CURRICULUM VITAE

PERSONAL INFORMATION

Family name, First name: YALCIN Binnaz

Researcher unique identifier: ORCID 0000-0002-1924-6807

Date of birth: 20th July 1977 Nationality: French-Turkish

URL for web site: https://blog.u-bourgogne.fr/yalcingroup/

EDUCATION

2014 HDR in Neurosciences (<u>Accreditation to Direct Research</u>, Highest French Academic degree)

Institute of Genetics and Molecular and Cellular Biology, University of Strasbourg, France

2007 **PhD** in Genetics (Supervision: Prof Richard Mott)

Wellcome Centre for Human Genetics, University of Oxford, Exeter College, UK

• CURRENT POSITION

2019-present **Tenured Assistant Professor** in Neurogenetics

Dijon Centre of Translational and Molecular Medicine, INSERM Unit 1231, Dijon, France

• PREVIOUS POSITIONS

2016-2019	Junior Team Leader
	Institute of Genetics and Molecular and Cellular Biology, University of Strasbourg, France
2011-2016	Independent Research Fellow
	Centre for Integrative Genomics, Switzerland
2007-2011	Postdoctoral Fellow (Advisor: Prof Jonathan Flint)
	Wellcome Centre for Human Genetics, Oxford University, UK
2002-2005	PhD entitled "Gene Mapping in Animal Models for Anxiety" (Advisor: Prof Richard Mott)
	Wellcome Centre for Human Genetics, Oxford University, UK

• SELECTED FELLOWSHIPS, AWARDS AND APPOINTMENTS

2023-2027	ANR Researcher Laureate, University of Bourgogne Franche-Comté, France
2022-2023	Nominated Scientific Evaluation Member, European Joint Programme on Rare Diseases
2022-2026	Nominated Scientific Evaluation Member, INSERM
2021	INSERM International Researcher Award (first laureate of the program)
2020	European Commission Solve-RD Award (first laureate of the seeding program)
2019	INSERM Permanent Researcher position in Genetics (ranked first)
2019	Mary Lyon Award in recognition for an early-stage independent female researcher
2018-2021	ANR Young Researcher Laureate, University of Strasbourg, France
2016-2019	Chair of Excellence Award, University of Strasbourg, France
2016	Gutenberg Mobility Award, University of Strasbourg, France
2015	Tremplin Fellowship for Female Researchers, University of Lausanne, Switzerland
2013-2016	SNSF Ambizione Fellowship, University of Lausanne, Switzerland
2013	Korner Mobility Fellowship Fund, University of Sussex, UK
2011-2014	ANR Postdoctoral Researcher Laureate, University of Strasbourg
2011-2013	EMBO Long-Term Researcher Fellowship, University of Lausanne, Switzerland
2008-2011	Wellcome Trust Postdoctoral Fellowship, University of Oxford, UK
2008	Academic Merit Award, University of Oxford, UK
2002	Wellcome PhD Studentship Award, University of Oxford, UK

• INVITED TEACHING ACTIVITIES

2023	1-hour broad audience lecture, <u>Topic</u> : Neurobiology of Cohen syndrome, Clermont, France
2021-present	2-hour invited lecture, <u>Topic</u> : Neuro-anatomy, UBFC, Besançon, France
2020-present	2-hour invited lecture, <u>Topic</u> : Neuro-signalisation, University of Burgundy, Dijon, France
2019	4-hour broad audience lecture, <u>Topic</u> : Brain Development in Children, Dijon, France
2012-present	Annual 2-hour invited lecture, <u>Topic</u> : Mouse Genetics, Pasteur Institute, Paris, France
2018	2-hour invited lecture, <u>Topic</u> : Next Generation Sequencing, University of Cambridge, UK
2016	3-days ITMO School, <u>Topic</u> : Translational Research in Neurosciences, Bordeaux, France
2015	1-week European Advanced School, <u>Topic</u> : Mouse Phenogenomics, Alsace, France
2013	2-hour invited lecture, <u>Topic</u> : Mouse Genomics, ESBS, University of Strasbourg, France

ORGANISATION OF SCIENTIFIC MEETINGS

2023	Chair IMGS conference, Session: Human Disease Model, Japan (200 participants)
2022	Chair IRC5 conference, Session: Human Genetics, online (<u>250 participants</u>)
2021-present	Coordinator of Webinars on Brain Research, IRC5 consortium (50 participants)
2020-present	Organizer of Internal Seminars, INSERM Unit 1231, Dijon, France (50 participants)
2020	Organizer of Symposium on Brain Imaging, University of Burgundy, Dijon (50 participants)
2018	Organizer of Symposium on Statistics, University of Strasbourg, France (50 participants)
2017	Chair IMGS conference, Session: Human Disease Model, Germany (150 participants)

• INSTITUTIONAL RESPONSIBILITIES

2023	Nominated member at the European Joint Programme on Rare Diseases
2022-2027	Nominated member at the INSERM Scientific Evaluation Board
2022-present	Scientific Advisor Panel Member of the PhD School (University of Milan)
2020	Nomination as President for the International Mouse Genome Society (IMGS)
2019-present	Ambassador, eLIFE Early Career Researchers Community
2016-present	PhD thesis examiner (N=13)
2016-present	Member of examination board for Masters and PhD students (N=4)
2016-present	HDR (Accreditation to Direct Research, Highest French Academic degree) examiner (N=2)

• REVIEWING ACTIVITIES

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2016-present	Expert Reviewer of Journals: Nature, Nature Genetics, Nucleic Acids Research, Trends in						
	Genetics, Cell Reports, Theranostics, eLife, Neurobiology of Disease, Frontiers Cell &						
	Developmental Biology, PLOS Genetics, International Journal of						
	Neuropsychopharmacology, Progress in Neuropsychopharmacology & Biological Psychiatry,						
	Frontiers in Neuroscience, Human Mutation, Molecular Neurobiology, Molecular Genetics &						
	Metabolism, BMC Genomics (x2), Biomolecules, PLOS One (x2), Genes Brain & Behaviour,						
	Journal of Clinical Medicine, Mammalian Genome (x3), BMC Research Notes, Journal of						
	Metabolomics & Systems Biology, Genes Genomes & Genetics (x3), Life Science Alliance,						
	Brain Research (x3), Journal of Paediatric Genetics (1 every 2 months)						
2016-present	Expert Reviewer of Grants: Sorbonne University Emergence Grant, European Joint						
	Programme on Rare Diseases, MRC UK, French Foundation for Medical Research Line-						
	Pomaret, IDEX Marseille, Research Foundation Flanders, Rare Disease Foundation						

2016-present **Editorial Board Member**: Frontiers in Genetics

• MEMBERSHIPS OF SCIENTIFIC SOCIETIES

2021-present	Member of various societies including NeuroFrance, FENS, ISTT, CFATG and ESHG
2020	Nominated Member to take the leadership of the International Mammalian Genome Society
2019-present	Co-director, International Research Consortium for the Corpus Callosum and Connectivity
2016-present	Elected Member, International Mammalian Genome Society
2005-2011	Elected Member, Complex Trait Consortium

• CAREER BREAKS (account for the temporary dip in publications)

03/18-04/19	6-month pregnancy-related sick leave + 6-month maternity leave
04/12-09/12	6-month sick leave
08/07-09/08	6-month pregnancy-related sick leave + 6-month maternity leave
07/05-08/06	6-month pregnancy-related sick leave + 6-month maternity leave

OTHER

The High Council for the Evaluation of Research and Higher Education Institutions (HCERES) has scored my team as "excellent to outstanding with great future prospects".

• COVID-19 IMPACT TO SCIENTIFIC PRODUCTIVITY

During the lockdown periods, my laboratory interrupted mouse experiments and reduced colony size to a strict minimum as we were not allowed to breed mice anymore. Also, during these periods, I experienced increased caring responsibilities for my dependent father and my children with home schooling.

Current research grants:

Project Title	Funding source	Amount (Euros)	Period	Role of the PI	Relation to current proposal
Chromatin remodeler CHD1L in neurogenesis and in 1q21.1 Copy Number Variant-associated neurodevelopmental phenotypes (CHROMATISM)	ANR	450 000	4 years (2022-2026)	Co- Coordinator	None
An integrated approach to uncover genetic, molecular, and cellular mechanisms underlying severe neurodevelopmental disorders (NeuroGen)	CEFIPRA	200 000	3 years (2022- 2025)	Co- Coordinator	None
Genetic and functional characterization of WDR genes in neurodevelopmental disorders (WDR)	ANR	250 000	5 years (2018- 2023)	Coordinator	None
Deciphering physio-pathological mechanisms to better understand comorbidities between neurodevelopmental disorders and obesity (DIOBE)	ANER	50 000	2 years (2022- 2024)	Coordinator	None
Mouse models for CTNNA2 heterozygous de novo missense variants (CTNNA2)	Solve-RD	20 000	3 years (2021- 2023)	Coordinator	None

On-going and submitted grant applications:

Project Title	Funding source	Amount (Euros)	Period	Role of the PI	Relation to current proposal
Investigation of the cerebellum as a target for the treatment of autism spectrum disorder	ANR	500 000	4 years (2024-2027)	Co- Coordinator	None
Dysregulation of Midasin 1 AAA ATPase impacts pre-60S particles export to the cytosol and causes a novel multisystem disease with cerebellar atrophy	Telethon	200 000	2 years (2024-2025)	Co- Investigator	None
In-depth functional characterization of new CTNNA2 variants involved in pediatric neurodevelopmental disorders to reduce the diagnostic odyssey	NIH	100 000	2 years (2024-2025)	Co- Coordinator	None
Genetics and physiopathology of rare disorders of the developing brain	BQR	100 000	2 years (2024-2025)	Coordinator	None
An integrated approach to uncover genetic, molecular, and cellular mechanisms underlying severe neurodevelopmental disorders	INSERM	75 000	5 years (2024-2028)	Coordinator	None

EARLY ACHIEVEMENTS TRACK-RECORD

• SHORT BIOSKETCH

I have contributed to important aspects of neurological disease genetics using high-throughput genetic, genomic methodologies and mice as genetic models. My ambition has always been to pursue my career in functional brain disease genetics at the highest level. In 2011, I secured an EMBO Young Investigator Fellowship allowing me to move to Switzerland where I obtained an Ambizione Fellowship to work on the genetics of neurodevelopmental diseases. I set up my own laboratory in 2016 at the IGBMC (Strasbourg, France) through an international call in the framework of a Chair of excellence in research program. One year later, the High Council for the Evaluation of Research and Higher Education scored my team as "excellent to outstanding with great future prospects". I presented my work at 49 International Conferences of which 12 were under a formal invitation. I have received 17 awards/prizes. To fund my research, I raised and managed more than three million Euros and employed since 2016 four PhD students, three postdoctoral fellows, one engineer, six research assistants. I hosted in my laboratory more than 60 students worldwide at all levels (MSc, BSc, Intern, Assistants and ERASMUS). My first three PhD students have graduated each with one significant first-author paper (Kannan et al. 2017 PNAS, Mikhaleva et al. 2019 Nature Communications, Kretz et al. 2023 Genome Biology In revision). In 2019, I secured a tenured INSERM researcher position after being ranked Major in a highly competitive national selection. I have built an international reputation for the contribution to the field of brain biology with the identification of several important genes associated with neuroanatomical phenotypes and biomarkers. Driving the field forward, my leadership was recognized by an invitation to be part of the Governance Board of the International Research Consortium for the Corpus Callosum and Cerebral Connectivity (IRC5) and the International Mammalian Genome Society (IMGS).

• OFFICIAL SUPERVISION OF GRADUATE STUDENTS AND POSTDOCTORAL FELLOWS

2019-present 1 Postdoc, 2 PhDs, 1 Engineer, 10 Masters, 18 Undergraduates, 2 Research Assistants

Inserm Unit 1231, University of Bourgogne Franche-Comté, France

2016-2019 2 Postdocs, 2 PhDs, 11 Masters, 16 Undergraduates, 4 Research Assistants

Institute of Genetics and Molecular and Cellular Biology, University of Strasbourg, France

PUBLICATIONS

Total of 44 publications: Biometrics (January 2023):

17 corresponding/last-author Category Normalised Citation Impact 3.2

9 first-author 11% of publications in **Top 1%**, 40% in **Top 10%**

8 invited H-index 23, Sum of the times cited 5532

Articles published in: Cell (x1), Nature (x2), Nature Genetics (x2), Nature Communications (x3), Genome Biology (x4), MBE (x1), Science Advances (x1), Biological Psychiatry (x2), AJHG (x3), PNAS (x2), Genome Research (x2), GIM (x1), Journal of Neuroscience (x1), PLoS Genetics (x3), Human Genetics (x1), HMG (x1), IJMS (x1), Epilepsia (x1), Frontiers in Genetics (x1), Genetics (x2), Behavioural Brain Research (x1), PLoS ONE (x1), JANAT (x1), Mammalian Genome (x2) and CPMB (x3)

Selected publications

- 1. Kretz P, Wagner C, Mikhaleva A, Hugel S, Morella I, Montillot C, Kannan M, Fischer MC, Milhau M, Yalcin I, Brambilla R, Selloum M, Herault Y, Reymond A, Collins SC, <u>Yalcin B*</u>. Dissecting the autism-associated 16p11.2 locus identifies sex-specific drivers in neuroanatomical phenotypes and unveils a new role for the major vault protein (in revision in **Genome Biology**). [*Correspondence].
- 2. Szpak M, Collins SC, Li Y, Liu X, Ayub Q, Vancollie VE, Lelliott CJ, Xue Y, <u>Yalcin B*</u>, Yang H, Tyler-Smith C*. A positively-selected MAGEE2 LoF allele is associated with sexual dimorphism in human brain size, and shows similar phenotypes in Magee2 null mice. **Molecular Biology & Evolution**. 2021 Aug 31. [*Co-Correspondence].
- 3. Duncan AR, Vitobello A, Collins SC, Vancollie VE, Lelliott CJ, Rodan L, Shi J, Seman AR, Agolini E, Novelli A, Prontera P, Guillen Sacoto MJ, Santiago-Sim T, Trimouille A, Goizet C, Nizon M, Bruel AL, Philippe C, Grant PE, Wojcik MH, Stoler J, Genetti CA, van Dooren MF, Maas SM, Alders M, Faivre L, Sorlin A, Yoon G, <u>Yalcin B*</u>, Agrawal PB*. Heterozygous Variants in KDM4B Lead to Global Developmental Delay and Neuroanatomical Defects. AJHG. 2020 Dec;107(6):1170-1177 [*Co-Correspondence].
- 4. Collins SC, Mikhaleva A, Vrcelj K, Vancollie VE, Wagner C, Demeure N, Whitley H, Kannan M, Balz R, Anthony LFE, Edwards A, Moine H, White JK, Adams DJ, Reymond A, Lelliott CJ, Webber C, <u>Yalcin B*</u>. Large-scale neuroanatomical study uncovers 198 gene associations in mouse brain morphogenesis. Nature Communications. 2019 Aug;10(1):3465 [*Correspondence].
- 5. Kannan M, Bayam E, Wagner C, Rinaldi B, Kretz PF, Tilly P, Roos M, McGillewie L, Bär S, Minocha S, Chevalier C, Po C, Chelly J, Mandel JL, Borgatti R, Piton A, Kinnear C, Loos B, Adams DJ, Hérault Y, Collins SC, Friant S, Godin JD, <u>Yalcin B*</u>. WD40-repeat 47, a microtubule-associated protein, is essential for brain development and autophagy. PNAS. 2017 Oct 31;114(44) [*Correspondence].

- 6. <u>Yalcin B*</u>, Wong K, Bhomra A, Goodson M, Keane T, Adams D, Flint J. The fine-scale architecture of structural variants in 17 mouse genomes. **Genome Biology**. 2012, 13(3):R18 [*Correspondence].
- 7. <u>Yalcin B</u>, Wong K, Agam A, Goodson M, Keane T, Gan X, Nellåker C, Goodstadt L, Nicod J, Bhomra A, Whitley H, Cleak J, Dutton R, Mott R, Adams D, Flint J. Sequence based characterization of structural variation in the mouse genome. **Nature**. 2011, 477(7364):326-9.
- 8. <u>Yalcin B</u>, Nicod J, Bhomra A, Davidson S, Cleak J, Farinelli L, Østerås M, Yuan W, Whitley A, Gan X, Goodson M, Klenerman P, Satpathy A, Benoist C, Adams DJ, Mott R, Flint J. Commercially available outbred mice for genome-wide association studies. **Plos Genetics**. 2010, 2;6(9).
- 9. <u>Yalcin B</u>, Willis-Owen SA, Fullerton J, Meesaq A, Deacon RM, Rawlins JNP, Copley RR, Morris AP, Flint J, Mott R. Genetic dissection of a behavioural quantitative trait locus shows that *Rgs2* modulates anxiety in mice. **Nature Genetics**. 2004, 36:1197-1202.
- 10. <u>Yalcin B</u>, Fullerton J, Miller S, Keays DA, Brady SA, Bhomra A, Jefferson A, Volpi E, Copley RR, Flint J, Mott R. Unexpected complexity in the haplotypes of commonly used inbred strains of laboratory mice. **PNAS**. 2004, 101:9734-9739.

RESEARCH MONOGRAPHS

I have built an international reputation for the careful assessment of neuroanatomical phenotypes. I have been formally invited to publish our collection of standard operating procedures (PMIDs: 29944194, 27584555 and 27584555) and to contribute to a chapter on innovation in neurosciences published by Springer Nature.

INVITED CONFERENCES

	CONTENENCES
2023	4 th NeuroFrance Meeting, Lyon, Talk
2022	20 th Anniversary ICS, Strasbourg, France, Talk
2021	International Research Consortium on Corpus Callosum, Brazil, Talk (online)
2019	3 rd Meeting Gene Expression and Epigenome, Strasbourg, France, Talk
2018	33 rd International Mouse Genome Conference, USA, Mary Lyon Award Talk
2018	International Research Consortium on Corpus Callosum, California, USA, Talk
2017	Advancing Personalised Medicine with Animal Models, Athens, Greece, Talk
2017	2 nd Functional annotation of genome-wide variants, Lausanne, Switzerland, Talk
2016	11 th International Meeting on CNVc and Genes in Autism, Troina, Italy, Talk
2016	5 th International Mouse Phenotyping Consortium, Strasbourg, France, Talk
2013	6 th Colloque Belles Souris, Montpellier, France, Talk
2005	11 th International School on Neurogenetics, Bordeaux, France, Talk

• SELECTED CONFERENCES

2023	36 th International Mouse Genome Conference, Tsukuba, Japan, Talk
2022	35 th International Mouse Genome Conference, Vancouver, Canada, Talk
2022	International Research Consortium on Corpus Callosum, USA, Talk (online)
2022	54 th European Society of Human Genetics, Vienna, Austria, Poster and Talk
2022	11 th Assises de Génétique Humaine et Médicale, Rennes, France, Talk
2017	50 th European Society of Human Genetics, Copenhagen, Denmark, Talk
2016	1 st Genomics of Brain Disorders, Hinxton, United Kingdom, Poster & Talk
2015	65 th American Society of Human Genetics, USA, Poster & Talk (+ Award)
2015	17th International Cognitive Disorders Workshop, Strasbourg, France, Talk
2015	48 th European Society of Human Genetics, Glasgow, Scotland, Talk
2015	2 nd Gencodys Conference, Chania, Crete, Talk
2015	10 th International Meeting on CNV and Genes in Autism, Italy, Talk
2014	7 th Personal Genomes: Discovery, Treatment & Outcomes, CSH, USA, Talk
2014	28 th International Mouse Genome Conference, Maine, USA, Talk
2014	10 th Neurogenetics Symposium, Strasbourg, France, Talk
2014	6 th Meeting of the International Society for CSF Disorders, Bristol, UK, Talk
2013	1 st Neurosciences Meeting, Dijon, France, Talk
2012	11th Complex Trait Consortium Meeting, Pasteur Institute, Paris, France, Talk
2009	8 th Complex Trait consortium Meeting, Manchester, United Kingdom, Talk
2003	17 th International Mouse Genome Conference, Braunschweig, Germany, Talk
2003	2 nd Complex Trait Consortium Meeting, Oxford, United Kingdom, Talk