

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

Peer-reviewed research articles

- 1) Fisher SE, Black GCM, Lloyd SE, Hatchwell E, Wrong O, Thakker RV, Craig IW (1994) Isolation and partial characterization of a chloride channel which is expressed in kidney and is a candidate for Dent's disease (an X-linked hereditary nephrolithiasis). *Hum Mol Genet* 3:2053-9
- 2) Blair HJ, Ho M, Monaco AP, Fisher S, Craig IW, Boyd Y (1995) High-resolution comparative mapping of the proximal region of the mouse X chromosome. *Genomics* 28:305-10
- 3) Fisher SE, Hatchwell E, Chand A, Ockendon N, Monaco AP, Craig IW (1995) Construction of two YAC contigs in human Xp11.23-p11.22, one encompassing the loci OATL1, GATA, TFE3 and SYP, the other linking DXS255 to DXS146. *Genomics* 29:496-502
- 4) Fisher SE, van Bakel I, Lloyd SE, Pearce SHS, Thakker RV, Craig IW (1995) Cloning and characterization of CLCN5, the human kidney chloride channel gene implicated in Dent disease (an X-linked hereditary nephrolithiasis). *Genomics* 29:598-606
- 5) Shipley JM, Birdsall S, Clark J, Crew J, Gill S, Linehan M, Gnarra J, Fisher S, Craig IW, Cooper CS (1995) Mapping the chromosome X breakpoint in two papillary renal cell carcinoma cell lines with a t(X;1) (p11.2; q21.2) and the first report of a female case. *Cytogenet Cell Genet* 71:280-4
- 6) Lloyd SE, Pearce SHS, Fisher SE, Steinmeyer K, Schwappach B, Scheinman SJ, Harding B, Bolino A, Devoto M, Goodyer P, Rigden SPA, Wrong O, Jentsch TJJ, Craig IW, Thakker RV (1996) A common molecular basis for three inherited kidney stone diseases. *Nature* 379:445-9
- 7) Weterman MAJ, Wilbrink M, Janssen I, Janssen HAP, van den Berg E, Fisher SE, Craig I, Geurts van Kessel A (1996) Molecular cloning of the papillary renal cell carcinoma-associated translocation (X;1) (p11;q21) breakpoint. *Cytogenet Cell Genet* 75:2-6
- 8) Lloyd SE, Günther W, Pearce SHS, Thomson A, Bianchi ML, Bosio M, Craig IW, Fisher SE, Scheinman SJ, Wrong O, Jentsch TJ, Thakker RV (1997) Characterization of renal chloride channel CLCN5 mutations in hypercalciuric nephrolithiasis (kidney stones) disorders. *Hum Mol Genet* 6:1233-9
- 9) Fisher SE, Ciccodicola A, Tanaka K, Curci A, Desicato S, D'Urso M, Craig IW (1997) Sequence-based exon prediction around the SYP locus reveals a gene rich area containing novel genes in human proximal Xp. *Genomics* 45:340-7
- 10) Fisher SE, Vargha-Khadem F, Watkins KE, Monaco AP, Pembrey ME (1998) Localisation of a gene implicated in a severe speech and language disorder. *Nature Genet* 18:168-70
- 11) Fisher SE, Marlow AJ, Lamb J, Maestrini E, Williams DF, Richardson AJ, Weeks DE, Stein JF, Monaco AP (1999) A quantitative-trait locus on chromosome 6p influences different aspects of developmental dyslexia. *Am J Hum Genet* 64:146-56
- 12) Tanaka K, Fisher SE and Craig IW (1999) Characterisation and genomic organisation of the mouse *Cln5* gene and identification of putative promoter and enhancer regions. *Genomics* 58:281-92
- 13) Lai CSL¹, Fisher SE¹, Hurst JA, Levy ER, Hodgson S, Fox M, Jeremiah S, Povey S, Jamison DC, Green ED, Vargha-Khadem F, Monaco AP (2000) The SPCH1 region on human 7q31: genomic characterization of the critical interval and localization of translocations associated with speech and language disorder. *Am J Hum Genet* 67:357-68; ¹joint first authors
- 14) Siddiqui MR, Meisner S, Tosh K, Balakrishnan K, Ghei S, Fisher SE, Golding M, Narayan NPS, Sitaraman T, Sengupta U, Pitchappan RM, Hill AVS (2001) A major susceptibility locus for leprosy in India maps to chromosome 10p13. *Nature Genet* 27:439-41
- 15) Marlow AJ, Fisher SE, Richardson AJ, Francks C, Talcott JB, Monaco AP, Stein JF, Cardon LR (2001) Investigation of quantitative measures related to reading disability in a large sample of sib-pairs from the UK. *Behav Genet* 31:219-30
- 16) Lai CSL¹, Fisher SE¹, Hurst JA, Vargha-Khadem F, Monaco AP (2001) A forkhead-domain gene is mutated in a severe speech and language disorder. *Nature* 413:519-23; ¹joint first authors

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 17)** Fisher SE, Francks C, Marlow AJ, MacPhie IL, Newbury DF, Cardon LR, Ishikawa-Brush Y, Richardson AJ, Talcott JB, Gayán J, Olson RK, Pennington BF, Smith SD, DeFries JC, Stein JF, Monaco AP (2002) Independent genome-wide scans identify a chromosome 18 quantitative-trait locus influencing dyslexia. *Nature Genet* 30:86-91
- 18)** The SLI Consortium (2002) A genome-wide scan identifies two novel loci involved in Specific Language Impairment (SLI). *Am J Hum Genet* 70:384-98
- 19)** Francks C, Fisher SE, MacPhie IL, Richardson AJ, Marlow AJ, Stein JF, Monaco AP (2002) A genome wide linkage screen for relative hand skill in sibling pairs. *Am J Hum Genet* 70:800-5
- 20)** Francks C, Fisher SE, Olson RK, Pennington BF, Smith SD, DeFries JC, Monaco AP (2002) Quantitative association analysis within the chromosome 2p12-16 dyslexia susceptibility region: Microsatellite markers and candidate genes SEMA4F and OTX1. *Psych Genet* 12:35-41
- 21)** Fisher SE, Francks C, McCracken JT, McGough JT, Marlow AJ, MacPhie IL, Newbury DF, Crawford LR, Palmer CJS, Woodward JA, Del'Homme M, Cantwell D, Nelson SF, Monaco AP, Smalley SL (2002) A genomewide scan for loci involved in Attention-Deficit/Hyperactivity Disorder. *Am J Hum Genet* 70:1183-96
- 22)** Newbury DF, Bonora E, Lamb JA, Fisher SE, Lai CSL, Baird G, Jannoun L, Slonims V, Stott CM, Merricks MJ, Bolton PF, Bailey A, Monaco AP and the International Molecular Genetic Study of Autism Consortium (2002) FOXP2 is not a major susceptibility gene for autism or Specific Language Impairment (SLI). *Am J Hum Genet* 70:1318-27
- 23)** Enard W, Przeworski M, Fisher SE, Lai CSL, Wiebe V, Kitano T, Monaco AP, Pääbo S (2002) Molecular evolution of FOXP2, a gene involved in speech and language. *Nature* 418:869-72
- 24)** Smalley SL, Kustanovich V, Minassian SL, Stone J, Ogdie M, McGough JJ, McCracken, MacPhie IL, Francks C, Fisher SE, Cantor RM, Monaco AP, Nelson SF (2002) Genetic linkage of Attention-Deficit/Hyperactivity Disorder (ADHD) on chromosome 16p13 in a region implicated in autism. *Am J Hum Genet* 71:959-63
- 25)** Francks C, DeLisi LE, Fisher SE, Laval SH, Rue JE, Stein JF, Monaco AP (2003) Confirmatory evidence for linkage of relative hand skill to 2p12-q11. *Am J Hum Genet* 72:499-502
- 26)** Marlow AJ, Fisher SE, Francks C, MacPhie IL, Richardson AJ, Talcott JB, Stein JF, Monaco AP, Cardon LR (2003) Use of multivariate linkage analysis for dissection of a complex cognitive trait. *Am J Hum Genet* 72:561-70
- 27)** Ogdie MN, Macphie IL, Minassian SL, Yang M, Fisher SE, Francks C, Cantor RM, McCracken JT, McGough JJ, Nelson SF, Monaco AP, Smalley SL (2003) A genome-wide scan for Attention-Deficit/Hyperactivity Disorder in an extended sample: suggestive linkage on 17p11. *Am J Hum Genet* 72:1268-79
- 28)** Lai CSL, Gerrelli D, Monaco AP, Fisher SE¹, Copp AJ¹ (2003) FOXP2 expression during brain development coincides with adult sites of pathology in a severe speech and language disorder. *Brain* 126:2455-62; ¹joint corresponding authors
- 29)** Francks C, Fisher SE, Marlow AJ, MacPhie IL, Taylor KE, Richardson AJ, Stein JF, Monaco AP (2003) Familial and genetic effects on motor coordination, laterality, and reading-related cognition. *Am J Psych* 160:1970-7
- 30)** Francks C, DeLisi LE, Shaw SH, Fisher SE, Richardson AJ, Stein JF, Monaco AP (2003) Parent-of-origin effects on handedness and schizophrenia susceptibility on chromosome 2p12-q11. *Hum Mol Genet* 12:3225-30
- 31)** Loo SK, Fisher SE, Francks C, Ogdie MN, MacPhie IL, Yang M, McCracken JT, McGough JJ, Nelson SF, Monaco AP, Smalley SL (2004) Genome-wide scan of reading ability in affected sibling pairs with attention-deficit/hyperactivity disorder: unique and shared genetic effects. *Mol Psychiatry* 9:485-93
- 32)** Ogdie MN, Fisher SE, Yang M, Ishii J, Francks C, RM, McCracken JJ, McGough JT, Smalley SL, Nelson SF (2004) Attention Deficit Hyperactivity Disorder: fine mapping supports linkage to 5p13, 6q12, 16p13, and 17p11. *Am J Hum Genet* 75:661-8

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 33)** The SLI Consortium (2004) Highly significant linkage to the SLI1 locus in an expanded sample of individuals affected by specific language impairment. *Am J Hum Genet* 74:1225-38
- 34)** Scerri TS, Fisher SE, Francks C, MacPhie IL, Paracchini S, Richardson AJ, Stein JF, Monaco AP (2004) Putative functional alleles of DYX1C1 are not associated with dyslexia susceptibility in a large sample of sibling pairs from the UK. *J Med Genet* 41:853-7
- 35)** Francks C, Paracchini S, Smith SD, Richardson AJ, Scerri TS, Cardon LR, Marlow AJ, MacPhie IL, Walter J, Pennington BF, Fisher SE, Olson RK, DeFries JC, Stein JF, Monaco AP (2004) A 77 kilobase region of chromosome 6p22.2 is associated with dyslexia in families from the United Kingdom and from the United States. *Am J Hum Genet* 75:1046-58
- 36)** MacDermot KD, Bonora E, Sykes N, Coupe AM, Lai CSL Vernes SC, Vargha-Khadem F, McKenzie F, Smith RL, Monaco AP, Fisher SE (2005) Identification of FOXP2 truncation as a novel cause of developmental speech and language deficits. *Am J Hum Genet* 76:1074-80
- 37)** Gayán J, Willcutt EG, Fisher SE, Francks C, Cardon LR, Olson RK, Pennington BF, Smith SD, Monaco AP, DeFries JC (2005) Bivariate linkage scan for reading disability and attention-deficit/hyperactivity disorder localizes pleiotropic loci. *J Child Psychol Psychiatr* 46:1045-56
- 38)** Ogdie MN, Bakker SC, Fisher SE, Francks C, Yang MH, Cantor RM, Loo SK, van der Meulen E, Pearson P, Buitelaar J, Monaco A, Nelson SF, Sinke RJ, Smalley SL (2006) Pooled genome-wide linkage data on 424 ADHD ASPs suggests genetic heterogeneity and a common risk locus at 5p13. *Mol Psychiatry* 11:5-8
- 39)** Vernes SC, Nicod J, Elahi FM, Coventry JA, Kenny N, Coupe A-M, Bird LE, Davies KE, Fisher SE (2006) Functional genetic analysis of mutations implicated in a human speech and language disorder. *Hum Mol Genet* 15:3154-67
- 40)** French CA, Groszer M, Preece C, Coupe A-M, Rajewsky K, Fisher SE (2007) Generation of mice with a conditional Foxp2 null allele. *Genesis* 45:440-6
- 41)** Monaco AP & The SLI Consortium (2007) Multivariate linkage analysis of Specific Language Impairment (SLI). *Ann Hum Genet* 71:660-73
- 42)** Francks C, Maegawa S, Laurén J, Abrahams B, Velayos-Baeza A, Medland SE, Colella S, Groszer M, McAuley EZ, Caffrey TM, Timmusk T, Pruunsild P, Koppel I, Lind PA, Matsumoto-Itaba N, Nicod J, Xiong L, Joober R, Enard W, Krinsky B, Nanba E, Richardson AJ, Riley BP, Martin NG, Strittmatter SM, Möller H-J, Rujescu D, St Clair D, Muglia P, Roos JL, Fisher SE, Wade-Martins R, Rouleau GA, Stein JF, Karayiorgou M, Geschwind DH, Ragoussis J, Kendler KS, Airaksinen MS, Oshimura M, DeLisi LE, Monaco AP (2007) LRRTM1 on chromosome 2p12 is a maternally suppressed gene that is associated paternally with handedness and schizophrenia. *Mol Psychiatry* 12:1129-39, 1057
- 43)** Vernes SC, Spiteri E, Nicod J, Groszer M, Taylor JM, Davies KE, Geschwind DH, Fisher SE (2007) High-throughput analysis of promoter occupancy identifies direct neural targets of FOXP2, a gene mutated in speech and language disorders. *Am J Hum Genet* 81:1232-50
- 44)** Spiteri E, Konopka G, Coppola G, Bomar J, Oldham M, Ou J, Vernes SC, Fisher SE, Ren B, Geschwind DH (2007) Identification of the transcriptional targets of FOXP2, a gene involved in speech and language, in developing human brain. *Am J Hum Genet* 81:1144-57
- 45)** Groszer M, Keays DA, Deacon RM, de Bono JP, Prasad-Mulcare S, Gaub S, Baum MG, French CA, Nicod J, Coventry JA, Enard W, Fray M, Brown SD, Nolan PM, Pääbo S, Channon KM, Costa RM, Eilers J, Ehret G, Rawlins JN, Fisher SE (2008) Impaired synaptic plasticity and motor learning in mice with a point mutation implicated in human speech deficits. *Curr Biol* 18:354-62
- 46)** Falcaro M, Pickles A, Newbury DF, Addis L, Banfield E, Fisher SE, Monaco AP, Simkin Z, Conti-Ramsden G, and the SLI Consortium (2008) Genetic and phenotypic effects of phonological short-term memory and grammatical morphology in Specific Language Impairment. *Genes Brain Behav* 7:393-402
- 47)** Vernes SC, Newbury DF, Abrahams BS, Winchester L, Nicod J, Groszer M, Alarcón M, Oliver PL, Davies KE, Geschwind DH, Monaco AP, Fisher SE (2008) A functional genetic link between distinct developmental language disorders. *N Engl J Med* 359:2337-45

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 48)** Vernes SC, MacDermot KD, Monaco AP, [Fisher SE](#) (2009) Assessing the impact of FOXP1 mutations on developmental verbal dyspraxia. *Eur J Hum Genet* 17:1354-8
- 49)** Enard W, Gehre S, Hammerschmidt K, Hölter SM, Blass T, Somel M, Brückner MK, Schreiweis C, Winter C, Sohr R, Becker L, Wiebe V, Nickel B, Giger T, Müller U, Groszer M, Adler T, Aguilar A, Bolle I, Calzadawack J, Claudia C, Ehrhardt N, Favor J, Fuchs H, Gailus-Durner V, Hans W, Hözlwimmer G, Javaheri A, Kalaydjiev S, Kallnik M, Kling E, Kunder S, Moßbrugger I, Naton B, Racz I, Rathkolb B, Rozman J, Schrewe A, Busch DH, Graw J, Ivandic B, Klingenspor M, Klopstock T, Ollert M, Quintanilla-Martinez L, Schulz H, Wolf E, Wurst W, Zimmer A, [Fisher SE](#), Morgenstern R, Arendt T, Hrabé de Angelis M, Fischer J, Schwarz J, Pääbo S (2009) A humanized version of Foxp2 affects cortico-basal ganglia circuits in mice. *Cell* 137:961-71
- 50)** Kurt S, Groszer M, [Fisher SE](#), Ehret G (2009) Modified sound-evoked brainstem potentials in Foxp2 mutant mice. *Brain Res* 1289:30-6
- 51)** Newbury DF, Winchester L, Addis L, Paracchini S, Buckingham LL, Clark A, Cohen W, Cowie H, Dworzynski K, Everitt A, Goodyer IM, Hennessy E, Kindley AD, Miller LL, Nasir J, O'Hare A, Shaw D, Simkin Z, Simonoff E, Slonims V, Watson J, Ragoussis J, [Fisher SE](#), Seckl J, Helms PJ, Bolton PF, Pickles A, Conti-Ramsden G, Baird G, Bishop DVM, Monaco AP, SLI Consortium (2009) CMIP and ATP2C2 modulate phonological short-term memory in language impairment. *Am J Hum Genet* 85:264-72
- 52)** Gaub S, Groszer M, [Fisher SE](#), Ehret G (2010) The structure of innate vocalizations in Foxp2 deficient mouse pups. *Genes Brain Behav* 9:390-401
- 53)** Roll P, Vernes SC, Bruneau N, Cillario J, Ponsolle-Lenfant M, Massacrier A, Rudolf G, Khalife M, Hirsch E, [Fisher SE](#), Szepetowski P (2010) Molecular networks implicated in speech-related disorders: FOXP2 regulates the SRPX2/uPAR complex. *Hum Mol Genet* 19:4848-60
- 54)** Whitehouse AJO, Bishop DVM, Ang QW, Pennell CE, [Fisher SE](#) (2011) CNTNAP2 variants affect early language development in the general population. *Genes Brain Behav* 10:451-6
- 55)** O'Roak BJ, Deriziotis P, Lee C, Vives L, Schwartz JJ, Girirajan S, Karakoc E, MacKenzie AP, Ng S, Baker C, Rieder M, Nickerson D, Bernier R, [Fisher SE](#), Shendure J, Eichler EE (2011) Exome sequencing in sporadic autism spectrum disorders identifies severe de novo mutations. *Nature Genet* 43:585-9
- 56)** Vernes SC, Oliver PL, Spiteri E, Lockstone HE, Puliyadi R, Taylor JM, Ho J, Mombereau C, Brewer A, Lowy E, Nicod J, Groszer M, Baban D, Sahgal N, Cazier J-B, Ragoussis J, Davies KE, Geschwind DH, [Fisher SE](#) (2011) Foxp2 regulates gene networks implicated in neurite outgrowth in the developing brain. *PLoS Genet* 7:e1002145
- 57)** French CA, Jin X, Campbell TG, Gerfen E, Groszer M, [Fisher SE](#), Costa RM (2012) An aetiological Foxp2 mutation causes aberrant striatal activity and alters plasticity during skill learning. *Mol Psychiatry* 17:1077-85
- 58)** Kurt S, [Fisher SE](#), Ehret G (2012) Foxp2 mutations impair auditory-motor-association learning. *PLoS One* 7:e33130
- 59)** Walker RM, Hill AE, Newman AC, Hamilton G, Torrance HS, Anderson SM, Ogawa F, Derizioti P, Nicod J, Vernes SC, [Fisher SE](#), Thomson PA, Porteous DJ, Evans KL (2012) The DISC1 promoter: characterisation and regulation by FOXP2. *Hum Mol Genet* 21:2862-72
- 60)** Newbury DF, Mari F, Sadighi Akha E, Hurst J, MacDermot KD, Canitano R, Monaco AP, Taylor JC, Renieri A, [Fisher SE](#), Knight SJL (2013) Dual copy number variants involving 16p11 and 6q22 in a case of childhood apraxia of speech and pervasive developmental disorder. *Eur J Hum Genet* 21:361-5
- 61)** Gialluisi A, Dediu D, Francks C, [Fisher SE](#) (2013) Persistence and transmission of recessive deafness and sign language: new insights from village sign languages. *Eur J Hum Genet* 21:894-6
- 62)** Gregersen PK, Kowalsky E, Lee A, Baron-Cohen S, [Fisher SE](#), Asher JE, Ballard D, Freudenberg J, Li W (2013) Absolute pitch exhibits phenotypic and genetic overlap with synesthesia. *Hum Mol Genet* 22:2097-104
- 63)** Ayub Q, Yngvadottir B, Chen Y, Xue Y, Hu M, Vernes SC, [Fisher SE](#), Tyler-Smith C (2013) FOXP2 targets show evidence of positive selection in European populations. *Am J Hum Genet* 92:696-706

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 64)** Brandler WM, Morris AP, Evans DM, Scerri TS, Kemp JP, Timpson NJ, St Pourcain B, Davey Smith G, Ring SM, Stein J, Monaco AP, Talcott JB, [Fisher SE](#), Webber C, Paracchini S (2013) Common variants in left/right asymmetry genes and pathways are associated with relative hand skill. *PLoS Genet* 9:e1003751
- 65)** Baron-Cohen S, Johnson D, Asher J, Wheelwright S, [Fisher SE](#), Gregersen PK, Allison C (2013) Is synaesthesia more common in autism? *Mol Autism* 4:40
- 66)** Simpson NH, Addis L, Brandler WM, Slonims V, Clark A, Watson J, Scerri TS, Hennessy ER, Stein J, Talcott J, Conti-Ramsden G, O'Hare A, Baird G, Fairfax BP, Knight JC, Paracchini S, [Fisher SE](#), Newbury DF, the SLI Consortium (2014) Increased prevalence of sex chromosome aneuploidies in Specific Language Impairment and dyslexia. *Dev Med Child Neurol* 56:346-53
- 67)** Guadalupe T, Zwiers MP, Teumer A, Wittfeld K, Arias Vasquez A, Hoogman M, Hagoort P, Fernández G, Buitelaar J, Hegenscheid K, Völzke H, Franke B, [Fisher SE](#), Grabe HJ, Francks C (2014) Measurement and genetics of human subcortical and hippocampal asymmetries in large datasets. *Hum Brain Mapp* 35:3277-89
- 68)** Nudel R, Simpson NH, Baird G, O'Hare A, Conti-Ramsden G, Bolton PF, Hennessy ER, SLI Consortium, Monaco AP, Fairfax BP, Knight JC, Winney B, [Fisher SE](#), Newbury DF (2014) Associations of HLA alleles with specific language impairment. *J Neurodev Disord* 6:1
- 69)** Derizioti P, Graham SA, Busquets Estruch S, [Fisher SE](#) (2014) Investigating protein-protein interactions in live cells using Bioluminescence Resonance Energy Transfer. *J Vis Exp* 87:e51438
- 70)** Ceroni F, Simpson NH, Francks C, Baird G, Conti-Ramsden G, Clark A, Bolton PF, Hennessy ER, Donnelly P, Bentley DR, Martin H, IMGSAC, SLI Consortium, WGS500 Consortium, Parr J, Pagnamenta AT, Maestrini E, Bacchelli E, [Fisher SE](#), Newbury DF (2014) Homozygous microdeletion of exon 5 in ZNF277 in a girl with specific language impairment. *Eur J Hum Genet* 22:1165-71
- 71)** Nudel R, Simpson NH, Francks C, Baird G, O'Hare A, Conti-Ramsden G, Bolton PF, Hennessy ER, SLI Consortium, ALSPAC, Paracchini S, Monaco AP, [Fisher SE](#), Newbury DF (2014) Genome-wide association analyses of imprinting effects in specific language impairment identify loci shared with other neurodevelopmental disorders. *Genes Brain Behav* 13:418-29
- 72)** Guadalupe T, Willems RM, Zwiers MP, Arias Vasquez A, Hoogman M, Hagoort P, Fernández G, Buitelaar J, Franke B, [Fisher SE](#), Francks C (2014) Differences in cerebral cortical anatomy of left- and right-handers. *Front Psychol* 5:261
- 73)** Baron-Cohen S, Murphy L, Chakrabarti B, Craig I, Mallya U, Lakatošová S, Rehnstrom K, Peltonen L, Wheelwright S, Allison C, [Fisher SE](#), Warrier V (2014) A genome wide association study of mathematical ability reveals an association at chromosome 3q29, a locus associated with autism and learning difficulties: a preliminary study. *PLoS One* 9:e96374
- 74)** Hoogman M, Guadalupe T, Zwiers MP, Klarenbeek P, Francks C, [Fisher SE](#) (2014) Assessing the effects of common variation in the FOXP2 gene on human brain structure. *Front Hum Neurosci* 8:473
- 75)** Cai D, Fonteijn H, Guadalupe T, Zwiers M, Wittfeld K, Teumer A, Hoogman M, Arias-Vásquez A, Yang Y, Buitelaar J, Fernández G, Brunner HG, van Bokhoven H, Franke B, Hegenscheid K, Homuth G, [Fisher SE](#), Grabe HJ, Francks C, Hagoort P (2014) A genome wide search for quantitative trait loci affecting the cortical surface area and thickness of Heschl's gyrus. *Genes Brain Behav* 13:675-85
- 76)** Gialluisi A, Newbury DF, Wilcutt EG, Olson RK, DeFries JC, Brandler WM, Pennington BF, Smith SD, Scerri TS, Simpson NH, The SLI Consortium, Luciano M, Evans DM, Bates TC, Stein JF, Talcott JB, Monaco AP, Paracchini S, Francks C, [Fisher SE](#) (2014) Genome-wide screening for DNA variants associated with reading and language traits. *Genes Brain Behav* 13:686-701
- 77)** Brucato N, DeLisi LE, [Fisher SE](#), Francks C (2014) Hypomethylation of the paternally inherited LRRTM1 promoter linked to schizophrenia. *Am J Med Genet B* 165B:555-63
- 78)** Deriziotis P, O'Roak BJ, Graham SA, Estruch SB, Dimitropoulou D, Bernier RA, Gerds J, Shendure JA, Eichler EE, [Fisher SE](#) (2014) De novo TBR1 mutations in sporadic autism disrupt protein functions. *Nature Commun* 5:4954

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 79)** Schreiweis C, Bornschein U, Burguière E, Kerimoglu C, Schreiter S, Dannemann M, Goyal S, Rea E, French CA, Puliyadi R, Groszer M, Fisher SE, Mundry R, Winter C, Hevers W, Pääbo S, Enard W, Graybiel AM (2014) Humanized Foxp2 accelerates learning by enhancing transitions from declarative to procedural performance. *Proc Natl Acad Sci USA* 111:14253-8
- 80)** Cousijn H, Eissing M, Zwiers M, Fernández G, Fisher SE, Franke B, Harrison PJ, Arias-Vasquez A (2014) No effect of schizophrenia risk genes MIR137, TCF4, and ZNF804A on macroscopic brain structure. *Schizophr Res* 159:329-32
- 81)** Guadalupe T, Zwiers MP, Wittfeld K, Teumer A, Arias-Vásquez A, Hoogman M, Hagoort P, Fernández G, Buitelaar J, van Bokhoven H, Hegenscheid K, Völzke H, Franke B, Fisher SE, Grabe HJ, Francks C (2015) Asymmetry within and around the planum temporale is sexually dimorphic and influenced by genes involved in steroid biology. *Cortex* 62:41-55
- 82)** Gupta CN, Calhoun VD, Rachkonda S, Chen J, Liu J, Segall J, Franke B, Zwiers MP, Arias-Vasquez A, Buitelaar J, Fisher SE, Fernández G, van Erp TGM, Potkin S, Ford J, Mathalon D, McEwen S, Lee HJ, Mueller BA, Greve DN, Andreassen O, Agartz I, Gollub RL, Sponheim SR, Ehrlich S, Wang L, Pearlson G, Glahn DC, Sprooten E, Mayer AR, Stephen J, Jung RE, Canive J, Bustillo J, Turner JA (2015) Patterns of gray matter abnormalities in schizophrenia based on an international mega-analysis. *Schizophr Bull* 41:1133-42
- 83)** Zhao H, Zhou W, Yao Z, Wan Y, Cao J, Zhang L, Zhao J, Li H, Zhou R, Li B, Wei G, Zhang Z, French CA, Dekker JD, Yang Y, Yao Z, Fisher SE, Tucker HO, Guo X (2015) Foxp1/2/4 regulate osteogenesis and chondrocyte hypertrophy during endochondral ossification. *Dev Biol* 398:242-54
- 84)** Simpson NH, Ceroni F, Reader RH, Covill L, Fairfax BP, Knight JC, the SLI Consortium, Hennessy ER, Bolton PF, Conti-Ramsden G, O'Hare A, Baird G, Fisher SE, Newbury DF (2015) Genome-wide analysis identifies a role for common copy number variants in specific language impairment. *Eur J Hum Genet* 23:1370-7
- 85)** Hibar DP *et al* (2015) Common genetic variants influence human subcortical brain structures. *Nature* 520:224-9
- 86)** Villanueva P, Nudel R, Hoischen A, Fernández MA, Simpson NH, Gilissen C, Reader RH, Jara L, Echeverry MM, Francks C, Baird G, Conti-Ramsden G, O'Hare A, Bolton PF, Hennessy ER, the SLI Consortium, Palomino H, Carvajal-Carmona L, Veltman JA, Cazier J-B, De Barbieri Z, Fisher SE¹, Newbury DF¹ (2015) Exome sequencing in an admixed isolated population indicates NFXL1 variants confer a risk for Specific Language Impairment. *PLoS Genet* 11:e1004925; ¹joint senior authors
- 87)** Brucato N, Guadalupe T, Franke B, Fisher SE, Francks C (2015) A schizophrenia-associated HLA locus affects thalamus volume and asymmetry. *Brain Behav Immun* 46:311-8
- 88)** Pettigrew KA, Fajutrao Valles SF, Moll K, Northstone K, Ring S, Pennell C, Wang C, Leavett R, Hayiou-Thomas ME, Thompson P, Simpson NH, Fisher SE, the SLI Consortium, Whitehouse AJO, Snowling MJ, Newbury DF, Paracchini S (2015) Lack of replication for the myosin-18B association with mathematical ability in independent cohorts. *Genes Brain Behav* 14:369-76
- 89)** Spaeth JM, Hunter CS, Bonatakis L, Guo M, French CA, Slack I, Hara M, Fisher SE, Ferrer J, Morrissey EE, Stanger BZ, Stein R (2015) The FOXP1, FOXP2 and FOXP4 transcription factors are required for islet alpha cell proliferation and function in mice. *Diabetologia* 58:1836-44
- 90)** Gascoyne DM, Spearman H, Lyne L, Puliyadi R, Pérez-Alcántara M, Coulton L, Fisher SE, Croucher PI, Banham AH (2015) The forkhead transcription factor FOXP2 is required for regulation of p21WAF1/CIP1 in 143B osteosarcoma cell growth arrest. *PLoS ONE* 10:e0128513
- 91)** Warriar V, Chakrabarti B, Murphy L, Chan A, Craig I, Mallya U, Lakatošová S, Rehnstrom K, Peltonen L, Wheelwright S, Allison C, Fisher SE, Simon Baron-Cohen S (2015) A pooled genome-wide association study of Asperger Syndrome. *PLoS ONE* 10:e0131202
- 92)** Chen J, Calhoun VD, Arias-Vasquez A, Zwiers MP, van Hulzen K, Fernández G, Fisher SE, Franke B, Turner JA, Liu J (2015) G-protein genomic association with normal variation in gray matter density. *Hum Brain Mapp* 36:4272-86

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 93)** Lozano R, Vino A, Cristina Lozano C, [Fisher SE](#), Deriziotis P (2015) A de novo FOXP1 variant in a patient with autism, intellectual disability and language impairment. *Eur J Hum Genet* 23:1702-7
- 94)** Becker M, Devanna P, [Fisher SE](#), Vernes SC (2015) A chromosomal rearrangement in a child with severe speech and language disorder separates FOXP2 from a functional enhancer. *Mol Cytogenet* 8:69
- 95)** Fedorenko E, Morgan A, Murray E, Cardinaux A, Mei C, Tager-Flusberg H, [Fisher SE](#), Kanwisher N (2016) A highly penetrant form of childhood apraxia of speech due to deletion of 16p11.2. *Eur J Hum Genet* 24:302-6
- 96)** Gaub S, [Fisher SE](#), Ehret G (2016) Ultrasonic vocalizations of adult male Foxp2-mutant mice: behavioral contexts of arousal and emotion. *Genes Brain Behav* 15:243-59
- 97)** Sollis E, Graham SA, Vino A, Froehlich H, Vreeburg M, Dimitropoulou D, Gilissen C, Pfundt R, Rappold GA, Brunner HG, Deriziotis P, [Fisher SE](#) (2016) Identification and functional characterization of de novo FOXP1 variants provides novel insights into the etiology of neurodevelopmental disorder. *Hum Mol Genet* 25:546-57
- 98)** Estruch SB, Graham SA, Deriziotis P, [Fisher SE](#) (2016) The language-related transcription factor FOXP2 is post-translationally modified with small ubiquitin-like modifiers. *Sci Rep* 6:20911
- 99)** Becker M, Guadalupe T, Franke B, Hibar DP, Renteria ME, Stein JL, Thompson PM, Francks C, Vernes SC, [Fisher SE](#) (2016) Early developmental gene enhancers affect subcortical volumes in the adult human brain. *Hum Brain Mapp* 37:1788-800
- 100)** Gialluisi A, Visconti A, Wilcutt EG, Smith SD, Pennington BF, Falchi M, DeFries JC, Olson RK, Francks C, [Fisher SE](#) (2016) Investigating the effects of copy number variants on reading and language performance. *J Neurodev Disord* 8:17
- 101)** Carrion-Castillo A, van Bergen E, Vino A, van Zuijlen T, de Jong PF, Francks C, [Fisher SE](#) (2016) Evaluation of results from genome-wide studies of language and reading in a novel independent dataset. *Genes Brain Behav* 15:531-41
- 102)** Woo YJ, Wang T, Guadalupe T, Nebel RA, Vino A, Del Bene VA, Molholm S, Ross LA, Zwiers MP, [Fisher SE](#), Foxe JJ, Abrahams BS (2016) A common CYFIP1 variant at the 15q11.2 disease locus is associated with structural variation at the language-related left supramarginal gyrus. *PLoS ONE* 11:e0158036
- 103)** Dias C, Estruch SB, Graham SA, McRae J, Sawiak SJ, Hurst JA, Joss SK, Holder SE, Morton JEV, Turner C, Thevenon J, Mellul K, Sanchez-Andrade G, Ibarra-Soria X, Deriziotis P, Santos RF, Lee S-C, Faivre L, Kleefstra T, Liu P, Hurles ME, DDD Study, [Fisher SE](#)¹, Logan DW¹ (2016) BCL11A haploinsufficiency causes an intellectual disability syndrome and dysregulates transcription *Am J Hum Genet* 99:253-74;
- ¹joint corresponding authors**
- 104)** Li S, Morley M, Lu M, Zhou S, Stewart K, French CA, Tucker HO, [Fisher SE](#), Morrisey EE (2016) Foxp transcription factors suppress a non-pulmonary gene expression program to permit proper lung development. *Dev Biol* 416:338-46
- 105)** Chabout J¹, Sarkar A, Patel SR, Raiden T, Dunson DB, [Fisher SE](#)¹, Jarvis ED¹ (2016) A Foxp2 mutation implicated in human speech deficits alters sequencing of ultrasonic vocalizations in adult male mice. *Front Behav Neurosci* 10:197; **¹joint corresponding authors**
- 106)** Estruch SB, Graham SA, Chinnappa SM, Deriziotis P, [Fisher SE](#) (2016) Functional characterization of rare FOXP2 variants in neurodevelopmental disorder. *J Neurodev Disord* 8:44
- 107)** Adams HH *et al* (2016) Novel genetic loci underlying human intracranial volume identified through genome-wide association. *Nature Neurosci* 19:1569-82
- 108)** Uddén J, Snijders TM, [Fisher SE](#), Hagoort P (2017) A common variant of the CNTNAP2 gene is associated with structural variation in the left superior occipital gyrus. *Brain Lang* 172:16-21
- 109)** Kavaklioglu T, Guadalupe T, Zwiers M, Marquand AF, Onnink M, Shumskaya E, Brunner H, Fernandez G, [Fisher SE](#), Francks C (2017) Structural asymmetries of the human cerebellum in relation to cerebral cortical asymmetries and handedness. *Brain Struct Funct* 222:1611-23

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 110)** Gialluisi A, Guadalupe T, Francks C, [Fisher SE](#) (2017) Neuroimaging genetic analyses of novel candidate genes associated with reading and language. *Brain Lang* 172:9-15
- 111)** Guadalupe T, Mathias SR, vanErp TG, Whelan CD, Zwiers MP, Abe Y, Abramovic L, Agartz I, Andreassen OA, Arias-Vásquez A, Aribisala BS, Armstrong NJ, Arolt V, Artiges E, Ayesa-Arriola R, Baboyan VG, Banaschewski T, Barker G, Bastin ME, Baune BT, Blangero J, Bokde AL, Boedhoe PS, Bose A, Brem S, Brodaty H, Bromberg U, Brooks S, Büchel C, Buitelaar J, Calhoun VD, Cannon DM, Cattrell A, Cheng Y, Conrod PJ, Conzelmann A, Corvin A, Crespo-Facorro B, Crivello F, Dannlowski U, de Zubicaray GI, de Zwarte SM, Deary IJ, Desrivières S, Doan NT, Donohoe G, Dørum ES, Ehrlich S, Espeseth T, Fernández G, Flor H, Fouche JP, Frouin V, Fukunaga M, Gallinat J, Garavan H, Gill M, Suarez AG, Gowland P, Grabe HJ, Grotegerd D, Gruber O, Hagenaars S, Hashimoto R, Hauser TU, Heinz A, Hibar DP, Hoekstra PJ, Hoogman M, Howells FM, Hu H, Hulshoff Pol HE, Huyser C, Ittermann B, Jahanshad N, Jönsson EG, Jurk S, Kahn RS, Kelly S, Kraemer B, Kugel H, Kwon JS, Lemaitre H, Lesch KP, Lochner C, Luciano M, Marquand AF, Martin NG, Martínez-Zalacáin I, Martinot JL, Mataix-Cols D, Mather K, McDonald C, McMahon KL, Medland SE, Menchón JM, Morris DW, Mothersill O, Maniega SM, Mwangi B, Nakamae T, Nakao T, Narayanaswamy JC, Nees F, Nordvik JE, Onnink AM, Opel N, Ophoff R, Paillère Martinot ML, Papadopoulos Orfanos D, Pauli P, Paus T, Poustka L, Reddy JY, Renteria ME, Roiz-Santiañez R, Roos A, Royle NA, Sachdev P, Sánchez-Juan P, Schmaal L, Schumann G, Shumskaya E, Smolka MN, Soares JC, Soriano-Mas C, Stein DJ, Strike LT, Toro R, Turner JA, Tzourio-Mazoyer N, Uhlmann A, Hernández MV, van den Heuvel OA, van der Meer D, van Haren NE, Veltman DJ, Venkatasubramanian G, Vetter NC, Vuletic D, Walitza S, Walter H, Walton E, Wang Z, Wardlaw J, Wen W, Westlye LT, Whelan R, Wittfeld K, Wolfers T, Wright MJ, Xu J, Xu X, Yun JY, Zhao J, Franke B, Thompson PM, Glahn DC, Mazoyer B, [Fisher SE](#), Francks C (2017) Human subcortical brain asymmetries in 15847 people worldwide reveal effects of age and sex. *Brain Imaging Behav* 11:1497-1514
- 112)** Hibar DP *et al* (2017) Novel genetic loci associated with hippocampal volume. *Nature Commun* 8:13624
- 113)** Carrion-Castillo A, Maassen B, Franke B, Heister A, Naber M, van der Leij A, Francks C, [Fisher SE](#) (2017) Association analysis of dyslexia candidate genes in a Dutch longitudinal sample. *Eur J Hum Genet* 25:452-60
- 114)** De Kovel CGF, Lisgo S, Karlebach G, Ju J, Cheng G, [Fisher SE](#), Francks C (2017) Left-right asymmetry of maturation rates in human embryonic neural development. *Biol Psychiatry* 82:204-12
- 115)** Acuna-Hidalgo R, Deriziotis P, Steehouwer M, Gilissen C, Graham SA, Hoover-Fong J, Telegrafi AB, Destree A, Smigiel R, Lambie LA, Kayserili H, Altunoglu U, Lapi E, Uzielli ML, Aracena M, Nur BG, Mihci E, Moreira LM, Ferreira VB, Horovitz DD, da Rocha KM, Jezela-Stanek A, Brooks A, Reutter H, Cohen JS, Fatemi A, Smitka M, Grebe T, DiDonato N, Deshpande C, Vandersteen A, Lourenço CM, Dufke A, Rossier E, Andre G, Baumer A, Spencer C, McGaughan J, Veltman JA, De Vries BBA, Schinzel A, [Fisher SE](#), Hoischen A, van Bon BW (2017) Overlapping SETBP1 gain-of-function mutations in Schinzel-Giedion syndrome and hematologic malignancies. *PLoS Genet* 13:e1006683
- 116)** Stergiakouli E, Davey Smith G, Martin J, Skuse DH, Viechtbauer W, Ring SM, Ronald A, Evans DM, [Fisher SE](#), Thapar A, St Pourcain B (2017) Shared genetic influences between dimensional ASD and ADHD symptoms during child and adolescent development. *Mol Autism* 8:18
- 117)** Chen XS, Reader RH, Hoischen A, Veltman JA, Simpson NH, Francks C, Newbury DF, [Fisher SE](#) (2017) Next-generation DNA sequencing identifies novel gene variants and pathways involved in specific language impairment. *Sci Rep* 7:46105
- 118)** Sollis E, Deriziotis P, Saitou H, Miyake N, Matsumoto N, Hoffer MJV, Ruivenkamp CAL, Alders M, Okamoto N, Bijlsma EK, Plomp AS, [Fisher SE](#) (2017) Equivalent missense variant in the FOXP2 and FOXP1 transcription factors causes distinct neurodevelopmental disorders. *Hum Mutat* 38:1542-54
- 119)** St Pourcain B, Robinson EB, Anttila V, Sullivan BB, Maller J, Golding J, Skuse D, Ring S, Evans DM, Zammit S, [Fisher SE](#), Neale BM, Anney R, Ripke S, Hollegaard MV, Werge T, iPSYCH-SSI-Broad Autism Group, Ronald A, Grove J, Hougaard DM, Børglum AD, Mortensen PB, Daly M, Smith GD (2018) ASD and schizophrenia show distinct developmental profiles in common genetic overlap with population-based social communication difficulties. *Mol Psychiatry* 23:263-70

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 120)** Devanna P, Chen XS, Ho J, Gajewski D, Smith SD, Gialluisi A, Francks C, Fisher SE, Newbury DF, Vernes SC (2018) Next-gen sequencing identifies non-coding variation disrupting miRNA binding sites in neurological disorders. *Mol Psychiatry* 23:1375-84
- 121)** St Pourcain B, Eaves LJ, Ring SM, Fisher SE, Medland S, Evans DM, Davey Smith G (2018) Developmental changes within the genetic architecture of social communication behavior: a multivariate study of genetic variance in unrelated individuals. *Biol Psychiatry* 83:598-606
- 122)** Morgan AT, Haaften LV, van Hulst K, Edley C, Mei C, Tan TY, Amor D, Fisher SE, Koolen DA (2018) Early speech development in Koolen de Vries syndrome limited by oral praxis and hypotonia. *Eur J Hum Genet* 26:75-84
- 123)** Kuerbitz J, Arnett M, Ehrman S, Williams MT, Vorhees CV, Fisher SE, Garratt AN, Muglia LJ, Waclaw RR, Campbell K (2018) Loss of intercalated cells (ITCs) in the mouse amygdala of Tshz1 mutants correlates with fear, depression, and social interaction phenotypes. *J Neurosci* 38:1160-77
- 124)** Becker M, Devanna P, Fisher SE, Vernes SC (2018) Mapping of human FOXP2 enhancers reveals complex regulation. *Front Mol Neurosci* 11:47
- 125)** Estruch SB, Graham SA, Quevedo M, Vino A, Dekkers DHW, Deriziotis P, Sollis E, Demmers J, Poot RA, Fisher SE (2018) Proteomic analysis of FOXP proteins reveals interactions between cortical transcription factors associated with neurodevelopmental disorders. *Hum Mol Genet* 27:1212-27
- 126)** Tilot AK, Kucera KS, Vino A, Asher JE, Baron-Cohen S, Fisher SE (2018) Rare variants in axonogenesis genes connect three families with sound-color synesthesia. *Proc Natl Acad Sci USA* 115:3168-73
- 127)** Mei C, Fedorenko E, Amor DJ, Boys A, Hoeflin C, Carew P, Burgess T, Fisher SE, Morgan AT (2018) Deep phenotyping of speech and language skills in individuals with 16p11.2 deletion. *Eur J Hum Genet* 26:676-86
- 128)** Snijders Blok L, Hiatt SM, Bowling KM, Prokop JW, Engel KL, Cochran JN, Bebin EM, Bijlsma EK, Ruivenkamp CAL, Terhal P, Simon MEH, Smith R, Hurst JA; DDD study, McLaughlin H, Person R, Crunk A, Wangler MF, Streff H, Symonds JD, Zuberi SM, Elliott KS, Sanders VR, Masunga A, Hopkin RJ, Dubbs HA, Ortiz-Gonzalez XR, Pfundt R, Brunner HG, Fisher SE, Kleefstra T, Cooper GM (2018) De novo mutations in MED13, a component of the Mediator complex, are associated with a novel neurodevelopmental disorder. *Hum Genet* 137:375-88
- 129)** Kong XZ, Mathias SR, Guadalupe T; ENIGMA Laterality Working Group, Glahn DC, Franke B, Crivello F, Tzourio-Mazoyer N, Fisher SE, Thompson PM, Francks C (2018) Mapping cortical brain asymmetry in 17,141 healthy individuals worldwide via the ENIGMA Consortium *Proc Natl Acad Sci USA* 115:E5154-63
- 130)** de Kovel CGF, Lisgo SN, Fisher SE, Francks C (2018) Subtle left-right asymmetry of gene expression profiles in embryonic and foetal human brains. *Sci Rep* 8:12606
- 131)** Xu S, Liu P, Chen Y, Chen Y, Zhang W, Zhao H, Cao Y, Wang F, Jiang N, Lin S, Li B, Zhang Z, Wei Z, Fan Y, Jin Y, He L, Zhou R, Dekker JD, Tucker HO, Fisher SE, Yao Z, Liu Q, Xia X, Guo X (2018) Foxp2 regulates anatomical features that may be relevant for vocal behaviors and bipedal locomotion. *Proc Natl Acad Sci USA* 115:8799-804
- 132)** Snijders Blok L, Rousseau J, Twist J, Ehresmann S, Takaku M, Venselaar H, Rodan LH, Nowak CB, Douglas J, Swoboda KJ, Steeves MA, Sahai I, Stumpel CTRM, Stegmann APA, Wheeler P, Willing M, Fiala E, Kochhar A, Gibson WT, Cohen ASA, Agbahovbe R, Innes AM, Au PYB, Rankin J, Anderson IJ, Skinner SA, Louie RJ, Warren HE, Afenjar A, Keren B, Nava C, Buratti J, Isapof A, Rodriguez D, Lewandowski R, Propst J, van Essen T, Choi M, Lee S, Chae JH, Price S, Schnur RE, Douglas G, Wentzensen IM, Zweier C, Reis A, Bialer MG, Moore C, Koopmans M, Brilstra EH, Monroe GR, van Gassen KLI, van Binsbergen E, Newbury-Ecob R, Bownass L, Bader I, Mayr H, Wortmann SB, Jakielski KJ, Strand EA, Kloth K, Bierhals T, the DDD study, Roberts JD, Petrovich RM, Machida S, Kurumizaka H, Lelieveld S, Pfundt R, Jansen S, Deriziotis P, Faivre L, Thevenon J, Assoum M, Shriberg L, Kleefstra T, Brunner HG, Wade PA, Fisher SE¹, Campeau PM¹ (2018) CHD3 helicase domain mutations cause a neurodevelopmental syndrome with macrocephaly and impaired speech and language. *Nature Commun* 9:4619; ¹joint corresponding authors

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 133)** den Hoed J, Sollis E, Venselaar H, Estruch SB, Deriziotis P, [Fisher SE](#) (2018) Functional characterization of TBR1 variants in neurodevelopmental disorder. *Sci Rep* 8:14279
- 134)** van Rhijn JR, [Fisher SE](#), Vernes SC, Nadif Kasri N (2018) Foxp2 loss of function increases striatal direct pathway inhibition via increased GABA release. *Brain Struct Funct* 223:4211-26
- 135)** French CA, Vinueza Veloz MF, Zhou K, Peter S, [Fisher SE](#)¹, Costa RM¹, De Zeeuw CI (2019) Differential effects of Foxp2 disruption in distinct motor circuits. *Mol Psychiatry* 24:447-62; ¹**joint corresponding authors**
- 136)** Eising E, Carrion-Castillo A, Vino A, Strand EA, Jakielski KJ, Scerri TS, Hildebrand MS, Webster R, Ma A, Mazoyer B, Francks C, Bahlo M, Scheffer IE, Morgan AT, Shriberg LD, [Fisher SE](#) (2019). A set of regulatory genes co-expressed in embryonic human brain is implicated in disrupted speech development. *Mol Psychiatry* 24:1065-78
- 137)** Haworth S, Shapland CY, Hayward C, Prins BP, Felix JF, Medina-Gomez C, Rivadeneira F, Wang C, Ahluwalia TS, Vrijheid M, Guxens M, Sunyer J, Tachmazidou I, Walter K, Iotchkova V, Jackson A, Cleal L, Huffmann J, Min JL, Sass L, Timmers PRHJ, UK10K consortium, Davey Smith G, [Fisher SE](#), Wilson JF, Cole TJ, Fernandez-Orth D, Bønnelykke K, Bisgaard H, Pennell CE, Jaddoe VWV, Dedoussis G, Timpson N, Zeggini E, Vitart V, St Pourcain B (2019) Low-frequency variation in TP53 has large effects on head circumference and intracranial volume. *Nature Commun* 10:357
- 138)** Gunz P, Tilot AK, Wittfeld K, Teumer A, Shapland CY, van Erp TGM, Dannemann M, Vernot B, Neubauer S, Guadalupe T, Fernandez G, Brunner H, Enard W, Fallon J, Hosten N, Völker U, Profico A, Di Vincenzo F, Manzi G, Kelso J, St Pourcain B, Hublin J-J, Franke B, Pääbo S, Macciardi F, Grabe HJ, [Fisher SE](#) (2019) Neandertal introgression sheds light on modern human endocranial globularity. *Curr Biol* 29:120-7
- 139)** Verhoef E, Demontis D, Burgess S, Shapland CY, Dale PS, Okbay A, Neale BM, Faraone SV; iPSYCH-Broad-PGC ADHD Consortium, Stergiakouli E, Davey Smith G, [Fisher SE](#), Børnglum AD, St Pourcain B (2019) Disentangling polygenic associations between attention-deficit/hyperactivity disorder, educational attainment, literacy and language. *Transl Psychiatry* 9:35
- 140)** Gialluisi A, Andlauer TFM, Mirza-Schreiber N, Moll K, Becker J, Hoffmann P, Ludwig KU, Czamara D, St Pourcain B, Brandler W, Honbolygó F, Tóth D, Csépe V, Huguet G, Morris AP, Hulshander J, Willcutt EG, DeFries JC, Olson RK, Smith SD, Pennington BF, Vaessen A, Maurer U, Lyytinen H, Peyrard-Janvid M, Leppänen PHT, Brandeis D, Bonte M, Stein JF, Talcott JB, Fauchereau F, Wilcke A, Francks C, Bourgeron T, Monaco AP, Ramus F, Landerl K, Kere J, Scerri TS, Paracchini S, [Fisher SE](#), Schumacher J, Nöthen MM, Müller-Myhsok B, Schulte-Körne G (2019) Genome-wide association scan identifies new variants associated with a cognitive predictor of dyslexia. *Transl Psychiatry* 9:77
- 141)** Castells-Nobau A, Eidhof I, Fenckova M, Brenman-Suttner DB, Scheffer-de Gooyert JM, Christine S, Schellevis RL, van der Laan K, Quentin C, van Nihuijs L, Hofmann F, Ejsmont R, [Fisher SE](#), Kramer JM, Sigrist SJ, Simon AF, Schenck A (2019) Conserved regulation of neurodevelopmental processes and behavior by FoxP in *Drosophila*. *PLoS ONE* 14:e0211652
- 142)** Truong DT, Adams AK, Paniagua S, Frijters JC, Boada R, Hill DE, Lovett MW, Mahone EM, Willcutt EG, Wolf M, Defries JC, Gialluisi A, Francks C, [Fisher SE](#), Olson RK, Pennington BF, Smith SD, Bosson-Heenan J, Gruen JR; Pediatric, Imaging, Neurocognition, and Genetics Consortium (2019) Multivariate genome-wide association study of rapid automatized naming and rapid alternating stimulus in Hispanic American and African-American youth. *J Med Genet* 56:557-66
- 143)** Snijders Blok L, Kleefstra T, Venselaar H, Maas S, Kroes HY, Lachmeijer AMA, van Gassen KLI, Firth HV, Tomkins S, Bodek S; DDD Study, Ōunap K, Wojcik MH, Cunniff C, Bergstrom K, Powis Z, Tang S, Shinde DN, Au C, Iglesias AD, Izumi K, Leonard J, Abou Tayoun A, Baker SW, Tartaglia M, Niceta M, Dentici ML, Okamoto N, Miyake N, Matsumoto N, Vitobello A, Faivre L, Philippe C, Gilissen C, Wiel L, Pfundt R, Deriziotis P, Brunner HG, [Fisher SE](#) (2019) De novo variants disturbing the transactivation capacity of POU3F3 cause a characteristic neurodevelopmental disorder. *Am J Hum Genet* 105:403-12
- 144)** Uddén J, Hultén A, Bendtz K, Mineroff Z, Kucera KS, Vino A, Fedorenko E, Hagoort P, [Fisher SE](#) (2019) Toward robust functional neuroimaging genetics of cognition. *J Neurosci* 39:8778-87

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 145)** Carrion-Castillo A, Van der Haegen L, Tzourio-Mazoyer N, Kavaklioglu T, Badillo S, Chavent M, Saracco J, Brysbaert M, [Fisher SE](#), Mazoyer B, Francks C (2019) Genome sequencing for rightward hemispheric language dominance. *Genes Brain Behav* 18:e12572
- 146)** Tilot AK, VINO A, Kucera KS, Carmichael DA, van den Heuvel L, den Hoed J, Sidoroff-Dorso AV, Campbell A, Porteous DJ, St Pourcain B, van Leeuwen TM, Ward J, Rouw R, Simner J, [Fisher SE](#) (2019) Investigating genetic links between grapheme-colour synaesthesia and neuropsychiatric traits. *Phil Trans Roy Soc B* 374:20190026
- 147)** Ioumpa K, Graham SA, Clausner T, [Fisher SE](#), van Lier R, van Leeuwen TM (2019) Enhanced self-reported affect and prosocial behaviour without differential physiological responses in mirror-sensory synaesthesia. *Phil Trans Roy Soc B* 374:20190395
- 148)** Satizabal CL *et al* (2019) Genetic architecture of subcortical brain structures in 38,851 individuals. *Nature Genet* 51:1624-36
- 149)** Postema MC, van Rooij D, Anagnostou E, Arango C, Auzias G, Behrmann M, Filho GB, Calderoni S, Calvo R, Daly E, Deruelle C, Di Martino A, Dinstein I, Duran FLS, Durston S, Ecker C, Ehrlich S, Fair D, Fedor J, Feng X, Fitzgerald J, Floris DL, Freitag CM, Gallagher L, Glahn DC, Gori I, Haar S, Hoekstra L, Jahanshad N, Jalbrzikowski M, Janssen J, King JA, Kong XZ, Lazaro L, Lerch JP, Luna B, Martinho MM, McGrath J, Medland SE, Muratori F, Murphy CM, Murphy DGM, O'Hearn K, Oranje B, Parellada M, Puig O, Retico A, Rosa P, Rubia K, Shook D, Taylor MJ, Tosetti M, Wallace GL, Zhou F, Thompson PM, [Fisher SE](#), Buitelaar JK, Francks C (2019) Altered structural brain asymmetry in autism spectrum disorder in a study of 54 datasets. *Nature Commun* 10:4958
- 150)** van der Meer D, Rokicki J, Kaufmann T, Córdova-Palomera A, Moberget T, Alnæs D, Bettella F, Frei O, Doan NT, Søndersby IE, Smeland OB, Agartz I, Bertolino A, Bralten J, Brandt CL, Buitelaar JK, Djurovic S, van Donkelaar M, Dørum ES, Espeseth T, Faraone SV, Fernández G, [Fisher SE](#), Franke B, Haatveit B, Hartman CA, Hoekstra PJ, Håberg AK, Jönsson EG, Kolskår KK, Le Hellard S, Lund MJ, Lundervold AJ, Lundervold A, Melle I, Monereo Sánchez J, Norbom LC, Nordvik JE, Nyberg L, Oosterlaan J, Papalino M, Papassotiropoulos A, Pergola G, de Quervain DJF, Richard G, Sanders AM, Selvaggi P, Shumskaya E, Steen VM, Tønnesen S, Ulrichsen KM, Zwiers MP, Andreassen OA, Westlye LT; Alzheimer's Disease Neuroimaging Initiative; Pediatric Imaging, Neurocognition and Genetics Study (2020) Brain scans from 21,297 individuals reveal the genetic architecture of hippocampal subfield volumes. *Mol Psychiatry* 25:3053-65.
- 151)** Søndersby IE, Gústafsson Ó, Doan NT, Hibar DP, Martin-Brevet S, Abdellaoui A, Ames D, Amunts K, Andersson M, Armstrong NJ, Bernard M, Blackburn N, Blangero J, Boomsma DI, Bralten J, Brattbak HR, Brodaty H, Brouwer RM, Bülow R, Calhoun V, Caspers S, Cavalleri G, Chen CH, Cichon S, Ciufolini S, Corvin A, Crespo-Facorro B, Curran JE, Dale AM, Dalvie S, Dazzan P, de Geus EJC, de Zubicaray GI, de Zwarte SMC, Delanty N, den Braber A, Desrivières S, Donohoe G, Draganski B, Ehrlich S, Espeseth T, [Fisher SE](#), Franke B, Frouin V, Fukunaga M, Gareau T, Glahn DC, Grabe H, Groenewold NA, Haavik J, Håberg A, Hashimoto R, Hehir-Kwa JY, Heinz A, Hillegers MHJ, Hoffmann P, Holleran L, Hottenga JJ, Hulshoff HE, Ikeda M, Jahanshad N, Jernigan T, Jockwitz C, Johansson S, Jonsdottir GA, Jönsson EG, Kahn R, Kaufmann T, Kelly S, Kikuchi M, Knowles EEM, Kolskår KK, Kwok JB, Hellard SL, Leu C, Liu J, Lundervold AJ, Lundervold A, Martin NG, Mather K, Mathias SR, McCormack M, McMahon KL, McRae A, Milaneschi Y, Moreau C, Morris D, Mothersill D, Mühleisen TW, Murray R, Nordvik JE, Nyberg L, Olde Loohuis LM, Ophoff R, Paus T, Pausova Z, Penninx B, Peralta JM, Pike B, Prieto C, Pudas S, Quinlan E, Quintana DS, Reinbold CS, Marques TR, Raymond A, Richard G, Rodriguez-Herreros B, Roiz-Santiañez R, Rokicki J, Rucker J, Sachdev P, Sanders AM, Sando SB, Schmaal L, Schofield PR, Schork AJ, Schumann G, Shin J, Shumskaya E, Sisodiya S, Steen VM, Stein DJ, Steinberg S, Strike L, Teumer A, Thalamuthu A, Tordesillas-Gutierrez D, Turner J, Ueland T, Uhlmann A, Ulfarsson MO, van 't Ent D, van der Meer D, van Haren NEM, Vaskinn A, Vassos E, Walters GB, Wang Y, Wen W, Whelan CD, Wittfeld K, Wright M, Yamamori H, Zayats T, Agartz I, Westlye LT, Jacquemont S, Djurovic S, Stefánsson H, Stefánsson K, Thompson P, Andreassen OA; 16p11.2 European Consortium, for the ENIGMA-CNV working group (2020) Dose response of the 16p11.2 distal copy number variant on intracranial volume and basal ganglia. *Mol Psychiatry* 25:584-602

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 152)** Postema MC, Carrion-Castillo A, [Fisher SE](#), Vingerhoets G, Francks C (2020) The genetics of situs inversus without primary ciliary dyskinesia *Sci Rep* 10:3677
- 153)** Writing Committee for the ENIGMA-CNV Working Group, van der Meer D, S nderby IE, Kaufmann T, Walters GB, Abdellaoui A, Ames D, Amunts K, Andersson M, Armstrong NJ, Bernard M, Blackburn NB, Blangero J, Boomsma DI, Brodaty H, Brouwer RM, B low R, Cahn W, Calhoun VD, Caspers S, Cavalleri GL, Ching CRK, Cichon S, Ciufolini S, Corvin A, Crespo-Facorro B, Curran JE, Dalvie S, Dazzan P, de Geus EJC, de Zubicaray GI, de Zwarte SMC, Delanty N, den Braber A, Desrivieres S, Di Forti M, Doherty JL, Donohoe G, Ehrlich S, Eising E, Espeseth T, [Fisher SE](#), Fladby T, Frei O, Frouin V, Fukunaga M, Gareau T, Glahn DC, Grabe HJ, Groenewold NA, G stafsson  , Haavik J, Haberg AK, Hashimoto R, Hehir-Kwa JY, Hibar DP, Hillegers MHJ, Hoffmann P, Holleran L, Hottenga JJ, Hulshoff Pol HE, Ikeda M, Jacquemont S, Jahanshad N, Jockwitz C, Johansson S, J nsson EG, Kikuchi M, Knowles EEM, Kwok JB, Le Hellard S, Linden DEJ, Liu J, Lundervold A, Lundervold AJ, Martin NG, Mather KA, Mathias SR, McMahon KL, McRae AF, Medland SE, Moberget T, Moreau C, Morris DW, M hleisen TW, Murray RM, Nordvik JE, Nyberg L, Olde Loohuis LM, Ophoff RA, Owen MJ, Paus T, Pausova Z, Peralta JM, Pike B, Prieto C, Quinlan EB, Reinbold CS, Reis Marques T, Rucker JJH, Sachdev PS, Sando SB, Schofield PR, Schork AJ, Schumann G, Shin J, Shumskaya E, Silva AI, Sisodiya SM, Steen VM, Stein DJ, Strike LT, Tamnes CK, Teumer A, Thalamuthu A, Tordesillas-Guti rrez D, Uhlmann A,  lfarsson M , van 't Ent D, van den Bree MBM, Vassos E, Wen W, Wittfeld K, Wright MJ, Zayats T, Dale AM, Djurovic S, Agartz I, Westlye LT, Stef nsson H, Stef nsson K, Thompson PM, Andreassen OA (2020) Association of copy number variation of the 15q11.2 BP1-BP2 region with cortical and subcortical morphology and cognition. *JAMA Psychiatry* 77:420-30
- 154)** Carrion-Castillo A, Pepe A, Kong XZ, [Fisher SE](#), Mazoyer B, Tzourio-Mazoyer N, Crivello F, Francks C (2020) Genetic effects on planum temporale asymmetry and their limited relevance to neurodevelopmental disorders, intelligence or educational attainment. *Cortex* 124:137-53
- 155)** Kong X, Tzourio-Mazoyer N, Joliot M, Fedorenko E, Liu J, [Fisher SE](#), Francks C (2020) Gene expression correlates of the cortical network underlying sentence processing. *Neurobiology of Language* 1:77-103
- 156)** Hildebrand MS, Jackson VE, Scerri TS, Van Reyk O, Coleman M, Braden R, Turner S, Rigbye KA, Boys A, Barton S, Webster R, Fahey M, Saunders K, Parry-Fielder B, Paxton G, Hayman M, Coman D, Goel H, Baxter A, Ma A, Davis N, Reilly S, Delatycki M, Li geois FJ, Connelly A, Gecz J, [Fisher SE](#), Amor DJ, Scheffer IE, Bahlo M, Morgan AT (2020) Severe childhood speech disorder: Gene discovery highlights transcriptional dysregulation. *Neurology* 94:e2148-67
- 157)** Grasby KL *et al* (2020) The genetic architecture of the human cerebral cortex. *Science* 367:eaay6690
- 158)** Doust C, Gordon SD, Garden N, [Fisher SE](#), Martin NG, Bates TC, Luciano M (2020) The association of dyslexia and developmental speech and language disorder candidate genes with reading and language abilities in adults. *Twin Res Hum Genet* 23:23-32
- 159)** Connaughton DM, Dai R, Owen DJ, Marquez J, Mann N, Graham-Paquin AL, Nakayama M, Coyaud E, Laurent EMN, St-Germain JR, Blok LS, Vino A, Kl mbt V, Deutsch K, Wu CW, Kolvenbach CM, Kause F, Ottlewski I, Schneider R, Kitzler TM, Majmundar AJ, Buerger F, Onuchic-Whitford AC, Youying M, Kolb A, Salmanullah D, Chen E, van der Ven AT, Rao J, Ityel H, Seltzsam S, Rieke JM, Chen J, Vivante A, Hwang DY, Kohl S, Dworschak GC, Hermle T, Alders M, Bartolomeus T, Bauer SB, Baum MA, Brilstra EH, Challman TD, Zyskind J, Costin CE, Dipple KM, Duijkers FA, Ferguson M, Fitzpatrick DR, Fick R, Glass IA, Hulick PJ, Kline AD, Krey I, Kumar S, Lu W, Marco EJ, Wentzensen IM, Mefford HC, Platzer K, Povolotskaya IS, Savatt JM, Shcherbakova NV, Senguttuvan P, Squire AE, Stein DR, Thiffault I, Voinova VY, Somers MJG, Ferguson MA, Traum AZ, Daouk GH, Daga A, Rodig NM, Terhal PA, van Binsbergen E, Eid LA, Tasic V, Rasouly HM, Lim TY, Ahram DF, Gharavi AG, Reutter HM, Rehm HL, MacArthur DG, Lek M, Laricchia KM, Lifton RP, Xu H, Mane SM, Sanna-Cherchi S, Sharrocks AD, Raught B, [Fisher SE](#), Bouchard M, Khokha MK, Shril S, Hildebrandt F (2020) Mutations of the transcriptional corepressor ZMYM2 cause syndromic urinary tract malformations. *Am J Hum Genet* 107:727-42
- 160)** Urbanus BHA, Peter S, [Fisher SE](#), De Zeeuw CI (2020) Region-specific Foxp2 deletions in cortex, striatum or cerebellum cannot explain vocalization deficits observed in spontaneous global knockouts. *Sci Rep* 10:21631.

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 161)** Henson RN, Suri S, Knights E, Rowe JB, Kievit RA, Lyall DM, Chan D, Eising E, [Fisher SE](#) (2020) Effect of APOE polymorphism on cognition and brain in the CamCAN cohort. *Brain and Neuroscience Advances* 4:2398212820961704.
- 162)** Thompson PA, Bishop DVM, Eising E, [Fisher SE](#), Newbury DF (2020) Generalized Structured Component Analysis in candidate gene association studies: applications and limitations. *Wellcome Open Res* 4:142
- 163)** Verhoef E, Shapland CY, [Fisher SE](#), Dale PS, St Pourcain B (2021) The developmental origins of genetic factors influencing language and literacy: Associations with early-childhood vocabulary. *J Child Psychol Psychiatry* 62:728-38
- 164)** Smeets CJLM, Ma KY, [Fisher SE](#), Verbeek DS (2021) Cerebellar developmental deficits underlie neurodegenerative disorder Spinocerebellar Ataxia Type 23. *Brain Pathol* 31:239-52
- 165)** Gialluisi A, Andlauer TFM, Mirza-Schreiber N, Moll K, Becker J, Hoffmann P, Ludwig KU, Czamara D, St Pourcain B, Honbolygó F, Tóth D, Csépe V, Huguet H, Chaix Y, Iannuzzi S, Demonet J-F, Morris AP, Hulslander J, Willcutt EG, DeFries JC, Olson RK, Smith SD, Pennington BF, Vaessen A, Maurer U, Lyytinen H, Peyrard-Janvid M, Leppänen PHT, Brandeis D, Bonte M, Stein JF, Talcott JB, Fauchereau F, Wilcke A, Kirsten H, Müller B, Francks C, Bourgeron T, Monaco AP, Ramus F, Landerl K, Kere J, Scerri TS, Paracchini S, [Fisher SE](#), Schumacher J, Nöthen MM, Müller-Myhsok B, Schulte-Körne G (2021) Genome-wide association study reveals new insights into the heritability and genetic correlates of developmental dyslexia. *Mol Psychiatry* 26:3004-17
- 166)** Snijders Blok L, Vino A, den Hoed J, Underhill HR, Monteil D, Hong L, Reynoso Santos FJ, Chung WK, Amaral MD, Schnur RE, Santiago-Sim T, Sinds Y, Brunner HG, Kleefstra T, [Fisher SE](#) (2021) Heterozygous variants that disturb the transcriptional repressor activity of FOXP4 cause a developmental disorder with speech/language delays and multiple congenital abnormalities. *Genet Med* 23:534-42
- 167)** Tilot AK, Khramtsova EA, Liang D, Grasby KL, Jahanshad N, Painter J, Colodro-Conde L, Bralten J, Hibar DP, Lind PA, Liu S, Brotman SM, Thompson PM, Medland SE, Macciardi F, Stranger BE, Davis LK, [Fisher SE](#)¹, Stein JL¹ (2021) The evolutionary history of common genetic variants influencing human cortical surface area *Cereb Cortex* 31:1873-87 ¹joint corresponding authors
- 168)** den Hoed J, de Boer E, Voisin N, Dingemans AJM, Guex N, Wiel L, Nellaker C, Amudhavalli SM, Banka S, Bena FS, Ben-Zeev B, Bonagura VR, Bruel AL, Brunet T, Brunner HG, Chew HB, Chrast J, Cimbalistiené L, Coon H; DDD Study, Délot EC, Démurger F, Denommé-Pichon AS, Depienne C, Donnai D, Dymont DA, Elpeleg O, Faivre L, Gilissen C, Granger L, Haber B, Hachiya Y, Abedi YH, Hanebeck J, Hehir-Kwa JY, Horist B, Itai T, Jackson A, Jewell R, Jones KL, Joss S, Kashii H, Kato M, Kattentidt-Mouravieva AA, Kok F, Kotzaeridou U, Krishnamurthy V, Kučinskás V, Kuechler A, Lavillaureix A, Liu P, Manwaring L, Matsumoto N, Mazel B, McWalter K, Meiner V, Mikati MA, Miyatake S, Mizuguchi T, Moey LH, Mohammed S, Mor-Shaked H, Mountford H, Newbury-Ecob R, Odent S, Orec L, Osmond M, Palculict TB, Parker M, Petersen AK, Pfundt R, Preikšaitienė E, Radtke K, Ranza E, Rosenfeld JA, Santiago-Sim T, Schwager C, Sinnema M, Snijders Blok L, Spillmann RC, Stegmann APA, Thiffault I, Tran L, Vaknin-Dembinsky A, Vedovato-Dos-Santos JH, Schrier Vergano SA, Vilain E, Vitobello A, Wagner M, Waheeb A, Willing M, Zuccarelli B, Kini U, Newbury DF, Kleefstra T, Reymond A, [Fisher SE](#)¹, Vissers LELM (2021) Mutation-specific pathophysiological mechanisms define different neurodevelopmental disorders associated with SATB1 dysfunction. *Am J Hum Genet* 108:346-56 ¹corresponding author
- 169)** Verhoef E, Shapland CY, [Fisher SE](#), Dale PS, St Pourcain B (2021) The developmental genetic architecture of vocabulary skills during the first three years of life: Capturing emerging associations with later-life reading and cognition. *PLoS Genet* 17:e1009144
- 170)** Sha Z, Schijven D, Carrion-Castillo A, Joliot M, Mazoyer B, [Fisher SE](#), Crivello F, Francks C (2021) The genetic architecture of structural left-right asymmetry of the human brain. *Nature Hum Behav* 5:1226-1239
- 171)** Kong XZ, Postema M, Schijven D, Castillo AC, Pepe A, Crivello F, Joliot M, Mazoyer B, [Fisher SE](#), Francks C (2021) Large-Scale phenomic and genomic analysis of brain asymmetrical skew. *Cereb Cortex* 31:4151-68

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 172)** Carrion-Castillo A, Estruch SB, Maassen B, Franke B, Francks C, [Fisher SE](#) (2021) Whole-genome sequencing identifies functional noncoding variation in SEMA3C that cosegregates with dyslexia in a multigenerational family. *Hum Genet* 140:1183-1200
- 173)** Postema MC, Hoogman M, Ambrosino S, Asherson P, Banaschewski T, Bandeira CE, Baranov A, Bau CHD, Baumeister S, Baur-Streubel R, Bellgrove MA, Biederman J, Bralten J, Brandeis D, Brem S, Buitelaar JK, Busatto GF, Castellanos FX, Cercignani M, Chaim-Avancini TM, Chantiluke KC, Christakou A, Coghill D, Conzelmann A, Cubillo AI, Cupertino RB, de Zeeuw P, Doyle AE, Durston S, Earl EA, Epstein JN, Ethofer T, Fair DA, Fallgatter AJ, Faraone SV, Frodl T, Gabel MC, Gogberashvili T, Grevet EH, Haavik J, Harrison NA, Hartman CA, Heslenfeld DJ, Hoekstra PJ, Hohmann S, Høvik MF, Jernigan TL, Kardatzki B, Karkashadze G, Kelly C, Kohls G, Konrad K, Kuntsi J, Lazaro L, Lera-Miguel S, Lesch KP, Louza MR, Lundervold AJ, Malpas CB, Mattos P, McCarthy H, Namazova-Baranova L, Nicolau R, Nigg JT, Novotny SE, Oberwelland Weiss E, O'Gorman Tuura RL, Oosterlaan J, Oranje B, Paloyelis Y, Pauli P, Picon FA, Plessen KJ, Ramos-Quiroga JA, Reif A, Reneman L, Rosa PGP, Rubia K, Schranke A, Schweren LJS, Seitz J, Shaw P, Silk TJ, Skokauskas N, Soliva Vila JC, Stevens MC, Sudre G, Tamm L, Tovar-Moll F, van Erp TGM, Vance A, Vilarroya O, Vives-Gilabert Y, von Polier GG, Walitza S, Yoncheva YN, Zanetti MV, Ziegler GC, Glahn DC, Jahanshad N, Medland SE; ENIGMA ADHD Working Group, Thompson PM, [Fisher SE](#), Franke B, Francks C (2021) Analysis of structural brain asymmetries in attention-deficit/hyperactivity disorder in 39 datasets. *J Child Psychol Psychiatry* 62:1202-19
- 174)** Søndersby IE, van der Meer D, Moreau C, Kaufmann T, Walters GB, Ellegaard M, Abdellaoui A, Ames D, Amunts K, Andersson M, Armstrong NJ, Bernard M, Blackburn NB, Blangero J, Boomsma DI, Brodaty H, Brouwer RM, Bülow R, Bøen R, Cahn W, Calhoun VD, Caspers S, Ching CRK, Cichon S, Ciufolini S, Crespo-Facorro B, Curran JE, Dale AM, Dalvie S, Dazzan P, de Geus EJC, de Zubicaray GI, de Zwarte SMC, Desrivieres S, Doherty JL, Donohoe G, Draganski B, Ehrlich S, Eising E, Espeseth T, Fejgin K, [Fisher SE](#), Fladby T, Frei O, Frouin V, Fukunaga M, Gareau T, Ge T, Glahn DC, Grabe HJ, Groenewold NA, Gústafsson Ó, Haavik J, Haberg AK, Hall J, Hashimoto R, Hehir-Kwa JY, Hibar DP, Hillegers MHJ, Hoffmann P, Holleran L, Holmes AJ, Homuth G, Hottenga JJ, Hulshoff Pol HE, Ikeda M, Jahanshad N, Jockwitz C, Johansson S, Jönsson EG, Jørgensen NR, Kikuchi M, Knowles EEM, Kumar K, Le Hellard S, Leu C, Linden DEJ, Liu J, Lundervold A, Lundervold AJ, Maillard AM, Martin NG, Martin-Brevet S, Mather KA, Mathias SR, McMahon KL, McRae AF, Medland SE, Meyer-Lindenberg A, Moberget T, Moderato C, Sánchez JM, Morris DW, Mühleisen TW, Murray RM, Nielsen J, Nordvik JE, Nyberg L, Loohuis LMO, Ophoff RA, Owen MJ, Paus T, Pausova Z, Peralta JM, Pike GB, Prieto C, Quinlan EB, Reinbold CS, Marques TR, Rucker JJH, Sachdev PS, Sando SB, Schofield PR, Schork AJ, Schumann G, Shin J, Shumskaya E, Silva AI, Sisodiya SM, Steen VM, Stein DJ, Strike LT, Suzuki IK, Tamnes CK, Teumer A, Thalamuthu A, Tordesillas-Gutiérrez D, Uhlmann A, Ulfarsson MO, van 't Ent D, van den Bree MBM, Vanderhaeghen P, Vassos E, Wen W, Wittfeld K, Wright MJ, Agartz I, Djurovic S, Westlye LT, Stefansson H, Stefansson K, Jacquemont S, Thompson PM, Andreassen OA; ENIGMA-CNV working group (2021) 1q21.1 distal copy number variants are associated with cerebral and cognitive alterations in humans. *Transl Psychiatry* 11:182
- 175)** Jansen NA, Braden RO, Srivastava S, Otness EF, Lesca G, Rossi M, Nizon M, Bernier RA, Quelin C, Haeringen A, Kleefstra T, Wong MMK, Whalen S, [Fisher SE](#), Morgan AT, van Bon BW (2021) Clinical delineation of SETBP1 haploinsufficiency disorder. *Eur J Hum Genet* 29:1198-1205
- 176)** Morgan A, Braden R, Wong MMK, Colin E, Amor D, Liégeois F, Srivastava S, Vogel A, Bizaoui V, Ranguin K, [Fisher SE](#), van Bon BW (2021) Speech and language deficits are central to SETBP1 haploinsufficiency disorder. *Eur J Hum Genet* 29:1216-25
- 177)** Braden RO, Amor DJ, [Fisher SE](#), Mei C, Myers CT, Mefford H, Gill D, Srivastava S, Swanson LC, Goel H, Scheffer IE, Morgan AT (2021) Severe speech impairment is a distinguishing feature of FOXP1-related disorder. *Dev Med Child Neurol* 63:1417-26
- 178)** Snijders Blok L, Goosen YM, van Haaften L, van Hulst K, [Fisher SE](#), Brunner HG, Egger JIM, Kleefstra T (2021) Speech-language profiles in the context of cognitive and adaptive functioning in SATB2-associated syndrome. *Genes Brain Behav* 20:e12761

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 179)** Shapland CY, Verhoef E, Davey Smith G, [Fisher SE](#), Verhulst B, Dale PS, St Pourcain B (2021) Multivariate genome-wide covariance analyses of literacy, language and working memory skills reveal distinct etiologies. *NPJ Sci Learn* 6:23.
- 180)** Sha Z, Pepe A, Schijven D, Carrión-Castillo A, Roe JM, Westerhausen R, Joliot M, [Fisher SE](#), Crivello F, Francks C (2021) Handedness and its genetic influences are associated with structural asymmetries of the cerebral cortex in 31,864 individuals. *Proc Natl Acad Sci USA* 118:e2113095118
- 181)** Kong XZ; ENIGMA Laterality Working Group, Francks C (2022) Reproducibility in the absence of selective reporting: An illustration from large-scale brain asymmetry research. *Hum Brain Mapp* 43:244-54
- 182)** Wierenga LM, Doucet GE, Dima D, Agartz I, Aghajani M, Akudjedu TN, Albajes-Eizagirre A, Alnaes D, Alpert KI, Andreassen OA, Anticevic A, Asherson P, Banaschewski T, Bargallo N, Baumeister S, Baur-Streubel R, Bertolino A, Bonvino A, Boomsma DI, Borgwardt S, Bourque J, den Braber A, Brandeis D, Breier A, Brodaty H, Brouwer RM, Buitelaar JK, Busatto GF, Calhoun VD, Canales-Rodríguez EJ, Cannon DM, Caseras X, Castellanos FX, Chaim-Avancini TM, Ching CR, Clark VP, Conrod PJ, Conzelmann A, Crivello F, Davey CG, Dickie EW, Ehrlich S, Van't Ent D, [Fisher SE](#), Fouché JP, Franke B, Fuentes-Claramonte P, de Geus EJ, Giorgio AD, Glahn DC, Gotlib IH, Grabe HJ, Gruber O, Gruner P, Gur RE, Gur RC, Gurholt TP, de Haan L, Haatveit B, Harrison BJ, Hartman CA, Hatton SN, Heslenfeld DJ, van den Heuvel OA, Hickie IB, Hoekstra PJ, Hohmann S, Holmes AJ, Hoogman M, Hosten N, Howells FM, Hulshoff Pol HE, Huysen C, Jahanshad N, James AC, Jiang J, Jönsson EG, Joska JA, Kalnin AJ; Karolinska Schizophrenia Project (KaSP) Consortium, Klein M, Koenders L, Kolskår KK, Krämer B, Kuntsi J, Lagopoulos J, Lazaro L, Lebedeva IS, Lee PH, Lochner C, Machielsen MW, Maingault S, Martin NG, Martínez-Zalacaín I, Mataix-Cols D, Mazoyer B, McDonald BC, McDonald C, McIntosh AM, McMahon KL, McPhilemy G, van der Meer D, Menchón JM, Naaijen J, Nyberg L, Oosterlaan J, Paloyelis Y, Pauli P, Pergola G, Pomarol-Clotet E, Portella MJ, Radua J, Reif A, Richard G, Roffman JL, Rosa PG, Sacchet MD, Sachdev PS, Salvador R, Sarró S, Satterthwaite TD, Saykin AJ, Serpa MH, Sim K, Simmons A, Smoller JW, Sommer IE, Soriano-Mas C, Stein DJ, Strike LT, Szeszko PR, Temmingh HS, Thomopoulos SI, Tomyshev AS, Trollor JN, Uhlmann A, Veer IM, Veltman DJ, Voineskos A, Völzke H, Walter H, Wang L, Wang Y, Weber B, Wen W, West JD, Westlye LT, Whalley HC, Williams SC, Wittfeld K, Wolf DH, Wright MJ, Yoncheva YN, Zanetti MV, Ziegler GC, de Zubicaray GI, Thompson PM, Crone EA, Frangou S, Tamnes CK (2022) Greater male than female variability in regional brain structure across the lifespan. *Hum Brain Mapp* 43:470-99
- 183)** Guadalupe T, Kong XZ, Akkermans SEA, [Fisher SE](#), Francks C (2022) Relations between hemispheric asymmetries of grey matter and auditory processing of spoken syllables in 281 healthy adults. *Brain Struct Funct* 227(2):561-72
- 184)** Schlag F, Allegrini AG, Buitelaar J, Verhoef E, van Donkelaar M, Plomin R, Rimfeld K, [Fisher SE](#), St Pourcain B (2022) Polygenic risk for mental disorder reveals distinct association profiles across social behaviour in the general population. *Mol Psychiatry* 27:1588-98
- 185)** Sha Z, van Rooij D, Anagnostou E, Arango C, Auzias G, Behrmann M, Bernhardt B, Bolte S, Busatto GF, Calderoni S, Calvo R, Daly E, Deruelle C, Duan M, Duran FLS, Durston S, Ecker C, Ehrlich S, Fair D, Fedor J, Fitzgerald J, Floris DL, Franke B, Freitag CM, Gallagher L, Glahn DC, Haar S, Hoekstra L, Jahanshad N, Jalbrzikowski M, Janssen J, King JA, Lazaro L, Luna B, McGrath J, Medland SE, Muratori F, Murphy DGM, Neufeld J, O'Hearn K, Oranje B, Parellada M, Pariente JC, Postema MC, Remnelius KL, Retico A, Rosa PGP, Rubia K, Shook D, Tammimies K, Taylor MJ, Tosetti M, Wallace GL, Zhou F, Thompson PM, [Fisher SE](#), Buitelaar JK, Francks C (2022) Subtly altered topological asymmetry of brain structural covariance networks in autism spectrum disorder across 43 datasets from the ENIGMA consortium. *Mol Psychiatry* 27:2114-25
- 186)** Boyce JO, Jackson VE, van Reyk O, Parker R, Vogel AP, Eising E, Horton SE, Gillespie NA, Scheffer IE, Amor DJ, Hildebrand MS, [Fisher SE](#), Martin NG, Reilly S, Bahlo M, Morgan AT (2022) Self-reported impact of developmental stuttering across the lifespan *Dev Med Child Neurol* 64:1297-1306
- 187)** Brouwer RM *et al* (2022) Genetic variants associated with longitudinal changes in brain structure across the lifespan. *Nature Neurosci* 25:421-32

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

188) Frangou S, Modabbernia A, Williams SCR, Papachristou E, Doucet GE, Agartz I, Aghajani M, Akudjedu TN, Albajes-Eizagirre A, Alnaes D, Alpert KI, Andersson M, Andreasen NC, Andreassen OA, Asherson P, Banaschewski T, Bargallo N, Baumeister S, Baur-Streubel R, Bertolino A, Bonvino A, Boomsma DI, Borgwardt S, Bourque J, Brandeis D, Breier A, Brodaty H, Brouwer RM, Buitelaar JK, Busatto GF, Buckner RL, Calhoun V, Canales-Rodríguez EJ, Cannon DM, Caseras X, Castellanos FX, Cervenka S, Chaim-Avancini TM, Ching CRK, Chubar V, Clark VP, Conrod P, Conzelmann A, Crespo-Facorro B, Crivello F, Crone EA, Dale AM, Dannlowski U, Davey C, de Geus EJC, de Haan L, de Zubicaray GI, den Braber A, Dickie EW, Di Giorgio A, Doan NT, Dørum ES, Ehrlich S, Erk S, Espeseth T, Fatouros-Bergman H, [Fisher SE](#), Fouche JP, Franke B, Frodl T, Fuentes-Claramonte P, Glahn DC, Gotlib IH, Grabe HJ, Grimm O, Groenewold NA, Grotegerd D, Gruber O, Gruner P, Gur RE, Gur RC, Hahn T, Harrison BJ, Hartman CA, Hatton SN, Heinz A, Heslenfeld DJ, Hibar DP, Hickie IB, Ho BC, Hoekstra PJ, Hohmann S, Holmes AJ, Hoogman M, Hosten N, Howells FM, Hulshoff Pol HE, Huyser C, Jahanshad N, James A, Jernigan TL, Jiang J, Jönsson EG, Joska JA, Kahn R, Kalnin A, Kanai R, Klein M, Klyushnik TP, Koenders L, Koops S, Krämer B, Kuntsi J, Lagopoulos J, Lázaro L, Lebedeva I, Lee WH, Lesch KP, Lochner C, Machielsen MWJ, Maingault S, Martin NG, Martínez-Zalacaín I, Mataix-Cols D, Mazoyer B, McDonald C, McDonald BC, McIntosh AM, McMahon KL, McPhilemy G, Meinert S, Menchón JM, Medland SE, Meyer-Lindenberg A, Naaijen J, Najt P, Nakao T, Nordvik JE, Nyberg L, Oosterlaan J, de la Foz VO, Paloyelis Y, Pauli P, Pergola G, Pomarol-Clotet E, Portella MJ, Potkin SG, Radua J, Reif A, Rinker DA, Roffman JL, Rosa PGP, Sacchet MD, Sachdev PS, Salvador R, Sánchez-Juan P, Sarró S, Satterthwaite TD, Saykin AJ, Serpa MH, Schmaal L, Schnell K, Schumann G, Sim K, Smoller JW, Sommer I, Soriano-Mas C, Stein DJ, Strike LT, Swagerman SC, Tamnes CK, Temmingh HS, Thomopoulos SI, Tomyshev AS, Tordesillas-Gutiérrez D, Trollor JN, Turner JA, Uhlmann A, van den Heuvel OA, van den Meer D, van der Wee NJA, van Haren NEM, van 't Ent D, van Erp TGM, Veer IM, Veltman DJ, Voineskos A, Völzke H, Walter H, Walton E, Wang L, Wang Y, Wassink TH, Weber B, Wen W, West JD, Westlye LT, Whalley H, Wierenga LM, Wittfeld K, Wolf DH, Worker A, Wright MJ, Yang K, Yoncheva Y, Zanetti MV, Ziegler GC; Karolinska Schizophrenia Project, Thompson PM, Dima D (2022) Cortical thickness across the lifespan: Data from 17,075 healthy individuals aged 3-90 years. *Hum Brain Mapp* 43:431-51

189) Dima D, Modabbernia A, Papachristou E, Doucet GE, Agartz I, Aghajani M, Akudjedu TN, Albajes-Eizagirre A, Alnaes D, Alpert KI, Andersson M, Andreasen NC, Andreassen OA, Asherson P, Banaschewski T, Bargallo N, Baumeister S, Baur-Streubel R, Bertolino A, Bonvino A, Boomsma DI, Borgwardt S, Bourque J, Brandeis D, Breier A, Brodaty H, Brouwer RM, Buitelaar JK, Busatto GF, Buckner RL, Calhoun V, Canales-Rodríguez EJ, Cannon DM, Caseras X, Castellanos FX, Cervenka S, Chaim-Avancini TM, Ching CRK, Chubar V, Clark VP, Conrod P, Conzelmann A, Crespo-Facorro B, Crivello F, Crone EA, Dannlowski U, Dale AM, Davey C, de Geus EJC, de Haan L, de Zubicaray GI, den Braber A, Dickie EW, Di Giorgio A, Doan NT, Dørum ES, Ehrlich S, Erk S, Espeseth T, Fatouros-Bergman H, [Fisher SE](#), Fouche JP, Franke B, Frodl T, Fuentes-Claramonte P, Glahn DC, Gotlib IH, Grabe HJ, Grimm O, Groenewold NA, Grotegerd D, Gruber O, Gruner P, Gur RE, Gur RC, Hahn T, Harrison BJ, Hartman CA, Hatton SN, Heinz A, Heslenfeld DJ, Hibar DP, Hickie IB, Ho BC, Hoekstra PJ, Hohmann S, Holmes AJ, Hoogman M, Hosten N, Howells FM, Hulshoff Pol HE, Huyser C, Jahanshad N, James A, Jernigan TL, Jiang J, Jönsson EG, Joska JA, Kahn R, Kalnin A, Kanai R, Klein M, Klyushnik TP, Koenders L, Koops S, Krämer B, Kuntsi J, Lagopoulos J, Lázaro L, Lebedeva I, Lee WH, Lesch KP, Lochner C, Machielsen MWJ, Maingault S, Martin NG, Martínez-Zalacaín I, Mataix-Cols D, Mazoyer B, McDonald C, McDonald BC, McIntosh AM, McMahon KL, McPhilemy G, Meinert S, Menchón JM, Medland SE, Meyer-Lindenberg A, Naaijen J, Najt P, Nakao T, Nordvik JE, Nyberg L, Oosterlaan J, de la Foz VO, Paloyelis Y, Pauli P, Pergola G, Pomarol-Clotet E, Portella MJ, Potkin SG, Radua J, Reif A, Rinker DA, Roffman JL, Rosa PGP, Sacchet MD, Sachdev PS, Salvador R, Sánchez-Juan P, Sarró S, Satterthwaite TD, Saykin AJ, Serpa MH, Schmaal L, Schnell K, Schumann G, Sim K, Smoller JW, Sommer I, Soriano-Mas C, Stein DJ, Strike LT, Swagerman SC, Tamnes CK, Temmingh HS, Thomopoulos SI, Tomyshev AS, Tordesillas-Gutiérrez D, Trollor JN, Turner JA, Uhlmann A, van den Heuvel OA, van den Meer D, van der Wee NJA, van Haren NEM, Van't Ent D, van Erp TGM, Veer IM, Veltman DJ, Voineskos A, Völzke H, Walter H, Walton E, Wang L, Wang Y, Wassink TH, Weber B, Wen W, West JD, Westlye LT, Whalley H, Wierenga LM, Williams SCR, Wittfeld K, Wolf DH, Worker A, Wright MJ, Yang K, Yoncheva Y, Zanetti MV, Ziegler GC, Thompson PM, Frangou S; Karolinska Schizophrenia Project (2022) Subcortical volumes across the lifespan: Data from 18,605 healthy individuals aged 3-90 years. *Hum Brain Mapp* 43:452-69

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 190)** Park BY, Larivière S, Rodríguez-Cruces R, Royer J, Tavakol S, Wang Y, Caciagli L, Caligiuri ME, Gambardella A, Concha L, Keller SS, Cendes F, Alvim MKM, Yasuda C, Bonilha L, Gleichgerrcht E, Focke NK, Kreilkamp BAK, Domin M, von Podewils F, Langner S, Rummel C, Rebsamen M, Wiest R, Martin P, Kotikalapudi R, Bender B, O'Brien TJ, Law M, Sinclair B, Vivash L, Kwan P, Desmond PM, Malpas CB, Lui E, Alhusaini S, Doherty CP, Cavalleri GL, Delanty N, Kälviäinen R, Jackson GD, Kowalczyk M, Mascalchi M, Semmelroch M, Thomas RH, Soltanian-Zadeh H, Davoodi-Bojd E, Zhang J, Lenge M, Guerrini R, Bartolini E, Hamandi K, Foley S, Weber B, Depondt C, Absil J, Carr SJA, Abela E, Richardson MP, Devinsky O, Severino M, Striano P, Parodi C, Tortora D, Hatton SN, Vos SB, Duncan JS, Galovic M, Whelan CD, Bargalló N, Pariente J, Conde-Blanco E, Vaudano AE, Tondelli M, Meletti S, Kong XZ, Francks C, [Fisher SE](#), Caldairou B, Ryten M, Labate A, Sisodiya SM, Thompson PM, McDonald CR, Bernasconi A, Bernasconi N, Bernhardt BC (2022) Topographic divergence of atypical cortical asymmetry and atrophy patterns in temporal lobe epilepsy. *Brain* 145:1285-98
- 191)** van der Spek J, den Hoed J, Snijders Blok L, Dingemans AJM, Schijven D, Nellaker C, Venselaar H, Astuti GDN, Barakat TS, Bebin EM, Beck-Wödl S, Beunders G, Brown NJ, Brunet T, Brunner HG, Campeau PM, Čuturilo G, Gilissen C, Haack TB, Hüning I, Husain RA, Kamien B, Lim SC, Lovrecic L, Magg J, Maver A, Miranda V, Monteil DC, Ockeloen CW, Pais LS, Plaiasu V, Raiti L, Richmond C, Rieß A, Schwaibold EMC, Simon MEH, Spranger S, Tan TY, Thompson ML, de Vries BBA, Wilkins EJ, Willemsen MH, Francks C, Vissers LELM, [Fisher SE](#), Kleefstra T (2022) Inherited variants in CHD3 show variable expressivity in Snijders Blok-Campeau syndrome. *Genet Med* 24:1283-96
- 192)** de Boer E, Ockeloen CW, Kampen RA, Hampstead JE, Dingemans AJM, Rots D, Lütje L, Ashraf T, Baker R, Barat-Houari M, Angle B, Chatron N, Denommé-Pichon AS, Devinsky O, Dubourg C, Elmslie F, Elloumi HZ, Faivre L, Fitzgerald-Butt S, Geneviève D, Goos JAC, Helm BM, Kini U, Lasa-Aranzasti A, Lesca G, Lynch SA, Mathijssen IMJ, McGowan R, Monaghan KG, Odent S, Pfundt R, Putoux A, van Reeuwijk J, Santen GWE, Sasaki E, Sorlin A, van der Spek PJ, Stegmann APA, Swagemakers SMA, Valenzuela I, Viora-Dupont E, Vitobello A, Ware SM, Wéber M, Gilissen C, Low KJ, [Fisher SE](#), Vissers LELM, Wong MMK, Kleefstra T (2022) Missense variants in ANKRD11 cause KBG syndrome by impairment of stability or transcriptional activity of the encoded protein. *Genet Med* 24:2051-64
- 193)** Chormai P, Pu Y, Hu H, [Fisher SE](#), Francks C, Kong XZ (2022) Machine learning of large-scale multimodal brain imaging data reveals neural correlates of hand preference. *Neuroimage* 262:119534
- 194)** Eising E, Mirza-Schreiber N, de Zeeuw EL, Wang CA, Truong DT, Allegrini AG, Shapland CY, Zhu G, Wigg KG, Gerritse ML, Molz B, Alagöz G, Gialluisi A, Abbondanza F, Rimfeld K, van Donkelaar M, Liao Z, Jansen PR, Andlauer TFM, Bates TC, Bernard M, Blokland K, Bonte M, Børglum AD, Bourgeron T, Brandeis D, Ceroni F, Csépe V, Dale PS, de Jong PF, DeFries JC, Démonet JF, Demontis D, Feng Y, Gordon SD, Guger SL, Hayiou-Thomas ME, Hernández-Cabrera JA, Hottenga JJ, Hulme C, Kere J, Kerr EN, Koomar T, Landerl K, Leonard GT, Lovett MW, Lyytinen H, Martin NG, Martinelli A, Maurer U, Michaelson JJ, Moll K, Monaco AP, Morgan AT, Nöthen MM, Pausova Z, Pennell CE, Pennington BF, Price KM, Rajagopal VM, Ramus F, Richer L, Simpson NH, Smith SD, Snowling MJ, Stein J, Strug LJ, Talcott JB, Tiemeier H, van der Schroeff MP, Verhoef E, Watkins KE, Wilkinson M, Wright MJ, Barr CL, Boomsma DI, Carreiras M, Franken MJ, Gruen JR, Luciano M, Müller-Myhsok B, Newbury DF, Olson RK, Paracchini S, Paus T, Plomin R, Reilly S, Schulte-Körne G, Tomblin JB, van Bergen E, Whitehouse AJO, Willcutt EG, St Pourcain B, Francks C, [Fisher SE](#) (2022) Genome-wide analyses of individual differences in quantitatively assessed reading- and language-related skills in up to 34,000 people. *Proc Natl Acad Sci USA* 119:e2202764119
- 195)** Alagöz G, Molz B, Eising E, Schijven D, Francks C, Stein JL, [Fisher SE](#) (2022) Using neuroimaging genomics to investigate the evolution of human brain structure. *Proc Natl Acad Sci USA* 119:e2200638119
- 196)** Doust C, Fontanillas P, Eising E, Gordon SD, Wang Z, Alagöz G, Molz B; 23andMe Research Team; Quantitative Trait Working Group of the GenLang Consortium; Pourcain BS, Francks C, Marioni RE, Zhao J, Paracchini S, Talcott JB, Monaco AP, Stein JF, Gruen JR, Olson RK, Willcutt EG, DeFries JC, Pennington BF, Smith SD, Wright MJ, Martin NG, Auton A, Bates TC, [Fisher SE](#), Luciano M (2022) Discovery of 42 genome-wide significant loci associated with dyslexia. *Nature Genet* 54:1621-9

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 197)** Tielbeek JJ, Uffelmann E, Williams BS, Colodro-Conde L, Gagnon É, Mallard TT, Levitt BE, Jansen PR, Johansson A, Sallis HM, Pistis G, Saunders GRB, Allegrini AG, Rimfeld K, Konte B, Klein M, Hartmann AM, Salvatore JE, Nolte IM, Demontis D, Malmberg ALK, Burt SA, Savage JE, Sugden K, Poulton R, Harris KM, Vrieze S, McGue M, Iacono WG, Mota NR, Mill J, Viana JF, Mitchell BL, Morosoli JJ, Andlauer TFM, Ouellet-Morin I, Tremblay RE, Côté SM, Gouin JP, Brendgen MR, Dionne G, Vitaro F, Lupton MK, Martin NG; COGA Consortium; Spit for Science Working Group; Castelao E, Rääkkönen K, Eriksson JG, Lahti J, Hartman CA, Oldehinkel AJ, Snieder H, Liu H, Preisig M, Whipp A, Vuoksimaa E, Lu Y, Jern P, Rujescu D, Giegling I, Palviainen T, Kaprio J, Harden KP, Munafò MR, Morneau-Vaillancourt G, Plomin R, Viding E, Boutwell BB, Aliev F, Dick DM, Popma A, Faraone SV, Børghlum AD, Medland SE, Franke B, Boivin M, Pingault JB, Glennon JC, Barnes JC, Fisher SE, Moffitt TE, Caspi A, Polderman TJC, Posthuma D (2022) Uncovering the genetic architecture of broad antisocial behavior through a genome-wide association study meta-analysis. *Mol Psychiatry* 27:4453-63
- 198)** Kaspi A, Hildebrand MS, Jackson VE, Braden R, van Reyk O, Howell T, Debono S, Lauretta M, Morison L, Coleman MJ, Webster R, Coman D, Goel H, Wallis M, Dabscheck G, Downie L, Baker EK, Parry-Fielder B, Ballard K, Harrold E, Ziegenfusz S, Bennett MF, Robertson E, Wang L, Boys A, Fisher SE, Amor DJ, Scheffer IE, Bahlo M, Morgan AT (2023) Genetic aetiologies for childhood speech disorder: novel pathways co-expressed during brain development. *Mol Psychiatry* 28:1647-63
- 199)** Morison LD, Meffert E, Stampfer M, Steiner-Wilke I, Vollmer B, Schulze K, Briggs T, Braden R, Vogel A, Thompson-Lake D, Patel C, Blair E, Goel H, Turner S, Moog U, Riess A, Liegeois F, Koolen DA, Amor DJ, Kleefstra T, Fisher SE, Zweier C, Morgan AT (2023) In-depth characterisation of a cohort of individuals with missense and loss-of-function variants disrupting FOXP2. *J Med Genet* 60:597-607
- 200)** Snijders Blok L, Verseput J, Rots D, Venselaar H, Innes AM, Stumpel C, Öunap K, Reinson K, Seaby EG, McKee S, Burton B, Kim K, van Hagen JM, Waisfisz Q, Joset P, Steindl K, Rauch A, Li D, Zackai EH, Sheppard SE, Keena B, Hakonarson H, Roos A, Kohlschmidt N, Cereda A, Iacone M, Rebessi E, Kernohan KD, Campeau PM, Millan F, Taylor JA, Lochmüller H, Higgs MR, Goula A, Bernhard B, Velasco DJ, Schmanski AA, Stark Z, Gallacher L, Pais L, Marcogliese PC, Yamamoto S, Raun N, Jakub TE, Kramer JM, den Hoed J, Fisher SE, Brunner HG, Kleefstra T (2023) A clustering of heterozygous missense variants in the crucial chromatin modifier WDR5 defines a new neurodevelopmental disorder. *Hum Genet Genom Advances* 4:100157
- 201)** Lemaitre H, Le Guen Y, Tilot AK, Stein JL, Philippe C, Mangin JF, Fisher SE, Frouin V (2023) Genetic variations within human gained enhancer elements affect human brain sulcal morphology. *Neuroimage* 265:119773
- 202)** Sollis E, den Hoed J, Quevedo M, Estruch SB, Vino A, Dekkers DHW, Demmers JA, Poot R, Deriziotis P, Fisher SE (2023) Characterization of the TBR1 interactome: variants associated with neurodevelopmental disorders disrupt novel protein interactions. *Hum Mol Genet* 32:1497-1510
- 203)** Abbondanza F, Dale PS, Wang CA, Hayiou-Thomas ME, Toseeb U, Koomar TS, Wigg KG, Feng Y, Price KM, Kerr EN, Guger SL, Lovett MW, Strug LJ, van Bergen E, Dolan CV, Tomblin JB, Moll K, Schulte-Körne G, Neuhoff N, Warnke A, Fisher SE, Barr CL, Michaelson JJ, Boomsma DI, Snowling MJ, Hulme C, Whitehouse AJO, Pennell CE, Newbury DF, Stein J, Talcott JB, Bishop DVM, Paracchini S (2023) Language and reading impairments are associated with increased prevalence of non-right-handedness. *Child Dev* 94:970-84
- 204)** Sha Z, Schijven D, Fisher SE, Francks C (2023) Genetic architecture of the white matter connectome of the human brain. *Sci Adv* 9:eadd2870
- 205)** Heim F, Fisher SE, Scharff C, Ten Cate C, Riebel K (2023) Effects of Cortical FoxP1 Knockdowns on Learned Song Preference in Female Zebra Finches. *eNeuro* 10:ENEURO.0328-22.2023
- 206)** Schijven D *et al* (2023) Large-scale analysis of structural brain asymmetries in schizophrenia via the ENIGMA consortium. *Proc Natl Acad Sci USA* 120:e2213880120. doi: 10.1073/pnas.2213880120
- 207)** Oliveira-Stahl G, Farboud S, Sterling ML, Heckman JJ, van Raalte B, Lenferink D, van der Stam A, Smeets CJLM, Fisher SE¹, Englitz B¹ (2023) High-precision spatial analysis of mouse courtship vocalization behavior reveals sex and strain differences. *Sci Rep* 13:5219 ¹joint senior authors

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 208)** Horton S, Jackson V, Boyce J, Franken MC, Siemers S, John MS, Hearps S, van Reyk O, Braden R, Parker R, Vogel AP, Eising E, Amor DJ, Irvine J, Fisher SE, Martin NG, Reilly S, Bahlo M, Scheffer I, Morgan A (2023) Self-Reported Stuttering Severity Is Accurate: Informing Methods for Large-Scale Data Collection in Stuttering. *J Speech Lang Hear Res* doi: 10.1044/2023_JSLHR-23-00081. Online ahead of print.
- 209)** Boen R *et al* (2024) Beyond the Global Brain Differences: Intraindividual Variability Differences in 1q21.1 Distal and 15q11.2 BP1-BP2 Deletion Carriers. *Biol Psychiatry* 95:147-60
- 210)** Verhoef E, Allegrini AG, Jansen PR, Lange K, Wang CA, Morgan AT, Ahluwalia TS, Symeonides C; EAGLE Working Group; Eising E, Franken MC, Hypponen E, Mansell T, Ollislagers M, Omerovic E, Rimfeld K, Schlag F, Selzam S, Shapland CY, Tiemeier H, Whitehouse AJO, Saffery R, Bønnelykke K, Reilly S, Pennell CE, Wake M, Cecil CAM, Plomin R, Fisher SE, St Pourcain B (2024) Genome-Wide Analyses of Vocabulary Size in Infancy and Toddlerhood: Associations With Attention-Deficit/Hyperactivity Disorder, Literacy, and Cognition-Related Traits. *Biol Psychiatry* 95:859-69
- 211)** Bignardi G, Smit DJA, Vessel EA, Trupp MD, Ticini LF, Fisher SE, Polderman TJC (2024) Genetic effects on variability in visual aesthetic evaluations are partially shared across visual domains. *Commun Biol* 7:55
- 212)** Ge R, Yu Y, Qi YX, Fan YN, Chen S, Gao C, Haas SS, New F, Boomsma DI, Brodaty H, Brouwer RM, Buckner R, Caseras X, Crivello F, Crone EA, Erk S, Fisher SE, Franke B, Glahn DC, Dannlowski U, Grotegerd D, Gruber O, Hulshoff Pol HE, Schumann G, Tamnes CK, Walter H, Wierenga LM, Jahanshad N, Thompson PM, Frangou S; ENIGMA Lifespan Working Group (2024) Normative modelling of brain morphometry across the lifespan with CentileBrain: algorithm benchmarking and model optimisation. *Lancet Digit Health* 6:e211-e221
- 213)** de Hoyos L, Barendse MT, Schlag F, van Donkelaar MMJ, Verhoef E, Shapland CY, Klassmann A, Buitelaar J, Verhulst B, Fisher SE, Rai D, St Pourcain B (2024) Structural models of genome-wide covariance identify multiple common dimensions in autism. *Nature Commun* 15:1770
- 214)** Goltermann O, Alagöz G, Molz B, Fisher SE (2024) Neuroimaging genomics as a window into the evolution of human sulcal organization. *Cereb Cortex* 34:bhae078
- 215)** Wesseldijk LW, Henechowicz TL, Baker DJ, Bignardi G, Karlsson R, Gordon RL, Mosing MA, Ullén F, Fisher SE (2024) Notes from Beethoven's genome. *Curr Biol* 34:R233-R234
- 216)** Schijven D, Soheili-Nezhad S, Fisher SE, Francks C (2024) Exome-wide analysis implicates rare protein-altering variants in human handedness. *Nature Commun* 15:2632
- 217)** Eising E, VINO A, Mabie HL, Campbell TF, Shriberg LD, Fisher SE (2024) Genome sequencing of idiopathic speech delay. *Hum Mutat* 2024: 9692863. doi:10.1155/2024/9692863
- 218)** Engelen MM, Franken MJ, Stipdonk LW, Horton SE, Jackson VE, Reilly S, Morgan AT, Fisher SE, van Dulmen S, Eising E (2024) The Association Between Stuttering Burden and Psychosocial Aspects of Life in Adults. *J Speech Lang Hear Res* 67:1385-99
- 219)** Heim F, Scharff C, Fisher SE, Riebel K, Ten Cate C (2024) Auditory discrimination learning and acoustic cue weighing in female zebra finches with localized FoxP1 knockdowns. *J Neurophysiol* 131:950-63
- 220)** Kurth F, Schijven D, Van den Heuvel OA, Hoogman M, Van Rooij D, Stein DJ, Buitelaar JK, Bölte S, Auzias G, Kushki A, Venkatasubramanian G, Rubia K, Bollmann S, Isaksson J, Jaspers-Fayer F, Marsh R, Batistuzzo MC, Arnold PD, Bressan RA, Stewart ES, Gruner P, Sorensen L, Pan PM, Silk TJ, Gur RC, Cubillo AI, Haavik J, O'Gorman Tuura RL, Hartman CA, Calvo R, McGrath J, Calderoni S, Jackowski A, Chantiluke KC, Satterthwaite TD, Busatto GF, Nigg JT, Gur RE, Retico A, Tosetti M, Gallagher L, Szeszko PR, Neufeld J, Ortiz AE, Ghisleni C, Lazaro L, Hoekstra PJ, Anagnostou E, Hoekstra L, Simpson B, Plessen JK, Deruelle C, Soreni N, James A, Narayanaswamy J, Reddy JYC, Fitzgerald J, Bellgrove MA, Salum GA, Janssen J, Muratori F, Vila M, Garcia Giral M, Ameis SH, Bosco P, Lundin Remnélius K, Huyser C, Pariente JC, Jalbrzikowski M, Rosa PGP, O'Hearn KM, Ehrlich S, Mollon J, Zugman A, Christakou A, Arango C, Fisher SE, Kong X-Z, Franke B, Medland SE, Thomopoulos SI, Jahanshad N, Glahn DC, Thompson PM, Francks C, Luders E (in press) Large-scale analysis of structural brain asymmetries during neurodevelopment: Age effects and sex differences in 4,265 children and adolescents. *Hum Brain Mapp*
- 221)** García-Marín LM *et al* (in press) Genomic analysis of intracranial and subcortical brain volumes yields polygenic scores accounting for brain variation across ancestries. *Nature Genet*

Review/perspective articles in journals

[Includes peer-reviewed articles as indicated]

- 1) Fisher SE, Stein JF, Monaco AP (1999) A genome-wide search strategy for identifying quantitative trait loci involved in reading and spelling disability (developmental dyslexia). *Eur Child & Adol Psych* 8-S3:47-51**
- 2) Francks C, Fisher SE, Marlow AJ, Richardson AJ, Stein JF, Monaco AP (2000) A sibling-pair based approach for mapping genetic loci that influence quantitative measures of reading disability. *Prostaglandins, Leukotrienes & Essential Fatty Acids* 63:27-31**
- 3) Fisher SE, DeFries JC (2002) Developmental dyslexia: genetic dissection of a complex cognitive trait. *Nature Rev Neurosci* 3:767-80 [peer-reviewed]**
- 4) Marcus GF, Fisher SE (2003) FOXP2 in focus: what can genes tell us about speech and language? *Trends Cogn Sci* 7:257-62 [peer-reviewed]**
- 5) Fisher SE, Lai CSL, Monaco AP (2003) Deciphering the genetic basis of speech and language disorders. *Annu Rev Neurosci* 26:57-80**
- 6) Fisher SE (2005) Dissection of molecular mechanisms underlying speech and language disorders. *Appl Psycholing* 26:111-28**
- 7) Fisher SE (2005) On genes, speech, and language. *N Engl J Med* 353:1655-7**
- 8) Fisher SE, Marcus GF (2006) The eloquent ape: genes, brains and the evolution of language. *Nature Rev Genet* 7:9-20 [peer-reviewed]**
- 9) Fisher SE, Francks C (2006) Genes, cognition and dyslexia: learning to read the genome. *Trends Cogn Sci* 10:250-7 [peer-reviewed]**
- 10) Fisher SE (2006) Tangled webs: tracing the connections between genes and cognition. *Cognition* 101:270-97 [peer-reviewed]**
- 11) White SA, Fisher SE, Geschwind DH, Scharff C, Holy TE (2006) Singing mice, songbirds, and more: models for FOXP2 function and dysfunction in human speech and language. *J Neurosci* 26:10376-9 [peer-reviewed]**
- 12) Fisher SE (2007) Molecular windows into speech and language disorders. *Folia Phoniatica et Logopaedica* 59:130-40**
- 13) Fisher SE, Scharff C (2009) FOXP2 as a molecular window into speech and language. *Trends Genet* 25:166-77 [peer-reviewed]**
- 14) Vernes SC, Fisher SE (2009) Unravelling neurogenetic networks implicated in developmental language disorders. *Biochem Soc Trans* 37:1263-9**
- 15) Newbury DF, Fisher SE, Monaco AP (2010) Recent advances in the genetics of language impairment. *Genome Medicine* 2:6**
- 16) Fisher SE (2010) Genetic susceptibility to stuttering. *N Engl J Med* 362:750-2**
- 17) Fisher SE, Ridley M (2013) Culture, genes and the human revolution. *Science* 340:929-30 [peer-reviewed]**
- 18) Graham SA, Fisher SE (2013) Decoding the genetics of speech and language. *Curr Opin Neurobiol* 23:43-51 [peer-reviewed]**
- 19) Deriziotis P, Fisher SE (2013) Neurogenomics of speech and language disorders: the road ahead. *Genome Biol* 14:204**
- 20) Carrion-Castillo A, Franke B, Fisher SE (2013) Molecular genetics of dyslexia: an overview. *Dyslexia* 19:214-40 [peer-reviewed]**
- 21) Willems RM, Van der Haegen L, Fisher SE, Francks C (2014) On the other hand: Left-handers in cognitive neuroscience and neurogenetics. *Nature Rev Neurosci* 15:193-201 [peer-reviewed]**

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

- 22)** Thompson PM *et al* (2014) The ENIGMA Consortium: large-scale collaborative analyses of neuroimaging and genetic data. *Brain Imaging Behav* 8:153-82
- 23)** French CA, Fisher SE (2014) What can mice tell us about Foxp2 function? *Curr Opin Neurobiol* 28C:72-9 [peer-reviewed]
- 24)** Fisher SE, Vernes SC (2015) Genetics and the language sciences *Annu Rev Linguist* 1:6.1–6.22
- 25)** Gingras B, Peretz I, Huron D, Honing H, Trainor L, Fisher SE (2015) Defining the biological bases of individual differences in musicality. *Phil Trans Roy Soc B* 370:20140092 [peer-reviewed]
- 26)** Graham SA, Deriziotis P, Fisher SE (2015) Insights into the genetic foundations of human communication. *Neuropsychol Rev* 25:3-26 [peer-reviewed]
- 27)** Graham SA, Fisher SE (2015) Understanding language from a genomic perspective. *Annu Rev Genet* 49:131-60
- 28)** Morgan A, Fisher SE, Scheffer I, Hildebrand M (2016) FOXP2-related speech and language disorders. In: *GeneReviews* (eds. Adam MP, Ardinger HH, Pagon RA, Wallace SE, Bean LJH, Stephens K, Amemiya A) <http://www.ncbi.nlm.nih.gov/books/NBK368474/> [updated 2017 & 2023] [peer-reviewed]
- 29)** Thompson PM, Andreassen OA, Arias-Vasquez A, Bearden CE, Boedhoe PS, Brouwer RM, Buckner RL, Buitelaar JK, Bulaeva KB, Cannon DM, Cohen RA, Conrod PJ, Dale AM, Deary IJ, Dennis EL, de Reus MA, Desrivieres S, Dima D, Donohoe G, Fisher SE, Fouché J-P, Francks C, Frangou S, Franke B, Ganjgahi H, Garavan H, Glahn DC, Grabe HJ, Guadalupe T, Gutman BA, Hashimoto R, Hibar DP, Holland D, Hoogman M, Pol HEH, Hosten N, Jahanshad N, Kelly S, Kochunov P, Kremen WS, Lee PH, Mackey S, Martin NG, Mazoyer B, McDonald C, Medland SE, Morey RA, Nichols TE, Paus T, Pausova Z, Schmaal L, Schumann G, Shen L, Sisodiya SM, Smit DJ, Smoller JW, Stein DJ, Stein JL, Toro R, Turner JA, van den Heuvel M, van den Heuvel OA, van Erp TG, van Rooij D, Veltman DJ, Walter H, Wang Y, Wardlaw JM, Whelan CD, Wright MJ, Ye J (2017) ENIGMA and the individual: Predicting factors that affect the brain in 35 countries worldwide. *Neuroimage* 145:389-408 [peer-reviewed]
- 30)** Fisher SE (2017) Evolution of language: Lessons from the genome. *Psychon Bull Review* 24:34-40 [peer-reviewed]
- 31)** Deriziotis P, Fisher SE (2017) Speech and language: Translating the genome. *Trends Genet* 33:642-56 [peer-reviewed]
- 32)** Zubicaray G, Fisher SE (2017) Genes, Brain, and Language: A brief introduction to the Special Issue. *Brain Lang* 172:1-2
- 33)** Fisher SE (2019) Human genetics: The evolving story of FOXP2. *Curr Biol* 29:R65-7
- 34)** Fisher SE, Tilot AK (2019) Bridging senses: novel insights from synaesthesia. *Phil Trans Roy Soc B* 374:20190022
- 35)** den Hoed J, Fisher SE (2020) Genetic pathways involved in human speech disorders. *Curr Opin Genet Dev* 65:103-111 [peer-reviewed]
- 36)** Thompson PM *et al* (2020) ENIGMA and global neuroscience: A decade of large-scale studies of the brain in health and disease across more than 40 countries. *Transl Psychiatry* 10:100 [peer-reviewed]
- 37)** den Hoed J, Devaraju K, Fisher SE (2021) Molecular networks of the FOXP2 transcription factor in the brain. *EMBO Rep* 22:e52803 [peer-reviewed]
- 38)** Kong XZ, Postema MC, Guadalupe T, de Kovel C, Boedhoe PSW, Hoogman M, Mathias SR, van Rooij D, Schijven D, Glahn DC, Medland SE, Jahanshad N, Thomopoulos SI, Turner JA, Buitelaar J, van Erp TGM, Franke B, Fisher SE, van den Heuvel OA, Schmaal L, Thompson PM, Francks C (2022) Mapping brain asymmetry in health and disease through the ENIGMA consortium. *Hum Brain Mapp* 43:167-81 [peer-reviewed]
- 39)** Nayak S, Coleman PL, Ladányi E, Nitin R, Gustavson DE, Fisher SE, Magne CL, Gordon RL (2022) The Musical Abilities, Pleiotropy, Language, and Environment (MAPLE) Framework for Understanding Musicality-Language Links Across the Lifespan. *Neurobiology of Language* 3:615-64 [peer-reviewed]

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

40) Søndery IE, Ching CRK, Thomopoulos SI, van der Meer D, Sun D, Villalon-Reina JE, Agartz I, Amunts K, Arango C, Armstrong NJ, Ayesa-Arriola R, Bakker G, Bassett AS, Boomsma DI, Bülow R, Butcher NJ, Calhoun VD, Caspers S, Chow EWC, Cichon S, Ciufolini S, Craig MC, Crespo-Facorro B, Cunningham AC, Dale AM, Dazzan P, de Zubicaray GI, Djurovic S, Doherty JL, Donohoe G, Draganski B, Durdle CA, Ehrlich S, Emanuel BS, Espeseth T, Fisher SE, Ge T, Glahn DC, Grabe HJ, Gur RE, Gutman BA, Haavik J, Håberg AK, Hansen LA, Hashimoto R, Hibar DP, Holmes AJ, Hottenga JJ, Hulshoff Pol HE, Jalbrzikowski M, Knowles EEM, Kushan L, Linden DEJ, Liu J, Lundervold AJ, Martin-Brevet S, Martínez K, Mather KA, Mathias SR, McDonald-McGinn DM, McRae AF, Medland SE, Moberget T, Modenato C, Monereo Sánchez J, Moreau CA, Mühleisen TW, Paus T, Pausova Z, Prieto C, Ragothaman A, Reinbold CS, Reis Marques T, Repetto GM, Reymond A, Roalf DR, Rodriguez-Herreros B, Rucker JJ, Sachdev PS, Schmitt JE, Schofield PR, Silva AI, Stefansson H, Stein DJ, Tamnes CK, Tordesillas-Gutiérrez D, Ulfarsson MO, Vajdi A, van 't Ent D, van den Bree MBM, Vassos E, Vázquez-Bourgon J, Vila-Rodriguez F, Walters GB, Wen W, Westlye LT, Wittfeld K, Zackai EH, Stefansson K, Jacquemont S, Thompson PM, Bearden CE, Andreassen OA; ENIGMA-CNV Working Group; ENIGMA 22q11.2 Deletion Syndrome Working Group. (2022) Effects of copy number variations on brain structure and risk for psychiatric illness: Large-scale studies from the ENIGMA working groups on CNVs. *Hum Brain Mapp* 43:300-28 [peer-reviewed]

SIMON E. FISHER - PUBLICATIONS (JUNE 2024)

Book chapters

- 1) Fisher SE, Smith SD (2001) Progress towards the identification of genes influencing developmental dyslexia. In: *Dyslexia: Theory and good practice* (ed. Fawcett AJ) 39-64 (Whurr, London, UK)
- 2) Fisher SE (2002) Isolation of the genetic factors underlying speech and language disorders. In: *Behavioral Genetics in the Postgenomic Era* (eds. Plomin R, DeFries JC, Craig IW, McGuffin P) 205-26 (APA Books, Washington DC, USA)
- 3) Fisher SE (2003) The genetic basis of a severe speech and language disorder. In: *Neurosciences in the postgenomic era* (eds. Mallet J, Christen Y) 125-34 (Springer Verlag, Germany)
- 4) Fisher SE (2006) How can animal studies help to uncover the roles of genes implicated in human speech and language disorders? In: *Transgenic and Knockout Models of Neuropsychiatric Disorders* (eds. Fisch GS, Flint J) 127-49 (Humana press, USA)
- 5) Ramus F, Fisher SE (2009) Genetics of language. In: *The Cognitive Neurosciences 4th Edition* (ed. Gazzaniga MS) 855-71 (MIT Press, Cambridge, MA, USA)
- 6) Marcus GF, Fisher SE (2011) Genes and language. In: *Cambridge Encyclopedia of the Language Sciences* (ed. Hogan PC) 341-4 (Cambridge University Press, New York, USA)
- 7) Vernes SC, Fisher SE (2011) Functional genomic dissection of speech and language disorders. In: *Genomics, Proteomics, and the Nervous System* (ed. Clelland JD) 253-78 (Springer)
- 8) Fisher SE (2012) Building bridges between genes, brains and language. In: *Birdsong, Speech and Language. Converging mechanisms* (eds. Bolhuis JJ, Everaert M) 425-54 (MIT Press, Cambridge, MA, USA).
- 9) Vernes SC, Fisher SE (2013) Genetic pathways implicated in speech and language. In: *Animal Models of Speech and Language Disorders* (ed. Helekar S) 13-40 (Springer)
- 10) Fisher SE (2014) Translating the Genome in Human Neuroscience. In: *The Future of The Brain: Essays By The World's Leading Neuroscientists* (eds. Marcus G, Freeman J) 149-58 (Princeton University Press)
- 11) Fisher SE (2015) A molecular genetic perspective on speech and language. In: *The Neurobiology of Language* (eds. Hickok G, Small SL) 13-24 (Elsevier)
- 12) De Kovel CGF, Fisher SE (2018) Molecular genetic methods. In: *Research Methods in Psycholinguistics and the Neurobiology of Language: A Practical Guide* (eds. De Groot AMB, Hagoort P) 330-353 (Hoboken: Wiley-Blackwell)
- 13) Gingras B, Honing H, Peretz I, Trainor LJ, Fisher SE (2018) Defining the biological bases of individual differences in musicality. In *The Origins of Musicality* (ed. Honing H) 221-250 (MIT Press).
- 14) Fisher SE (2019) Genes and language: Key issues and ways forward. In *Human Language: from Genes and Brains to Behavior* (ed. Hagoort P) 609-620 (MIT Press)
- 15) Burenkova OV, Fisher SE (2019) Genetic insights into the neurobiology of speech and language. In *All About Language: Science, Theory, and Practice* (eds. Grigorenko E, Shtyrov Y, McCardle P) (Paul Brookes Publishing, Inc, Baltimore, MD)