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**Degrees**

Ph. D., Behavior Genetics (Emphasis: Quantitative Genetics), Department of Psychology, University of Colorado, Boulder, USA, 1994 [Dissertation title: "Structural Models for the Development of Specific Cognitive Abilities in the Colorado Adoption Project"]

M. A., Behavior Genetics (Emphasis: Quantitative Genetics), Department of Psychology, University of Colorado, Boulder, USA, 1992

B. A. (Honours), Psychology (Emphasis: Quantitative Psychology), The University of Winnipeg, Canada, 1989 [Thesis title: "Self-Deception and Impression Management in Approval Motivated Behaviour"]

**Academic appointments**

**2009–present Assistant Professor, Department of Psychiatry, The University of Hong Kong**

**2007–present Principal Investigator, The State Key Laboratory of Brain and Cognitive Sciences, The University of Hong Kong**

**2007–2009** Honorary Associate Professor, Department of Psychiatry, The University of Hong Kong

**2006–2009** Research Assistant Professor, Department of Psychiatry and Genome Research Centre, The University of Hong Kong

**2000–2006** Head of Statistical Genetics Applications, Wellcome Trust Centre for Human Genetics, University of Oxford

**2000–2006** Honorary Senior Lecturer in Statistical Genetics, Social, Genetic and Developmental Psychiatry Research Centre, Institute of Psychiatry, King's College London

**1999–2000** Senior Statistical Geneticist, Wellcome Trust Centre for Human Genetics, University of Oxford

**1999–2000** Senior Lecturer in Statistical Genetics, Social, Genetic and Developmental Psychiatry Research Centre, Institute of Psychiatry, King's College London

**1996–1999** Lecturer in Statistical Genetics, Social, Genetic and Developmental Psychiatry Research Centre, Institute of Psychiatry, King's College London

**1995–1999** Research Associate, Institute for Behavioral Genetics, University of Colorado

**1991** Teaching Assistant, two Introduction to Behavior Genetics courses, University of Colorado

**1989–1990** Lecturer, Introduction to Computing course, University of Colorado

**1989–1994** Predoctoral Trainee and Graduate Research Assistant, Institute for Behavioral Genetics, University of Colorado

#### Professional activities

**2012–present** Member, Editorial Board (Academic Editor), PeerJ

**2010–present** Member, Editorial Board (Academic Editor), PLoS One

**2005–2008** Elected Treasurer, Behavior Genetics Association

**2004–present** Member, Advisory Board, Journal of Child Psychology & Psychiatry

**2004** Guest instructor at the Cambridge Complex Traits Conference, Hinxton, United Kingdom, 4–7 April.

**2002–2005** Elected Member-at-Large, Behavior Genetics Association

**1999** Science Advisor, EINSHAC workshop on Science and Cases on Neuro and Behavioral Genetics, Airlie Conference Center, Virginia, 30 June–3 July 1999.

**1997–present** Associate Editor, Behavior Genetics

**1997** Guest faculty member, 18th Wellcome Trust Summer School on statistical genetics, Hinxton, United Kingdom, July.

**1993–present** Faculty, International Workshops on Methodology of Twin and Family Studies

#### University committees

**2013–present** Chair, IT Committee, Department of Psychiatry, The University of Hong Kong

**2013–2014** Faculty Human Resources Committee, LKS Faculty of Medicine, The University of Hong Kong

**2011–2013** Medical Loan Funds Committee, LKS Faculty of Medicine, The University of Hong Kong

**2000–2005** Member, IT Committee, Wellcome Trust Centre for Human Genetics, University of Oxford

**1994–1999** Member, Computing Committee, Institute for Behavioral Genetics, University of Colorado, Boulder

#### Honors and awards

**2000** Fulker Award (best paper of the year in *Behavior Genetics*), Behavior Genetics Association, for "Neale, M. C., Cherny, S. S., Sham, P. C., Whitfield, J. B., Heath, A. C., Birley, A. C., and Martin, N. G. (1999) Distinguishing population stratification from genuine allelic effects with Mx: Association of ADH2 with alcohol consumption. *Behavior Genetics*, 29, 233–244; doi:10.1023/A:1021638122693."

**1994** Dozier Award, Department of Psychology, University of Colorado

**1989** Academic Proficiency Scholarship, University of Winnipeg

**1989** Dr. C. J. Robson Scholarship in Psychology, University of Winnipeg

**1988–1989** Member of the University of Winnipeg Honours Society

**1988–1989** "Student of Highest Distinction", University of Winnipeg

**Grants and fellowships**

1. Natural Sciences and Engineering Research Council of Canada Postgraduate Scholarships (PGS-2 and PGS-3), 1/6/90–31/5/93.
2. NHLBI: “Biobehavioral Studies of Cardiovascular Disease” (HL-40962), 1/3/94–28/2/99, PI: Stephen Manuck, TDC: US\$4,148,124. Role on project: Consultant, 0% AY.
3. Natural Sciences and Engineering Research Council of Canada Postdoctoral Fellow, 1/1/95–31/12/95.
4. NIMH: “Workshop on Methodology of Twin Studies” (R25MH019918), 1/9/98–31/8/2008, PI: John K. Hewitt, TDC: US\$218,469. Role on project: Faculty member.
5. NIAAA: “Genes for Ethanol Sensitivity in the Rat — QTL” (AA-10556), 1/4/96–31/3/2000, PI: Richard A. Deitrich, TDC: US\$717,148. Role on project: Co-Principal Investigator, 25% AY.
6. Medical Research Council of Great Britain: “Genetics and Anxiety/Depression in 25,000 Sibs,” 1/9/97–31/8/2004, PI: Pak Sham, YDC: US\$110,204. Role on project: Co-Investigator/consultant.
7. NIDA: “Antisocial Drug Dependence: Genetics and Treatment, Component I: Adolescent Drug/Alcohol Dependence: Chromosomal Loci” (DA-11015), 1/8/97–30/6/2002, PI: John K. Hewitt, YDC: US\$47,018. Role on project: Co-Investigator/consultant.
8. NIDA: “Antisocial Drug Dependence: Genetics and Treatment, Component IV: Heritable Early Indicators of Risk for Drug Dependence” (DA-11015), 1/8/97–30/6/2002, PI: John K. Hewitt, YDC: US\$155,638. Role on project: Co-Investigator/consultant.
9. NEI: “Variance Components Models for Mapping QTLs” (EY-12562), 1/12/1998–30/9/2008, PI: John K. Hewitt, TDC: US\$1,419,799 (1/10/2002–30/9/2008). Role on project: Co-I, 1/12/1998–31/10/1999; PI of Oxford subcontract, 10% AY, 1/11/1999–31/3/2006; Co-PI of University of Hong Kong subcontract, 1/10/2006–30/9/2008 (HK\$636,082 direct costs to HKU).
10. Seed Funding Programme for Basic Research (HKU): “Whole genome family based association study to search for RET-dependent modifiers in Hirschsprung’s disease” (200611159152), 01/05/2007–30/04/2009, PI: Prof PKH Tam, TDC: HK\$120,000. Role on project: Co-I.
11. URC/CRCG — Conference Grants for Teaching Staff (HKU): “37th Annual Meeting of the Behavior Genetics Association: *QTL methodology in behavior genetics: from linkage to genomewide association*” (200703170087), 03/06/2007–06/06/2007, TDC: HK\$13,500. Role on project: PI.
12. University Grants Council (Hong Kong)/Competitive Earmarked Research Grants (CERG): “Genetic dissection of Hirschsprung’s disease” (HKU 7759/07M), 01/01/2008–01/01/2011, TDC: HK\$1,781,813, PI: Prof PKH Tam. Role on project: Co-I.
13. University Grants Council (Hong Kong)/Competitive Earmarked Research Grants (CERG): “Genome-wide association study of schizophrenia” (HKU 7747/07M), 01/01/2008–01/01/2011, TDC: HK\$1,798,990, PI: Prof PC Sham. Role on project: Co-I.
14. Seed Funding Programme for Basic Research (HKU): “Fine mapping of a Hirschsprung’s disease locus on the 3p21 candidate region” (200709159003), 30/6/2008–29/6/2010, TDC: HK\$80,000, PI: Dr MM Garcia-Barceló. Role on project: Co-I.
15. Seed Funding Programme for Basic Research (HKU): “Genome-wide association mapping of epilepsy using DNA pooling” (200711159114), 1/4/2008–31/3/2010, TDC: HK\$80,000. Role on project: PI.
16. URC/CRCG — Conference Grants for Teaching Staff (HKU): “Behavior Genetics Association Conference 2008: *Genomewide association study of Hirschsprung’s disease*” (200703170522), 25/06/2008–28/06/2008, TDC: HK\$13,500. Role on project: PI.

17. University Grants Council (Hong Kong)/General Research Fund (GRF): "Genome-wide association mapping of susceptibility loci for symptomatic epilepsy" (HKU 7623/08M), 1/1/2009–31/12/2010, TDC: HK\$1,570,295. Role on project: PI.
18. University Grants Council (Hong Kong)/General Research Fund (GRF): "Fine mapping of Hirschsprung's disease loci on the 3p21 and 9q31 candidate regions" (HKU 7650/08M), 1/1/2009–30/6/2010, TDC: HK\$566,774, PI: Dr MM Garcia-Barceló. Role on project: Co-I.
19. University Grants Council (Hong Kong)/General Research Fund (GRF): "Genome-wide association study for the identification of genes underlying anorectal malformations" (HKU 7756/08M), 1/1/2009–31/12/2011, TDC: HK\$1,266,337, PI: Prof PKH Tam. Role on project: Co-I.
20. URC/CRCG — Conference Grants for Teaching Staff (HKU): "39th Annual Meeting of the Behavior Genetics Association: *Selection strategies for sib-pair association studies*" (200807170340), 17/06/2009–20/06/2009, TDC: HK\$16,500. Role on project: PI.
21. University Grants Council (Hong Kong)/General Research Fund (GRF): "Identification of functional variants in Neuregulin-1 (NRG1), a newly discovered Hirschsprung's disease gene" (HKU 7656/09M), 01/10/2009–30/09/2011, TDC: HK\$802,995, PI: Dr MM Garcia-Barceló. Role on project: Co-I.
22. URC/CRCG — Conference Grants for Teaching Staff (HKU): "59th Annual Meeting of the American Society of Human Genetics: *Fine mapping of Hirschsprung disease loci in 9q31*" (200807170544), 20/10/2009–24/10/2009, TDC: HK\$16,500. Role on project: PI.
23. Small Project Funding (HKU): "Sequencing of the neuregulin-1 (NRG1) gene in schizophrenia patients" (200907176047), 1/8/2009–31/1/2011, TDC: HK\$40,000. Role on project: Co-I.
24. University Grants Council (Hong Kong)/General Research Fund (GRF): "Genotype by risk factor interactions in cognitive decline: The Guangzhou Biobank Cohort Study" (HKU 7623/09M), 1/1/2010–31/12/2012, TDC: HK\$197,650. Role on project: PI.
25. Seed Funding Programme for Basic Research (HKU): "Exon sequencing of Semaphorin, a novel Hirschsprung's Disease susceptibility locus" (200911159190), 15/3/2010–14/3/2012, TDC: HK\$67,000. Role on project: PI.
26. National Institute of Neurological Disorders and Stroke (NINDS/NIH): "Validation of clinical assessment tools for population genetic studies of epilepsy" (R21NS069223), 1/7/2010–30/6/2012, TDC: US\$152,930, PI: Prof P Kwan. Role on project: Co-PI.
27. University Grants Council (Hong Kong)/Innovation and Technology Support Programme (Tier 2): "Next-Generation High-throughput and Targeted DNA Sequencing Platforms for Systematic Discovery of Novel Diagnostic Markers of Common Disease" (ITS/245/09FP), 1/8/2010–31/1/2012, TDC: HK\$5,856,986, PI: Prof S Lok/Dr AHY Tong. Role on project: Co-I.
28. University Grants Council (Hong Kong)/General Research Fund (GRF): "Deep Re-Sequencing of Hirschsprung's Disease Candidate Genes" (HKU 7686/10M), 1/10/2010–30/9/2013, TDC: HK\$1,000,000, PI: Prof PKH Tam. Role on project: Co-I.
29. URC/CRCG — Conference Grants for Teaching Staff (HKU): "60th Annual Meeting of the American Society of Human Genetics: *A Genome-Wide Association Study of Symptomatic Epilepsy in Han Hong Kong Chinese Detects Multiple Variants*" (201007170081), 2/11/2010–7/11/2010, TDC: HK\$16,500. Role on project: PI.
30. University Grants Council (Hong Kong)/General Research Fund (GRF): "Host genetic determinants of human influenza infection" (HKU 7675/10M), 1/1/2011–30/6/2012, TDC: HK\$598,995, PI: Dr DKM Ip. Role on project: Co-I.
31. University Grants Council (Hong Kong)/General Research Fund (GRF): "Are life long sex-steroids causally related to cardiovascular disease in men and women?" (HKU 7697/10M), 1/1/2011–30/6/2012, TDC: HK\$640,900, PI: Prof TH Lam. Role on project: Co-I.

32. Seed Funding Programme for Basic Research (HKU): "Meta-analysis of Chinese and European genome-wide association studies of Hirschsprung's disease and confirmation of novel loci" (201011159160), 15/3/2011–14/3/2013, TDC: HK\$43,000. Role on project: PI.
33. Seed Funding Programme for Basic Research (HKU): "Investigation of structural variations in Caudal Regression Syndrome Patients" (201010159012), 1/4/2011–31/3/2013, TDC: HK\$98,590. Role on project: Co-I.
34. URC/CRCG — Conference Grants for Teaching Staff (HKU): "12th International Congress of Human Genetics: RET *mutational spectrum in Hirschsprungs disease: evaluation of 601 Chinese patients*" (201007170629), 11/10/2011–15/10/2011, TDC: HK\$16,500. Role on project: PI.
35. University Grants Council (Hong Kong)/General Research Fund (GRF): "Whole-exome sequencing to identify genes underlying Caudal Regression Syndrome" (HKU 7666/11M), 1/12/2011–30/11/2013, TDC: HK\$750,000, PI: Dr MM Garcia-Barceló. Role on project: Co-I.
36. University Grants Council (Hong Kong)/General Research Fund (GRF): "Method and Software for Personal Risk Profiling of Complex Diseases" (HKU 7775/11M), 1/1/2012–31/12/2014, TDC: HK\$945,000, PI: Prof PC Sham. Role on project: Co-I.
37. University Grants Council (Hong Kong)/Theme-Based Research Scheme (TBRS): "Personalized Medicine for Cardiovascular Diseases: From Genomic Testing and Biomarkers to Human Pluripotent Stem Cell Platform" (T12-705/11), 1/1/2012–31/12/2016, TDC: HK\$34,944,000, PC: Prof HK Tse. Role on project: Co-I.
38. URC/CRCG — Conference Grants for Teaching Staff (HKU): "Annual meeting of the American Society of Human Genetics: *Genome-wide copy number variation in anorectal malformations*" (201107170657), 05/11/2012–11/11/2012, TDC: HK\$16,500. Role on project: PI.
39. University Grants Council (Hong Kong)/Theme-Based Research Scheme (TBRS): "Functional analyses of how genomic variation affects personal risk for degenerative skeletal disorders" (T12-708/12-N), 1/1/2013–31/12/2017, TDC: HK\$64,990,000, PC: Prof KSE Cheah. Role on project: Co-I.
40. University Grants Council (Hong Kong)/General Research Fund (GRF): "Exome sequencing of mesial temporal lobe epilepsy with hippocampal sclerosis in parent-offspring trios" (HKU 7630/12M), 1/1/2013–31/12/2014, TDC: HK\$850,000. Role on project: PI.
41. University Grants Council (Hong Kong)/General Research Fund (GRF): "Whole-exome sequencing to uncover causative genes for perinatal biliary atresia" (HKU 7661/12M), 1/1/2013–31/12/2014, TDC: HK\$850,000, PI: Dr MM Garcia-Barceló. Role on project: Co-I.
42. University Grants Council (Hong Kong)/General Research Fund (GRF): "Bioinformatics tools for identifying disease loci from exome sequencing data" (HKU 7764/12M), 1/1/2013–31/12/2015, TDC: HK\$1,098,000, PI: Prof PC Sham. Role on project: Co-I.
43. University Grants Council (Hong Kong)/General Research Fund (GRF): "Study on genomic structural variations in Hirschsprung disease" (HKU 7776/12M), 1/1/2013–31/12/2015, TDC: HK\$1,000,000, PI: Prof PKH Tam. Role on project: Co-I.
44. University Grants Council (Hong Kong)/General Research Fund (GRF): "Congenital dilatation of the bile ducts (CCD): a genetic study" (HKU 7669/13M), 1/9/2013–31/08/2015, TDC: HK\$1,472,255, PI: Dr MM Garcia-Barceló. Role on project: Co-I.
45. University Grants Council (Hong Kong)/General Research Fund (GRF): "Statistical methods for characterizing the genetic component of polygenic diseases" (HKU 7765/13M), 1/1/2014–31/12/2015, TDC: HK\$690,089, PI: Prof PC Sham. Role on project: Co-I.
46. University Grants Council (Hong Kong)/General Research Fund (GRF): "Uncovering the genetic lesions underlying persistent cloaca by whole genome sequencing" (HKU 7782/13M), 1/1/2014–31/12/2016, TDC: HK\$822,731, PI: Prof PKH Tam. Role on project: Co-I.

47. Health and Medical Research Fund - Full Grant (HMRF): "Elucidating the genetic basis for early-age onset nasopharyngeal carcinoma in Hong Kong" (01121496), 1/11/2013–30/4/2015, TDC: HK\$1,000,000, PI: Prof ML Lung. Role on project: Co-I.
48. Health and Medical Research Fund - Mini Grant (HMRF): "A transethnic meta-analysis of genome-wide association studies for Hirschsprung disease" (01121796), 1/1/2014–31/12/2015, TDC: HK\$78,800, PI: Dr SM Tang. Role on project: Co-I.
49. Health and Medical Research Fund - Full Grant (HMRF): "Uncovering the genetic lesions underlying the most severe form of Hirschsprung (HSCR) disease by whole genome sequencing (WGS): a pilot study in 8 family trios" (01121516), 1/4/2014–31/3/2016, TDC: HK\$1,000,000, PI: Dr MM Garcia-Barcelo. Role on project: Co-I.

#### Articles

1. Becker, G. and **Cherny, S. S.** (1992). A five-factor nuclear model of socially desirable responding. *Social Behavior and Personality*, *20*, 163–192; doi:10.2224/sbp.1992.20.3.163.
2. **Cherny, S. S.**<sup>1</sup>, Cardon, L. R., Fulker, D. W., and DeFries, J. C. (1992). Differential heritability across levels of cognitive ability. *Behavior Genetics*, *22*, 153–162; doi:10.1007/BF01066994.
3. **Cherny, S. S.**<sup>1</sup>, DeFries, J. C., and Fulker, D. W. (1992). Multiple regression analysis of twin data: A model-fitting approach. *Behavior Genetics*, *22*, 489–497; doi:10.1007/BF01066617.
4. Benson, J. B., **Cherny, S. S.**, Haith, M. M., and Fulker, D. W. (1993). Rapid assessment of infant predictors of adult IQ: The midtwin/midparent approach. *Developmental Psychology*, *29*, 434–447; doi:10.1037/0012-1649.29.3.434.
5. Rodriguez, L. R., Fulker, D. W., and **Cherny, S. S.** (1993). A maximum-likelihood model-fitting approach to conducting a Hayman analysis of diallel tables with complete or missing data. *Behavior Genetics*, *23*, 69–76; doi:10.1007/BF01067555.
6. Becker, G. and **Cherny, S. S.** (1994). Gender-controlled measures of socially desirable responding. *Journal of Clinical Psychology*, *50*, 746–752; doi:10.1002/1097-4679(199409)50:5<746::AID-JCLP2270500512>3.0.CO;2-V.
7. **Cherny, S. S.**<sup>1</sup>, Fulker, D. W., Corley, R. P., Plomin, R., and DeFries, J. C. (1994) Continuity and change in infant shyness from 14 to 20 months. *Behavior Genetics*, *24*, 365–379; doi:10.1007/BF01067538.
8. **Cherny, S. S.**<sup>1</sup>, Fulker, D. W., Emde, R. N., Robinson, J., Corley, R. P., Reznick, S. J., Plomin, R., and DeFries, J. C. (1994) A developmental-genetic analysis of continuity and change in the Bayley Mental Development Index from 14 to 24 months: The MacArthur Longitudinal Twin Study. *Psychological Science*, *5*, 354–360; doi:10.1111/j.1467-9280.1994.tb00285.x.
9. Schmitz, S., **Cherny, S. S.**, Fulker, D. W., and Mrazek, D. (1994) Genetic and environmental influences on early childhood behavior. *Behavior Genetics*, *24*, 25–34; doi:10.1007/BF01067926.
10. Cardon, L. R., Fulker, D. W., and **Cherny, S. S.** (1995) Linkage analysis of a common oligogenic disease using selected sib pairs. *Genetic Epidemiology*, *12*, 741–746; doi:10.1002/gepi.1370120635.
11. Fulker, D. W. and **Cherny, S. S.** (1995) Genetic and environmental influences on cognition during childhood. *Population Research and Policy Review*, *14*, 283–300; doi:10.1007/BF01074393.
12. Fulker, D. W., **Cherny, S. S.**, and Cardon, L. R. (1995) Multipoint interval mapping of quantitative trait loci, using sib pairs. *American Journal of Human Genetics*, *56*, 1224–1233; PMID:7726180.
13. Hu, S., Pattatucci, A. M. L., Patterson, C., Li, L., Fulker, D. W., **Cherny, S. S.**, Kruglyak, L., and Hamer, D. H. (1995) Linkage between Xq28 and sexual orientation in males but not in females. *Nature Genetics*, *11*, 248–256; doi:10.1038/ng1195-248.

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14. Fulker, D. W. and **Cherny, S. S.** (1996) An improved multipoint sib-pair analysis of quantitative traits. *Behavior Genetics*, *26*, 527–532; doi:10.1007/BF02359758.
15. Petrill, S. A., Saudino, K., **Cherny, S. S.**, Emde, R. N., Hewitt, J. K., Fulker, D. W., and Plomin, R. (1997) Exploring the genetic etiology of low general cognitive ability from 14 to 36 months. *Developmental Psychology*, *33*, 544–548; doi:10.1037/0012-1649.33.3.544.
16. Stallings, M. C., **Cherny, S. S.**, Young, S. E., Miles, D. R., Hewitt, J. K., and Fulker, D. W. (1997) The familial aggregation of depressive symptoms, antisocial behavior, and alcohol abuse. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, *74*, 183–191; doi:10.1002/(SICI)1096-8628(19970418)74:2<183::AID-AJMG14>3.0.CO;2-E.
17. Petrill, S. A., Saudino, K., **Cherny, S. S.**, Emde, R. N., Fulker, D. W., Hewitt, J. K., and Plomin, R. (1998) Exploring the genetic and environmental etiology of high general cognitive ability in fourteen- to thirty-six-month-old twins. *Child Development*, *69*, 68–74; doi:10.1111/j.1467-8624.1998.tb06133.x.
18. Schmitz, S., **Cherny, S. S.**, and Fulker, D. W. (1998) Increase in power through multivariate analyses. *Behavior Genetics*, *28*, 357–363; doi:10.1023/A:1021669602220.
19. Fulker, D. W., **Cherny, S. S.**<sup>1</sup>, Sham, P. C., and Hewitt, J. K. (1999) Combined linkage and association sib-pair analysis for quantitative traits. *American Journal of Human Genetics*, *64*, 259–267; doi:10.1086/302193.
20. Gayán, J., Smith, S. D., **Cherny, S. S.**, Cardon, L. R., Fulker, D. W., Brower, A. M., Olson, R. K., Pennington, B. F., and DeFries, J. C. (1999) Quantitative trait locus for specific language and reading deficits on chromosome 6p. *American Journal of Human Genetics*, *64*, 157–164; doi:10.1086/302191.
21. Neale, M. C., **Cherny, S. S.**, Sham, P. C., Whitfield, J. B., Heath, A. C., Birley, A. C., and Martin, N. G. (1999) Distinguishing population stratification from genuine allelic effects with Mx: Association of ADH2 with alcohol consumption. *Behavior Genetics*, *29*, 233–244; doi:10.1023/A:1021638122693.
22. Talbot, C. J., Nicod, A., **Cherny, S. S.**, Fulker, D. W., Collins, A. C., and Flint, J. (1999) High-resolution mapping of quantitative trait loci in outbred mice. *Nature Genetics*, *21*, 305–308; doi:10.1038/6825.
23. Saudino, K. J., **Cherny, S. S.**, and Plomin, R. (2000) Parent ratings of temperament in twins: Explaining the 'too low' DZ correlations. *Twin Research*, *3*, 224–233; doi:10.1375/136905200320565193.
24. Sham, P. C., **Cherny, S. S.**, Purcell, S., and Hewitt, J. K. (2000) Power of linkage versus association analysis of quantitative traits, by use of variance-components models, for sibship data. *American Journal of Human Genetics*, *66*, 1616–1630; doi:10.1086/302891.
25. Sham, P. C., Sterne, A., Purcell, S., **Cherny, S.**, Webster, M., Rijdsdijk, F., Asherson, P., Ball, D., Craig, I., Eley, T., Goldberg, D., Gray, J., Mann, A., Owen, M., and Plomin, R. (2000) GENESIS: creating a composite index of the vulnerability to anxiety and depression in a community-based sample of siblings. *Twin Research*, *3*, 316–322; doi:10.1375/136905200320565292.
26. Sham, P. C., Zhao, J. H., **Cherny, S. S.**, and Hewitt, J. K. (2000) Variance-components QTL linkage analysis of selected and non-normal samples: conditioning on trait values. *Genetic Epidemiology*, *19*, S22–S28; doi:10.1002/1098-2272(2000)19:1+<::AID-GEPI4>3.0.CO;2-S.
27. Abecasis, G. R., Cardon, L. R., Cookson, W. O. C., Sham, P. C., and **Cherny, S. S.**<sup>1</sup> (2001) Association analysis in a variance components framework. *Genetic Epidemiology*, *21*, S341–S346; <http://www.sph.umich.edu/csg/abecasis/publications/pdf/Genet.Epidemiol.vol.21-pp.S341-S346.pdf>.
28. Abecasis, G. R., **Cherny, S. S.**, and Cardon, L. R. (2001) The impact of genotyping error on family-based analysis of quantitative traits. *European Journal of Human Genetics*, *9*, 130–134; doi:10.1038/sj.ejhg.5200594.
29. Abecasis, G. R., **Cherny, S. S.**, Cookson, W. O. C., and Cardon, L. R. (2001) GRR: Graphical representation of relationship errors. *Bioinformatics*, *17*, 742–743; doi:10.1093/bioinformatics/17.8.742.

30. Beekman, M., Lakenberg, N., **Cherny, S. S.**, de Knijff, P., Vogler, G. P., Frants, R. R., Boomsma, D. I., and Slagboom, P. E. (2001) A powerful and rapid approach to human genome scanning using small quantities of genomic DNA. *Genetical Research*, *77*, 129–134; doi:10.1017/S001667230100492X.
31. **Cherny, S. S.**<sup>1</sup>, Abecasis, G. R., Cookson, W. O. C., Sham, P. C., and Cardon, L. R. (2001) The effect of genotype and pedigree error on linkage analysis: Analysis of three asthma genome scans. *Genetic Epidemiology*, *21*, S117-S122; <http://www.sph.umich.edu/csg/abecasis/publications/pdf/Genet.Epidemiol.vol.21-pp.S117-S122.pdf>.
32. Lessem, J. M. and **Cherny, S. S.**<sup>1</sup> (2001) DeFries-Fulker multiple regression of sibship QTL data: a SAS<sup>®</sup> macro. *Bioinformatics*, *17*, 371–372; doi:10.1093/bioinformatics/17.4.371.
33. Purcell, S., **Cherny, S. S.**, Hewitt, J. K., and Sham, P. C. (2001) Optimal sibship selection for genotyping in quantitative trait locus linkage analysis. *Human Heredity*, *52*, 1–13; doi:10.1159/000053350.
34. Rijdsdijk, F. V., Sham, P. C., Sterne, A., Purcell, S., McGuffin, P., Farmer, A., Goldberg, D., Mann, A., **Cherny, S. S.**, Webster, M., Ball, D., Eley, T. C., and Plomin, R. (2001) Life events and depression in a community sample of siblings. *Psychological Medicine*, *31*, 401–410; doi:10.1017/S0033291701003361.
35. Abecasis, G. R., **Cherny, S. S.**, Cookson, W. O., and Cardon, L. R. (2002) MERLIN — rapid analysis of dense genetic maps using sparse gene flow trees. *Nature Genetics*, *30*, 97–101; doi:10.1038/ng786.
36. Jawaid, A., Bader, J. S., Purcell, S., **Cherny, S. S.**, and Sham, P. (2002) Optimal selection strategies for QTL mapping using pooled DNA samples. *European Journal of Human Genetics*, *10*, 125–132; doi:10.1038/sj.ejhg.5200771.
37. Sham, P. C., Purcell, S., **Cherny, S. S.**, and Abecasis, G. R. (2002) Powerful regression-based quantitative-trait linkage analysis of general pedigrees. *American Journal of Human Genetics*, *71*, 238–253; doi:10.1086/341560.
38. Bishop, E. G., **Cherny, S. S.**, Corley, R., Plomin, R., DeFries, J. C., and Hewitt, J. K. (2003) Developmental genetic analysis of general cognitive ability from 1 to 12 years in a sample of adoptees, biological siblings, and twins. *Intelligence*, *31*, 31–49; doi:10.1016/S0160-2896(02)00112-5.
39. Cader, Z. M., Noble-Topham, S., Dyment, D. D., **Cherny, S. S.**, Brown, J. D., Rice, G. P. A., and Ebers, G. C. (2003) Significant linkage to migraine with aura on chromosome 11q24. *Human Molecular Genetics*, *12*, 2511–2517; doi:10.1093/hmg/ddg252.
40. Marlow, A. J., Fisher, S. E., Francks, C., MacPhie, I. L., **Cherny, S. S.**, Richardson, A. J., Talcott, J. B., Stein, J. F., Monaco, A. P., and Cardon, L. R. (2003) Use of multivariate linkage analysis for dissection of a complex cognitive trait. *American Journal of Human Genetics*, *72*, 561–570; doi:10.1086/368201.
41. Purcell, S., **Cherny, S. S.**, and Sham, P. C. (2003) Genetic Power Calculator: design of linkage and association genetic mapping studies of complex traits. *Bioinformatics*, *19*, 149–150; doi:10.1093/bioinformatics/19.1.149.
42. **Cherny, S. S.**<sup>1</sup>, Sham, P. C., and Cardon, L. R. (2004) Introduction to the Special Issue on Variance Components Methods for Mapping Quantitative Trait Loci. *Behavior Genetics*, *34*, 125–126; doi:10.1023/B:BEGE.0000013938.54772.42.
43. Nash, M. W., Huezo-Diaz, P., Williamson, R. J., Sterne, A., Purcell, S., Hoda, F., **Cherny, S. S.**, Abecasis, G. R., Prince, M., Gray, J. A., Ball, D., Asherson, P., Mann, A., Goldberg, D., McGuffin, P., Farmer, A., Plomin, R., Craig, I. W., and Sham, P. C. (2004) Genome-wide linkage analysis of a composite index of neuroticism and mood-related scales in extreme selected sibships. *Human Molecular Genetics*, *13*, 2173–2182; doi:10.1093/hmg/ddh239.
44. Petrill, S. A., Lipton, P. A., Hewitt, J. K., Plomin, R., **Cherny, S. S.**, Corley, R., and DeFries, J. C. (2004) Genetic and environmental contributions to general cognitive ability through the first 16 years of life. *Developmental Psychology*, *40*, 805–812; doi:10.1037/0012-1649.40.5.805.



45. The U.S.-Venezuela Collaborative Research Project<sup>2</sup> and Wexler, N. S. (2004) Venezuelan kindreds reveal that genetic and environmental factors modulate Huntington's disease age of onset. *Proceedings of the National Academy of Sciences of the United States of America*, *101*, 3498–3503; doi:10.1073/pnas.0308679101.
46. Posthuma, D., **Cherny, S. S.**, and Boomsma, D. I. (2006) Introduction to the Special Issue: Human linkage studies for behavioral traits. *Behavior Genetics*, *36*, 1–3; doi:10.1007/s10519-005-9010-2.
47. Andresen, J. M., Gayán, J., **Cherny, S. S.**, Brocklebank, D., Alkorta-Aranburu, G., Addis, E. A., The US-Venezuela Collaborative Research Group<sup>2</sup>, Cardon, L. R., Housman, D. E., and Wexler, N. S. (2007) Replication of twelve association studies for Huntington's disease residual age of onset in large Venezuelan kindreds. *Journal of Medical Genetics*, *44*, 44–50; doi:10.1136/jmg.2006.045153.
48. Andresen, J. M., Gayán, J., Djoussé, L., Roberts, S., Brocklebank, D., **Cherny, S. S.**, The US-Venezuela Collaborative Research Group<sup>2</sup>, The HD MAPS Collaborative Research Group<sup>3</sup>, Cardon, L. R., Gusella, J. F., MacDonald, M. E., Myers, R. H., Housman, D. E., and Wexler, N. S. (2007) The relationship between CAG repeat length and age of onset differs for Huntington's Disease patients with juvenile onset or adult onset. *Annals of Human Genetics*, *71*, 295–301; doi:10.1111/j.1469-1809.2006.00335.x.
49. Goode, E. L., **Cherny, S. S.**, Christian, J. C., Jarvik, G. P., and de Andrade, M. (2007) Heritability of longitudinal measures of body mass index and lipid and lipoprotein levels in aging twins. *Twin Research and Human Genetics*, *10*, 703–711; doi:10.1375/twin.10.5.703.
50. Hopper, C. J., Lessem, J. M., Hartman, C. A., Stallings, M. C., **Cherny, S. S.**, Corley, R. P., Hewitt, J. K., Krauter, K. S., Mikulich-Gilbertson, S. K., Rhee, S. H., Smolen, A., Young, S. E., and Crowley, T. J. (2007) A genome-wide scan for loci influencing adolescent cannabis dependence symptoms: Evidence for linkage on chromosomes 3 and 9. *Drug and Alcohol Dependence*, *89*, 34–41; doi:10.1016/j.drugalcdep.2006.11.015.
51. Willer, C. J., Dymont, D. A., **Cherny, S.**, Ramagopalan, S. V., Herrera, B. M., Morrison, K. M. E., Sadovnick, A. D., Risch, N. J., and Ebers, G. C. (2007) A genome-wide scan in forty large pedigrees with multiple sclerosis. *Journal of Human Genetics*, *52*, 955–962; doi:10.1007/s10038-007-0194-6.
52. **Cherny, S. S.** (2008) Variance components and related methods for mapping quantitative trait loci. *Sociological Methods and Research*, *37*, 227–250; doi:10.1177/0049124108324524.
53. Dymont, D. A., Cader, M. Z., Datta, A., Broxholme, S. J., **Cherny, S. S.**, Willer, C. J., Ramagopalan, S., Herrera, B. M., Orton, S., Chao, M., Sadovnick, A. D., Hader, M., Hader, W., and Ebers, G. C. (2008) A first stage genome-wide screen for regions shared identical-by-descent in Hutterite families with multiple sclerosis. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, *147B*, 467–472; doi:10.1002/ajmg.b.30620.
54. Garcia-Barceló, M. M., Fong, P. Y., Tang, C. S., Miao, X. P., So, M. T., Yuan, Z. W., Li, L., Guo, W. H., Liu, L., Wang, B., Sun, X. B., Huang, L. M., Tou, J. F., Wong, K. K. Y., Ngan, E. S. W., Lui, V. C. H., **Cherny, S. S.**, Sham, P. C., and Tam, P. K. H. (2008) Mapping of a Hirschsprung's disease locus in 3p21. *European Journal of Human Genetics*, *16*, 833–840; doi:10.1038/ejhg.2008.18.

<sup>2</sup> N S Wexler, J Lorimer, J Porter, F Gomez, C Moskowitz, E Shackell, K Marder, G Penchaszadeh, S A Roberts, J Gayán, D Brocklebank, **S S Cherny**, L R Cardon, J Gray, S R Dlouhy, S Wiktorski, M E Hodes, P M Conneally, J B Penney, J Gusella, J-H Cha, M Irizarry, D Rosas, S Hersch, Z Hollingsworth, M MacDonald, A B Young, J M Andresen, D E Housman, M Mieja de Young, E Bonilla, T Stillings, A Negrette, S R Snodgrass, M D Martinez-Jaurrieta, M A Ramos-Arroyo, J Bickham, J S Ramos, F Marshall, I Shoulson, G J Rey, A Feigin, M Arnheim, A Acevedo-Cruz, L Acosta, J Alvir, K Fischbeck, L M Thompson, A Young, L Dure, C J O'Brien, J Paulsen, A Brickman, D Krch, S Peery, P Hogarth, D S Higgins, Jr., and B Landwehrmeyer

<sup>3</sup> M R Hayden, E W Almquist, R R Brinkman, O Suchowersky, A Durr, C Dodé, F Squitieri, P J Morrison, M Nance, C A Ross, R L Margolis, A Rosenblatt, E Gómez-Tortosa, D M Cabrero, R J A Trent, E McCusker, A Novelletto, M Frontali, J S Paulsen, R Jones, A Zanko, T Ashizawa, A Lazzarini, J-L Li, V C Wheeler, A L Russ, G Xu, J S Mysore, T Gillis, M Hakky, L A Cupples, M Saint-Hilaire, J-H J Cha, S M Hersch, J B Penney, M Harrison, K Marder, R K Abramson, P M Conneally, J F Gusella, M E MacDonald, R H Myers

55. Garcia-Barceló, M. M., Lui, V. C. H., Miao, X., So, M. T., Leon, T. Y. Y., Yuan, Z. W., Li, L., Liu, L., Wang, B., Sun, X. B., Huang, L. M., Tou, J. F., Ngan, E. S. W., **Cherny, S. S.**, Chan, K. W., Lee, K. H., Wang, W., Wong, K. K. Y., and Tam, P. K. H. (2008) Mutational analysis of *SHH* and *GLI3* in anorectal malformations. *Birth Defects Research (Part A): Clinical and Molecular Teratology*, *82*, 644–648; doi:10.1002/bdra.20482.
56. Gayán, J., Brocklebank, D., Andresen, J. M., Alkorta-Aranburu, G., The US-Venezuela Collaborative Research Group<sup>2</sup>, Cader, M. Z., Roberts, S. A., **Cherny, S. S.**, Wexler, N. S., Cardon, L. R., and Housman, D. E. (2008) Genomewide linkage scan reveals novel loci modifying age of onset of Huntington's disease in the Venezuelan HD kindreds. *Genetic Epidemiology*, *32*, 445–453; doi:10.1002/gepi.20317.
57. Hur, Y.-M., Kaprio, J., Iacono, W. G., Boomsma, D. I., McGue, M., Silventoinen, K., Martin, N. G., Luciano, M., Visscher, P. M., Rose, R. J., He, M., Ando, J., Ooki, S., Nonaka, K., Lin, C. C. H., Lajunen, H. R., Cornes, B. K., Bartels, M., van Beijsterveldt, C. E., **Cherny, S. S.**, and Mitchell, K. (2008) Genetic influences on the difference in variability of height, weight, and body mass index between Caucasian and East Asian adolescent twins. *International Journal of Obesity*, *32*, 1455–1467; doi:10.1038/ijo.2008.144.
58. Ong, K. L., Leung, R. Y. H., Wong, L. Y. F., **Cherny, S. S.**, Sham, P. C., Lam, T. H., Lam, K. S. L., and Cheung, B. M. Y. (2008) Association of a polymorphism in the lipin 1 gene with systolic blood pressure in men. *American Journal of Hypertension*, *21*, 539–545; doi:10.1038/ajh.2008.21.
59. Ong, K. L., Leung, R. Y. H., Wong, L. Y. F., **Cherny, S. S.**, Sham, P. C., Lam, T. H., Lam, K. S. L., and Cheung, B. M. Y. (2008) Association of F11 receptor gene polymorphisms with central obesity and blood pressure. *Journal of Internal Medicine*, *263*, 322–332; doi:10.1111/j.1365-2796.2007.01886.x.
60. Sham, P. C., **Cherny, S. S.**, Kao, P. Y. P., Song, Y.-Q., Chan, D., and Cheung, K. M. C. (2008) (iii) Whole-genome association studies of complex diseases. *Current Orthopaedics*, *22*, 251–258; doi:10.1016/j.cuor.2008.05.006.
61. Yeung, J. M. Y., Sham, P. C., Chan, A. S. W., and **Cherny, S. S.**<sup>1</sup> (2008) OpenADAM: an open source genome-wide association data management system for Affymetrix SNP arrays. *BMC Genomics*, *9*, 636; doi:10.1186/1471-2164-9-636.
62. Brocklebank, D., Gayán, J., Andresen, J. M., Roberts, S. A., The International-Venezuela Collaborative Research Group<sup>2</sup>, Young, A. B., Snodgrass, S. R., Penney, J. B., Ramos-Arroyo, M. A., Cha, J. J., Rosas, H. D., Hersch, S. M., Feigin, A., **Cherny, S. S.**, Wexler, N. S., Housman, D. E., and Cardon, L. R. (2009) Repeat instability in the 27-39 CAG range of the HD gene in the Venezuelan kindreds: Counseling implications. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, *150B(3)*, 425–429; doi:10.1002/ajmg.b.30826.
63. Cheung, B. M. Y., Ong, K. L., **Cherny, S. S.**, Sham, P. C., Tso, A. W. K., and Lam, K. S. L. (2009) Diabetes prevalence and therapeutic target achievement in the United States, 1999–2006. *The American Journal of Medicine*, *122(5)*, 443–453; doi:10.1016/j.amjmed.2008.09.047.
64. Garcia-Barceló, M. M., Tang, C. S. M., Ngan, E. S. W., Lui, V. C. H., Chen, Y., So, M. T., Leon, T. Y. Y., Miao, X. P., Shum, C. K. Y., Liu, F. Q., Yeung, M. Y., Yuan, Z. W., Guo, W. H., Liu, L., Sun, X. B., Huang, L. M., Tou, J. F., Song, Y. Q., Chan, D., Cheung, K. M. C., Wong, K. K. Y., **Cherny, S. S.**<sup>1</sup>, Sham, P. C.<sup>1</sup>, and Tam, P. K. H.<sup>1</sup> (2009) Genome-wide association study identifies *NRG1* as a susceptibility locus for Hirschsprung's disease. *Proceedings of the National Academy of Sciences of the United States of America*, *106(8)*, 2694–2699; doi:10.1073/pnas.0809630105.
65. Kwan, J. S. H., **Cherny, S. S.**, Kung, A. W. C., and Sham, P. C. (2009) Novel sib pair selection strategy increases power in quantitative association analysis. *Behavior Genetics*, *39(5)*, 571–579; doi:10.1007/s10519-009-9284-x.
66. Ngan, E. S. W., Lang, B. H. H., Liu, T., Shum, C. K. Y., So, M.-T., Lau, D. K. C., Leon, T. Y. Y., **Cherny, S. S.**, Tsai, S. Y., Lo, C.-Y., Khoo, U.-S., Tam, P. K. H., and Garcia-Barceló, M. M. (2009) A

- germline mutation (A339V) in Thyroid Transcription Factor-1 (TTF-1/NKX2.1) in patients with multinodular goiter and papillary thyroid carcinoma. *Journal of the National Cancer Institute*, 101(3), 162–175; doi:10.1093/jnci/djn471.
67. Ong, K. L., Leung, R. Y. H., Babinska, A., Salifu, M. O., Ehrlich, Y. H., Kordecki, E., Wong, L. Y. F., Tso, A. W. K., **Cherny, S. S.**, Sham, P. C., Lam, T. H., Lam, K. S. L., and Cheung, B. M. Y. (2009) Elevated plasma level of soluble F11 receptor / junctional adhesion molecule-A (F11R/JAM-A) in hypertension. *American Journal of Hypertension*, 22(5), 500–505; doi:10.1038/ajh.2009.23.
68. Sham, P. C., **Cherny, S. S.**, and Purcell, S. (2009) Application of genome-wide SNP data for uncovering pairwise relationships and quantitative trait loci. *Genetica*, 136(2), 237–243; doi:10.1007/s10709-008-9349-4.
69. Cornes, B. K., Tang, C. S., Leon, T. Y. Y., Hui, K. J. W. S., So, M.-T., Miao, X., **Cherny, S. S.**, Sham, P. C., Tam, P. K. H., and Garcia-Barceló, M. M. (2010) Haplotype analysis reveals a possible founder effect of RET mutation R114H for Hirschsprung's disease in the Chinese population. *PLoS ONE*, 5(6), e10918; doi:10.1371/journal.pone.0010918.
70. Garcia-Barceló, M. M., Yeung, M. Y., Miao, X. P., Tang, C. S. M., Cheng, G., So, M. T., Ngan, E. S. W., Lui, V. C. H., Chen, Y., Liu, X. L., Hui, K. J. W. S., Li, L., Guo, W. H., Sun, X. B., Tou, J. F., Chan, K. W., Wu, X. Z., Song, Y. Q., Chan, D., Cheung, K., Chung, P. H. Y., Wong, K. K. Y., Sham, P. C., **Cherny, S. S.**<sup>1</sup>, and Tam, P. K. H.<sup>1</sup> (2010) Genome-wide association study identifies a susceptibility locus for biliary atresia on 10q24.2. *Human Molecular Genetics*, 19(14), 2917–2925 ; doi:10.1093/hmg/ddq196.
71. Kung, A. W. C., Xiao, S.-M., **Cherny, S.**, Li, G. H. Y., Gao, Y., Tso, G., Lau, K. S., Luk, K. D. K., Liu, J.-M., Cui, B., Zhang, M.-J., Zhang, Z.-L., He, J.-W., Yue, H., Xia, W.-B., Luo, L.-M., He, S.-L., Kiel, D. P., Karasik, D., Hsu, Y.-H., Cupples, L. A., Demissie, S., Stykarsdottir, U., Halldorsson, B. V., Sigurdsson, G., Thorsteinsdottir, U., Stefansson, K., Richards, J. B., Zhai, G., Soranzo, N., Valdes, A., Spector, T. D., and Sham, P. C. (2010) Association of *JAG1* with bone mineral density and osteoporotic fractures: a genome-wide association study and follow-up replication studies. *American Journal of Human Genetics*, 86, 229–239; doi:10.1016/j.ajhg.2009.12.014.
72. Li, M.-X., Sham, P. C., **Cherny, S. S.**, and Song, Y.-Q. (2010) A knowledge-based weighting framework to boost the power of genome-wide association studies. *PLoS ONE*, 5(12), e14480; doi:10.1371/journal.pone.0014480.
73. Ong, K. L., Li, M., Tso, A. W. K., Xu, A., **Cherny, S. S.**, Sham, P. C., Tse, H. T., Lam, T. H., Cheung, B. M. Y., Lam, K. S. L., and the Investigators of the Hong Kong Cardiovascular Risk Factor Prevalence Study (2010) Association of genetic variants in the adiponectin gene with adiponectin level and hypertension in Hong Kong Chinese. *European Journal of Endocrinology*, 163, 251–257; doi:10.1530/EJE-10-0251.
74. Ong, K. L., Tso, A. W. K., **Cherny, S. S.**, Sham, P. C., Lam, K. S. L., Jiang, C. Q., Thomas, G. N., Lam, T. H., and Cheung, B. M. Y. (2010) A genetic variant in the gene encoding fibrinogen beta chain predicted development of hypertension in Chinese men. *Thrombosis and Haemostasis*, 103(4), 728–735; doi:10.1160/TH09-10-0692.
75. Ong, K. L., Tso, A. W. K., Lam, K. S. L., **Cherny, S. S.**, Sham, P. C., and Cheung, B. M. Y. (2010) Using glycosylated hemoglobin to define the metabolic syndrome in United States adults. *Diabetes Care*, 33(8), 1856–1858; doi:10.2337/dc10-0190.
76. So, H.-C., Fong, P. Y., Chen, R. Y. L., Hui, T. C. K., Ng, M. Y. M., **Cherny, S. S.**, Mak, W. W. M., Cheung, E. F. C., Chan, R. C. K., Chen, E. Y. H., Li, T., and Sham, P. C. (2010) Identification of neuroglycan C and interacting partners as potential susceptibility genes for schizophrenia in a Southern Chinese population. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, 153B, 103–113; doi:10.1002/ajmg.b.30961.

77. Styrkarsdottir, U., Halldorsson, B. V., Gudbjartsson, D. F., Tang, N. L. S., Koh, J.-M., Xiao, S.-M., Kwok, T. C. Y., Kim, G. S., Chan, J. C. N., **Cherny, S.**, Lee, S. H., Kwok, A., Ho, S., Gretarsdottir, S., Kostic, J. P., Palsson, S. Th., Sigurdsson, G., Sham, P. C., Kim, B.-J., Kung, A. W. C., Kim, S.-Y., Woo, J., Leung, P.-C., Kong, A., Thorsteinsdottir, U., and Stefansson, K. (2010) European bone mineral density loci are also associated with BMD in East-Asian populations. *PLoS One*, *5*(10), e13217; doi:10.1371/journal.pone.0013217.
78. Tang, C. S., Sribudiani, Y., Miao, X. P., de Vries, A. R., Burzynski, G., So, M. T., Leon, Y. Y., Yip, B. H., Osinga, J., Hui, K. J., Verheij, J. B., **Cherny, S. S.**, Tam, P. K., Sham, P. C., Hofstra, R. M., and Garcia-Barceló, M. M. (2010) Fine mapping of the 9q31 Hirschsprung's disease locus. *Human Genetics*, *127*(6), 675–683; doi:10.1007/s00439-010-0813-8.
79. Toulopoulou, T., Goldberg, T. E., Mesa, I. R., Picchioni, M., Rijdsdijk, F., Stahl, D., **Cherny, S. S.**, Sham, P., Faraone, S. V., Tsuang, M., Weinberger, D. R., Seidman, L. J., and Murray, R. M. (2010) Impaired intellect and memory: A missing link between genetic risk and schizophrenia? *Archives of General Psychiatry*, *67*(9), 905–913; doi:10.1001/archgenpsychiatry.2010.99.
80. Cheung, B. M. Y., Ong, K. L., Tso, A. W. K., **Cherny, S. S.**, Sham, P. C., Lam, T. H., and Lam, K. S. L. (2011) Gamma-glutamyl transferase level predicts the development of hypertension in Hong Kong Chinese. *Clinica Chimica Acta*, *412*, 1326–1331; doi:10.1016/j.cca.2011.03.030.
81. Cheung, B. M. Y., Ong, K. L., Tso, A. W. K., Leung, R. Y. H., Xu, A., **Cherny, S. S.**, Sham, P. C., Lam, T. H., and Lam, K. S. L. on behalf of the Investigators of the Hong Kong Cardiovascular Risk Factor Prevalence Study (2011) Plasma adrenomedullin level is related to a single nucleotide polymorphism in the adrenomedullin gene. *European Journal of Endocrinology*, *165*, 571–577; doi:10.1530/EJE-11-0513.
82. Cheung, B. M. Y., Ong, K. L., Tso, A. W. K., Leung, R. Y. H., **Cherny, S. S.**, Sham, P. C., Thomas, G. N., Lam, T. H., and Lam, K. S. L. on behalf of the Investigators of the Hong Kong Cardiovascular Risk Factor Prevalence Study (2011) Relationship of plasma interleukin-6 and its genetic variants with hypertension in Hong Kong Chinese. *American Journal of Hypertension*, *24*(12), 1331–1337; doi:10.1038/ajh.2011.141.
83. Garcia-Barceló, M. M., Miao, X., Tang, C. S., So, H. C., Tang, W., Leon, T. Y., So, M., Yip, B., Chen, R. Y., Cheung, E. F., Chen, E. Y., Li, T., Tam, P., **Cherny, S. S.**, and Sham, P. C. (2011) No NRG1 V266L mutations in Chinese schizophrenic patients. *Psychiatric Genetics*, *21*(1), 47–49; doi:10.1097/YPG.0b013e328341355b.
84. Gui, H., Li, M., Sham, P. C., and **Cherny, S. S.**<sup>1</sup> (2011) Comparisons of seven algorithms for pathway analysis using the WTCCC Crohn's Disease Dataset. *BMC Research Notes*, *4*, 386; doi:10.1186/1756-0500-4-386.
85. Ho, J. W., Choi, S.-C., Lee, Y.-F., Hui, T. C., **Cherny, S. S.**, Garcia-Barceló, M.-M., Carvajal-Carmona, L., Liu, R., To, S.-H., Yau, T.-K., Chung, C. C., Yau, C. C., Hui, S. M., Lau, P. Y., Yuen, C.-H., Wong, Y. W., Ho, S., Fung, S. S., Tomlinson, I. P., Houlston, R. S., Cheng, K. K., and Sham, P. C. (2011) Replication study of SNP associations for colorectal cancer in Hong Kong Chinese. *British Journal of Cancer*, *104*, 369–375; doi:10.1038/sj.bjc.6605977.
86. Ngan, E. S.-W., Garcia-Barceló, M. M., Yip, B. H.-K., Poon, H.-C., Lau, S.-T., Kwok, C. K.-M., Sat, E., Sham, M.-H., Wong, K. K.-Y., Wainwright, B. J., **Cherny, S. S.**, Hui, C.-C., and Sham, P. C., Lui, V. C.-H., and Tam, P. K.-H. (2011) Hedgehog/Notch-induced premature gliogenesis represents a new disease mechanism for Hirschsprung disease in mice and humans. *The Journal of Clinical Investigation*, *121*(9), 3467–3478; doi:10.1172/JCI43737.
87. Ong, K. L., Tso, A. W. K., **Cherny, S. S.**, Sham, P. C., Lam, T. H., Lam, K. S. L., and Cheung, B. M. Y. on behalf of the Investigators of the Hong Kong Cardiovascular Risk Factor Prevalence Study (2011) Role of genetic variants in the gene encoding lipocalin-2 in the development of elevated blood pressure. *LCEH: Clinical and Experimental Hypertension*, *33*(7), 484–491; doi:10.3109/10641963.2010.549276.

88. Ong, K. L., Tso, A. W. K., Leung, R. Y. H., **Cherny, S. S.**, Sham, P. C., Lam, T. H., Cheung, B. M. Y., and Lam, K. S. L. (2011) A genetic variant in the gene encoding adrenomedullin predicts the development of dysglycemia over 6.4 years in Chinese. *Clinica Chimica Acta*, *412*, 353–357; doi:10.1016/j.cca.2010.11.007.
89. Sham, P. C., **Cherny, S. S.**, and Hall, M.-H. (2011) Statistical issues and approaches in endophenotype research. *Chinese Science Bulletin*, *56*, 3403–3408; doi:10.1007/s11434-011-4746-y.
90. So, H. C., Gui, A. H., **Cherny, S. S.**, and Sham, P. C. (2011) Evaluating the heritability explained by known susceptibility variants: a survey of ten complex diseases. *Genetic Epidemiology*, *35*, 310–317; doi:10.1002/gepi.20579.
91. So, M.-T., Leon, T. Y.-Y., Cheng, G., Tang, C. S.-M., Miao, X.-P., Cornes, B. K., Ngo, D. N., Cui, L., Ngan, E. S.-W., Lui, V. C.-H., Wu, X.-Z., Wang, B., Wang, H., Yuan, Z.-W., Huang, L.-M., Li, L., Xia, H., Zhu, D., Liu, J., Nguyen, T. L., Chan, I. H.-Y., Chung, P. H.-Y., Liu, X.-L., Zhang, R., Wong, K. K.-Y., Sham, P.-C., **Cherny, S. S.**, Tam, P. K.-H., Garcia-Barcelo, M.-M. (2011) *RET* mutational spectrum in Hirschsprung disease: evaluation of 601 Chinese patients. *PLoS One*, *6*(12), e28986; doi:10.1371/journal.pone.0028986.
92. So, H. C., Kwan, J. S. H., **Cherny, S. S.**, and Sham, P. C. (2011) Risk prediction of complex diseases from family history and known susceptibility loci, with applications for cancer screening. *American Journal of Human Genetics*, *88*(5), 548–565; doi:10.1016/j.ajhg.2011.04.001.
93. Tang, C. S.-M., Tang, W.-K., So, M.-T., Miao, X.-P., Leung, B. M.-C., Yip, B. H.-K., Leon, T. Y.-Y., Ngan, E. S.-W., Lui, V. C.-H., Chen, Y., Chan, I. H.-Y., Chung, P. H.-Y., Liu, X.-L., Wu, X.-Z., Wong, K. K.-Y., Sham, P.-C., **Cherny, S. S.**<sup>1</sup>, Tam, P. K.-H., and Garcia-Barceló, M.-M.<sup>1</sup> (2011) Fine mapping of the *NRG1* Hirschsprung's disease locus. *PLoS ONE*, *6*(1), e16181; doi:10.1371/journal.pone.0016181.
94. Yang, J., Yang, W., Hirankarn, N., Ye, D. Q., Zhang, Y., Pan, H. F., Mok, C. C., Chan, T. M., Wong, R. W., Mok, M. Y., Lee, K. W., Wong, S. N., Leung, A. M., Li, X. P., Avihingsanon, Y., Rianthavorn, P., Deekajorndej, T., Suphapeetiporn, K., Shotlersuk, V., Baum, L., Kwan, P., Lee, T. L., Ho, M. H., Lee, P. P., Wong, W. H., Zeng, S., Zhang, J., Wong, C. M., Ng, I. O., Garcia-Barceló, M. M., **Cherny, S. S.**, Tam, P. K., Sham, P. C., Lau, C. S., and Lau, Y. L. (2011) *ELF1* is associated with systemic lupus erythematosus in Asian populations. *Human Molecular Genetics*, *20*(3), 601–607; doi:10.1093/hmg/ddq474.
95. Zhang, L., Yang, W., Ying, D., **Cherny, S. S.**, Hildebrandt, F., Sham, P. C., and Lau, Y. L. (2011) Homozygosity mapping on a single patient—identification of homozygous regions of recent common ancestry by using population data. *Human Mutation*, *32*(3), 345–353; doi:10.1002/humu.21432.
96. Zhang, Y., Yang, W., Mok, C. C., Chan, T. M., Wong, R. W., Mok, M. Y., Lee, K. W., Wong, S. N., Leung, A. M., Lee, T. L., Ho, M. H., Lee, P. P., Wong, W. H., Yang, J., Zhang, J., Wong, C. M., Ng, I. O., Garcia-Barceló, M. M., **Cherny, S. S.**, Tam, P. K., Sham, P. C., Lau, C. S., and Lau, Y. L. (2011) Two missense variants in UHRF1BP1 are independently associated with systemic lupus erythematosus in Hong Kong Chinese. *Genes and Immunity*, *12*, 231–234; doi:10.1038/gene.2010.66.
97. Cheung, B. M. Y., Ong, K. L., Tso, A. W. K., Leung, R. Y. H., Xu, A., **Cherny, S. S.**, Sham, P. C., Lam, T. H., and Lam, K. S. L. on behalf of the Investigators of the Hong Kong Cardiovascular Risk Factor Prevalence Study (2012) C-reactive protein as a predictor of hypertension in the Hong Kong Cardiovascular Risk Factor Prevalence Study (CRISPS) cohort. *Journal of Human Hypertension*, *26*, 108–116; doi:10.1038/jhh.2010.125.
98. Guo, Y., Baum, L. W., Sham, P. C., Wong, V., Ng, P. W., Lui, C. H. T., Sin, N. C., Tsoi, T. H., Tang, C. S. M., Kwan, J. S. H., Yip, B. H. K., Xiao, S. M., Thomas, G. N., Lau, Y. L., Yang, W., **Cherny, S. S.**<sup>1</sup>, and Kwan, P.<sup>1</sup> (2012) Two-stage genome-wide association study identifies variants in *CAMSAP1L1* as susceptibility loci for epilepsy in Chinese. *Human Molecular Genetics*, *21*(5), 1184–1189; doi:10.1093/hmg/ddr550.

99. Guo, Y., Tomlinson, B., Chu, T., Fang, Y. J., Gui, H., Tang, C. S., Yip, B. H., **Cherny, S. S.**<sup>1</sup>, Hur, Y.-M., Sham, P. C., Lam, T. H.<sup>1</sup>, Thomas, G. N.<sup>1</sup> (2012) A genome-wide linkage and association scan reveals novel loci for hypertension and blood pressure traits. *PLoS One*, *7*(2), e31489; doi:10.1371/journal.pone.0031489.
100. Hur, Y.-M., **Cherny, S. S.**, and Sham, P. C. (2012) Heritability of hallucinations in adolescent twins. *Psychiatry Research*, *199*(2), 98–101; doi:10.1016/j.psychres.2012.04.024.
101. Li, M. X., Yeung, J. M. Y., **Cherny, S. S.**, and Sham, P. C. (2012) Evaluating the effective number of independent tests and significant *p*-value thresholds in commercial genotyping arrays and public imputation reference datasets. *Human Genetics*, *131*, 747–756; doi:10.1007/s00439-011-1118-2.
102. Li, R., Yang, W., Zhang, J., Hirankarn, N., Pan, H. F., Mok, C. C., Chan, T. M., Wong, R. W., Mok, M. Y., Lee, K. W., Wong, S. N., Leung, A. M., Li, X. P., Avihingsanon, Y., Lee, T. L., Ho, M. H., Lee, P. P., Wong, W. H., Wong, C. M., Ng, I. O., Yang, J., Li, P. H., Zhang, Y., Zhang, L., Li, W., Baum, L., Kwan, P., Rianthavorn, P., Deekajorndej, T., Suphapeetiporn, K., Shotelersuk, V., Garcia-Barceló, M. M., **Cherny, S. S.**, Tam, P. K., Sham, P. C., Lau, C. S., Shen, N., Lau, Y. L., and Ye, D. Q. (2012) Association of CD247 with systemic lupus erythematosus in Asian populations. *Lupus*, *21*, 75–83; doi:10.1177/0961203311422724.
103. Ngo, D.-N., So, M.-T., Gui, H., Tran, A.-Q., Bui, D.-H., **Cherny, S.**, Tam, P. K.-H., Nguyen, T.-L., and Garcia-Barceló, M.-M. (2012) Screening of the *RET* gene of Vietnamese Hirschsprung patients identifies 2 novel missense mutations. *Journal of Pediatric Surgery*, *47*(10), 1859–1864; doi:10.1016/j.jpedsurg.2012.05.020.
104. Pang, X., Liu, C., Shi, L., Liu, R., Liang, D., Li, H., **Cherny, S. S.**, Chen, S. (2012) Utility of the trnH-psbA intergenic spacer region and its combinations as plant DNA barcodes: a meta-analysis. *PLoS One*, *7*(11), e48833; doi:10.1371/journal.pone.0048833.
105. Tang, C. S.-M., Cheng, G., So, M.-T., Yip, B. H.-K., Miao, X.-P., Wong, E. H.-M., Ngan, E. S.-W., Lui, V. C.-H., Song, Y.-Q., Chan, D., Cheung, K., Yuan, Z.-W., Lei, L., Chung, P. H.-Y., Liu, X.-L., Wong, K. K.-Y., Marshall, C. R., Scherer, S., **Cherny, S. S.**, Sham, P.-C., Tam, P. K.-H., and Garcia-Barceló, M.-M. (2012) Genome-wide copy number analysis uncovers a new HSCR gene: *NRG3*. *PLoS Genetics*, *8*(5), e1002687; doi:10.1371/journal.pgen.1002687.
106. Tang, C. S.-M., Ngan, E. S.-W., Tang, W.-K., So, M.-T., Cheng, G., Miao, X.-P., Leon, T. Y.-Y., Leung, B. M.-C., Hui, K.-J. W. S., Lui, V. C.-H., Chen, Y., Chan, I. H.-Y., Chung, P. H.-Y., Liu, X.-L., Wong, K. K.-Y., Sham, P.-C., **Cherny, S. S.**<sup>1</sup>, Tam, P. K.-H.<sup>1</sup>, and Garcia-Barceló, M.-M.<sup>1</sup> (2012) Mutations in the *NRG1* gene are associated with Hirschsprung disease. *Human Genetics*, *131*, 67–76; doi:10.1007/s00439-011-1035-4.
107. Yang, H.-C., Liang, Y.-J., Chen, J.-W., Chiang, K.-M., Chung, C.-M., Ho, H.-Y., Ting, C.-T., Lin, T.-H., Sheu, S.-H., Tsai, W.-C., Chen, J.-H., Leu, H.-B., Yin, W.-H., Chiu, T.-Y., Chern, C.-I., Lin, S.-J., Tomlinson, B., Guo, Y., Sham, P. C., **Cherny, S. S.**, Lam, T. H., Thomas, G. N., Pan, W.-H. (2012) Identification of *IGF1*, *SLC4A4*, *WWOX*, and *SFMBT1* as hypertension susceptibility genes in Han Chinese with a genome-wide gene-based association study. *PLoS One*, *7*(3), e32907; doi:10.1371/journal.pone.0032907.
108. Cheng, G., Tang, C. S.-M., Wong, E. H.-M., Cheng, W. W.-C., So, M.-T., Miao, X., Zhang, R., Cui, L., Liu, X., Ngan, E. S.-W., Lui, V. C.-H., Chung, P. H.-Y., Chan, I. H.-Y., Liu, J., Zhong, W., Xia, H., Yu, J., Qiu, X., Wu, X.-Z., Wang, B., Dong, X., Tou, J., Huang, L., Yi, B., Ren, H., Chan, E. K.-W., Ye, K., O'reilly, P. F., Wong, K. K.-Y., Sham, P.-C., **Cherny, S. S.**, Tam, P. K.-H., and Garcia-Barceló, M.-M. (2013) Common genetic variants regulating *ADD3* gene expression alter biliary atresia risk. *Journal of Hepatology*, *59*(6), 1285–1291; doi:10.1016/j.jhep.2013.07.021.
109. Cui, L., Wong, E. H.-M., Cheng, G., Firmato de Almeida, M., So, M.-T., Sham, P.-C., **Cherny, S. S.**, Tam, P. K.-H., and Garcia-Barceló, M.-M. (2013) Genetic analyses of a three generation family segregating Hirschsprung Disease and iris heterochromia. *PLoS ONE*, *8*(6), e66631; doi:10.1371/journal.pone.0066631.

110. Gui, H., Tang, W. K., So, M. T., Proitsi, P., Sham, P. C., Tam, P. K., Ngan, E. S.-W.<sup>1</sup>, **Cherny, S. S.**<sup>1</sup>, and Garcia-Barceló, M. M.<sup>1</sup> (2013) RET and NRG1 interplay in Hirschsprung disease. *Human Genetics*, 132(5), 591–600; doi:10.1007/s00439-013-1272-9.
111. Ma, R. C., Hu, C., Tam, C. H., Zhang, R., Kwan, P., Leung, T. F., Thomas, G. N., Go, M. J., Hara, K., Sim, X., Ho, J. S., Wang, C., Li, H., Lu, L., Wang, Y., Li, J. W., Wang, Y., Lam, V. K., Wang, J., Yu, W., Kim, Y. J., Ng, D. P., Fujita, H., Panoutsopoulou, K., Day-Williams, A. G., Lee, H. M., Ng, A. C., Fang, Y. J., Kong, A. P., Jiang, F., Ma, X., Hou, X., Tang, S., Lu, J., Yamauchi, T., Tsui, S. K., Woo, J., Leung, P. C., Zhang, X., Tang, N. L., Sy, H. Y., Liu, J., Wong, T. Y., Lee, J. Y., Maeda, S., Xu, G., **Cherny, S. S.**, Chan, T. F., Ng, M. C., Xiang, K., Morris, A. P., DIAGRAM Consortium, Keildson, S., The MuTHER Consortium, Hu, R., Ji, L., Lin, X., Cho, Y. S., Kadowaki, T., Tai, E. S., Zeggini, E., McCarthy, M. I., Hon, K. L., Baum, L., Tomlinson, B., So, W. Y., Bao, Y., Chan, J. C., and Jia, W. (2013) Genome-wide association study in a Chinese population identifies a susceptibility locus for type 2 diabetes at 7q32 near *PAX4*. *Diabetologia*, 56, 1291–1305; doi:10.1007/s00125-013-2874-4.
112. Song, Y.-Q., Karasugi, T., Cheung, K. M.-C., Chiba, K., Ho, D. W.-H., Miyake, A., Kao, P. Y.-P., Sze, K. L., Yee, A., Takahashi, A., Kawaguchi, Y., Mikami, Y., Matsumoto, M., Togawa, D., Kanayama, M., Shi, D., Dai, J., Jiang, Q., Wu, C., Tian, W., Wang, N., Leong, J. C.-Y., Luk, K. D.-K., Yip, S.-P., **Cherny, S. S.**, Wang, J., Mundlos, S., Kelempisioti, A., Eskola, P. J., Männikkö, M., Mäkelä, P., Karppinen, J., Järvelin, M.-R., O'Reilly, P. F., Kubo, M., Kimura, T., Kubo, T., Toyama, Y., Mizuta, H., Cheah, K. S.-E., Tsunoda, T., Sham, P.-C., Ikegawa, S., Chan, D. (2013) Lumbar disc degeneration is linked to a carbohydrate sulfotransferase 3 variant. *The Journal of Clinical Investigation*, 123(11), 4909–4917; doi:10.1172/JCI69277.
113. Wong, E. H.-M., Cui, L., Ng, C.-L., Tang, C. S.-M., Liu, X.-L., So, M.-T., Yip, B. H.-K., Cheng, G., Zhang, R., Tang, W.-K., Yang, W., Lau, Y.-L., Baum, L., Kwan, P., Sun, L.-D., Zuo, X.-B., Ren, Y.-Q., Yin, X.-Y., Miao, X.-P., Liu, J., Lui, V. C.-H., Ngan, E. S.-W., Yuan, Z.-W., Zhang, S.-W., Xia, J., Wang, H., Sun, X.-B., Wang, R., Chang, T., Chan, I. H.-Y., Chung, P. H.-Y., Zhang, X.-J., Wong, K. K.-Y., **Cherny, S. S.**, Sham, P.-C., Tam, P. K.-H., and Garcia-Barceló, M.-M. (2013) Genome-wide copy number variation study in anorectal malformations. *Human Molecular Genetics*, 22(3), 621–631; doi:10.1093/hmg/dd5451.
114. Wong, E. H. M., Ng, C.-L., Lui, V. C.-H., So, M.-T., **Cherny, S. S.**, Sham, P.-C., Tam, P. K.-H., and Garcia-Barceló, M.-M. (2013) Gene network analysis of candidate loci for human anorectal malformations. *PLoS One*, 8(8), e69142; doi:10.1371/journal.pone.0069142.
115. Wong, E. H.-M., So, H.-C., Li, M., Wang, Q., Butler, A. W., Paul, B., Wu, H.-M., Hui, T. C.-K., Choi, S. C., So, M.-T., Garcia-Barceló, M.-M., McAlonan, G. M., Chen, E. Y.-H., Cheung, E. F.-C., Chan, R. C.-K., Purcell, S. M., **Cherny, S. S.**, Chen, R. R.-L., Li, T., Sham, P.-C. (2013) Common variants on Xq28 conferring risk of schizophrenia in Han Chinese. *Schizophrenia Bulletin*, in press; doi:10.1093/schbul/sbt104.
116. Yang, W., Tang, H., Zhang, Y., Tang, X., Zhang, J., Sun, L., Yang, J., Cui, Y., Zhang, L., Hirankarn, N., Cheng, H., Pan, H. F., Gao, J., Lee, T. L., Sheng, Y., Lau, C. S., Li, Y., Chan, T. M., Yin, X., Ying, D., Lu, Q., Leung, A. M., Zuo, X., Chen, X., Tong, K. L., Zhou, F., Diao, Q., Tse, N. K., Xie, H., Mok, C. C., Hao, F., Wong, S. N., Shi, B., Lee, K. W., Hui, Y., Ho, M. H., Liang, B., Lee, P. P., Cui, H., Guo, Q., Chung, B. H., Pu, X., Liu, Q., Zhang, X., Zhang, C., Chong, C. Y., Fang, H., Wong, R. W., Sun, Y., Mok, M. Y., Li, X.-P., Avihingsanon, Y., Zhai, Z., Rianthavorn, P., Deekajorndej, T., Suphapeetiporn, K., Gao, F., Shotelersuk, V., Kang, X., Ying, S. K., Zhang, L., Wong, W. H., Zhu, D., Fung, S. K., Zeng, F., Lai, W. M., Wong, C. M., Ng, I. O., Garcia-Barceló, M.-M., **Cherny, S. S.**, Shen, N., Tam, P. K., Sham, P. C., Ye, D.-Q., Yang, S., Zhang, X., and Lau, Y. L. (2013) Meta-analysis followed by replication identifies loci in or near *CDKN1B*, *TET3*, *CD80*, *DRAM1*, and *ARID5B* as associated with systemic lupus erythematosus in Asians. *American Journal of Human Genetics*, 92(1), 41–51; doi:10.1016/j.ajhg.2012.11.018.
117. Zhang, J., Zhang, Y., Yang, J., Zhang, L., Sun, L., Pan, H.-F., Hirankarn, N., Ying, D., Zeng, S., Lee, T. L., Lau, C. S., Chan, T. M., Leung, A. M., Mok, C. C., Wong, S. N., Lee, K. W., Ho, M. H., Lee,

P. P., Chung, B. H., Chong, C. Y., Wong, R. W., Mok, M. Y., Wong, W. H., Tong, K. L., Tse, N. K., Li, X.-P., Avihingsanon, Y., Rianthavorn, P., Deekajorndej, T., Suphapeetiporn, K., Shotelersuk, V., Ying, S. K., Fung, S. K., Lai, W. M., Garcia-Barceló, M.-M., **Cherny, S. S.**, Tam, P. K., Cui, Y., Sham, P. C., Yang, S., Ye, D. Q., Zhang, X.-J., Lau, Y. L., Yang, W. (2013) Three SNPs in chromosome 11q23.3 are independently associated with systemic lupus erythematosus in Asians. *Human Molecular Genetics*, in press; doi:10.1093/hmg/ddt424.

### Book Reviews

1. **Cherny, S. S.** (2001) Review of *Advances in Twin and Sib-Pair Analysis*. *American Journal of Human Genetics*, 69, 668; doi:10.1086/322935.

### Book chapters

1. Fulker, D. W., **Cherny, S. S.**, and Cardon, L. R. (1993) Continuity and change in cognitive development. In R. Plomin and G. E. McClearn (eds.), *Nature, Nurture, and Psychology* (pp. 77–97). American Psychological Association, Washington, D. C.; <http://books.google.com/books?id=J2nMPgAACAAJ>.
2. Cardon, L. R. and **Cherny, S. S.** (1994) Adoption design methodology. In J. C. DeFries, R. Plomin, and D. W. Fulker, *Nature and Nurture During Middle Childhood* (pp. 26–45). Blackwell, Oxford; <http://books.google.com/books?id=7G-tQgAACAAJ>.
3. **Cherny, S. S.** (1994) Home environmental influences on general cognitive ability. In J. C. DeFries, R. Plomin, and D. W. Fulker, *Nature and Nurture During Middle Childhood* (pp. 262–280). Blackwell, Oxford; <http://books.google.com/books?id=7G-tQgAACAAJ>.
4. **Cherny, S. S.** and Cardon, L. R. (1994) General cognitive ability. In J. C. DeFries, R. Plomin, and D. W. Fulker, *Nature and Nurture During Middle Childhood* (pp. 46–56). Blackwell, Oxford; <http://books.google.com/books?id=7G-tQgAACAAJ>.
5. Whitfield, K. E. and **Cherny, S. S.** (1994) Motor development. In J. C. DeFries, R. Plomin, and D. W. Fulker, *Nature and Nurture During Middle Childhood* (pp. 173–180). Blackwell, Oxford; <http://books.google.com/books?id=7G-tQgAACAAJ>.
6. **Cherny, S. S.**, Fulker, D. W., and Hewitt, J. K. (1997) Cognitive development from infancy to middle childhood. In R. J. Sternberg and E. L. Grigorenko (eds.), *Intelligence, Heredity, and Environment* (pp. 463–482). Cambridge University Press, Cambridge; <http://books.google.com/books?id=BMXVZzvLXz8C>.
7. **Cherny, S. S.**, Fulker, D. W., Plomin, R., and DeFries, J. C. (2001) Continuity and change in general cognitive ability from 14 to 36 months. In R. N. Emde and J. K. Hewitt (eds.), *Infancy to Early Childhood: Genetic and Environmental Influences on Developmental Change* (pp. 206–220). Oxford University Press, New York; <http://books.google.com/books?id=VCfjfcLf-uUC>.
8. **Cherny, S. S.**, Saudino, K. J., Fulker, D. W., Plomin, R., Corley, R. P., and DeFries, J. C. (2001) The development of observed shyness from 14 to 20 months: Shyness in context. In R. N. Emde and J. K. Hewitt (eds.), *Infancy to Early Childhood: Genetic and Environmental Influences on Developmental Change* (pp. 269–282). Oxford University Press, New York; <http://books.google.com/books?id=VCfjfcLf-uUC>.
9. Saudino, K. J. and **Cherny, S. S.** (2001) Parental ratings of temperament in twins. In R. N. Emde and J. K. Hewitt (eds.), *Infancy to Early Childhood: Genetic and Environmental Influences on Developmental Change* (pp. 73–88). Oxford University Press, New York; <http://books.google.com/books?id=VCfjfcLf-uUC>.
10. Saudino, K. J. and **Cherny, S. S.** (2001) Sources of continuity and change in observed temperament. In R. N. Emde and J. K. Hewitt (eds.), *Infancy to Early Childhood: Genetic and Environmental Influences on Developmental Change* (pp. 89–110). Oxford University Press, New York; <http://books.google.com/books?id=VCfjfcLf-uUC>.



11. Bishop, E. G., **Cherny, S. S.**, and Hewitt, J. K. (2003) Developmental Analysis of IQ. In S. A. Petrill, R. Plomin, J. C. DeFries, and J. K. Hewitt (eds.), *Nature, Nurture, and the Transition to Early Adolescence*, (pp. 13–27). Oxford University Press, New York; <http://books.google.com/books?id=bQMeaLG-ByYC>.
12. Gagne, J. R., Saudino, K. J., and **Cherny, S. S.** (2003) Genetic influences on temperament in early adolescence: A multimethod perspective. In S. A. Petrill, R. Plomin, J. C. DeFries, and J. K. Hewitt (eds.), *Nature, Nurture, and the Transition to Early Adolescence*, (pp. 166–184). Oxford University Press, New York; <http://books.google.com/books?id=bQMeaLG-ByYC>.
13. **Cherny, S. S.** (2005) Cholesky decomposition. In B. S. Everitt and D. C. Howell (eds.), *Encyclopedia of Statistics in Behavioral Science*, (Vol 1, pp. 262–263). John Wiley & Sons, Chichester, U.K.; doi:10.1002/0470013192.bsa090.
14. **Cherny, S. S.** (2005) Correlation issues in genetics research. In B. S. Everitt and D. C. Howell (eds.), *Encyclopedia of Statistics in Behavioral Science*, (Vol 1, pp. 402–403). John Wiley & Sons, Chichester, U.K.; doi:10.1002/0470013192.bsa139.
15. **Cherny, S. S.** (2008) Chapter 9: Regression methods for linkage analysis. In B. M. Neale, M. A. R. Ferreira, S. E. Medland, and D. Posthuma (eds.), *Statistical genetics: Gene Mapping through Linkage and Association*, (pp. 153–180). Taylor and Francis, U.K.; <http://genemapping.org>.
16. **Cherny, S. S.** (2009) QTL methodology in behavior genetics. In Y. K. Kim (ed.), *Handbook of Behavioral Genetics*, (pp. 35–45). Springer, New York; doi:10.1007/978-0-387-76727-7\_3.
17. Sham, P. C. and **Cherny, S. S.** (2010) Genetic architecture of complex diseases. In E. Zeggini and A. Morris (eds.), *Analysis of Complex Disease Association Studies: A Practical Guide*, (pp. 1–13). Elsevier, Oxford; [http://books.google.com/books?id=jD8ST0\\_kwakC](http://books.google.com/books?id=jD8ST0_kwakC).

#### Abstracts

1. **Cherny, S. S.**, Cardon, L. R., and Fulker, D. W. (1990) Multivariate genetic analysis of three situation-specific composite measures of shyness in 14-month-old twins from the MacArthur Longitudinal Twin Study. *Behavior Genetics*, *20*, 710.
2. Bishop, E. G., **Cherny, S. S.**, DiLalla, L. F., and Fulker, D. W. (1991) The infant Visual Expectation Paradigm as a predictor of adult IQ: The Twin Infant Project. *Behavior Genetics*, *21*, 561-562.
3. **Cherny, S. S.**, Bishop, E. G., Williams, N. D., DiLalla, L. F., and Fulker, D. W. (1991) Infant predictors of adult IQ: The Twin Infant Project. *Behavior Genetics*, *21*, 565.
4. **Cherny, S. S.** and Fulker, D. W. (1991) A longitudinal hierarchical model of infant shyness: The MacArthur Longitudinal Twin Study. *Behavior Genetics*, *21*, 564-565.
5. **Cherny, S. S.** and Fulker, D. W. (1992) Continuity and change in general intelligence from ages 1 through 9 years. *Behavior Genetics*, *22*, 715.
6. Fulker, D. W. and **Cherny, S. S.** (1992) Continuity and change in general cognitive ability from 1 to 9 years of age. *Abstracts of the Seventh International Congress of Twin Studies*, p. 89.
7. Schmitz, S., **Cherny, S. S.**, and Fulker, D. W. (1992) Genetic and environmental influences on the etiology of problem behavior in early childhood. *Behavior Genetics*, *22*, 751.
8. Whitfield, K. E., **Cherny, S. S.**, Fulker, D. W., and Reznick, J. S. (1992) A multivariate analysis of cognitive measures at 14 months: The MacArthur Longitudinal Twin Study. *Behavior Genetics*, *22*, 762.
9. **Cherny, S. S.** (1993) A genetic analysis of continuity and change in general cognitive ability from 14 to 48 months: The MacArthur Longitudinal Twin Study. *Abstracts of the 60th Anniversary Meeting of the Society for Research in Child Development*, *9*, 290.

10. **Cherny, S. S.** and Fulker, D. W. (1993) Continuity and change in specific cognitive abilities from ages 3 through 9 years. *Behavior Genetics*, 23, 549.
11. Finkel, D., Whitfield, K. E., **Cherny, S. S.**, and McGue, M. (1993) A quantitative genetic analysis of information processing in later life. *The Gerontologist*, 33(Special Issue), 222.
12. Schmitz, S., **Cherny, S. S.**, and Fulker, D. W. (1993) Assessing genetic and environmental influences on problem behavior: Twin analysis of the Child Behavior Checklist. *Abstracts of the 60th Anniversary Meeting of the Society for Research in Child Development*, 9, 587.
13. Cardon, L. R., **Cherny, S. S.**, and Fulker, D. W. (1994) Multipoint methods for linkage analysis of quantitative trait loci in sib pairs. *American Journal of Human Genetics*, 55(Supplement), A147.
14. **Cherny, S. S.** and Fulker, D. W. (1994) The structure of specific cognitive abilities in early and middle childhood. *Behavior Genetics*, 24, 509.
15. **Cherny, S. S.**, Stallings, M. C., Young, S. E., Miles, D. R., Carey, G., Hewitt, J. K., and Fulker, D. W. (1994) Familial aggregation of depressive symptoms, antisocial personality, and substance abuse. *Behavior Genetics*, 24, 509.
16. **Cherny, S. S.**, Stallings, M. C., Young, S. E., Miles, D. R., Carey, G., Hewitt, J. K., and Fulker, D. W. (1994) Familial transmission of depressive symptoms in adolescent substance abusing males. *Behavior Genetics*, 24, 509.
17. Stallings, M. C., **Cherny, S. S.**, Young, S. E., Miles, D. R., Carey, G., Hewitt, J. K., and Fulker, D. W. (1994) Familial aggregation of alcohol and drug problems in adolescent males. *Behavior Genetics*, 24, 531-532.
18. Stallings, M. C., **Cherny, S. S.**, Young, S. E., Miles, D. R., Carey, G., Hewitt, J. K., and Fulker, D. W. (1994) Personality risk factors in adolescent substance abuse. *Behavior Genetics*, 24, 531.
19. Wilson, S. M., **Cherny, S. S.**, and Fulker, D. W. (1994) Continuity and change in general intelligence from ages 1 through 7 years. *Behavior Genetics*, 24, 537.
20. **Cherny, S. S.**, Corley, R. P., Fulker, D. W., DeFries, J. C., Goodman, D. W., Markel, P. D., and Johnson, T. E. (1995) Regression approaches to QTL analyses in selected  $F_2$  samples. *Alcoholism: Clinical and Experimental Research*, 19, 48A.
21. **Cherny, S. S.** and Fulker, D. W. (1995) Continuity and change in specific cognitive abilities in the Colorado Adoption Project and Colorado Twin Study. *Behavior Genetics*, 25, 259.
22. **Cherny, S. S.**, Fulker, D. W., and Corley, R. P. (1995) Continuity and change in general cognitive ability from 14 months to 7 years in the Colorado Twin Study. *Abstracts of the Eighth International Congress of Twin Studies*, p. 28.
23. **Cherny, S. S.**, Fulker, D. W., Hu, S., Pattatucci, A. M. L., Patterson, C., and Hamer, D. H. (1995) Multipoint interval mapping of a quantitative trait locus for sexual orientation using selected sib pairs. *Behavior Genetics*, 25, 259.
24. Corley, R. P., **Cherny, S. S.**, Fulker, D. W., DeFries, J. C., Goodman, D. W., Flint, J., and Collins, A. C. (1995) Comparison of regression and maximum-likelihood approaches to quantitative trait loci interval mapping in  $F_2$  experiments involving selected samples. *Behavior Genetics*, 25, 260.
25. Corley, R. P., **Cherny, S. S.**, Stallings, M. C., and Young, S. E. (1995) Temperamental and familial factors in adolescent substance use. *Abstracts of the Biennial Meeting of the Society for Research in Child Development*, 10, p. 344.
26. Fulker, D. W., Young, S. E., **Cherny, S. S.**, and Schmitz, S. (1995) Sources of individual differences in the development of language and cognition. *Abstracts of the Biennial Meeting of the Society for Research in Child Development*, 10, p. 202.

27. Saudino, K. J., **Cherny, S. S.**, and Plomin, R. (1995) Components of behavioral inhibition: Exploring the genetic and environmental relations between Emotionality, Shyness, and Behavioral Inhibition. *Behavior Genetics*, 25, 286.
28. Stallings, M. C., **Cherny, S. S.**, Young, S. E., Carey, G., Hewitt, J. K., and Fulker, D. W. (1995) Personality factors in early-onset adolescent substance abuse. *Behavior Genetics*, 25, 289.
29. Stallings, M. C., **Cherny, S. S.**, Young, S. E., Corley, R. P., Hewitt, J. K., and Fulker, D. W. (1995) The coaggregation of alcohol abuse, depressive symptoms, and antisocial personality in the families of multiple-problem adolescents. *Alcoholism: Clinical and Experimental Research*, 19, 71A.
30. Young, S. E., Stallings, M. C., **Cherny, S. S.**, Carey, G., Hewitt, J. K., and Fulker, D. W. (1995) Familial transmission in substance dependence and comorbid psychiatric problems in severely affected adolescent males and their relatives. *Behavior Genetics*, 25, 294–295.
31. **Cherny, S. S.**, Corley, R. P., Fulker, D. W., Plomin, R., and DeFries, J. C. (1996) Parent-offspring resemblance for adult general cognitive ability in the Colorado Adoption Project. *Behavior Genetics*, 26, 583.
32. **Cherny, S. S.** and Fulker, D. W. (1996) Continuity and change in specific cognitive abilities in twins, siblings, and adoptees. *Proceedings of the Biennial International Society for the Study of Development Conference*, 374.
33. Corley, R. P., **Cherny, S. S.**, Schmitz, S., and Fulker, D. W. (1996) The Colorado Twin Study: Design and methods. *Proceedings of the Biennial International Society for the Study of Development Conference*, 47.
34. Fulker, D. W. and **Cherny, S. S.** (1996) The Analysis of Quantitative Trait Loci for Complex Behaviors. *Abstracts of the 9th International Congress of Human Genetics*.
35. Fulker, D. W., Schmitz, S., and **Cherny, S. S.** (1996) Overview of the Colorado Adoption Project (CAP). *Proceedings of the Biennial International Society for the Study of Development Conference*, 46.
36. Pogue-Geile, M. F., Manuck, S. B., Kamarck, T., **Cherny, S. S.**, and Debski, T. (1996) Cardiovascular Reactivity to Stress: A Twin Study. *Behavior Genetics*, 26, 594.
37. **Cherny, S. S.** and Fulker, D. W. (1997) Effects of non-normal errors on the efficacy of selection strategies for mapping quantitative trait loci using sib pairs. *Behavior Genetics*, 27, 586.
38. Knopik, V. S., Olson, R. K., Alarcón, M., **Cherny, S. S.**, and DeFries, J. C. (1997) Etiology of covariation between measures of speeded and perceptual processes and reading performance: A twin study. *Behavior Genetics*, 27, 596.
39. Saudino, K. J. and **Cherny, S. S.** (1997) Ratings of temperament in twins: A comparison of alternative models. *Behavior Genetics*, 27, 604–605.
40. Talbot, C. J., Nicod, A., **Chern[e]y, S.**, Fulker, D., Collins, A., and Flint, J. (1998) Fine mapping of QTLs for emotionality in mice as a model of human anxiety. *American Journal of Human Genetics*, 61, 1731.
41. **Cherny, S. S.**, Fulker, D. W., and Sham, P. (1998) Maximum-likelihood methods of association and linkage. *Behavior Genetics*, 28, 467.
42. **Cherny, S. S.**, Fulker, D. W., and Sham, P. (1998) Maximum-likelihood methods of association and linkage. *Twin Research*, 1, 84.
43. **Cherny, S. S.**, Fulker, D. W., Sham, P. C., and Hewitt, J. K. (1998) Maximum-likelihood variance components combined linkage and association sibship analysis for quantitative traits. *American Journal of Human Genetics*, 63, A284.

44. Gayán, J., Smith, S. D., **Cherny, S. S.**, Cardon, L. R., Fulker, D. W., Kimberling, W. J., Olson, R. K., Pennington, B. F., and DeFries, J. C. (1998) Quantitative trait locus for specific language and reading deficits on chromosome 6p. *Behavior Genetics*, *28*, 468–469.
45. Talbot, C. J., Nicod, A., **Chern[e]y, S.**, Fulker, D., Collins, A., and Flint, J. (1998) Ultra-fine mapping of quantitative trait loci (QTLs). *Brain Research*, *809*, P119.
46. **Cherny, S. S.**, Purcell, S., Rijdsdijk, F., Hewitt, J. K., and Sham, P. C. (1999) Selecting maximally informative sibships for QTL linkage analysis. *Behavior Genetics*, *29*, 352.
47. Purcell, S., **Cherny, S. S.**, Rijdsdijk, F., Hewitt, J. K., and Sham, P. C. (1999) Selecting maximally informative sibships for QTL association analysis. *Behavior Genetics*, *29*, 367.
48. Purcell, S., **Cherny, S. S.**, and Sham, P. C. (1999) Selecting maximally informative sibships for QTL linkage and association analysis. *Molecular Psychiatry*, *4*, S11.
49. Sham, P. C., **Cherny, S. S.**, Purcell, S., and Hewitt, J. K. (1999) Power of QTL linkage and association analysis. *Behavior Genetics*, *29*, 369–370.
50. Abecasis, G. R., **Chern[e]y, S. S.**, Cookson, W. O. C., and Cardon, L. R. (2000) MERLIN — Multipoint Engine for Rapid Likelihood Inference. *American Journal of Human Genetics*, *67*, Supplement 2, 327.
51. Cardon, L. R., Abecasis, G. R., and **Cherny, S. S.** (2000) The effect of genotype error on the power to detect linkage and association with quantitative traits. *American Journal of Human Genetics*, *67*, Supplement 2, 1713.
52. Andresen, J. M., Cardon, L. R., **Cherny, S. S.**, Gusella, J. F., MacDonald, M. E., Housman, D. E., and Wexler, N. S. (2001) Variance in age of onset of Huntington's disease that is independent of CAG repeat length remains highly heritable. *American Journal of Human Genetics*, *69*, Supplement 4, 2145.
53. **Cherny, S. S.**, Cardon, L. R., and Abecasis, G. R. (2001) The effects of genotype errors on mapping complex traits: Detection and treatment. *Behavior Genetics*, *31*, 449–450.
54. Hewitt, J. K., Stallings, M., Krauter, K. S., Crowley, T. J., Corley, R., Riggs, P., Young, S., Mikulich, S. K., **Cherny, S. S.**, Rhea, S., and Ehlers, K. (2001) Finding genes that predispose to substance dependence in adolescents: The Colorado Center on Antisocial Drug Dependence (CADD). *Drug and Alcohol Dependence*, *63*, S66.
55. Jawaid, A., Purcell, S., **Cherny, S.**, and Sham, P. (2001) A generalised threshold defined case-control selection strategy for QTL association mapping in pooled DNA samples. *American Journal of Medical Genetics*, *105*, 12.
56. Jawaid, A., Purcell, S., **Cherny, S. S.**, and Sham, P. C. (2001) Optimal selection strategies for QTL mapping using pooled DNA samples. *Behavior Genetics*, *31*, 456.
57. Lessem, J. M., **Cherny, S. S.**, and Abecasis, G. R. (2001) Optimum methods of linkage analysis in a selected sample. *Drug and Alcohol Dependence*, *63*, S90.
58. Lessem, J. M., **Cherny, S. S.**, Abecasis, G. R., Sham, P. C., and Purcell, S. (2001) An overview of regression methods of linkage analysis in selected samples. *Behavior Genetics*, *31*, 458.
59. Sham, P. C., Purcell, S., **Cherny, S. S.**, and Abecasis, G. R. (2002) A powerful regression-based method of quantitative-trait linkage analysis for general pedigrees. *Behavior Genetics*, *32*, 485.
60. Rhea, S. A., Haberstick, B. C., Corley, R. P., and **Cherny, S. S.** (2004) Friends and families: Combined twin-adoption analysis of parenting and peer relationships. *Behavior Genetics*, *34*, 658.
61. Lessem, J. M., Hopfer, C. J., **Cherny, S. S.**, Corley, R. P., Hewitt, J. K., Krauter, K. S., Mikulich, S. K., Rhee, S. H., Smolen, A., Stallings, M. C., Young, S. E., and Crowley, T. J. (2005) Linkage for cannabis dependence on chromosomes 3 and 9. *Behavior Genetics*, *35*, 811.

62. Neale, B. M., **Cherny, S. S.**, and Sham, P. C. (2005) Genome-wide association: Practical application of current technology to overcome multiple testing. *American Journal of Medical Genetics, Part B: Neuropsychiatric Genetics*, 138B, 19–20.
63. Andresen, J. M., Gayán, J., Djoussé, L., Roberts, S., **Cherny, S. S.**, The US-Venezuela Collaborative Research Group<sup>2</sup>, The HD MAPS Collaborative Research Group<sup>3</sup>, Cardon, L. R., Gusella, J. F., MacDonald, M. E., Myers, R. H., Housman, D. E., and Wexler, N. S. (2006) The relationship between CAG repeat length and age of onset differs for adult-onset and juvenile-onset subtypes of Huntington's Disease. *Abstracts of HD 2006: Changes, Advances and Good News (CAG)<sub>n</sub>*, #50.
64. Andresen, J. M., Gayán, J., Brocklebank, D., Alkorta-Aranburu, G., Addis, E., Severins, R., Thole, A., The US-Venezuela Collaborative Research Group<sup>2</sup>, **Cherny, S. S.**, Cardon, L. R. Housman, D. E., and Wexler, N. S. (2006) Genotyping studies in the large Venezuelan kindreds: Association studies for residual age of onset and proof of principle for RNA interference therapy. *Abstracts of HD 2006: Changes, Advances and Good News (CAG)<sub>n</sub>*, #51.
65. Brocklebank, D., Gayán, J., Andresen, J. M., The US-Venezuela Collaborative Research Group<sup>2</sup>, Roberts, S. A., **Cherny, S. S.**, Housman, D. E., Wexler, N. S., and Cardon, L. R. (2006) Transmission and instability of the Huntington's Disease CAG repeat length in a large, multigenerational Venezuelan pedigree. *Abstracts of HD 2006: Changes, Advances and Good News (CAG)<sub>n</sub>*, #49.
66. Gayán, J., Brocklebank, D., Andresen, J. M., Alkorta-Aranburu, G., The US-Venezuela Collaborative Research Group<sup>2</sup>, Cader, Z. M., Roberts, S. A., **Cherny, S. S.**, Wexler, N. S., Cardon, L. R., and Housman, D. E. (2006) Linkage genome scan reveals loci modifying age of onset of Huntington's Disease. *Abstracts of HD 2006: Changes, Advances and Good News (CAG)<sub>n</sub>*, #8.
67. **Cherny, S. S.** (2007) QTL Methodology in behavior genetics: from linkage to genomewide association. *Behavior Genetics*, 37, 744.
68. Garcia-Barceló, M., Tang, C., **Cherny, S.**, Sham, P., and Tam, P. (2007) Genome-wide association study identifies new HSCR loci. Poster presented at the 57th Annual meeting of the American Society of Human Genetics, Program Number 1211/T, San Diego, October 2007.
69. Gayán, J., Brocklebank, D., Andresen, J. M., Alkorta-Aranburu, G., The US-Venezuela Collaborative Research Group<sup>2</sup>, Cader, Z. M., Roberts, S. A., **Cherny, S. S.**, Wexler, N. S., Cardon, L. R., and Housman, D. E. (2007) Genomewide linkage analysis of age of onset of Huntington's Disease. *Behavior Genetics*, 37, 755.
70. Tam, P. K. H., Tang, S. M., Garcia-Barceló, M. M., **Cherny, S. S.**, and Sham, P. C. (2007) Genome-wide association analysis of Hirschsprung's disease. *Abstracts of The 9th International Meeting on Human Genome Variation and Complex Genome Analysis*, Barcelona.
71. **Cherny, S. S.**, Garcia-Barceló, M. M., Tang, C. S., Sham, P. C., and Tam, P. K. (2008) Genome-wide association study of Hirschsprung's disease. *Behavior Genetics*, 38(6), 618.
72. Cheung, B. M. Y., Ong, K. L., Leung, R. Y. H., Wong, L. Y. F., **Cherny, S. S.**, Sham, P. C., Lam, T. H., and Lam, K. S. L. (2008) Association of a polymorphism in the lipin 1 gene with systolic blood pressure in men. *Journal of Hypertension*, 26(Suppl 1), S137.
73. Garcia-Barceló, M., Yeung, M. Y., Tang, C., Sham, P. C., **Cherny, S.**, and Tam, P. K. (2008) Genome-wide association study identifies susceptibility loci for biliary atresia. *Abstracts of the 58th Annual Meeting of the American Society of Human Genetics*, Abstract #1661T, p. 327. Poster presentation.
74. Tang, C. S., Garcia-Barceló, M. M., **Cherny, S. S.**, Sham, P. C., and Tam, P. K. H. (2008) Genetic variants on *NRG1* confer susceptibility to Hirschsprung's disease. *Abstracts of the 58th Annual Meeting of the American Society of Human Genetics*, Abstract #193, p. 73. Platform presentation, Friday, November 14.

75. **Cherny, S. S.**, Kwan, J. S. H., Kung, A. W. C., and Sham, P. C. (2009) Selection strategies for sib-pair association studies. *Behavior Genetics*, 39(6), 641.
76. **Cherny, S. S.**, Tang, C. S. M., Sribudiani, Y., Miao, X. P., So, M. T., Sham, P. C., Tam, P. K. H., Garcia-Barceló, M. M., and Hofstra, R. (2009) Fine mapping of Hirschsprung's disease loci in 9q31. Poster presented at the 59th Annual Meeting of the American Society of Human Genetics, Abstract 721/W, Poster Board #379, p. 224, Honolulu, Hawaii, USA, 21 October 2009.
77. Garcia-Barceló, M., Tang, W. K., Miao, X. P., Tang, C., So, M. T., Leon, Y. Y., Sham, P. C., **Cherny, S. S.**, and Tam, P. K. (2009) Identification of rare variants in the NRG1 gene of Hirschsprungs patients. Poster presented at the 59th Annual Meeting of the American Society of Human Genetics, Abstract 725/W, Poster Board #383, p. 225, Honolulu, Hawaii, USA, 21 October 2009.
78. Ong, K. L., Lam, K. S. L., **Cherny, S. S.**, Tso, A. W. K., Leung, R. Y. H., Sham, P. C., and Cheung, B. M. Y. (2009) Association of single nucleotide polymorphisms in the C-reactive protein gene with systolic blood pressure in Hong Kong Chinese women. *British Journal of Pharmacology*, 68(2), 303.
79. So, H. C., Chen, R. Y. L., Chen, E. Y. H., Cheung, E. F. C., **Cherny, S. S.**, Li, T., and Sham, P. C. (2009) Genome-wide association study of schizophrenia in a Chinese population. Poster presented at the 59th Annual Meeting of the American Society of Human Genetics, Abstract 2185/W, Poster Board #734, p. 622, Honolulu, Hawaii, USA, 22 October 2009.
80. So, H. C., Sham, P. C., and **Cherny, S. S.** (2009) Evaluating variance in liability explained by individual genetic variants and relationship to individualized risk prediction. *Genetic Epidemiology*, 33(8), 809. Paper presented at the 18th Annual Meeting of the International Genetic Epidemiology Society (Abstract 195), Honolulu, Hawaii, USA, 10-20 October 2009.
81. Tam, P. K. H., Tang, C. S. M., Ngan, E. S. W., Lui, V. C. H., Chen, Y., So, M. T., Leon, T. Y. Y., Miao, X. P., Shum, C. K. Y., Liu, F. Q., Yeung, M. Y., Yuan, Z. W., Guo, W. H., Liu, L., Sun, X. B., Huang, L. M., Tou, J. F., Song, Y. Q., Chan, D., Cheung, K. M. C., Wong, K. K. Y., **Cherny, S. S.**, Sham, P. C., and Garcia-Barceló, M. M. (2009) Genome-wide association study identifies NRG1 as a susceptibility locus for Hirschsprung's disease. *Neurogastroenterology and Motility*, 21(2), XXVII.
82. Tang, C. S., Garcia-Barceló, M. M., **Cherny, S. S.**, Sham, P. C., and Tam, P. K. H. (2009) Genome-wide profile of copy number variants for Hirschsprung disease. Poster presented at the 59th Annual Meeting of the American Society of Human Genetics, Abstract 965/W, Poster Board #623, p. 297, Honolulu, Hawaii, USA, 21 October 2009.
83. Toulopoulou, T., Goldberg, T., Weinberger, D. R., Rijdsdijk, F., Faraone, S., Tsuang, M., Stahl, D., Picchioni, M., Sham, P., **Chern[e]y, S.**, Murray, R., and Seidman, L. (2009) Are intelligence and memory good endophenotypes for schizophrenia? Genetic models in a Harvard, IoP, and NIH collaboration. *Schizophrenia Bulletin*, 35, 110. Invited paper presented at the 12th International Congress on Schizophrenia Research, San Diego, CA, Mar 28–Apr 1, 2009.
84. Yip, B. H., Tang, C. S., **Cherny, S. S.**, Tam, P. K., [Ngan], E. S., Garcia-Barceló, M. M., and Sham, P. C. (2009) Quantifying epistasis between two sets of signaling pathway genes by canonical correlation analysis. Poster presented at the 59th Annual Meeting of the American Society of Human Genetics, Abstract 1852/W, Poster Board #401, p. 533. Honolulu, Hawaii, USA, 22 October 2009.
85. **Cherny, S. S.**, Guo, Y. L., Sham, P. C., Baum, L., and Kwan, P. (2010) A genome-wide association study of symptomatic epilepsy in Han Chinese detects multiple variants. Poster presented at the 60th Annual Meeting of the American Society of Human Genetics, Abstract 2638/F, p. 764. Washington, D.C., USA, 5 November 2010.
86. Garcia-Barceló, M., Tang, C., Tang, W. K., So, M. T., Sham, P. C., **Cherny, S. S.**, and Tam, P. K. (2010) Fine mapping of the NRG1 Hirschsprungs-associated gene. Poster presented at the 60th Annual Meeting of the American Society of Human Genetics, Abstract 1155/F, p. 361. Washington, D.C., USA, 5 November 2010.

87. Kwan, J., **Cherny, S.**, and Sham, P. (2010) Assessing statistical significance in PLINK segmental sharing test. Poster presented at the 60th Annual Meeting of the American Society of Human Genetics, Abstract 2844/W, p. 818. Washington, D.C., USA, 3 November 2010.
88. Li, M., Sham, P., and **Cherny, C.**[sic] (2010) A powerful approach to efficiently combine p-values for gene-based association analysis. Poster presented at the 60th Annual Meeting of the American Society of Human Genetics, Abstract 2815/W, p. 811. Washington, D.C., USA, 3 November 2010.
89. Ong, K. L., Li, M. F., Tso, A. W. K., Xu, A. M., **Cherny, S. S.**, Sham, P. C., Tse, H. F., Cheung, B. M. Y., and Lam, K. S. L. (2010) Adiponectin gene polymorphisms, plasma adiponectin concentration and persistent hypertension in Hong Kong Chinese. *British Journal of Clinical Pharmacology*, 70(2), 303. Presented at the Meeting of the Clinical Pharmacology Section of the British-Pharmacological-Society London, ENGLAND, DEC 15–17, 2009, British Pharmacol Soc, Clin Pharmacol Sect.
90. Ong, K. L., Tso, A. W. K., Leung, R. Y. H., Xu, A., **Cherny, S. S.**, Sham, P. C., Lam, K. S. L., and Cheung, B. M. Y. (2010) C-reactive protein as a predictor of hypertension in the Hong Kong cardiovascular risk prevalence study (CRISPS) cohort. *European Heart Journal Supplements*, 12, S21. Presented at the International Congress of Cardiology, Hong Kong, 26–28 February 2010.
91. So, H. C., Li, M., Chen, R. Y., Cheung, E. F., Chen, E. Y., Cherny, S. S., Li, T., and Sham, P. C. (2010) Genome-wide association study of schizophrenia in a Chinese population. *International Journal of Neuropsychopharmacology*, 13, 171. Presented at the 27th CINP Congress Meeting, Hong Kong, 6–10 June 2010.
92. Wong, E. H. M., Tang, C. S. M., [Garcia-]Barceló, M. G.[sic], Zhang, X. J., Liu, J. J., **Cherny, S. S.**, Sham, P. C., and Tam, P. K. (2010) Genome-wide association study on anorectal malformations in the Chinese population. Poster presented at the 60th Annual Meeting of the American Society of Human Genetics, Abstract 2942/T, p. 846. Washington, D.C., USA, 5 November 2010.
93. Yang, H.-C., Liang, L.-J., Chiang, K.-M., Thomas, G., Tomlinson, B., **Cherny, S.**, Gu[o], Y., Lam, T.-H., Chen, J.-W., and Pan, W.-H. (2010) A genome-wide gene-based association study identifies SLC4A4, WWOX and COMMD7 as hypertension susceptibility genes in a Han Chinese population. Poster presented at the 60th Annual Meeting of the American Society of Human Genetics, Abstract 736/W, p. 240. Washington, D.C., USA, 3 November 2010.
94. Zhang, X. W., **Cherny, S. S.**, Huang, Y., Gao, X., Xiang, Y., Fu, Y. X., Meng, H. Q., Ma, X. H., Wang, Y. C., Li, T., and Sham, P. (2010) Effect of parental age on children's intelligence in the Southwestern China prospective twin registry. *Behavior Genetics*, 40(6), 820.
95. Cheung, B. M. Y., Ong, K. L., Tso, A. W. K., Leung, R. Y. H., **Cherny, S. S.**, Sham, P. C., Lam, T. H., and Lam, K. S. L. (2011) Plasma adrenomedullin level is related to plasma interleukin-6 and a polymorphism in the adrenomedullin gene. *Basic & Clinical Pharmacology & Toxicology*, 109(Suppl 1), 102–103. Abstract P146 appearing in *Abstracts of the 10th Congress of the European Association for Clinical Pharmacology and Therapeutics*, 26–29 June 2011, Budapest, Hungary.
96. **Cherny, S. S.**, Garcia-Barceló, M.-M., Leon, T. Y. Y., So, M. T., Sham, P. C., and Tam, P. H. (2011) *RET* mutational spectrum in Hirschsprungs disease: Evaluation of 601 Chinese patients. Poster presented at the 12th International Congress of Human Genetics / 61st Annual Meeting of the American Society of Human Genetics, Abstract 778F, p. 790. Montréal, Québec, Canada, 14 October 2011.
97. Garcia-Barceló, M., Tang, C. S. M., So, M. T., Marshall, C. R., Scherer, S., **Cherny, S.**, Sham, P., and Tam, P. (2011) Genome-wide copy number analysis uncovers a new HSCR gene: *NRG3*. Poster presented at the 12th International Congress of Human Genetics / 61st Annual Meeting of the American Society of Human Genetics, Abstract 415W, p. 25. Montréal, Québec, Canada, 12 October 2011.
98. Gui, H., Li, M., Sham, P., and **Cherny, S.** (2011) A systematic comparison of GWAS pathway analysis methods. Poster presented at the 12th International Congress of Human Genetics / 61st Annual Meeting of the American Society of Human Genetics, Abstract 619W, p. 83. Montréal, Québec, Canada, 12 October 2011.

99. **Cherny, S.** (2012) Using Twins, Adoptees, or Molecular Data to Separate Genetic from Environmental Causes of Variation Among Individuals. Invited lecture presented at the workshop Biological Basis of Individual Differences in Human Social Communicative Abilities—Quantitative Experiments, Computational Methods and Modeling. Hong Kong Baptist University, 2–5 October 2012; <http://cns.hkbu.edu.hk/workshop2012/handbook.pdf>, p. 20.
100. Cheng, G. Wong, E. H. M., Sham, P. C., **Cherny, S. S.**, Maas, S., Scherer, S. W., Marshall, C. R., Pereira, S. L., Tam, P. K. H., and Garcia-Barceló, M. (2012) Exome sequencing of a pedigree with Caudal Regression Syndrome (CRS). Poster presented at the 62nd Annual Meeting of the American Society of Human Genetics, Abstract 2932W. San Francisco, California, 7 November 2012.
101. **Cherny, S. S.**, Wong, E. H. M., Cui, L., Ng, C. L., Tang, C. S. M., So, M. T., Yip, B. H. K., Cheng, G., Liu, V. C. H., Sham, P. C., Tam, P. K. H., and Garcia-Barceló, M. M. (2012) Genome-wide copy number variation in anorectal malformations. Poster presented at the 62nd Annual Meeting of the American Society of Human Genetics, Abstract 2066F. San Francisco, California, 9 November 2012.
102. Garcia-Barceló, M.-M., Cheng, G., Tang, C. S. M., Liu, X. L., Zhang, R. Z., So, M. T., Wong, E. H. M., Chung, P. H. Y., Chan, I. H. Y., Liu, J., Zhong, W., Xia, H., Yu, J., Wong, K. K. Y., **Cherny, S. S.**, Sham, P. C., and Tam, P. K. H. (2012) Fine mapping on chromosome 10q24.2 implicates ADD3 in biliary atresia. Poster presented at the 62nd Annual Meeting of the American Society of Human Genetics, Abstract 1930T. San Francisco, California, 8 November 2012.
103. Wong, E., Butler, A. W., Wang, Q., **Cherny, S. S.**, Li, T., and Sham, P. C. (2012) Genome-wide copy number analysis on schizophrenia in Han Chinese. Poster presented at the 62nd Annual Meeting of the American Society of Human Genetics, Abstract 2094T. San Francisco, California, 8 November 2012.
104. **Cherny, S. S.**, Gui, H.-S., Xu, L., Sham, P.-C., Jiang, C.-Q., Lam, T.-H., Liu, B., Jin, Y.-L., Zhu, T., Zhang, W.-S., Thomas, G. N., Cheng, K.-K. (2013) Influence of Alzheimer's disease genes on cognitive decline: the Guangzhou Biobank Cohort Study. Poster presented at the 63rd Annual Meeting of the American Society of Human Genetics, Abstract 1190W. Boston, Massachusetts, 23 October 2013.
105. Garcia-Barceló, M., Wong, E. H.-M., Ng, C. H., Lui, V. C.-H., So, M.-T., **Cherny, S. S.**, Sham, P.-C., and Tam, P. K. (2013) Gene network analysis of candidate loci for human anorectal malformations. Poster presented at the 63rd Annual Meeting of the American Society of Human Genetics, Abstract 1027T. Boston, Massachusetts, 24 October 2013.
106. Tang, C. S., Wong, E., Gui, H., **Cherny, S.**, Sham, P., Tam, P., and Garcia-Barceló, M. (2013) Comprehensive comparison of copy number variations detection using Illumina Omni 2.5M and Affymetrix CytoScan<sup>®</sup> arrays. Poster presented at the 63rd Annual Meeting of the American Society of Human Genetics, Abstract 552T. Boston, Massachusetts, 24 October 2013.
107. Zhang, X.-W., Sham, P.-C., **Cherny, S. S.**, Meng, H.-Q., Fu, Y.-X., Huang, Y., and Li, T. (2013) Parents and teachers report on different aspect[s] of children's and adolescent's conduct disorder and hyperactivity/inattention behavior. Poster presented at the 63rd Annual Meeting of the American Society of Human Genetics, Abstract 1255F. Boston, Massachusetts, 25 October 2013.

#### Professional affiliations (past and present)

American Association for the Advancement of Science (past)

American Psychological Association (past)

American Psychological Society (past)

American Society of Human Genetics

Behavior Genetics Association

International Society for the Study of Behavioural Development (past)



**Scholarly meetings attended**

1. 20th Annual Meeting of the Behavior Genetics Association, Aussois, Modane, France, June 1990; presented one paper.
2. 21st Annual Meeting of the Behavior Genetics Association, Saint Louis, Missouri, June 1991; presented one paper and two posters.
3. Fourth International Workshop on the Methodology of Twin and Family Studies, Leuven, Belgium, September, 1991
4. 22nd Annual Meeting of the Behavior Genetics Association, Boulder, Colorado, July 1992; presented one paper and one poster and co-authored one paper.
5. 60th Anniversary Meeting of the Society for Research in Child Development, New Orleans, Louisiana, March 1993; presented two posters.
6. 23rd Annual Meeting of the Behavior Genetics Association, Sydney, N. S. W., Australia, July 1993; presented one paper.
7. 24th Annual Meeting of the Behavior Genetics Association, Barcelona, Catalonia, Spain, July 1994; presented one paper and five posters.
8. Genetic Analysis Workshop 9, Val Morin, Québec, Canada, October 1994; co-authored one paper.
9. 44th Annual Meeting of the American Society of Human Genetics, Montréal, Québec, Canada, October 1994; presented one poster.
10. Biennial Meeting of the Society for Research in Child Development, Indianapolis, Indiana, March 1995; presented one symposium paper and one poster.
11. Eighth International Twin Congress, Richmond, Virginia, May 1995; presented one paper.
12. 25th Annual Meeting of the Behavior Genetics Association, Richmond, Virginia, June 1995; presented one paper and five posters.
13. Research Society on Alcoholism Annual Scientific Meeting, Steamboat Springs, Colorado, June 1995; presented two posters.
14. XIVth Biennial Meetings of the International Society for the Study of Behavioural Development, Québec City, Québec, Canada, August 1996; presented one symposium paper and two posters.
15. 27th Annual Meeting of the Behavior Genetics Association, Toronto, Ontario, Canada, July 1997; presented one symposium paper and one poster and co-authored one paper.
16. World Congress on Psychiatric Genetics, Santa Fe, New Mexico, October 1997.
17. Workshop in the Genetics of Non-Linear Dynamic Systems, State College, Pennsylvania, May 1998.
18. Ninth International Twin Congress, Helsinki, Finland, June 1998; co-authored one paper.
19. 28th Annual Meeting of the Behavior Genetics Association, Stockholm, Sweden, June 1998; co-authored two papers.
20. 48th Annual Meeting of the American Society of Human Genetics, Denver, Colorado, October 1998; presented one poster.
21. S.A.G.E. Short Course, Seattle, Washington, December 1998.
22. 29th Annual Meeting of the Behavior Genetics Association, Vancouver, Canada, July 1999; presented one paper and co-authored two papers.

23. 49th Annual Meeting of the American Society of Human Genetics, San Francisco, California, October 1999.
24. NIH Statistical Genetics Initiative Workshop, Half Moon Bay, California, October 1999; co-authored one paper.
25. Second International Meeting on the Genetic Epidemiology of Complex Traits, Cambridge, United Kingdom, April 2000.
26. VIIIth CEPH Annual Conference, Paris, France, May 2000.
27. 30th Annual Meeting of the Behavior Genetics Association, Burlington, Vermont, June-July 2000.
28. 50th Annual Meeting of the American Society of Human Genetics, Philadelphia, Pennsylvania, October 2000; co-authored two posters.
29. EPOS Symposium on Uncovering the Genetic Basis of Anxiety and Depression, Amsterdam, Holland, October 2000.
30. Genetic Analysis Workshop 12, San Antonio, Texas, October 2000; co-authored two papers.
31. European Science Foundation Exploratory Workshop on Finding quantitative trait loci (QTLs) for complex traits in humans. Statistical analysis of data from selected groups of related individuals: An application to anxiety/depression, Oxford, United Kingdom, January 2001; co-authored one paper.
32. EMBO Workshop on Genetics of Osteoporosis: from Basic to Clinical Research, Sestri Levante, Italy, 31 March-3 April 2001; presented invited paper "The effects of pedigree and genotype error on linkage and association".
33. 31st Annual Meeting of the Behavior Genetics Association, Cambridge, United Kingdom, July 2001; presented one paper and one poster, co-authored one paper, co-chaired one symposium, and chaired one paper session.
34. 51st Annual Meeting of the American Society of Human Genetics, San Diego, California, October 2001; co-authored one poster.
35. 32nd Annual Meeting of the Behavior Genetics Association, Keystone, Colorado, July 2002; presented one paper.
36. 33rd Annual Meeting of the Behavior Genetics Association, Chicago, Illinois, June 2003.
37. 4th International Genetics of Complex Traits Conference, Cambridge, UK, April 2004; instructor.
38. 34th Annual Meeting of the Behavior Genetics Association, Aix-en-Provence, France, June 2004.
39. 35th Annual Meeting of the Behavior Genetics Association, Hollywood, California, June 2005.
40. Informatics Grand Rounds, Mayo Clinic, Rochester, Minnesota, July 2005. Presented invited paper "Variance components methods in quantitative genetic research"; [http://ndc.mayo.edu/mayo/research/bmi/grand\\_rounds05.cfm](http://ndc.mayo.edu/mayo/research/bmi/grand_rounds05.cfm).
41. 36th Annual Meeting of the Behavior Genetics Association, Storrs, Connecticut, June 2006.
42. HD 2006: Changes, Advances and Good News (CAG)<sub>n</sub>, Cambridge, Massachusetts, August 2006; co-authored one paper and four posters.
43. 37th Annual Meeting of the Behavior Genetics Association, Amsterdam, The Netherlands, June 2007; presented one symposium paper and co-authored one paper.
44. Biochemistry & Molecular Biology, Michigan State University, seminar series, 14 March 2008. Presented invited paper "Searching for Huntington's Disease modifier genes in the Venezuelan HD kindreds"; <http://www.bch.msu.edu/bchweekly/weeklyarchive/411108.pdf>.

45. 38th Annual Meeting of the Behavior Genetics Association, Louisville, Kentucky, June 2008; presented one paper.
46. HD 2008: The Milton Wexler Celebration of Life, Cambridge, Massachusetts, August 2008; co-authored one paper and two posters.
47. 39th Annual Meeting of the Behavior Genetics Association, Minneapolis, Minnesota, June 2009; presented one poster.
48. 59th Annual Meeting of the American Society of Human Genetics, Honolulu, Hawaii, October 2009; presented one poster and co-authored four additional posters.
49. 40th Annual Meeting of the Behavior Genetics Association, Seoul, Korea, June 2010; co-authored one paper.
50. 13th International Congress on Twin Studies, Seoul, Korea, June 2010.
51. 60th Annual Meeting of the American Society of Human Genetics, Washington, D. C., November 2010; presented one poster and co-authored five additional posters.
52. 41st Annual Meeting of the Behavior Genetics Association, Newport, Rhode Island, June 2011.
53. 12th International Congress of Human Genetics / 61st Annual Meeting of the American Society of Human Genetics, Montreal, Quebec, Canada, October 2011; presented three posters.
54. 42nd Annual Meeting of the Behavior Genetics Association, Edinburgh, United Kingdom, June 2012.
55. Biological Basis of Individual Differences in Human Social Communicative Abilities—Quantitative Experiments, Computational Methods and Modeling, Hong Kong Baptist University, 2–5 October 2012; presented invited paper.
56. 62nd Annual Meeting of the American Society of Human Genetics, San Francisco, California, November 2012; presented four posters.
57. 43rd Annual Meeting of the Behavior Genetics Association, Marseille, France, June 2013.
58. 63rd Annual Meeting of the American Society of Human Genetics, Boston, Massachusetts, October 2013; presented four posters.

#### **Refereed for journals**

*Addiction*

*American Journal of Human Genetics*

*American Journal of Medical Genetics, Part B: Neuropsychiatric Genetics*

*Annals of Human Genetics*

*Archives of General Psychiatry*

*Behavior Genetics*

*Bioinformatics*

*Biological Psychiatry*

*Biometrical Journal*

*BMC Genomics*

*Briefings in Bioinformatics*

*Canadian Journal of Behavioural Science*  
*Child Development*  
*Developmental Neuropsychology*  
*Developmental Science*  
*European Journal of Human Genetics*  
*Evolution and Human Behavior*  
*Genes, Brain and Behavior*  
*The Genes and Immunity Journal*  
*Genetic Epidemiology*  
*Genetics*  
*Heredity*  
*Human Genetics*  
*Human Heredity*  
*International Journal of Behavioral Development*  
*Journal of Child Psychology and Psychiatry*  
*The Journal of Clinical Investigation*  
*Journal of Consulting and Clinical Psychology*  
*Journal of Gerontology: Psychological Sciences*  
*Journal of Human Genetics*  
*Journal of Pediatric Biochemistry*  
*Journal of Statistical Planning and Inference*  
*The Lancet*  
*Molecular Biology Reports*  
*Molecular Psychiatry*  
*Personality and Individual Differences*  
*PLoS Genetics*  
*PLoS ONE*  
*Psychiatric Genetics*  
*Psychological Bulletin*  
*Psychological Methods*  
*Psychological Science*  
*Translational Psychiatry*  
*Twin Research and Human Development*  
*World Journal of Pediatrics*

**Grant reviews**

HKBU Faculty Research Grant, Hong Kong (2010)  
Health, Welfare and Food Bureau, Hong Kong (2007)  
Wellcome Trust, United Kingdom (2001; 2002; 2003)  
Foundation Fighting Blindness, USA (1999)

**Examination committees**

YU Zhuoyou, PhD, Department of Clinical Oncology, The University of Hong Kong, November 2013  
(Committee Chair)

ZENG Shuai, PhD, Department of Paediatrics and Adolescent Medicine, The University of Hong Kong,  
November 2013 (Committee Chair)

KWOK Ka Man, MPhil, Department of Surgery, The University of Hong Kong, October 2013 (Committee  
Chair)

ZHANG, Jing, PhD, Department of Paediatrics and Adolescent Medicine, The University of Hong Kong, June  
2013

FAN Yanhui, PhD, Department of Paediatrics and Adolescent Medicine, The University of Hong Kong,  
February 2013

CUI Long, MPhil, Department of Surgery, The University of Hong Kong, August 2012

ZHANG Lu, MPhil, Department of Paediatrics and Adolescent Medicine, The University of Hong Kong, June  
2012

ZHANG Yan, PhD, Department of Paediatrics and Adolescent Medicine, The University of Hong Kong, May  
2012 (Committee Chair)

TANG Wai Kiu, MRes(Med), Department of Surgery, The University of Hong Kong, July 2011

Eric PUN Cheuk Lun, BCogSc, Department of Psychology, The University of Hong Kong, May 2011

Patrick KAO Yu Ping, PhD, Department of Biochemistry, The University of Hong Kong, June 2010

LEE Yiu Fai, PhD, Department of Psychiatry, The University of Hong Kong, February 2010

LI Miaoxin, PhD, Department of Biochemistry, The University of Hong Kong, December 2009

XIAO Sumei, PhD, Department of Medicine, The University of Hong Kong, December 2009

Partha Bose, Transfer of status to DPhil, University of Oxford, September 2004

Clyde Francks, DPhil, University of Oxford, September 2001

Gabrielle Barnby, Transfer of status to DPhil, University of Oxford, May 2001

Yvonne Linney, PhD, King's College London, May 2001

Yvonne Linney, Upgrade from MPhil to PhD, King's College London, November 1999

Frühling Rijdsdijk, PhD, Free University, Amsterdam, January 1997

**Graduate student supervision**

Robert Milan PORSCH, Department of Psychiatry, The University of Hong Kong; primary supervisor; MPhil started September 2013

YUANG Yazhi, Department of Paediatrics and Adolescent Medicine, The University of Hong Kong; co-supervisor; PhD started September 2012

Sam CHOI Shing Wan, Department of Psychiatry, The University of Hong Kong; co-supervisor; MPhil started January 2012

John WONG Kwong Leong, Department of Psychiatry, The University of Hong Kong; primary supervisor; MPhil started September 2011; upgraded to PhD July 2013

Emily WONG Hoi Man, Department of Psychiatry, The University of Hong Kong; co-supervisor; PhD started January 2010; awarded June 2013.

Albert CHOI Siu Chung, Department of Psychiatry, The University of Hong Kong; primary supervisor; MPhil started January 2010; awarded 2012.

ZHANG Xiaowei, Department of Psychiatry, The University of Hong Kong; co-supervisor; PhD started January 2010

CHENG Guo, Department of Surgery, The University of Hong Kong; co-supervisor; MPhil started September 2009; upgraded to PhD September 2011; awarded 2013

GUI Hongsheng, Department of Psychiatry, The University of Hong Kong; primary supervisor; PhD started September 2009; awarded 2013

GUO Youling, Department of Psychiatry, The University of Hong Kong; primary supervisor; MPhil started September 2008; upgraded to PhD September 2010; awarded 2013.

Juilian YEUNG Ming Yiu, Department of Psychiatry, The University of Hong Kong; primary supervisor; MPhil started September 2007; awarded 2009; "Genome wide association studies of biliary atresia in Chinese."

ONG Kwok Leung, Department of Medicine, The University of Hong Kong; co-supervisor; PhD started 2007; awarded 2010; "Genetic variants of obesity-and inflammation-related genes in hypertension: genetic association studies using candidate gene approach."

Clara TANG Sze Man, Department of Psychiatry, The University of Hong Kong; co-supervisor; PhD started 2007; awarded 2010; "Genetic dissection of Hirschsprung's disease."

Ansar Jawaid, Institute of Psychiatry, King's College London; second supervisor; PhD awarded 2003

**References**

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