F. A., Thapar, A., Lesch, K. P., Faraone, S. V., Daly, M. J., Nguyen, T. T., Schafer, H. H., Steinhausen, H. C., Reif, A., Renner, T. J. & 13 others, Romanos, M., Romanos, J., Warnke, A., Walitza, S., Freitag, C. M., Meyer, J. D., Palmason, H. O., Rothenberger, A., Hawi, Z., Sergeant, J. A., Roeyers, H., Mick, E. O. & Biederman, J. J., 2010, In: *Journal of the American Academy of Child and Adolescent Psychiatry*. 49, 9, p. 906 - 920 15 p.

Evidence that genetic variation in the oxytocin receptor (OXTR) gene influences social cognition in ADHD

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Functional analysis of intron 8 and 3' UTR variable number of tandem repeats of SLC6A3: differential activity of intron 8 variants

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Haplotype association of calpain 10 gene variants with type 2 diabetes mellitus in an Irish sample

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Dopaminergic haplotype as a predictor of spatial inattention in children with attention-deficit/hyperactivity disorder

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Replication of an association of a promoter polymorphism of the dopamine transporter gene and Attention Deficit Hyperactivity Disorder

Doyle, C., Brookes, K. J., Simpson, J., Park, J., Scott, S. V., Coghill, D. R., Hawi, Z., Kirley, A., Gill, M. & Kent, L. S. W., 2009, In: *Neuroscience Letters*. 462, 2, p. 179 - 181 3 p.

Absence of the 7-repeat variant of the DRD4 VNTR is associated with drifting sustained attention in children with ADHD but not in controls

Johnson, K. A., Kelly, S. P., Robertson, I. H., Barry, E., Mulligan, A., Daly, M., Lambert, D. B., McDonnell, C., Connor, T. J., Hawi, Z., Gill, M. & Bellgrove, M. A., 2008, In: *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*. 147, 6, p. 927 - 937 11 p.

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Brookes, K. J., Hawi, Z., Kirley, A., Barry, E., Gill, M. & Kent, L. S. W., 2008, In: *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*. 147, 8, p. 1531 - 1535 5 p.

Creativity, psychosis, autism, and the social brain

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1/01/18 → 31/12/21

CogChip: development of a targeted genotyping chip for executive function

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20/12/17 → 19/12/21

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Department of Health (Australia)

20/01/20 → 30/06/24

How exercise modulates neural plasticity and memory consolidation

Coxon, J., Byblow, W. D. & Hawi, Z.

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