

Chunyu Liu, Ph.D.

Professor

Department of Psychiatry and Behavioral Sciences,
Department of Neuroscience and Physiology
Research Advisory Committee Chair
SUNY Upstate Medical University
Institute for Human Performance, room 3704
505 Irving Ave. Syracuse, NY 13210
(315) 464-3448 (office)

liuch@upstate.edu

ORCID: <https://orcid.org/0000-0002-5986-4415>

Scopus Author ID: 35315983600

ResearcherID: G-7561-2012

<https://publons.com/author/321349>

NIH Bibliography: [My Bibliography - NCBI \(nih.gov\)](#)
shorturl.at/joHXY

NSF ID: 000820124

NIH: CHUNYU

Scopus H-index: 41 [Scopus preview - Liu, Chunyu - Author details - Scopus](#)

Google H-index 47 [Chunyu Liu - Google Scholar](#)



Research Area

Methods: Psychiatric genetics; genomics and epigenomics; bioinformatics; psychological stress; iPSC/organoid cellular models

Questions: Molecular bases of psychiatric disorders and human behavioural traits

Education

Ph.D. in Genetics 1998

Hunan Medical University, National Laboratory of Medical Genetics of China, P.R. China
Dissertation: *Integrated data analysis for cloning and analyzing the novel human genes related to diseases*

Advisor: Jiahui Xia

M.S. in Cell Biology 1994

Xiamen University, P.R. China

Thesis: *Identification of hereditary variation of hybridization between domesticated silkworm and eri silkworm by RAPD assay*

Advisor: Yuanlin Chen

B.S. in Biochemistry 1991

Wuhan University, P.R. China

Advisor: Xiaojun Ma

Postdoctoral Training

Department of Psychiatry, the University of Chicago, USA 1998- 2000

Mentor: Elliot S. Gershon, M.D.

Professional training

Faculty Leadership Essentials for Academic Development (LEAD) course, SUNY Upstate Medical University. 2021-2022

Building Excellent Skills for Teaching (BEST) workshop, SUNY Upstate Medical University. 2021-2022

Professional Experience

Professor (tenured)	2017- present
Department of Psychiatry, SUNY Upstate Medical University, Syracuse, NY, USA	
Associate Professor (tenured in 2016)	2011-2017
Department of Psychiatry, University of Illinois at Chicago, USA	
Adjunct Professor	2014-present
State Key Laboratory of Medical Genetics, School of Life Science, Central South University, Changsha, China	
Visiting Professor	2017-2020
School of Psychology, Shaanxi Normal University, Xi'an, China	
Assistant Professor	2005- 2011
Department of Psychiatry and Behavioral Neuroscience, the University of Chicago, USA	
Research Associate (Assistant Professor)	2003- 2005
Department of Psychiatry, the University of Chicago, USA	
Research Associate	2000- 2003
Department of Psychiatry, the University of Chicago, USA	
Research Assistant & Research Associate	1994- 1998
Hunan Medical University, National Laboratory of Medical Genetics of China, P.R. China	

Registered Consultant at AlphaSights; GLG 2021 -

Completed Research Grants

1R01MH110920-01 (MPI: Chunyu Liu (Contact), Kevin P. White)	07/01/2016 - 06/30/2020
NIMH \$2,055,000 (total)	1 calendar months
1/2 Measuring translational dynamics and the proteome to identify potential brain biomarkers for psychiatric disease	
1 U01 MH103340-01 (MPIs: Chunyu Liu (Contact), Kevin White)	07/01/2014 – 06/31/2017
NIMH \$1,100,000 (each year)	2.4 calendar months
Genetic variants affect brain gene expression and risks of psychiatric disorders	
1R21AG045789-01A1 (MPIs: Elliot Gershon (Contact), Chunyu Liu, Geoff Faulkner)	07/01/2014 - 06/30/2017
NIA \$150,000	0.6 calendar months
Somatic Mutations in Brain in Alzheimer's Disease	
1P50AA022538-01 (PI: Subhash Pandey)	07/01/2014 – 06/30/2019
NIAAA \$1,137,842	0.6 calendar months
Center for Alcohol Research in Epigenetics.	
NIH 1R01MH094358-01A1 (PI: Rajiv P Sharma)	2012- 2017
The H3K9 Histone Switch 'Levels' in Schizophrenia Blood and Brain	
0.6 calendar months	
1R01ES024988 (PI: Chunyu Liu)	10/01/2014 – 09/30/2017
NIEHS \$325,549	1.8 calendar months
Integrating epigenomic maps to predict regulatory functions of genetic variants	

NIH 1R01MH094483-01 (MPIs: Craig, Edenberg, Gershon, Kelsoe, Nurnberger, and Schork) The Bipolar Genome Study	2012- 2015
NIH R01 MH080425-01 (direct costs \$1,720,459) <i>The Genetic and Genomic Study of MicroRNA in Bipolar and Schizophrenia</i> , PI: Chunyu Liu	2007- 2012
NIH R33 MH083521 (\$132,107) <i>A Human-Specific Gene (G72/G30) in Transgenic Mice</i> , PI: Chunyu Liu	2010- 2012
NIH R01 MH078151-01A1 (\$285,000) <i>Genomic Association Study of Bipolar Disorder</i> , PI: John Kelsoe	2007- 2010
NIH R21 MH083521-01 (direct costs \$198,057) <i>A Human-Specific Gene (G72/G30) in Transgenic Mice</i> , PI: Chunyu Liu	2008- 2010
NIH R01 MH61613-05A1 (\$1,221,410) <i>Genetic Linkage Studies in Bipolar Disorder</i> , PI: Elliot Gershon	2005- 2010
Brain Research Foundation Research Seed Grant (\$24,949) <i>Genetic Mapping of DNA Methylation Regulators in Human Cerebellum</i> , PI: Chunyu Liu	2008- 2009
NIH R01 MH65560-01 (\$1,789,320) <i>Fine Genomic Mapping of 13q32 in Bipolar Disorder</i> , PI: Elliot Gershon	2002- 2007
Brain Research Foundation Research Seed Grant (\$22,166) <i>Identification of Cis-regulatory Elements for Genes Expressed in Human Brain</i> , PI: Chunyu Liu	2006- 2007
Brain Research Foundation Research Seed Grant (\$25,000) <i>Neurogenesis and Plasticity in Bipolar Disorder</i> , PI: Chunyu Liu	2005- 2006
NARSAD Young Investigator Award (\$60,000) <i>Systematic Association Study of Candidate Genes in Bipolar Disorder</i> , PI: Chunyu Liu	2004- 2006
NARSAD Young Investigator Award (\$60,000) <i>Mutation Screening and Linkage Disequilibrium Analysis of Bipolar Disorder on 13q32</i> PI: Chunyu Liu	2001- 2003

Current Research Grant Support

U01 MH122591-01 (PI: Chunyu Liu)

.6 Calendar Months

NIH	\$99,476	04/01/20 -03/31/2025
1/3 High-resolution mapping of cell type-specific DNA (hydroxy)methylation in the human brain during postnatal development and in psychiatric disease		

- Single-base-pair resolution maps of DNA (hydroxy)methylation will be generated in sorted inhibitory and excitatory neurons at key time points of postnatal brain development as well as in the brains of psychiatrically normal individuals and patients with schizophrenia and bipolar disorder.
- Dr. Liu will coordinate data processing and perform analyses including QTL mapping, causal relationship, and GWAS integration.

1U01MH116489 (PI: Dan Geschwind) 8/17/2018 – 4/30/2023
 NIMH \$107,680 a year (for subward) **0.5 calendar month**
 2/2 Discovery and validation of neuronal enhancers associated with the development of psychiatric disorders.

R01MH122140 (PI: Chunyu Liu)
1.6 Calendar Months
 NIH \$494,902 a year 02/01/2022 - 03/30/2027
 Gene Expression Regulation in Brains of East Asian, African, and European Descent Explains Schizophrenia
 GWAS in Diverse Populations
 •This study aims to learn whether well-known racial differences in schizophrenia genetic risks are due to different risk genes between populations or due to the same risk genes represented by different genetic variants by studying genetic regulators of brain gene expression.
 • Dr. Liu will design the study, supervise data production and analysis, and organize publication and data release.

31871276 (PI: Chunyu Liu)
 National Natural Science Foundation of China 190,000 RMB (~\$30,000) a year 01/01/2019- 12/31/2022
 Gene expression regulatory networks and master regulators of the schizophrenia risk genes.
 No salary recovery

Pending

Center for Architecture of the Spliced Transcriptome in Human Brain Cells (CAST).
 Project Number: 1UM MH130974-01
 Name of PD/PI: Liu
 Project/Proposal Start and End Date: 07/01/22–06/30/2027
 Total Award Amount: \$23,817,271

An Atlas of RNA Elements and Their Regulatory Network Associated with ADRD
 Name of PD/PI: Xiajun Dong
 Primary Place of Performance: SUNY Upstate Medical University, New York State
 Project/Proposal Start and End Date: 07/01/22 – 06/30/2027
 Total Award Amount: \$407,150

Honors and Awards

Changjiang Scholar, Ministry of Education, China	2017
Chair’s Award for Most Outstanding Teaching 2007-2008	2008
Department of Psychiatry and Behavioral Neuroscience, the University of Chicago	
Southwest Florida Investigator Award 2002	2003
NARSAD Research Partners Program, National Alliance for Research on Schizophrenia and Depression, New York, USA	
National Outstanding Doctoral Dissertation Award	2000
Ministry of Education of P. R. China	
Cheung Kong Scholars Achievement Award	1999
Ministry of Education of P. R. China	

Professional Organizations

American College of Neuropsychopharmacology (Member since 2014)
(Education and Training Committee 2022 -)
Genetics Society of China
American Society of Human Genetics
Society of Biological Psychiatry
International Society of Psychiatric Genetics
(Global Diversity Taskforce, Committee Chair 2015 - 2021)
(IDEA committee 2022 -)
China International Exchange and Promotive Association for Medical and Health Care

Professional Service

Journal Academic Editor

PLoS One (2011 -)
Frontiers in Behavioral and Psychiatric Genetics (2014 -)
Genetics & Epigenetics (2014 -)
Journal of Psychiatry and Brain Science (2016 -)
American Journal of Medical Genetics, Part B (2017 -)

Ad hoc Reviewer

Genome Research
Molecular Psychiatry
Nature Communication
The American Journal of Human Genetics
The American Journal of Psychiatry
The New England Journal of Medicine
Biological Psychiatry
Schizophrenia Bulletin
Human Molecular Genetics
Nucleic Acids Research
Archives of General Psychiatry
Psychiatry Research
BioTechniques
Proteins: Structure, Function and Bioinformatics
Neuroscience Letters
Neurological Sciences
International Journal of Molecular Sciences
Progress in Neuro-Psychopharmacology & Biological Psychiatry
PLoS One
Bipolar Disorders
Journal of Psychopharmacology
BMC Research Notes
Journal of Cellular and Molecular Medicine

Neuroscience & Biobehavioral Reviews

Gene

Neurobiology of Disease

Pharmacogenomics

Frontiers in Neuroscience

Science

Grant Review

SUNY Upstate Medical University, Research Advisory Committee. 2018 - present

Permanent member: BGES 2016 - 2020

Ad hoc

Veteran Affair: Special Emphasis Panel on Million Veteran Program (SPLM) Subcommittee 2017

NIH Study section GHD 2016

NIH Special Emphasis Panel/Scientific Review Group 2010/01 ZRG1 BDCN-N (02); 2012/10 ZRG1 F08-Q (20); 2015 ZMH1-ERB-C; 2015 ZRG1 PSE-N

NIH Study section F08, 2012- 2013

NIH Study section MESH-BBBP-J 2014; BGES 2014, 2015;

NIH Study section GGG 2015

NIH Study section ZES1 2015

Mail reviewer for NIH 2013 ZRG1-BCMB-A 51 R.

ZRG1 MDCN-P (57) 2020

NIDA Core and Research Center of Excellence Grant Program (P30/50) 2020

NIH ZNS1 SRB-M (06) 2021

NIH Transformative Research Award (TRA) 2021

VA Mental Health & Behavior Sciences-A (MHBA) 2021

NIH ZRG1 PSE-H (70) R 2022

Medical Research Council (MRC, UK), Career Development Award Fellowship, NMHB Research Grant Ontario Mental Health Foundation (Canada), Innovational Research Incentives Scheme Vidi, WOTRO Science for Global Development (Netherlands), National Natural Science Foundation (China), French Funding Organization (France). PSI Foundation (Canada), Israel Science Foundation (Israel), Hong Kong Innovation and Technology Commission (China).

UIC, Fall 2012 Chancellor's Graduate Research Fellowship reviewer

Sessions Chaired at National and International Conferences/Symposia

- 1) Chair, Panel. Genetics of Intermediate Phenotypes in Psychosis. 60th American College of Neuropsychopharmacology. Annual Meeting. December 5-8, 2021.
- 2) Chair: PsychENCODE Updates: Discovery of Functional Elements for Psychiatric GWAS Signals. 2020 Virtual World Congress of Psychiatric Genetics. Virtual. October 16 – 22, 2020.
- 3) Chair, Education Session: microRNA: genetics, genomics and its relevance to psychiatry. XXIV. World Congress of Psychiatric Genetics. Jerusalem, Israel, Oct 28- Nov 3, 2016
- 4) Chair, Symposium: From Gene Expression to Disease Association. XXIV. World Congress of Psychiatric

- Genetics. Jerusalem, Israel, Oct 28- Nov 3, 2016
- 5) Co-chair, Symposium: Progress in Psychiatric Genetics in China. XXIV. World Congress of Psychiatric Genetics. Jerusalem, Israel, Oct 28- Nov 3, 2016
 - 6) Chair, Symposium: A Small Window to Psychiatric Genetics in China. XXII World Congress of Psychiatric Genetics. Copenhagen, Denmark, October 12-16, 2014
 - 7) Discussant, Symposium: Towards Translational Psychiatry: From Genomic Discoveries to Prediction of Treatment Response. XXII World Congress of Psychiatric Genetics. Copenhagen, Denmark, October 12-16, 2014
 - 8) Moderator, Chinese Psychiatric Genetics Summit, Changsha, Hunan, April 2-3, 2014
 - 9) Chair, Symposium: Integration of Genomics, Epigenomics and Genetics for a Biological Understanding of Psychiatric Diseases. Society of Biological Psychiatry. Philadelphia, May 3-5, 2012.
 - 10) Chair, Symposium: Functional Genomics in Post-GWAS of Neuropsychiatric Disease. XIX World Congress of Psychiatric Genetics. Washington DC, Sept 10-14, 2011.
 - 11) Chair, Symposium: DNA Methylation in Human Brain and Neuropsychiatric Diseases. XVII World Congress of Psychiatric Genetics. San Diego, November 3-8, 2009.
 - 12) Co-chair, Workshop: Critical Assessment of Massive Data Analysis (CAMDA 2009). Chicago. October 5-6, 2009.
 - 13) Chair, Symposium: microRNA in Human Brain and Neuropsychiatric Diseases. XVI World Congress of Psychiatric Genetics. Osaka, Japan. October 11-15, 2008.
 - 14) Co-chair, Symposium: Functional Studies of DAOA. 63rd Annual Meeting of Society of Biological Psychiatry. Washington DC, USA. May 1- 3, 2008.

Service to the community

Faculty Judge for the 2011, 2012 UIC Student Research Forum, College of Medicine
 Postdoc-Faculty Speed Mentoring Program
 Judge for the 2011 UIC Research Forum
 ACNP Education and Training Committee, 2022-

Teaching Experience

Course Developer and Instructor

Nanocourse: GS647-015 Nanocourse: Computer ABC for Biologists Graduate School, SUNY Upstate Medical University	2019-
Programming for Modern Biomedical Research (BIOS 20250) Biological Sciences Collegiate Division, the University of Chicago	2007- 2010

Course Lecturer

N631: Topics in Neuroscience (Genetics and epigenetics), Upstate	2022 -
PATH 512 Molecular Epidemiology And Biomarkers Of Disease: One lecture about microarray data analysis, University of Illinois at Chicago	2012
GCLS 504 Introduction to Research Methods: two lectures about microarray Graduation Education in Medical Sciences, University of Illinois at Chicago	2011 – 2012
Psychiatric Genetics Training Program course: Bioinformatics and Genomics Department of Psychiatry, the University of Chicago	2011 – 2013

Multidisciplinary Approaches to Psychiatric and Behavioral Genetics (PSYC 46900)	2003- 2009
Department of Psychiatry, the University of Chicago	
Residency Training Program in Psychiatry: Neurobiology sequence	2007- 2009
Department of Psychiatry, the University of Chicago Medical Center	

Research Mentor

Visiting Scientists

- Weihong Kuang, M.D., Associate Professor, Department of Psychiatry, West China Hospital of Sichuan University, Chengdu, P.R. China, June, 2007 - June, 2008
- Wei Luo, M.D., Ph.D., Associate Professor, Department of Neurology, Zhejiang University, 2nd affiliated hospital, Hangzhou, P.R. China, Aug, 2008 - Aug, 2009
- Ran Zhao, M.D., Ph.D., Professor, Department of Psychology, Central University of Finance and Economics, Beijing, P.R. China, Sept, 2011- Sept, 2012
- Zhengmao Hu, Ph.D. Associate Professor, Central South University, Changsha, China, Jan, 2013- Dec, 2013
- Jing Peng, M.D., Ph.D., Associate Professor, Central South University, Changsha, China, Dec, 2012- Oct, 2013
- Ning Yuan, M.D., Associate Professor, Hunan Brain Hospital, Changsha, China, 2016-2017
- ShunKai Lai, Ph.D. Candidate, Jinan University, Guangzhou, China, Feb – August, 2019
- Shishi Ming, Ph.D. Candidate, Central South University, Changsha, China, Oct 2019- Sept 2020
- Zhi Xu, Associate Professor, SouthEast University, Nanjing, China, Nov 2019 – Oct 2020

Postdoctoral Fellows

- Jiajun Shi, Ph.D. Oct. 2005 –Oct. 2008
- Lijun Cheng, Ph.D. Oct. 2005 –Oct. 2010
- Hongmin Zhu, Ph.D. Sept 2006 – Sept 2008
- Dandan Zhang, Ph.D. Nov. 2007.11- August, 2010
- Fabio Pibiri, Ph.D. Feb, 2009 –May, 2011
- Qi Chen, Ph.D. July, 2008–June, 2011
- Chao Chen, Ph.D. July, 2011- 2014
- Annie Shieh, Ph.D. August 2015 – 2018
- Shan Jiang, Ph.D. November 2015 - 2016
- Ramu Vadukapuram, Ph.D. 2016 - 2019

Graduate Students

Tang Haiyan, Meng Qingtuan, Bin Langman, Yan Jing, Central South University, 2014 – 2015
 Meng Qingtuan, Yan Xia, Chuan Jiao, Kangli Wang, Central South University, 2015 – 2019
 Cuihua Xia, Rujia Dai, Yi Jiang, Chaodong Ding, Central South University, 2015 – 2020

Visiting Graduate Students

Yongjun Wang, Ph.D. 2014-2015.
 Jinsong Tang, Ph.D. 2007-2008, Thesis: MRI Study on Brain Neuronetwork Connections in Early Onset Schizophrenia and DNA Copy Number Variations in Major Psychosis
 Chao Chen, Ph.D. 2008- 2011, Thesis: Evaluation of Six Batch Adjustment Methods in

Expression Microarray Data and Application of Gene Co-expression Network in Schizophrenia

Undergraduate Students

Christopher Armoskus, the University of Chicago, 2008

Karishma Furtado, the University of Chicago, 2009

Raj Patel, Loyola University, 2009

Lewie Zeng, the University of Chicago, 2009 - 2010

Kevin Zhang, the University of Chicago, 2009 - 2012

2012 Conte Center Summer Research Program

Ashwin Nayak, University of Illinois at Urbana-Champaign,

2012 Conte Center Summer Research Program

Alex Merlo, University of Illinois at Chicago, 2012

Noah Stephen Curtis, University of Illinois at Chicago, 2012

Lindsey Jay, University of Chicago, 2016

2016 Conte Center Summer Research Program

Pre-College Research Student Advisees, Illinois Mathematics and Science Academy

Ashley Ro, Nikita Veera, Shannon Tai, Kelly Yom, 2009 -2011

Summer interns

Jaden Tian (WashU), Zhanlin Chen (Yale), Lindsey Jay (U of C), Contessa Norris (Purdue) 2016

Basia Walenkiewicz (SUNY SURF program) 2018

Xiangyu Liao (UC Davis), Dina Silvestri (Syracuse University) 2021

Students that won awards:

Yan Xia (2016, ISPG Hugh Gurling Memorial Award)

Yan Xia (2017, ASHG Charles J. Epstein Awards)

Yan Xia (2017, ACGA Outstanding Graduate Student Presentation Award)

Chadong Ding (2018, ACGA Outstanding Graduate Student Presentation Award)

Rujia Dai (2018, ISPG Hugh Gurling Memorial Award)

Chuan Jia (2018, ASHG Charles J. Epstein Trainee Awards for Excellence in Human Genetics Research, Semifinalists)

Yu Chen (2021, ISPG Hugh Gurling Memorial Award)

Yu Chen (2021, ASHG Reviewer's Choice Award)

Current students at SUNY Upstate Medical University

Ph.D. candidates

- Rujia Dai (2019-)
- Gayathri Ganesh (2022 -)
- Yu Wei (2022 -)

MD/PhD candidates

- Tatiana Mikhailova (2022 -)

Extracurricular Education Program Created

BioMedical Programming Club: Biological Sciences Collegiate Division, the University of Chicago.

2007- 2010

Invited Presentations

1. Genetics and brain epigenetics of psychiatric disorders and human behavior. University of Connecticut. Animal Science Seminar. Sept 10, 2021. Online.
2. PsychENCODE and Psychiatric Genetics. WCPG. PGC Day. Online. Oct 12-14, 2021.
3. Basic Science Research of Psychiatry. Keynote speaker for the Annual meeting of Society of Basic Theory in Chinese Medicine. Online. Nov 3, 2020.
4. Functional Annotation of GWAS Findings in Psychiatric Disorders. Program in Quantitative Genomics Seminar. Harvard School of Public Health. Online. Nov 17. 2020.
5. Genetics and Epigenetics of Drug Response. SUNY Upstate Medical University, Department of Psychiatry. Grand Round. Feb 7, 2019
6. Studies of Genetics and Epigenetics of Drug Response. The University of Chicago HongKong Center. April 15, 2019
7. Open Science and Big Data. Zhejiang University. Hanzhou, China April 21. 2019.
8. Studies of Genetics and Epigenetics of Drug Response. Hunan Society of Psychiatrist 2019 Annual Meeting. Changsha, China, June 1st 2019
9. Current Psychiatric Genetics, Epigenetics, and Cellular Models. Chinese Society of Psychiatry 2019 Annual Meeting, Nanjing, China. August 29-31, 2019.
10. Psychiatric Genetics, Epigenetics, and Cellular Models in Coming Years. Beijing Normal University. Beijing China, Sept 6, 2019.
11. Psychiatric GWAS, PsychENCODE, and After. University of Texas at Houston. Oct 4th. 2019
12. Multi –omics of psychiatric disorders, SUNY Upstate Medical University, 18th annual Biomedical Sciences Retreat on September 11, 2018, Welch Allyn in Skaneateles, NY, USA
13. Aproaching Psychiatric Genetics. SURF Researrch Seminar, SUNY Upstate Medical University, June 28th, 2018, Syracuse, NY, USA
14. Precision Medicine on DSM5-based Psychiatric Disorders. 2018 Annual Sino-American Convention for the Diagnosis and Treatment of Major Mental Illnesses. May 25th, 2018, Beijing, China
15. Updates on PsychENCODE - a path to functional interpretation of GWAS signals. The Fifth Chinese Psychiatric Genetics Summit, Suzhou Mental Health Hospital, May 17th, 2018, Suzhou, Jiangsu, China.
16. Ten Years of Psychiatric GWAS and After. Vanderbilt Genetics Institute. Feb 21, 2018. Nashville, Tennessee. USA
17. Psychiatric GWAS and Clinical Implications. SUNY Upstate Psychiatry Grand Round. March 1, 2018. Syracuse, New York. USA
18. From Genetics to Regulatory Networks Underlying Psychiatric Disorders. Beijing Pediatric Research Institute. June 15, 2017. Beijing, China.
19. Brain Functional Annotation of Genetic Variants/Mutations. Xiangya International Summit of Pediatric Neurology. June 10, 2017. Changsha, China
20. Current Genetics of Schizophrenia from phenotype to gene, and drug targets. Xiangya Workshop on Schizophrenia. April 21, 2017. Changsha, China
21. Epigenetics of Psychiatric Disorders. 50th Winter Conference on Brain Research. January 28- Feb 2, 2017. Big Sky, Montana.

22. Coexpression Network Analyses Identify Regulatory Systems Associated With Psychiatric Disorders. American College of Neuropsychopharmacology Panel: Using Human Genetics GWAS and Expression Data to Drive Discovery. Dec 4-8, 2016. Hollywood, Florida,
23. From gene expression regulation networks to genetic association of psychiatric disorders. 14th Annual CBC Symposium. November 11, 2016. Chicago.
24. GWAS of Psychiatry: the told and untold. Retreat of the T32 Training Program. The University of Chicago. April 15, 2016. Chicago
25. Progress of genetics and epigenetics of psychiatric disorders. Forum for Development and Innovation of Mental Health Research. Huilongguan Hospital. May 8-9, 2015. Beijing, China.
26. Genetics and epigenetics of psychiatric disorders, today and future. Kangning Mental Hospital. May 7, 2015. Shenzhen, China.
27. Research progress of genetics and epigenetics of psychiatric disorders. Hunan Mental Hospital. April 30, 2015. Changsha, China.
28. Circadian rhythms of DNA methylation and gene expression. Global Chinese Congress of Genetics. October 29 to November 1st, 2014. Changsha, Hunan, China
29. Somatic mutation in Alzheimer's brain. Forum of Neuronal Circuitry of Emotion and Memory. November 8-9, 2014. Changsha, Hunan, China
30. De novo mutations in neuropsychiatric disorders. Xiangya Forum of Pediatric Neurology. November 8-9, 2014. Changsha, Hunan, China
31. Current psychiatric genetics and future perspective. Society of Chinese Traditional Medicine. August 21-23, 2014. Hanzhou, Zhejiang, China
32. Brain expression QTLs and GWAS signals of psychiatric diseases. International Conference on Functional and Comparative Genomics & Pharmacogenomics, November 12-14, 2013. North Shore, Chicago
33. Current genetics of Alzheimer's diseases. Neuropsychiatry Forum. July 6th, 2013. 148th Hospital. Zibo, Shangdong, China
34. Brain molecular QTL mapping and GWAS of psychiatric diseases, Symposium at Society of Biological Psychiatry, 68th Annual Scientific Convention, May 16-18, 2013. San Francisco, California
35. Sexually dimorphic DNA methylation sites are conserved across human brain and blood, Symposium at Society of Biological Psychiatry, 68th Annual Scientific Convention, May 16-18, 2013. San Francisco, California
36. Integrate genetics, epigenetics and genomics to understand psychiatric disorders. Psychiatry Institute Neuroscience Seminar. June 18th, 2012. University of Illinois at Chicago.
37. Brain eQTLs and function-based GWAS of bipolar disorder identified novel disease risk genes. American College of Neuropsychopharmacology Panel: Beyond genome-wide association studies - - new approaches to risk of psychiatric illness Dec 4-8, 2011. Waikoloa, Hawaii
38. Genetics of gene expression regulation and bipolar disorder. August 21, 2011. Beijing University. 6th Mental Health Hospital. Beijing, China
39. Genetics of gene expression regulation and bipolar disorder. March 21, 2011. Anschutz Medical Campus, University of Colorado, Colorado.
40. Genetics and epigenetics of schizophrenia and bipolar disorder. Nov. 24, 2009. Fudan University, State Key Laboratory of Genetic Engineering, Shanghai, China

41. Genetics and epigenetics of schizophrenia and bipolar disorder. Nov. 19, 2009. Kunming Primate Research Center, Chinese Academy of Sciences, Kunming, China
42. Current genetics and epigenetics of schizophrenia and bipolar disorder. Nov. 23, 2009. Wenzhou Medical College, Genomic Medicine Institute, Wenzhou, Zhejiang, China
43. Genetics and epigenetics of schizophrenia and bipolar disorder. Nov. 24, 2009. Bio-X Center, Shanghai Jiaotong University, Shanghai, China
44. MicroRNA genomic variations in psychiatric diseases. Symposium at the International Congress on Schizophrenia Research. March 28-31, 2009. San Diego, California
45. Genetics of schizophrenia and bipolar disorders. NCIBI (National Center for Integrative Biomedical informatics) Workshop on Translational Bioinformatics. July 29-30, 2009. Ann Arbor, Michigan
46. Sequencing in the study of psychiatric diseases. Symposium, Next Generation Sequencing Data Analysis and Exploring Next Generation Sequencing meetings. Sept. 21-23, 2009. Providence, Rhode Island
47. Genome-wide association study in psychiatric disorders. Lecture, Oct. 6, 2008. Institute of Mental Health, Central South University. Changsha, China
48. Association study of trace amine receptors in bipolar disorder. Symposium at the CINP International College of Neuropsychopharmacology. July 9-13, 2006. Chicago, Illinois

Patents Obtained:(Patent title: International Application Number: International Publication Number)

1. Human M6b1 gene: PCT/CN97/00107: WO 99/21982
2. Human Atrophin-1 related gene: PCT/CN97/00108: WO 99/21983
3. Exostosin-4 gene: PCT/CN97/00126: WO 99/25822
4. An Ataxin-2 like gene: PCT/CN98/00009: WO 99/36527
5. Isoform 1 of Dimethylglycine dehydrogenase-like gene: PCT/CN98/00040: WO 99/47559
6. Isoform 2 of Dimethylglycine dehydrogenase-like gene: PCT/CN98/00041: WO 99/47560
7. Connexin 31, Human Gap Junction Protein beta-3: PCT/CN98/00056: WO 99/51634
8. Connexin 31.1, Human Gap Junction Protein beta-4: PCT/CN98/00055: WO 99/51738

Copyright Held

SNP-Cruncher: SNP Data Mining Toolkit, the University of Chicago, 2001

Bioinformatics Web/Applications Developed

<http://bioinfo.psych.uic.edu/> for a group of tools

R package: Meth27QC, Teng L., Chen C., Liu C*. 2011.

Bibliography (H-index 45)

Peer-reviewed Journal Articles (updated 01/01/2022)

58 first or corresponding (*) author

1. Yu Chen, Jiacheng Dai, Longfei Tang, Tatiana Mikhailova, Qiuman Liang, Miao Li, Jiaqi Zhou, Jinghong Qiu, Richard Kopp, Cynthia Weickert, Chao Chen*, Chunyu Liu*. Neuroimmune transcriptome changes in patient brains of psychiatric and neurological disorders. 2022. (submitted)
2. Cuihua Xia, Rujia Dai, Jing Yu, Chunling Zhang, Ma-li Wong*, Chunyu Liu*. Dysregulated spliceosome gene expression may be a common process in brains of neurological and psychiatric disorders.

Discover Mental Health. 2022 (revision)

3. **Chunyu Liu***, Elliot S. Gershon. Disease-related Phenotypes for Genetic Studies of Psychiatric Disorders. *Molecular Psychiatry*. (revision)
4. Chao Chen, Qingtuan Meng, Wendiao Zhang, Xuan Wang, Chuan Jiao, Sheng Xu, Chunyu Liu, and Beisha Tang. Human forebrain organoids reveal connections between valproic acid exposure and autism risk. *Translational Psychiatry*. (accepted)
5. Yun Zhang, Chunyu Liu*. Evaluating the challenges and reproducibility of studies investigating DNA methylation signatures of psychological stress. *Epigenomics*. 2022. (Online)
6. Haixia Gu, Xue'er Ma, Jingjing Zhao, Chunyu Liu*. A Meta-analysis of Salivary Cortisol Responses in the Trier Social Stress Test to Evaluate the Effects of Speech Topics, Sex, and Sample Size. *Comprehensive Psychoneuroendocrinology*. Volume 10, May 2022.
7. Kangli Wang, Rujia Dai, Yan Xia, Jianghua Tian, Chuan Jiao, Tatiana Mikhailova, Chunling Zhang, Chao Chen, **Chunyu Liu***. Spatiotemporal specificity of correlated DNA methylation and gene expression pairs across different human tissues and stages of brain development. *Epigenetics*. 2021. (online).
8. Lulu Chen, Chiung-Ting Wu, Chia-Hsiang Lin, Rujia Dai, Chunyu Liu, Robert Clarke, Guoqiang Yu, Jennifer E Van Eyk, David M Herrington, Yue Wang. swCAM: estimation of subtype-specific expressions in individual samples with unsupervised sample-wise deconvolution. *Bioinformatics*. 2021 (in press)
9. Shishi Min; Zongchang Li; Annie Shieh; Gina Giase; Riyue Bao; Chunling Zhang; Liz Kuney; Richard Kopp; Huma Asif; Ney Alliey Rodriguez; Lixia Qin; David Wesley Craig; Geoffrey J. Faulkner; Elliot S. Gershon; Beisha Tang; Chao Chen; **Chunyu Liu***. Absence of coding somatic single nucleotide variants within well-known candidate genes in late-onset sporadic Alzheimer's Disease based on the analysis of multi-omics data. *Neurobiology of Aging*. 2021 (accepted)
10. Jiaqi Zhou, Miao Li, Xueying Wang, Yuwen He, Yan Xia, John A Sweeney, Richard F Kopp, **Chunyu Liu*** and Chao Chen*. Drug response-related DNA methylation changes in schizophrenia, bipolar disorder and major depressive disorder. *Frontiers in Neuroscience - Neurogenomics*. 2021.
11. Xianjun Dong*, **Chunyu Liu**, Mikhail Dozmorov*. Multi-omics Data Resources and Integration for Human Brain Disorders. *Briefings in Functional Genomics*. 2021
12. Todd Lencz, Max Lam, Chia-Yen Chen, tian ge, Xia Yan, David Hill, Joey Trampush, Jin Yu, Emma Knowles, Gail Davies, Eli Stahl, Laura Huckins, David Liewald, Srdjan Djurovic, Ingrid Melle, Andrea Christoforou, Ivar Reinvang, Pamela DeRosse, Astri Lundervold, Vidar Steen, Thomas Espeseth, Katri Raikonen, Elisabeth Widen, Aarno Palotie, Johan Eriksson, Ina Giegling, Bettina Konte, Annette Hartmann, Panos Roussos, Stella Giakoumaki, Katherine Burdick, Antony Payton, William Ollier, Ornit Chiba-Falek, Deborah Koltai, Anna Need, Elizabeth Cirulli, Aristotle Voineskos, Nikos Stefanis, Dimitrios Avramopoulos, Alex Hatzimanolis, Nikolaos Smyrnis, Robert Bilder, nelson freimer, Tyrone Cannon, Edythe London, Russell Poldrack, Fred Sabb, Eliza Congdon, Emily Drabant Conley, Matthew Scult, Dwight Dickinson, Richard Straub, Gary Donohoe, Derek Morris, Aiden Corvin, Michael Gill, Ahmad Hariri, Daniel Weinberger, Neil Pendleton, Panos Bitsios, Dan Rujescu, Jari Lahti, Stephanie Hellard, Matthew Keller, Ole Andreassen, Ian Deary, David Glahn, Hailiang Huang, **Chunyu Liu**, and Anil Maholtra. Identifying Nootropic Drug Targets via Large-Scale Cognitive GWAS and Transcriptomics. *Neuropsychopharmacology*. 2021
13. Jiacheng Dai, Chao Chen, Yu Chen, Rujia Dai, Yi Jiang, Jianghua Tian, Sihan Liu, Meng Xu, Miao Li, Jiaqi

- Zhou, **Chunyu Liu***. Agonal factors distort gene-expression patterns in human postmortem brains. *Frontiers Neuroscience*. 2021.
14. Chaodong Ding, Chunling Zhang, Richard Kopp, Liz Kuney, Qingtuan Meng, Le Wang, Yan Xia, Yi Jiang, Rujia Dai, Shishi Min, Wei-Dong Yao, Ma-Li Wong, Hongyu Ruan*, **Chunyu Liu***, Chao Chen*. Transcription factor POU3F2 regulates TRIM8 expression contributing to cellular functions implicated in schizophrenia. *Molecular Psychiatry*. 2020. (online)
 15. Qingtuan Meng, Le Wang, Rujia Dai, Jiawen Wang, Zongyao Ren, Sihan Liu, Yan Xia, Yi Jiang, Fangyuan Duan, Kangli Wang, **Chunyu Liu***, Chao Chen*. Integrative analyses prioritize GNL3 as a risk gene for bipolar disorder. *Molecular Psychiatry*. 2020. (online)
 16. Jiacheng Dai, Yu Chen, Cuihua Xia, Jiaqi Zhou, **Chunyu Liu***, Chao Chen*. Digital Sensory Phenotyping for Psychiatric Disorders. *Journal of Psychiatry and Brain Science* 2020;5(3):e200015
 17. Yi Jiang, Gina Giase, Kay Grennan, Annie Shieh, Yan Xia, Lide Han, Quan Wang, Qiang Wei, Rui Chen, Sihan Liu, Chao Chen*, Bingshan Li*, **Chunyu Liu***. DRAMS: A Tool to Detect and Re-Align Mixed-up Samples for Integrative Studies of Multi-omics Data. *PLOS Computational Biology*. 2020 (online)
 18. Dan Zhou, Yi Jiang, Xue Zhong, Nancy J Cox, **Chunyu Liu**, Eric R Gamazon. A unified framework for joint-tissue transcriptome-wide association and Mendelian randomization analysis. *Nat Genet*. 2020 Oct 5. doi: 10.1038/s41588-020-0706-2. Online ahead of print.
 19. Huma Asif, Ney Alliey-Rodriguez, Sarah Keedy, Carol Tamminga, John Sweeney, Godfrey Pearlson, Brett Clementz, Matcheri Keshavan, Peter Buckley, Chunyu Liu, Benjamin Neale, and Elliot Gershon. GWAS Significance Thresholds for Deep Phenotyping Studies Can Depend Upon Minor Allele Frequencies and Sample Size. *Molecular Psychiatry*. 2020. (online)
 20. Ney Alliey-Rodriguez, Tamar Grey, Rebecca Shafee, Huma Asif, Olivia Lutz, Nicolas Bolo, Jaya Padmanabhan, Neeraj Tandon, Madeline Klinger, Katherine Reis, Jonathan Spring, Lucas Coppes, Victor Zeng, Rachal Hegde, Dung Hoang, Deepthi Bannai, Uzma Nawaz, Philip Henson, Siyuan Liu, Diane Gage, Steven McCarroll, Jeffrey Bishop, Scot Hill, James Reilly, Rebekka Lencer, Brett Clementz, Peter Buckley, David Glahn, Shashwath Meda, Balaji Narayanan, Godfrey Pearlson, Matcheri Keshavan, Elena Ivleva, Carol Tamminga, John Sweeney, David Curtis, Judith Badner, Sarah Keedy, Judith Rapoport, Chunyu Liu, and Elliot Gershon. NRXN1 is associated with enlargement of the temporal horns of the lateral ventricles in psychosis. *Translational Psychiatry*. 2019. 9(1):230.
 21. **Chunyu Liu***; Stephen V. Faraone; Stephen J. Glatt. Psychiatric Genetics, Epigenetics, and Cellular Models in Coming Years. *Journal of Psychiatry and Brain Science* 2019;4(4):e190012
 22. Bryois J, Garrett ME, Song L, Safi A, Giusti-Rodriguez P, Johnson GD, Shieh AW, Buil A, Fullard JF, Roussos P, Sklar P, Akbarian S, Haroutunian V, Stockmeier CA, Wray GA, White KP, Liu C, Reddy TE, Ashley-Koch A, Sullivan PF, Crawford GE. Evaluation of chromatin accessibility in prefrontal cortex of individuals with schizophrenia. *Nat Commun*. 2018 Aug 7;9(1):3121. doi: 10.1038/s41467-018-05379-y. PubMed PMID: 30087329; PubMed Central PMCID: PMC6081462.
 23. Yan Xia, Rujia Dai, Kangli Wang, Chuan Jiao, Yuchen Xu, Honglei Li, Xi Jing, Chunling Zhang, Chao Chen* and **Chunyu Liu***. Sex-differential DNA methylation and associated regulation networks in human brain implicated in the sex-biased risks of psychiatric disorders. *Mol Psychiatry*. 2019. Apr 11. doi: 10.1038/s41380-019-0416-2
 24. Quan Wang , Rui Chen , Feixiong Cheng , Qiang Wei , Ying Ji , Dr. Hai Yang , Xue Zhong , Ran Tao , Dr. Zhexing Wen , Dr. James Sutcliffe , Dr. Chunyu Liu , Prof. Edwin Cook , Nancy Cox. A Bayesian framework that integrates multi-omics data and gene networks predicts risk genes from schizophrenia GWAS data. *Nature Neuroscience*. 2019. 22(5):691-699.
 25. Guoqiang Yu, David Miller, Chiung-Ting Wu, Eric Hoffman, Chunyu Liu, David Herrington, and Yue Wang. Asymmetric independence modeling identifies novel gene-environment interactions. *Scientific Reports*. 2019. 9(1):2455
 26. Yuan N, Chen Y, Xia Y, Dai J, **Liu C***. Inflammation-related biomarkers in major psychiatric disorders: a cross-disorder assessment of reproducibility and specificity in 43 meta-analyses. *Translational Psychiatry*. 2019. 9(1):233
 27. Gandal MJ*, Zhang P, Hadjimichael E, Walker RL, Chen C, Liu S, Won H, van Bakel H, Varghese M, Wang Y, Shieh AW, Haney J, Parhami S, Belmont J, Kim M, Moran Losada P, Khan Z, Mleczko J, Xia Y, Dai R, Wang D, Yang YT, Xu M, Fish K, Hof PR, Warrell J, Fitzgerald D, White K, Jaffe AE; PsychENCODE Consortium, Peters MA, Gerstein M, **Liu C***, Iakoucheva LM*, Pinto D*, Geschwind DH*. Transcriptome-wide isoform-level dysregulation in ASD, schizophrenia, and bipolar disorder. *Science*. 2018.
 28. Meng Q, Wang K, Brunetti T, Xia Y, Jiao C, Dai R, Fitzgerald D, Thomas A, Jay L, Eckart H, Grennan K,

- Imamura-Kawasawa Y, Li M, Sestan N, White KP, Chen C*, **Liu C***. (2018). The DGCR5 long noncoding RNA may regulate expression of several schizophrenia-related genes. *Science Translational Medicine*. 2018. 10(472)
29. Chen C*, Meng Q, Xia Y, Ding C, Wang L, Dai R, Cheng L, Gunaratne P, Gibbs RA, Min S, Coarfa C, Reid JG, Zhang C, Jiao C, Jiang Y, Giase G, Thomas A, Fitzgerald D, Brunetti T, Shieh A, Xia C, Wang Y, Wang Y, Badner JA, Gershon ES, White KP, **Liu C***. The transcription factor POU3F2 regulates a gene coexpression network in brain tissue from patients with psychiatric disorders. *Science Translational Medicine* 2018. 10(472)
 30. Li J, Hu S, Zhang K, Shi L, Zhang Y, Zhao T, Wang L, He X, Xia K, **Liu C***, Sun Z. A comparative study of the genetic components of three subcategories of autism spectrum disorder. *Mol Psychiatry*. 2018.
 31. Jiao C, Zhang C, Dai R, Xia Y, Wang K, Giase G, Chen C, **Liu C***. Positional effects revealed in Illumina methylation array and the impact on analysis. *Epigenomics*. 2018;10(5):643-59. PMC6021926.
 32. Jiao C, Yan P, Xia C, Shen Z, Tan Z, Tan Y, Wang K, Jiang Y, Huang L, Dai R, Wei Y, Xia Y, Meng Q, Ouyang Y, Yi L, Duan F, Dai J, Zhao S, **Liu C***, Chen C. BrainEXP: a database featuring with spatiotemporal expression variations and co-expression organizations in human brains. *Bioinformatics*. 2018.
 33. Horwitz T, Lam K, Chen Y, Xia Y, **Liu C***. A decade in psychiatric GWAS research. *Mol Psychiatry*. 2019 Mar;24(3):378-389.
 34. Gandal MJ, Haney JR, Parikshak NN, Leppa V, Ramaswami G, Hartl C, Schork AJ, Appadurai V, Buil A, Werge TM, Liu C, White KP, CommonMind C, Psych EC, i P-BWG, Horvath S, Geschwind DH. Shared molecular neuropathology across major psychiatric disorders parallels polygenic overlap. *Science*. 2018;359(6376):693-7. PMC5898828.
 35. Cen Z, Jiang Z, Chen Y, Zheng X, Xie F, Yang X, Lu X, Ouyang Z, Wu H, Chen S, Yin H, Qiu X, Wang S, Ding M, Tang Y, Yu F, Li C, Wang T, Ishiura H, Tsuji S, Jiao C, **Liu C**, Xiao J, Luo W. Intronic pentanucleotide TTTCA repeat insertion in the SAMD12 gene causes familial cortical myoclonic tremor with epilepsy type 1. *Brain*. 2018.
 36. Tang J, Fan Y, Li H, Xiang Q, Zhang DF, Li Z, He Y, Liao Y, Wang Y, He F, Zhang F, Shugart YY, **Liu C**, Tang Y, Chan RCK, Wang CY, Yao YG, Chen X. Whole-genome sequencing of monozygotic twins discordant for schizophrenia indicates multiple genetic risk factors for schizophrenia. *J Genet Genomics*. 2017;44(6):295-306.
 37. Li J, Wang L, Guo H, Shi L, Zhang K, Tang M, Hu S, Dong S, Liu Y, Wang T, Yu P, He X, Hu Z, Zhao J, **Liu C***, Sun ZS, Xia K. Targeted sequencing and functional analysis reveal brain-size-related genes and their networks in autism spectrum disorders. *Mol Psychiatry*. 2017;22(9):1282-90.
 38. Alliey-Rodriguez N, Grey TA, Shafee R, Padmanabhan J, Tandon N, Klinger M, Spring J, Coppes L, Reis K, Keshavan MS. Common variants of NRXN1, LRP1B and RORA are associated with increased ventricular volumes in psychosis-GWAS findings from the B-SNIP deep phenotyping study. *bioRxiv*. 2017:175489.
 39. Zarayeneh N, Oh JH, Kim D, Liu C, Gao J, Suh SC, Kang M, editors. Integrative Gene Regulatory Network inference using multi-omics data. *Bioinformatics and Biomedicine (BIBM)*, 2016 IEEE International Conference on; 2016: IEEE.
 40. **Liu C***. A Case for Core Genes. *Journal of Psychiatry and Brain Science*. 2017.
 41. **Liu C***, Grennan KS, Gershon ES. Current Practice of Genetics Research of Psychiatric Disorders. *Journal of Psychiatry and Brain Science*. 2016.
 42. Kim D, Kang M, Biswas A, **Liu C**, Gao J. Integrative approach for inference of gene regulatory networks using lasso-based random featurizing and application to psychiatric disorders. *BMC Med Genomics*. 2016.
 43. PsychENCODE Consortium, Akbarian S, **Liu C**, Knowles JA, Vaccarino FM, Farnham PJ, Crawford GE, Jaffe AE, Pinto D, Dracheva S, Geschwind DH, Mill J, Nairn AC, Abyzov A, Pochareddy S, Prabhakar S, Weissman S, Sullivan PF, State MW, Weng Z, Peters MA, White KP, Gerstein MB, Amiri A, Armoskus C, Ashley-Koch AE, Bae T, Beckel-Mitchener A, Berman BP, Coetzee GA, Coppola G, Francoeur N, Fromer M, Gao R, Grennan K, Herstein J, Kavanagh DH, Ivanov NA, Jiang Y, Kitchen RR, Kozlenkov A, Kundakovic M, Li M, Li Z, Liu S, Mangravite LM, Mattei E, Markenscoff-Papadimitriou E, Navarro FC, North N, Omberg L, Panchision D, Parikshak N, Poschmann J, Price AJ, Purcaro M, Reddy TE, Roussos P, Schreiner S, Scuderi S, Sebra R, Shibata M, Shieh AW, Skarica M, Sun W, Swarup V, Thomas A, Tsuji J, van Bakel H, Wang D, Wang Y, Wang K, Werling DM, Willsey AJ, Witt H, Won H, Wong CC, Wray GA, Wu EY, Xu X, Yao L, Senthil G, Lehner T, Sklar P, Sestan N. *Nat Neurosci*. 2015 Dec;18(12):1707-12. The PsychENCODE project.
 44. Wang N, Hoffman EP, Chen L, Chen L, Zhang Z, **Liu C**, Yu G, Herrington DM, Clarke R, Wang Y.

- Mathematical modelling of transcriptional heterogeneity identifies novel markers and subpopulations in complex tissues. *Sci Rep.* 2016;6:18909. PMC4703969.
45. **Liu C***, Saffen D, Schulze TG, Burmeister M, Sham PC, Yao YG, Kuo PH, Chen C, An Y, Dai J, Yue W, Li MX, Xue H, Su B, Chen L, Shi Y, Qiao M, Liu T, Xia K, Chan RCK. Psychiatric genetics in China: achievements and challenges. *Mol Psychiatry.* 2016;21(1):4-9. PMC4830695.
 46. Li M, Luo XJ, Landén M, Bergen SE, Hultman CM, Li X, Zhang W, Yao YG, Zhang C, Liu J, Mattheisen M, Cichon S, Mühleisen TW, Degenhardt FA, Nöthen MM, Schulze TG, Grigoriou-Serbanescu M, Li H, Fuller CK, Chen C, Dong Q, Chen C, Jamain S, Leboyer M, Bellivier F, Etain B, Kahn JP, Henry C, Preisig M, Kutalik Z, Castelao E, Wright A, Mitchell PB, Fullerton JM, Schofield PR, Montgomery GW, Medland SE, Gordon SD, Martin NG; MooDS Consortium; Swedish Bipolar Study Group, Rietschel M, **Liu C**, Kleinman JE, Hyde TM, Weinberger DR, Su B. Impact of a cis-associated gene expression SNP on chromosome 20q11.22 on bipolar disorder susceptibility, hippocampal structure and cognitive performance. *Br J Psychiatry.* 2015 Sep 3. [Epub ahead of print]
 47. Li J, Cai T, Jiang Y, Chen H, He X, Chen C, Li X, Shao Q, Ran X, Li Z, Xia K, Liu C, Sun ZS, Wu J. Genes with de novo mutations are shared by four neuropsychiatric disorders discovered from NPdenovo database. *Mol Psychiatry.* 2016;21(2):290-7. PMC4837654.
 48. **Liu C***, Chung M. Genetics and epigenetics of circadian rhythms and their potential roles in neuropsychiatric disorders. *Neurosci Bull.* 2015 Feb;31(1):141-59.
 49. Kang M, Zhang C, Chun H-W, Ding C, **Liu C**, Gao J. eQTL epistasis: detecting epistatic effects and inferring hierarchical relationships of genes in biological pathways. *Bioinformatics.* 2015;31(5):656-64.
 50. Chen C, Zhang C, Cheng L, Reilly JL, Bishop JR, Sweeney JA, Chen HY, Gershon ES, **Liu C***. Correlation between DNA methylation and gene expression in the brains of patients with bipolar disorder and schizophrenia. *Bipolar Disord.* 2014;16(8):790-9. PMC4302408.
 51. Zhao Z, Xu J, Chen J, Kim S, Reimers M, Bacanu SA, Yu H, **Liu C**, Sun J, Wang Q, Jia P, Xu F, Zhang Y, Kendler KS, Peng Z, Chen X. Transcriptome sequencing and genome-wide association analyses reveal lysosomal function and actin cytoskeleton remodeling in schizophrenia and bipolar disorder. *Mol Psychiatry.* 2015;20(5):563-72. PMC4326626.
 52. Kim DC, Wang J, **Liu C**, Gao J. Inference of SNP-gene regulatory networks by integrating gene expressions and genetic perturbations. *Biomed Res Int.* 2014;2014:629697. PMC4127230.
 53. Yu D, Mathews CA, Scharf JM, Neale BM, Davis LK, Gamazon ER, Derks EM, Evans P, Edlund CK, Crane J, Fagerness JA, Osiecki L, Gallagher P, Gerber G, Haddad S, Illmann C, McGrath LM, Mayerfeld C, Arepalli S, Barlassina C, Barr CL, Bellodi L, Benarroch F, Berrio GB, Bienvenu OJ, Black DW, Bloch MH, Brentani H, Bruun RD, Budman CL, Camarena B, Campbell DD, Cappi C, Silgado JC, Cavallini MC, Chavira DA, Chouinard S, Cook EH, Cookson MR, Coric V, Cullen B, Cusi D, Delorme R, Denys D, Dion Y, Eapen V, Egberts K, Falkai P, Fernandez T, Fournier E, Garrido H, Geller D, Gilbert DL, Girard SL, Grabe HJ, Grados MA, Greenberg BD, Gross-Tsur V, Grunblatt E, Hardy J, Heiman GA, Hemmings SM, Herrera LD, Hezel DM, Hoekstra PJ, Jankovic J, Kennedy JL, King RA, Konkashbaev AI, Kremeyer B, Kurlan R, Lanzagorta N, Leboyer M, Leckman JF, Lennertz L, **Liu C**, Lochner C, Lowe TL, Lupoli S, Macciardi F, Maier W, Manunta P, Marconi M, McCracken JT, Mesa Restrepo SC, Moessner R, Moorjani P, Morgan J, Muller H, Murphy DL, Naarden AL, Nurmi E, Ochoa WC, Ophoff RA, Pakstis AJ, Pato MT, Pato CN, Piacentini J, Pittenger C, Pollak Y, Rauch SL, Renner T, Reus VI, Richter MA, Riddle MA, Robertson MM, Romero R, Rosario MC, Rosenberg D, Ruhrmann S, Sabatti C, Salvi E, Sampaio AS, Samuels J, Sandor P, Service SK, Sheppard B, Singer HS, Smit JH, Stein DJ, Strengman E, Tischfield JA, Turiel M, Valencia Duarte AV, Vallada H, Veenstra-VanderWeele J, Walitza S, Wang Y, Weale M, Weiss R, Wendland JR, Westenberg HG, Shugart YY, Hounie AG, Miguel EC, Nicolini H, Wagner M, Ruiz-Linares A, Cath DC, McMahon W, Posthuma D, Oostra BA, Nestadt G, Rouleau GA, Purcell S, Jenike MA, Heutink P, Hanna GL, Conti DV, Arnold PD, Freimer NB, Stewart SE, Knowles JA, Cox NJ, Pauls DL. Cross-disorder genome-wide analyses suggest a complex genetic relationship between Tourette's syndrome and OCD. *Am J Psychiatry.* 2015;172(1):82-93. PMC4282594.
 54. Grennan, K. S., Chen, C., Gershon, E. S. and **Liu, C***. (2014), Molecular network analysis enhances understanding of the biology of mental disorders. *Bioessays*, 36: 606–616.
 55. Lencz T, Guha S, **Liu C**, Rosenfeld J, DeRosse P, John M, Kirtley A, Cheng L, Zhang C, Badner J, Ikeda M, Iwata N, Cichon S, Rietschel M, Nöthen M, Cheng A, Hodgkinson C, Yuan Q, Kane1 J, Lee A, Pisanté A, Gregersen P, Pe'er I, Malhotra A, Goldman A, Darvasi A. (2013) Genome-wide association study implicates NDST3 in Schizophrenia and Bipolar disorder. *Nature Communication.* 4:2739.
 56. Liu X, Chen Q, Tsai HJ; Wang G, Hong X, Zhou Y, Zhang C, **Liu C**, Liu R, Wang H, Zhang S, Yu Y,

- Mestan K, Pearson C, Otlans P, Zuckerman B, Wang X. (2013) Maternal Preconception Body Mass Index and Offspring Cord Blood DNA Methylation: Exploration of Early Life Origins of Disease. *Environmental and Molecular Mutagenesis*. 2013 Nov 15. doi: 10.1002/em.21827. [Epub ahead of print]
57. Davis LK, Yu D, Keenan CL, Gamazon ER, Konkashbaev AI, Derks EM, Neale BM, Yang J, Lee SH, Evans P, Barr CL, Bellodi L, Benarroch F, Berrio GB, Bienvenu OJ, Bloch MH, Blom RM, Bruun RD, Budman CL, Camarena B, Campbell D, Cappi C, Cardona Silgado JC, Cath DC, Cavallini MC, Chavira DA, Chouinard S, Conti DV, Cook EH, Coric V, Cullen BA, Deforce D, Delorme R, Dion Y, Edlund CK, Egberts K, Falkai P, Fernandez TV, Gallagher PJ, Garrido H, Geller D, Girard SL, Grabe HJ, Grados MA, Greenberg BD, Gross-Tsur V, Haddad S, Heiman GA, Hemmings SM, Hounie AG, Illmann C, Jankovic J, Jenike MA, Kennedy JL, King RA, Kremeyer B, Kurlan R, Lanzagorta N, Leboyer M, Leckman JF, Lennertz L, **Liu C**, Lochner C, Lowe TL, Macciardi F, McCracken JT, McGrath LM, Mesa Restrepo SC, Moessner R, Morgan J, Muller H, Murphy DL, Naarden AL, Ochoa WC, Ophoff RA, Osiecki L, Pakstis AJ, Pato MT, Pato CN, Piacentini J, Pittenger C, Pollak Y, Rauch SL, Renner TJ, Reus VI, Richter MA, Riddle MA, Robertson MM, Romero R, Rosàrio MC, Rosenberg D, Rouleau GA, Ruhrmann S, Ruiz-Linares A, Sampaio AS, Samuels J, Sandor P, Sheppard B, Singer HS, Smit JH, Stein DJ, Strengman E, Tischfield JA, Valencia Duarte AV, Vallada H, Van Nieuwerburgh F, Veenstra-Vanderweele J, Walitza S, Wang Y, Wendland JR, Westenberg HG, Shugart YY, Miguel EC, McMahon W, Wagner M, Nicolini H, Posthuma D, Hanna GL, Heutink P, Denys D, Arnold PD, Oostra BA, Nestadt G, Freimer NB, Pauls DL, Wray NR, Stewart SE, Mathews CA, Knowles JA, Cox NJ, Scharf JM. (2013) Partitioning the heritability of Tourette syndrome and obsessive compulsive disorder reveals differences in genetic architecture. *PLoS Genet*. 9(10):e1003864.
 58. Gershon ES*, Grennan K, Busnello J, Badner JA, Ovsiew F, Memon S, Alliey-Rodriguez N, Cooper J, Romanos B, **Liu C***. (2014) A rare mutation of CACNA1C in a patient with Bipolar disorder, and decreased gene expression associated with a Bipolar-associated common SNP of CACNA1C in brain. *Mol Psychiatry*. 19(8):890-4.
 59. Cross-Disorder Group of the Psychiatric Genomics Consortium*. (2013). Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. *Nature Genetics*. 45(9):984-94
 60. Qiu R, Chen C, Jiang H, Shen L, Wu M, **Liu C***. (2013). Large genomic region free of GWAS-based common variants contains fertility-related genes. *PLoS One*. 8(4):e61917
 61. Cheng L, Hattori E, Nakajima A, Woehrle N, Opal MD, Zhang C, Grennan K, Dulawa SC, Tang YP, Gershon ES, **Liu C***. (2014). Expression of the G72/G30 gene in transgenic mice induces behavioral changes. *Mol Psychiatry*. 19(2):175-83.
 62. Chen C, Cheng L, Grennan K, Pibiri F, Zhang C, Badner JA, Gershon ES, **Liu C***. (2013). Two Gene Co-expression Modules Robustly Differentiate Psychotics and Controls. *Mol Psychiatry*. 18(12):1308-14.
 63. Gamazon ER, Badner JA, Cheng L, Zhang C, Zhang D, Cox NJ, Gershon ES, Kelsoe JR, Greenwood TA, Nievergelt CM, Chen C, McKinney R, Shilling PD, Schork NJ, Smith EN, Bloss CS, Nurnberger JI, Edenberg HJ, Foroud T, Koller DL, Scheftner WA, Coryell W, Rice J, Lawson WB, Nwulia EA, Hipolito M, Byerley W, McMahon FJ, Schulze TG, Berrettini WH, Potash JB, Zandi PP, Mahon PB, McInnis MG, Zollner S, Zhang P, Craig DW, Szelinger S, Barrett TB, **Liu C***. (2013). Enrichment of cis-regulatory gene expression SNPs and methylation quantitative trait loci among bipolar disorder susceptibility variants. *Mol Psychiatry*. 18(3):340-6
 64. Gao Y, Galante M, El-Mallakh J, Nurnberger JI, Jr., Delamere NA, Lei Z, El-Mallakh RS, **BiGS Consortium**. (2012). BDNF expression in lymphoblastoid cell lines carrying BDNF SNPs associated with bipolar disorder. *Psychiatr Genet* 22: 253-255.
 65. Lahey BB, Michalska KJ, **Liu C**, Chen Q, Hipwell AE, Chronis-Tuscano A, Waldman ID, Decety J. (2012): Preliminary genetic imaging study of the association between estrogen receptor-alpha gene polymorphisms and harsh human maternal parenting. *Neurosci Lett* 525: 17-22.
 66. Davis LK, Gamazon ER, Kistner-Griffin E, Badner JA, **Liu C**, Cook EH, Sutcliffe JS, Cox NJ. (2012): Loci nominally associated with autism from genome-wide analysis show enrichment of brain expression quantitative trait loci but not lymphoblastoid cell line expression quantitative trait loci. *Mol Autism* 3: 3.
 67. Meier S, Mattheisen M, Vassos E, Strohmaier J, Treutlein J, Josef F, Breuer R, Degenhardt F, Muhleisen TW, Muller-Myhsok B, Steffens M, Schmael C, McMahon FJ, Kelsoe JR, Greenwood TA, Nievergelt CM, Barrett TB, McKinney R, Shilling PD, Schork NJ, Smith EN, Bloss CS, Nurnberger J, Edenberg HJ, Foroud T, Koller DL, Gershon ES, **Liu C**, Badner JA, Scheftner W, Lawson WB, Nwulia EA, Hipolito M, Coryell W, Rice J, Byerley W, McMahon F, Chen DT, Schulze TG, Berrettini W, Potash JB, Zandi PP, Mahon PB, McInnis M, Craig D, Szelinger S, Nothen MM, Cichon S, Schulze TG, Rietschel M. (2012): Genome-wide

- significant association between a 'negative mood delusions' dimension in bipolar disorder and genetic variation on chromosome 3q26.1. *Transl Psychiatry* 2: e165.
68. Nissen S, Liang S, Shekhtman T, Kelsoe JR, **BiGS Consortium**. (2012): Evidence for association of bipolar disorder to haplotypes in the 22q12.3 region near the genes stargazin, ift27 and parvalbumin. *Am J Med Genet B Neuropsychiatr Genet*.2012 Dec;159B(8):941-50. doi: 10.1002/ajmg.b.32099.
 69. Pirooznia M, Seifuddin F, Judy J, Mahon PB, **BiGS Consortium**, Potash JB, Zandi PP. (2012): Data mining approaches for genome-wide association of mood disorders. *Psychiatr Genet* 22: 55-61.
 70. Scharf JM, Yu D, Mathews CA, Neale BM, Stewart SE, Fagerness JA, Evans P, Gamazon E, Edlund CK, Service SK, Tikhomirov A, Osiecki L, Illmann C, Pluzhnikov A, Konkashbaev A, Davis LK, Han B, Crane J, Moorjani P, Crenshaw AT, Parkin MA, Reus VI, Lowe TL, Rangel-Lugo M, Chouinard S, Dion Y, Girard S, Cath DC, Smit JH, King RA, Fernandez TV, Leckman JF, Kidd KK, Kidd JR, Pakstis AJ, State MW, Herrera LD, Romero R, Fournier E, Sandor P, Barr CL, Phan N, Gross-Tsur V, Benarroch F, Pollak Y, Budman CL, Bruun RD, Erenberg G, Naarden AL, Lee PC, Weiss N, Kremeyer B, Berrio GB, Campbell DD, Cardona Silgado JC, Ochoa WC, Mesa Restrepo SC, Muller H, Valencia Duarte AV, Lyon GJ, Leppert M, Morgan J, Weiss R, Grados MA, Anderson K, Davarya S, Singer H, Walkup J, Jankovic J, Tischfield JA, Heiman GA, Gilbert DL, Hoekstra PJ, Robertson MM, Kurlan R, **Liu C**, Gibbs JR, Singleton A, for the North American Brain Expression Consortium, Hardy J, for the UK Human Brain Expression Database, Strengman E, Ophoff RA, Wagner M, Moessner R, Mirel DB, Posthuma D, Sabatti C, Eskin E, Conti DV, Knowles JA, Ruiz-Linares A, Rouleau GA, Purcell S, Heutink P, Oostra BA, McMahon WM, Freimer NB, Cox NJ, Pauls DL. (2013): Genome-wide association study of Tourette's syndrome. *Mol Psychiatry*. 18(6):721-8
 71. Stewart SE, Yu D, Scharf JM, Neale BM, Fagerness JA, Mathews CA, Arnold PD, Evans PD, Gamazon ER, Osiecki L, McGrath L, Haddad S, Crane J, Hezel D, Illman C, Mayerfeld C, Konkashbaev A, **Liu C**, Pluzhnikov A, Tikhomirov A, Edlund CK, Rauch SL, Moessner R, Falkai P, Maier W, Ruhrmann S, Grabe HJ, Lennertz L, Wagner M, Bellodi L, Cavallini MC, Richter MA, Cook EH, Jr., Kennedy JL, Rosenberg D, Stein DJ, Hemmings SM, Lochner C, Azzam A, Chavira DA, Fournier E, Garrido H, Sheppard B, Umana P, Murphy DL, Wendland JR, Veenstra-Vanderweele J, Denys D, Blom R, Deforce D, Van NF, Westenberg HG, Walitza S, Egberts K, Renner T, Miguel EC, Cappi C, Hounie AG, Conceicao do RM, Sampaio AS, Vallada H, Nicolini H, Lanzagorta N, Camarena B, Delorme R, Leboyer M, Pato CN, Pato MT, Voyiaziakis E, Heutink P, Cath DC, Posthuma D, Smit JH, Samuels J, Bienvenu OJ, Cullen B, Fyer AJ, Grados MA, Greenberg BD, McCracken JT, Riddle MA, Wang Y, Coric V, Leckman JF, Bloch M, Pittenger C, Eapen V, Black DW, Ophoff RA, Strengman E, Cusi D, Turiel M, Frau F, Macciardi F, Gibbs JR, Cookson MR, Singleton A, for the North American Brain Expression Consortium, Hardy J, for the UK Brain Expression Database, Crenshaw AT, Parkin MA, Mirel DB, Conti DV, Purcell S, Nestadt G, Hanna GL, Jenike MA, Knowles JA, Cox N, Pauls DL (2013): Genome-wide association study of obsessive-compulsive disorder. *Mol Psychiatry*. 18(7):788-98.
 72. Akula N, Baranova A, Seto D, Solka J, Nalls MA, Singleton A, Ferrucci L, Tanaka T, Bandinelli S, Cho YS, Kim YJ, Lee JY, Han BG, **BiGS Consortium**, WTCCC, McMahon FJ. (2011): A network-based approach to prioritize results from genome-wide association studies. *PLoS One* 6: e24220.
 73. Alliey-Rodriguez N, Zhang D, Badner JA, Lahey BB, Zhang X, Dinwiddie S, Romanos B, Plenys N, **Liu C**, Gershon ES. (2011): Genome-wide association study of personality traits in bipolar patients. *Psychiatr Genet* 21: 190-194.
 74. Badner JA, Koller D, Foroud T, Edenberg H, Nurnberger JI, Jr., Zandi PP, Willour VL, McMahon FJ, Potash JB, Hamshere M, Grozeva D, Green E, Kirov G, Jones I, Jones L, Craddock N, Morris D, Segurado R, Gill M, Sadovnick D, Remick R, Keck P, Kelsoe J, Ayub M, Maclean A, Blackwood D, **Liu CY**, Gershon ES, McMahon W, Lyon GJ, Robinson R, Ross J, Byerley W. (2012): Genome-wide linkage analysis of 972 bipolar pedigrees using single-nucleotide polymorphisms. *Mol Psychiatry*. 17(8):818-26. doi: 10.1038/mp.2011.89
 75. Belmonte MP, Pirooznia M, Goes FS, Seifuddin F, Steele J, Lee PH, Huang J, Hamshere ML, **BiGS Consortium**, Depaulo JR, Jr., Kelsoe JR, Rietschel M, Nothen M, Cichon S, Gurling H, Purcell S, Smoller JW, Craddock N, Schulze TG, McMahon FJ, Potash JB, Zandi PP. (2011): Genome-wide association analysis of age at onset and psychotic symptoms in bipolar disorder. *Am J Med Genet B Neuropsychiatr Genet* 156B: 370-378.
 76. Chen C, Grennan K, Badner J, Zhang D, Gershon E, Jin L, **Liu C***. (2011): Removing batch effects in analysis of expression microarray data: an evaluation of six batch adjustment methods. *PLoS One* 6: e17238.

77. Chen DT, Jiang X, Akula N, Shugart YY, Wendland JR, Steele CJ, Kassem L, Park JH, Chatterjee N, Jamain S, Cheng A, Leboyer M, Muglia P, Schulze TG, Cichon S, Nothen MM, Rietschel M, **BiGS Consortium**, McMahon FJ. (2011): Genome-wide association study meta-analysis of European and Asian-ancestry samples identifies three novel loci associated with bipolar disorder. *Mol Psychiatry*. 2011 Dec 20. doi: 10.1038/mp.2011.157.
78. Cichon S, Muhleisen TW, Degenhardt FA, Mattheisen M, Miro X, Strohmaier J, Steffens M, Meesters C, Herms S, Weingarten M, Priebe L, Haenisch B, Alexander M, Vollmer J, Breuer R, Schmal C, Tessmann P, Moebus S, Wichmann HE, Schreiber S, Muller-Myhsok B, Lucae S, Jamain S, Leboyer M, Bellivier F, Etain B, Henry C, Kahn JP, Heath S, **BiGS Consortium**, Hamshere M, O'Donovan MC, Owen MJ, Craddock N, Schwarz M, Vedder H, Kammerer-Ciernioch J, Reif A, Sasse J, Bauer M, Hautzinger M, Wright A, Mitchell PB, Schofield PR, Montgomery GW, Medland SE, Gordon SD, Martin NG, Gustafsson O, Andreassen O, Djurovic S, Sigurdsson E, Steinberg S, Stefansson H, Stefansson K, Kapur-Pojkic L, Oruc L, Rivas F, Mayoral F, Chuchalin A, Babadjanova G, Tiganov AS, Pantelejeva G, Abramova LI, Grigoriou-Serbanescu M, Diaconu CC, Czerski PM, Hauser J, Zimmer A, Lathrop M, Schulze TG, Wienker TF, Schumacher J, Maier W, Propping P, Rietschel M, Nothen MM. (2011): Genome-wide association study identifies genetic variation in neurocan as a susceptibility factor for bipolar disorder. *Am J Hum Genet* 88: 372-381.
79. Gershon ES, Iley-Rodriguez N, **Liu C**. (2011): After GWAS: searching for genetic risk for schizophrenia and bipolar disorder. *Am J Psychiatry* 168: 253-256.
80. **Liu C***. (2011): Brain expression quantitative trait locus mapping informs genetic studies of psychiatric diseases. *Neurosci Bull* 27: 123-133.
81. Nwulia EA, Hipolito MM, Aamir S, Lawson WB, Nurnberger JI, Jr., **BiGS Consortium**. (2011): Ethnic disparities in the perception of ethical risks from psychiatric genetic studies. *Am J Med Genet B Neuropsychiatr Genet* 156B: 569-580.
82. Sklar P, Ripke S, Scott LJ, Andreassen OA, Cichon S, Craddock N, Edenberg HJ, Nurnberger JI, Jr., Rietschel M, Blackwood D, Corvin A, Flickinger M, Guan W, Mattingsdal M, McQuillin A, Kwan P, Wienker TF, Daly M, Dudbridge F, Holmans PA, Lin D, Burmeister M, Greenwood TA, Hamshere ML, Muglia P, Smith EN, Zandi PP, Nievergelt CM, McKinney R, Shilling PD, Schork NJ, Bloss CS, Foroud T, Koller DL, Gershon ES, **Liu C**, Badner JA, Scheftner WA, Lawson WB, Nwulia EA, Hipolito M, Coryell W, Rice J, Byerley W, McMahon FJ, Schulze TG, Berrettini W, Lohoff FW, Potash JB, Mahon PB, McInnis MG, Zollner S, Zhang P, Craig DW, Szlinger S, Barrett TB, Breuer R, Meier S, Strohmaier J, Witt SH, Tozzi F, Farmer A, McGuffin P, Strauss J, Xu W, Kennedy JL, Vincent JB, Matthews K, Day R, Ferreira MA, O'Dushlaine C, Perlis R, Raychaudhuri S, Ruderfer D, Hyoun PL, Smoller JW, Li J, Absher D, Thompson RC, Meng FG, Schatzberg AF, Bunney WE, Barchas JD, Jones EG, Watson SJ, Myers RM, Akil H, Boehnke M, Chambert K, Moran J, Scolnick E, Djurovic S, Melle I, Morken G, Gill M, Morris D, Quinn E, Muhleisen TW, Degenhardt FA, Mattheisen M, Schumacher J, Maier W, Steffens M, Propping P, Nothen MM, Anjorin A, Bass N, Gurling H, Kandaswamy R, Lawrence J, McGhee K, McIntosh A, McLean AW, Muir WJ, Pickard BS, Breen G, St CD, Caesar S, Gordon-Smith K, Jones L, Fraser C, Green EK, Grozeva D, Jones IR, Kirov G, Moskvina V, Nikolov I, O'Donovan MC, Owen MJ, Collier DA, Elkin A, Williamson R, Young AH, Ferrier IN, Stefansson K, Stefansson H, Thornorgeirsson T, Steinberg S, Gustafsson O, Bergen SE, Nimgaonkar V, Hultman C, Landen M, Lichtenstein P, Sullivan P, Schalling M, Osby U, Backlund L, Frisen L, Langstrom N, Jamain S, Leboyer M, Etain B, Bellivier F, Petursson H, Sigur SE, Muller-Myhsok B, Lucae S, Schwarz M, Schofield PR, Martin N, Montgomery GW, Lathrop M, Oskarsson H, Bauer M, Wright A, Mitchell PB, Hautzinger M, Reif A, Kelsoe JR, Purcell SM, Psychiatric GWAS Consortium Bipolar Disorder Working Group. (2011): Large-scale genome-wide association analysis of bipolar disorder identifies a new susceptibility locus near ODZ4. *Nat Genet* 43: 977-983.
83. Smith EN, Koller DL, Panganiban C, Szlinger S, Zhang P, Badner JA, Barrett TB, Berrettini WH, Bloss CS, Byerley W, Coryell W, Edenberg HJ, Foroud T, Gershon ES, Greenwood TA, Guo Y, Hipolito M, Keating BJ, Lawson WB, **Liu C**, Mahon PB, McInnis MG, McMahon FJ, McKinney R, Murray SS, Nievergelt CM, Nurnberger JI, Jr., Nwulia EA, Potash JB, Rice J, Schulze TG, Scheftner WA, Shilling PD, Zandi PP, Zollner S, Craig DW, Schork NJ, Kelsoe JR. (2011): Genome-wide association of Bipolar disorder suggests an enrichment of replicable associations in regions near genes. *PLoS Genet* 7: e1002134.
84. Zhang D, Qian Y, Akula N, Iley-Rodriguez N, Tang J, Bipolar GS, Gershon ES, **Liu C***. (2011): Accuracy of CNV Detection from GWAS Data. *PLoS One* 6: e14511.
85. Dao DT, Mahon PB, Cai X, Kovacsics CE, Blackwell RA, Arad M, Shi J, Zandi PP, O'Donnell P, **BiGS Consortium**, Knowles JA, Weissman MM, Coryell W, Scheftner WA, Lawson WB, Levinson DF, Thompson SM, Potash JB, Gould TD (2010): Mood disorder susceptibility gene CACNA1C modifies mood-related

- behaviors in mice and interacts with sex to influence behavior in mice and diagnosis in humans. *Biol Psychiatry* 68: 801-810.
86. Liu C*, Cheng L, Badner JA, Zhang D, Craig DW, Redman M, Gershon ES. (2010): Whole-genome association mapping of gene expression in the human prefrontal cortex. *Mol Psychiatry* 15: 779-784.
 87. Piletz JE, Zhang X, Ranade R, Liu C. (2010): Database of genetic studies of bipolar disorder. *Psychiatr Genet*.
 88. Zhang D, Cheng L, Craig DW, Redman M, Liu C*. (2010): Cerebellar telomere length and psychiatric disorders. *Behav Genet* 40: 250-254.
 89. Zhang D, Cheng L, Badner JA, Chen C, Chen Q, Luo W, Craig DW, Redman M, Gershon ES, Liu C*. (2010): Genetic control of individual differences in gene-specific methylation in human brain. *Am J Hum Genet* 86: 411-419.
 90. Maheshwari M, Shi J, Badner JA, Skol A, Willour VL, Muzny DM, Wheeler DA, Gerald FR, Detera-Wadleigh S, McMahon FJ, Potash JB, Gershon ES, Liu C*, Gibbs RA. (2009): Common and rare variants of DAOA in bipolar disorder. *Am J Med Genet B Neuropsychiatr Genet* 150B: 960-966.
 91. Mahon PB, Payne JL, MacKinnon DF, Mondimore FM, Goes FS, Schweizer B, Jancic D, NIMH Genetics Initiative Bipolar Disorder Consortium, BiGS Consortium, Coryell WH, Holmans PA, Shi J, Knowles JA, Scheftner WA, Weissman MM, Levinson DF, Depaulo JR, Jr., Zandi PP, Potash JB. (2009): Genome-wide linkage and follow-up association study of postpartum mood symptoms. *Am J Psychiatry* 166: 1229-1237.
 92. Shi J, Badner JA, Willour VL, Potash JB, Gershon ES, Liu C*. (2009): Further evidence for an association of G72/G30 with schizophrenia in Chinese. *Schizophr Res* 107: 324-326.
 93. Smith EN, Bloss CS, Badner JA, Barrett T, Belmonte PL, Berrettini W, Byerley W, Coryell W, Craig D, Edenberg HJ, Eskin E, Foroud T, Gershon E, Greenwood TA, Hipolito M, Koller DL, Lawson WB, Liu C, Lohoff F, McInnis MG, McMahon FJ, Mirel DB, Murray SS, Nievergelt C, Nurnberger J, Nwulia EA, Paschall J, Potash JB, Rice J, Schulze TG, Scheftner W, Panganiban C, Zaitlen N, Zandi PP, Zollner S, Schork NJ, Kelsoe JR. (2009): Genome-wide association study of bipolar disorder in European American and African American individuals. *Mol Psychiatry* 14: 755-763.
 94. Zhang D, Cheng L, Qian Y, Aliey-Rodriguez N, Kelsoe JR, Greenwood T, Nievergelt C, Barrett TB, McKinney R, Schork N, Smith EN, Bloss C, Nurnberger J, Edenberg HJ, Foroud T, Scheftner W, Lawson WB, Nwulia EA, Hipolito M, Coryell W, Rice J, Byerley W, McMahon F, Schulze TG, Berrettini W, Potash JB, Belmonte PL, Zandi PP, McInnis MG, Zollner S, Craig D, Szlinger S, Koller D, Christian SL, Liu C*, Gershon ES. (2009): Singleton deletions throughout the genome increase risk of bipolar disorder. *Mol Psychiatry* 14: 376-380.
 95. Gershon ES, Liu C, Badner JA. (2008): Genome-wide association in bipolar. *Mol Psychiatry* 13: 1-2.
 96. Shi J, Badner JA, Liu C*. (2008): PDLIM5 and susceptibility to bipolar disorder: a family-based association study and meta-analysis. *Psychiatr Genet* 18: 116-121.
 97. Shi J, Wittke-Thompson JK, Badner JA, Hattori E, Potash JB, Willour VL, McMahon FJ, Gershon ES, Liu C*. (2008): Clock genes may influence bipolar disorder susceptibility and dysfunctional circadian rhythm. *Am J Med Genet B Neuropsychiatr Genet* 147B: 1047-1055.
 98. Shi J, Badner JA, Hattori E, Potash JB, Willour VL, McMahon FJ, Gershon ES, Liu C*. (2008): Neurotransmission and bipolar disorder: a systematic family-based association study. *Am J Med Genet B Neuropsychiatr Genet* 147B: 1270-1277.
 99. Shi J, Badner JA, Gershon ES, Liu C*. (2008): Allelic association of G72/G30 with schizophrenia and bipolar disorder: a comprehensive meta-analysis. *Schizophr Res* 98: 89-97.
 100. Shi J, Gershon ES, Liu C*. (2008): Genetic associations with schizophrenia: meta-analyses of 12 candidate genes. *Schizophr Res* 104: 96-107.
 101. Detera-Wadleigh SD, Liu C, Maheshwari M, Cardona I, Corona W, Akula N, Steele CJ, Badner JA, Kundu M, Kassem L, Potash JB, Gibbs R, Gershon ES, McMahon FJ, NIMH Genetics Initiative for Bipolar Disorder Consortium. (2007): Sequence variation in DOCK9 and heterogeneity in bipolar disorder. *Psychiatr Genet* 17: 274-286.
 102. Liu C*, Shi J, Badner JA, Zou H, Qian Y, Gershon ES. (2007): No association of trace amine receptor genes with bipolar disorder. *Mol Psychiatry* 12: 979-981.
 103. Shi J, Hattori E, Zou H, Badner JA, Christian SL, Gershon ES, Liu C*. (2007): No evidence for association between 19 cholinergic genes and bipolar disorder. *Am J Med Genet B Neuropsychiatr Genet* 144B: 715-723.
 104. Yao F, Zhang R, Zhu Z, Xia K, Liu C*. (2006): MutScreener: Primer design tool for PCR-direct sequencing. *Nucleic Acids Res* 34: W660-W664.

105. Ferraren DO, **Liu C**, Badner JA, Corona W, Rezvani A, Monje VD, Gershon ES, Bonner TI, Detera-Wadleigh SD. (2005): Linkage disequilibrium analysis in the LOC93081-KDEL1-BIVM region on 13q in bipolar disorder. *Am J Med Genet B Neuropsychiatr Genet* 133B: 12-17.
106. Hattori E, **Liu C**, Zhu H, Gershon ES. (2005): Genetic tests of biologic systems in affective disorders. *Mol Psychiatry* 10: 719-740.
107. Zhang R, Zhu Z, Zhu H, Nguyen T, Yao F, Xia K, Liang D, **Liu C***. (2005): SNP Cutter: a comprehensive tool for SNP PCR-RFLP assay design. *Nucleic Acids Res* 33: W489-W492.
108. Nguyen TH, **Liu C**, Gershon ES, McMahon FJ. (2004): Frequency Finder: a multi-source web application for collection of public allele frequencies of SNP markers. *Bioinformatics* 20: 439-443.
109. Hattori E, **Liu C**, Badner JA, Bonner TI, Christian SL, Maheshwari M, Detera-Wadleigh SD, Gibbs RA, Gershon ES. (2003): Polymorphisms at the G72/G30 gene locus, on 13q33, are associated with bipolar disorder in two independent pedigree series. *Am J Hum Genet* 72: 1131-1140.
110. **Liu C***, Bonner TI, Nguyen T, Lyons JL, Christian SL, Gershon ES. (2003): DNannotator: Annotation software tool kit for regional genomic sequences. *Nucleic Acids Res* 31: 3729-3735.
111. Christian SL, McDonough J, **Liu C**, Shaikh S, Vlamakis V, Badner JA, Chakravarti A, Gershon ES. (2002): An evaluation of the assembly of an approximately 15-Mb region on human chromosome 13q32-q33 linked to bipolar disorder and schizophrenia. *Genomics* 79: 635-656.
112. Maheshwari M, Christian SL, **Liu C**, Badner JA, Detera-Wadleigh S, Gershon ES, Gibbs RA. (2002): Mutation screening of two candidate genes from 13q32 in families affected with Bipolar disorder: human peptide transporter (SLC15A1) and human glypican5 (GPC5). *BMC Genomics* 3: 30.
113. **Liu C***, Badner JA, Christian SL, Guroff JJ, Detera-Wadleigh SD, Gershon ES. (2001): Fine mapping supports previous linkage evidence for a bipolar disorder susceptibility locus on 13q32. *Am J Med Genet* 105: 375-380.
114. Tang B, **Liu C**, Shen L, Dai H, Pan Q, Jing L, Ouyang S, Xia J. (2000): Frequency of SCA1, SCA2, SCA3/MJD, SCA6, SCA7, and DRPLA CAG trinucleotide repeat expansion in patients with hereditary spinocerebellar ataxia from Chinese kindreds. *Arch Neurol* 57: 540-544.
115. Li Y, Lu X, Xia J, Tang X, **Liu C**, Shi X. (1999): [STK11 gene mutation in Chinese with PJS]. *Zhonghua Yi Xue Za Zhi* 79: 425-427.
116. Tang B, Xia J, Wang D, Tang X, Shen L, **Liu C**, Dai H, Yan X, Pan Q, Xiao J, Zhang B, Ou Y. (1999): [CAG trinucleotide mutation detection in patients with hereditary spinocerebellar ataxia]. *Zhonghua Yi Xue Yi Chuan Xue Za Zhi* 16: 281-284.
117. Xia JH, **Liu C**, Ruan QG, Pan Q, Liao XD, Fu JJ, Cui F, Deng HX. (1999): [Molecular cloning of one splicing form of human M6b cDNA]. *Yi Chuan Xue Bao* 26: 439-446.
118. Xia JH, Yu KP, **Liu C**, Pan Q, Zheng D, Dai HP. (1999): [Molecular cloning of the human dimethylglycine dehydrogenase-like gene (DMGDHL1) from the sarcosinemia critical region at 9q34]. *Yi Chuan Xue Bao* 26: 591-597.
119. Xia JH, **Liu C**, Wang D, Ruan Q, Cui F, Xie W, Pan Q, Liao X, Dai HP, Deng H. (1999): Molecular cloning and localization of human atrophin-1-like gene. *Progress in Natural Science* 9: 203-210.
120. Cui F, **Liu C**, Xia J. (1998): Identification and localization of four retropseudogenes of the human polypeptide elongation factor 1alpha. *Hereditas (Beijing)* 20: 6-10.
121. **Liu C**, Chen Y, Gui M, Zhang C. (1998): Studies on the mechanism of variation of hybrids of domesticated silkworm and eri silkworm: RAPD analysis of genome. *Hereditas (Beijing)* 20: 5-8.
122. **Liu C**, Zhang C, Xia J. (1998): Identification and mapping of human auxilin by integrated data analysis. *Prog Biochem Biophys* 25: 434-439.
123. Xia J, Yang X, Ruan Q, Pan Q, **Liu C**, Xie W, Deng H. (1998): [Molecular cloning and characterization of novel protein kinase gene DYRK3]. *Zhonghua Yi Xue Yi Chuan Xue Za Zhi* 15: 327-332.
124. Xia J, Wang D, **Liu C**, Pan Q, Dai H, Deng H. (1998): Molecular cloning and Localization of human ataxin-2-like gene. *High Technology Letters* 4: 100-104.
125. Xia JH, **Liu C**, Tang BS, Pan Q, Huang L, Dai HP, Zhang BR, Xie W, Hu DX, Zheng D, Shi XL, Wang DA, Xia K, Yu KP, Liao XD, Feng Y, Yang YF, Xiao JY, Xie DH, Huang JZ. (1998): Mutations in the gene encoding gap junction protein beta-3 associated with autosomal dominant hearing impairment. *Nat Genet* 20: 370-373.
126. Zhang C, Chen Y, Gui M, **Liu C**. (1998): RAPD analysis of hereditary and variation of domesticated silkworm generated by introduction of eri silkworm DNA. *Hereditas (Beijing)* 20: 1-4.
127. Dong L, **Liu C**, Xia J. (1997): Progress of large scale sequencing of human genome. *Foreign Medical Sciences: Genetics* 20: 281-286.

128. **Liu C**, Zhang C, Chen Y. (1997): RAPD analysis of eri silkworm and domesticated silkworm. *Acta Sericologica Sinica* 23: 215-220.
129. Tang B, **Liu C**, Xia J, Pan Q, Long Z. (1997): The SCA1 mutation detection and analysis in the patients with hereditary spinocerebellar ataxia in one Chinese family. *Chinese J Neurol* 30: 110-113.
130. **Liu C**. (1996): Bioinformatics and human genome project in HGM'96. *Foreign Medical Sciences: Genetics* 19: 262-263.
131. **Liu C**, Tang B, Xia J. (1996): Progress in research on genes responsible for hereditary spinocerebellar ataxia. *Foreign Medical Sciences: Genetics* 19: 16-19.
132. **Liu C**, Zhang C, Xia J. (1996): Application of Random Primers in the Research of Molecular Biology. *Prog Biochem Biophys* 23: 517-520.

Note: * corresponding author

BiGS Consortium: Kelsoe JR, Greenwood TA, Nievergelt CM, McKinney R, Shilling PD, Schork NJ, Smith EN, Bloss CS, Nurnberger JI Jr, Edenberg HJ, Foroud T, Koller DL, Gershon ES, **Liu C**, Badner JA, Scheftner WA, Lawson WB, Nwulia EA, Hipolito M, Coryell W, Rice J, Byerley W, McMahon FJ, Schulze TG, Berrettini WH, Potash JB, Zandi PP, Mahon PB, McInnis MG, Zöllner S, Zhang P, Craig DW, Szlinger S, Barrett TB.

Invited Reviews/Editorials

1. Gershon ES, **Liu C**, Badner JA (2008): Genome-wide association in bipolar disorder. *Mol Psychiatry* 13: 1-2.
2. Tang J and **Liu C**. Genome-wide association studies of neuropsychiatric diseases. The 5th Symposium for Chinese Neuroscientists Worldwide. Progress of Neuroscience (5). 2008. 7-25-2008.

Book Chapters

1. Grennan K, Zhang F, **Liu C***.(2012): microRNAs and Their Potential Roles in Schizophrenia and Bipolar Disorder. miRNAs and human diseases. Kerala, India: Research Signpost. Editors: Junming Yue and Lu Lu.
2. **Liu C***. (2012): QTL Mapping of Molecular Traits for Studies of Human Complex Diseases. In Translational Bioinformatics: Computational Genomics. Springer. Editor: Yin Yao.
3. Shi J, Cheng L, Gershon ES, **Liu C***. (2008): G72/G30 in Neuropsychiatric Disorders. In Javitt, DC; Kantrowitz, J; Lajtha, A. editors. *Handbook of Neurochemistry and Molecular Neurobiology*: Springer.
4. **Liu C**, Piletz JE. (2006): Strategies and technologies for genetic study of complex diseases *Recent Developments in Nucleic Acids Research*. Kerala, India: Research Signpost, pp 267-305.
5. **Liu C***. (2004): DNannotator: Annotation software tool kit for regional genomic sequences. In Jurgen Fuchs, Maurizio Podda, editors. *Encyclopedia of Medical Genomics and Proteomics*. New York: Marcel Dekker, Inc.
6. **Liu C***, Zhou H. (2005) DNA genotyping. In *Experimental Protocols for Medical Molecular Biology in Chinese and English*. Edited by Wei Zheng. Peking Union Medical College Press. 215-244
7. **Liu C***, Chen C. (2014) Multi-gene inheritance: Medical Genetics (Chinese textbook).
8. Zhang D, **Liu C**. et al. (2014) DSM5 Desk Reference (Chinese version). Beijing University Press.
9. **Liu C***, Jiao C, Wang K, Yuan N. (2017) DNA Methylation and Psychiatric Disorders. In Epigenetics and Psychiatric Disease, Progress in Molecular Biology and Translational Science. volume 158. ELSEVIER. Editor: Dennis Grayson.

Abstracts

1. Haixia Gu, Chunyu Liu. New High-Throughput Method to Induce Robust Psychological Stress. 60th American College of Neuropsychopharmacology. Annual Meeting. December 5-8, 2021.
2. Longfei Tang, Yu Chen, Cuihua Xia, Chao Chen, Chunyu Liu. Comparing Gene Expression Changes in Blood and Brain of Schizophrenia. 2021 Virtual World Congress of Psychiatric Genetics, Oct 11-15. Virtual
3. Cindy Wen, Rujia Dai, Pawel F. Przytycki, Minsoo Kim, Arjun Bhattacharya, Pan Zhang, Rebecca L. Walker, Dalila Pinto, Katherine S. Pollard, Chunyu Liu, Michael J. Gandal. Large-Scale, Multi-Ethnic Resource of Gene, Isoform, and Splicing Regulation in the Developing Human Brain. 2021 Virtual

- World Congress of Psychiatric Genetics, Oct 11-15. Virtual
4. Yu Chen, Sihan Liu, Feiran Wang, Yi Jiang, Fangyuan Duan, Yan Xia, Miao Li, Wenying Qiu, Chao Ma, Jufang Huang, Shuhua Xu, Beisha Tang, Hailiang Huang, Chunyu Liu, Chao Chen. Brain eQTL of East Asian, African American, and European Descent Explains Schizophrenia GWAS in Diverse Populations. 2021 Virtual World Congress of Psychiatric Genetics, Oct 11-15. Virtual
 5. Cong Han, Yu Chen, Chao Chen, Chunyu Liu. Gene Body-Based eQTL Analysis Detects Signals Located in Genebody and Downstream of TES that are Missed by TSS-Centered Analysis for Long Transcripts. 2021 Virtual World Congress of Psychiatric Genetics, Oct 11-15. Virtual
 6. Rujia Dai, Ramu Vadukapuram, Chunling Zhang, Yan Xia, Chao Chen, Chunyu Liu. Transcription and Translation Regulated by microRNA in Human Brain and its Potential Contribution to Psychiatric Disorders. 59th ACNP. Virtual. Dec 5 – 9. 2020.
 7. Shishi Min, PsychENCODE Consortium, Daniel H. Geschwind, Chao Chen, Chunyu Liu. Detection of human viruses in the brains of patients with schizophrenia, bipolar disorder, and autism spectrum disorder. 2020 Virtual World Congress of Psychiatric Genetics. Virtual. October 16 – 22, 2020.
 8. Rujia Dai, Ramu Vadukapuram, Chunling Zhang, Yan Xia, Chao Chen, Chunyu Liu. Transcription and translation regulated by microRNA in human brain and its potential contribution to psychiatric disorders. 59th ACNP. Virtual.
 9. Amira Kefi, Lingling Huang, Chunyu Liu. A novel approach to identify and validate novel genes and transcripts using a probabilistic framework. ASHG 2020 Virtual Meeting. October 27-30, 2020.
 10. Yu Chen, Jiacheng Dai, Jinghong Qiu, Chao Chen, Chunyu Liu. Cross-Disorder Analyses of Immune-Related Gene Expressions in Brains of Major Neuropsychiatric Disorders. 58th ACNP. Orlando, FL, USA. Dec 8-11, 2019.
 11. Chuan Jiao, Richard Kopp, Elizabeth Kuney, Chao Chen, and Chunyu Liu. Sex-differences in brain development correspond to sex-differences in the age of onset for schizophrenia. **Oral presentation.** American Society of Human Genetics 2019. Oct 15- 19. 2019. Houston, TX, USA
 12. Yu Chen, Jiacheng Dai, Jinghong Qiu, Chunyu Liu, Chao Chen. The brain transcriptomic profiles of immunogenic systems in major neuropsychiatric disorders: a cross-disorder study. American Society of Human Genetics 2019. Oct 15- 19. 2019. Houston, TX, USA
 13. Sihan Liu, Yu Chen, Feiran Wang, Yi Jiang, Fangyuan Duan, Richard F. Kopp, PsychENCODE consortium, Chunyu Liu, Chao Chen. Interpret risk of schizophrenia using brain regulatory profiles across different populations. **Oral presentation.** American Society of Human Genetics 2019. Oct 15- 19. 2019. Houston, TX, USA
 14. Yi Jiang, Dan Zhou, Rui Chen, The PsychENCODE Consortium, Chao Chen, Bingshan Li, Chunyu Liu. Integrating multi-omics data for gene-level association test identifies schizophrenia risk genes. American Society of Human Genetics 2019. Oct 15- 19. 2019. Houston, TX, USA
 15. Meng Xu, Chuan Jiao, Yu Chen, Sihan Liu, Jianghua Tian, Chao Chen, Chunyu Liu. The relationship between subunits of a protein complex and gene co-expression networks in the human brain. American Society of Human Genetics 2019. Oct 15- 19. 2019. Houston, TX, USA
 16. Yan Xia, Yu Chen, Yi Jiang, Chunyu Liu, Chao Chen. Sex-specific brain transcriptome dysfunction burden in schizophrenia, autism spectrum disorder, and bipolar disorder. **Oral presentation.** American Society of Human Genetics 2019. Oct 15- 19. 2019. Houston, TX, USA
 17. Sihan Liu, Yu Chen, Feiran Wang, Yi Jiang, Fangyuan Duan, Richard F. Kopp, PsychENCODE consortium, Chunyu Liu, Chao Chen. Interpret risk of schizophrenia using brain regulatory profiles across different populations. 2019 World Congress of Psychiatric Genetics. October 26-31, 2019. Los Angeles, CA, USA
 18. Meng Xu, Chuan Jiao, Yu Chen, Sihan Liu, Jianghua Tian, Chao Chen, Chunyu Liu. The relationship between subunits of a protein complex and gene co-expression networks in the human brain. 2019 World Congress of Psychiatric Genetics. October 26-31, 2019. Los Angeles, CA, USA
 19. Yan Xia, Yu Chen, Yi Jiang, Chunyu Liu, Chao Chen. Sex-specific brain transcriptome dysfunction burden in schizophrenia, autism spectrum disorder, and bipolar disorder. 2019 World Congress of Psychiatric Genetics. October 26-31, 2019. Los Angeles, CA, USA
 20. Jeffrey R. Bishop, Bin Guo, Adam M. Lee, Paul D. Thuras, Joshua Baller, Chuan Jiao, Christopher R.

- Erbes, Melissa A. Polusny, Gregory J. Lamberty, Chunyu Liu, Baolin Wu and Kelvin O. Lim. Polygenic risk score associations with nonpharmacologic interventions for PTSD. American College of Neuropsychopharmacology 58th Annual Meeting. December 8 – 11, 2019. Orlando, FL, USA
21. Yan Xia, Chuan Jiao, Yu Chen, Yi Jiang, Chao Chen, Chunyu Liu. Sex-related brain transcriptome differences may participate in risks, symptoms and age-of-onset of psychiatric disorders. American College of Neuropsychopharmacology 58th Annual Meeting. December 8 – 11, 2019. Orlando, FL, USA
 22. Ding C. et al. Schizophrenia associated functional study of POU3F2-regulated TRIM8 signaling pathway in neural progenitor cells. American Society of Human Genetics 2018, October 18, 2018, San Diego, CA, USA
 23. Jiang Y. et al. DRAMS: A tool to Detect and Re-Align Mixed-up Samples leveraging multi-omics data, American Society of Human Genetics 2018, October 18, 2018, San Diego, CA, USA
 24. Dai R. et al. Cell type-specific alteration in schizophrenia and bipolar disorder, World Congress of Psychiatric Genetics. poster Oct 13, Glasgow, Scotland, UK
 25. Dai R. et al. Cell type-specific alteration in schizophrenia and bipolar disorder, **Oral presentation**. The American Society of Human Genetics 2018, October 18, 2018, San Diego, CA, USA
 26. Jiao, C. et al. Brain region- and developmental time-specific genes are enriched in psychiatric genetics signals. Abstract Program Number: 262, **Oral presentation**. The American Society of Human Genetics 2018, October 19, 2018, San Diego, CA, USA
 27. Xia Y, Dai R, Wang K, Jiao C, Xu Y, Li H, Jing X, Chen Y, The PsychENCODE Consortium, Chen C and **Liu C***. Sexually dimorphic DNA methylation in human brain and release to psychiatric disorders. **Oral presentation**. The American Society of Human Genetics 2017, Oct.17 - Oct.21, 2017, Orlando, USA
 28. Hu M, Xia Y, Zong X, Sweeney JA, Bishop JR, Giase G, Li B, Rubin LH, Wang Y, Li Z, He Y, Tang J, **Liu C**, Chen C, Chen X. Risperidone-induced DNA methylation alterations in first-episode drug-naïve schizophrenia patients and their relation with neuroimaging and cognitive phenotypes. **Oral presentation**. 2017 XXV World Congress of Psychiatric Genetics. Oct.13 - Oct.17, 2017, Orlando, USA
 29. Wang K, Dai R, Xia Y, Jiao C, Chen C, **Liu C***. Various relationships between DNA methylation and gene expression in different tissues and ages. **Oral presentation**. 2017 XXV World Congress of Psychiatric Genetics. Oct.13 - Oct.17, 2017, Orlando, USA
 30. Wang K, Dai R, Xia Y, Jiao C, Chen C, **Liu C***. Various relationships between DNA methylation and gene expression in different tissues and ages. The American Society of Human Genetics 2017, Oct.17 - Oct.21, 2017, Orlando, USA
 31. Dai R, Jiao C, Chen Y, Chen C, **Liu C***. Specificity and Co-expression of Cell Type-Specific Marker Genes in Brain. The American Society of Human Genetics 2017, Oct.17 - Oct.21, 2017, Orlando, USA
 32. Dai R, Jiao C, Chen Y, Chen C, **Liu C***. Specificity and Co-expression of Cell Type-Specific Marker Genes in Brain. 2017 XXV World Congress of Psychiatric Genetics. Oct.13 - Oct.17, 2017, Orlando, USA
 33. Hu M, Xia Y, Zong X, Sweeney JA, Bishop JR, Giase G, Li B, Rubin LH, Wang Y, Li Z, He Y, Tang J, Liu C, Chen C, Chen X. Risperidone-induced DNA methylation alterations in first-episode drug-naïve schizophrenia patients and their relation with neuroimaging and cognitive phenotypes. Poster presentation. The American Society of Human Genetics 2017, Oct.17 - Oct.21, 2017, Orlando, USA
 34. Horwitz T, Lam K, Chen Y, Xia Y, **Liu C***, A Decade in Psychiatric GWAS Research. American College of Neuropsychopharmacology 56th Annual Meeting. Palm Springs, CA, December 3-7, 2017.
 35. **Liu C**. Using Human Genetics Gwas and Expression Data to Drive Discovery. American College of Neuropsychopharmacology 55th Annual Meeting. Hollywood, Florida, December 4-8, 2016. (Symposium, oral presentation)
 36. Liu C. Global diversity task force: mission and vision. World Congress of Psychiatric Genetics. Toronto, Canada October 15-20, 2015. (oral presentation)
 37. Liu C, Tang H, Tang J, Chen H, Chen C. Circadian rhythms of DNA methylation and gene expression. American College of Neuropsychopharmacology. Phoenix, AZ, Dec 7-11, 2014.
 38. Liu C, Zhang C, Chen C, Badner J.A., Alliey-Rodriguez N., Gershon E.S., Gamazon E.R., IOCDF-GC,

- TSAICG, Cox N. Brain eQTLs shared by multiple psychiatric diseases. American College of Neuropsychopharmacology. Miami, FL, Dec 8-12, 2013.
39. Liu C, Zhang C, Chen C, Badner J.A., Alliey-Rodriguez N., Gershon E.S., Gamazon E.R., IOCDF-GC, TSAICG, Cox N. Using brain molecular QTLs to identify novel risk genes shared by multiple psychiatric diseases. The American Society of Human Genetics 61th Annual Meeting. Boston, MA, October 22-26, 2013. (oral presentation)
 40. Hause RJ. Jr., Archer CT., Cheng L, Liu C, and Jones RB. Genetic variation underlying protein levels in brain samples and lymphoblastoid cell lines. Boston, MA, October 22-26, 2013. (oral presentation)
 41. Liu C, Zhang C, Chen C, Badner J.A., Alliey-Rodriguez N., Gershon E.S., Gamazon E.R., IOCDF-GC, TSAICG, Cox N. Using brain molecular QTLs to identify novel risk genes shared by multiple psychiatric diseases. World Congress of Psychiatric Genetics. Boston, MA, October 17-21, 2013. (oral presentation)
 42. Liu C, Bishop J, Zhang C, Wong M, Patel SR, Leigh J, Pavuluri MN. Cell adhesion pathway is implicated in Lithium treatment for adolescent Mania via DNA Methylation alteration. American College of Neuropsychopharmacology. Miami, FL, Dec 2-6, 2012.
 43. Liu C, Zhang C, Chen Q, Liu X, Wang X, Chen C, Bishop J, Gershon E, Singleton A, Lipska B, Kleinman J, Sweeney J, Badner J, Cheng C. Sexually dimorphic DNA methylation are conserved across brain and blood. Univ of Illinois at Chicago, Department of Psychiatry Third Annual Research Forum Extravaganza. UIC west student center, Chicago. 2012, Sept 19, 2012.
 44. Chen C, Zhang C, Cheng C, Bishop J, Sweeney J, Liu C. DNA Methylation regulates gene expression differently across tissues. Univ of Illinois at Chicago, Department of Psychiatry Third Annual Research Forum Extravaganza. UIC west student center, Chicago. 2012, Sept 19, 2012.
 45. Liu C, Chen H, Chen C, Zhang C, Badner J, Gershon E, Cheng L. Interaction between SNPs and DNA methylation affects gene expression in human brain. Univ of Illinois at Chicago, Department of Psychiatry Third Annual Research Forum Extravaganza. UIC west student center, Chicago. 2012, Sept 19, 2012.
 46. Cheng L, Zhang C, Badner J, Chen C, Gershon E, Liu C. DNA Methylation in Cerebellum of Bipolar disorder, Schizophrenia and Depression patients. The American Society of Human Genetics 61th Annual Meeting & ICHG. Montreal, Canada. Oct. 11-15, 2011.
 47. Chen C, Cheng L, Pibiri F, Zhang C, Badner E, Gershon E, Liu C. A Metallothionein gene family co-expression module is differentially expressed in multiple studies of schizophrenia postmortem brains. The American Society of Human Genetics 61th Annual Meeting & ICHG. Montreal, Canada. Oct. 11-15, 2011.
 48. Liu C, Zhang C, Zhang K, Chen C, Gamazon E, Zhang X, Badner J. Common SNPs in expression Microarray probes impact eQTL studies. The American Society of Human Genetics 61th Annual Meeting & ICHG. Montreal, Canada. Oct. 11-15, 2011.
 49. Chen Q, Reed J, Cheng, Edenberg H, BiGS consortium, Gershon E, Liu C, Deep sequencing of brain expressed microRNAs and bipolar association studies of microRNA SNPs. The American Society of Human Genetics 61th Annual Meeting & ICHG. Montreal, Canada. Oct. 11-15, 2011.
 50. Cox N, Gamazon E, BadnerJ, Cheng L, Pibiri F, Zhang C, Chen C, Grennan K, Gershon E, Nicolae D, BiGS Consortium, Liu C. Brain eQTLs and function-based GWAS of Bipolar disorder identified novel disease risk genes. XIX World Congress of Psychiatric Genetics. Washington DC, Sept 10-14, 2011.
 51. Chen C, Cheng L, Pibiri F, Zhang C, Badner J, Gershon E, Liu C. Co-expression modules in Schizophrenia postmortem brains. XIX World Congress of Psychiatric Genetics. Washington DC, Sept 10-14, 2011.
 52. Liu C, Gershon ES, Zhang C, Badner JA, Cheng L, Zhang D. Age and sex effects on gene-specific methylation in human brain and its relevance to Psychiatry. (Symposium). 66th Annual Meeting of Society of Biological Psychiatry. San Francisco, CA. May 12, 2011.
 53. Liu C, Gamazon E, Cheng L, Pibiri F, Badner JA, Zhang C, Chen C, Grennan K, Gershon ES, Nicolae D, BiGS, Cox N. Brain eQTLs and function-based GWAS of Bipolar disorder. 13thInternational Congress on Schizophrenia Research. Colorado Spring, CO. April 2, 2011.
 54. Chen C, Zhang D, Badner JA, Gershon ES, and Liu C. Batch effects in microarray experiments: performance evaluation of six algorithms. The American Society of Human Genetics 60th Annual Meeting. Washington, DC. Nov. 2, 2010.
 55. Chen Q, Cheng L, Rodriguez A, Chen C, Zhang C, Gershon ES, Liu C. Brain DNA Methylation of major depressive Disorder and healthy control. The American Society of Human Genetics 60th Annual Meeting. Washington, DC. Nov. 2, 2010.
 56. Liu C, Zhang C, Chen C, Badner JA, Lipska BK, Rizig MA, Mcquillan A, Gurling H, and Gershon ES. Gene

- expression changes induced by haloperidol and clozapine in rodents. World Congress of Psychiatric Genetics. Athens, Greece. Oct. 3, 2010.
57. Liu C, Cheng L, Pibiri F, Badner JA, Chen C, Zhang C, Grennan K, and Gershon ES. eQTLs and sQTLs in human Cerebellum and Parietal cortex. The American Society of Human Genetics 60th Annual Meeting. Washington, DC. Nov. 2, 2010.
 58. Zhang D, Cheng L, Pibiri F, Chen C, Badner JA, Gershon ES, and Liu C. Rare CNVs affect gene expression in human brains. XVIII World Congress of Psychiatric Genetics. Athens, Greece. Oct. 3, 2010.
 59. Alliey N, Zhang D, Badner JA, Greenwood T, Kelsoe JR, Liu C, and Gershon ES. GWAS for personality traits in Bipolar patients. XVII World Congress of Psychiatric Genetics. San Diego, California. Nov. 4, 2009.
 60. Chen C, Zhang D, Cheng L, Badner JA, Gershon ES, and Liu C. Correlations between gene expression and DNA methylation in human brain. XVII World Congress of Psychiatric Genetics. San Diego, California. Nov. 4, 2009.
 61. Cheng L, Zhang D, Chen C, Badner JA, Luo W, and Liu C. Genome-wide methylation analysis of postmortem brains suggests gene differential methylation is associated with Schizophrenia and Major Depression. Emerging Evidence for Epigenomic Changes in Human Disease Workshop. Bethesda. MD. Mar. 16, 2009.
 62. Cheng L, Chen C, Badner JA, Zhang D, Gershon ES, and Liu C. Genome-wide Methylation analysis of postmortem brains suggests differential Methylation is associated with major depression and Bipolar disorder. XVII World Congress of Psychiatric Genetics. San Diego, California. Nov. 4, 2009.
 63. Liu C, Reid J, Cheng L, Alvi O, Lucas AS, Lewis L, Hall O, Nazareth L, Wheeler D, Muzny D, Zhang D, Gershon ES, and Gibbs RA. microRNA genomic variations in psychiatric diseases. International Congress on Schizophrenia Research. San Diego, California. Mar. 28, 2009.
 64. Smith EN, Badner JA, Barrett T, Belmonte PL, Berrettini W, Bloss CS, Byerley W, Coryell W, Craig D, Edenberg HJ, Eskin E, Foroud T, Gershon ES, Greenwood TA, Hipolito M, Koller DL, Lawson WB, Liu C, Lohoff F, McInnis MG, McMahon FJ, Mirel DB, Murray SS, Nievergelt C, Nurnberger J, Nwulia EA, Paschall J, Potash JB, Rice J, Schulze TG, Scheftner W, Panganiban C, Zaitlen N, Zandi PP, Zhang P, Zollner S, Schork NJ, and Kelsoe JR. Genome-wide association of Bipolar disorder. XVII World Congress of Psychiatric Genetics. San Diego, California. Nov. 4, 2009.
 65. Zhang D, Qian Y, Akula N, Gershon ES, and Liu C. Comparison of four CNV detection programs based on genome wide SNP array. XVII World Congress of Psychiatric Genetics. San Diego, California. Nov. 4, 2009.
 66. Cheng L, Tang Y, Gershon ES, and Liu C. Gene expression, biochemistry, and behavior study in a G72 transgenic mouse model. 63rd Annual Meeting of Society of Biological Psychiatry. Washington DC. May 1, 2008.
 67. Cheng L, Reid J, Alvi O, Lucas AS, Lewis L, Hall O, Nazareth L, Wheeler D, Muzny D, Zhang D, Gershon ES, Gibbs RA, and Liu C. microRNA genomic variations in psychiatric diseases. 58th Annual Meeting of the American Society of Human Genetics. Philadelphia, Pennsylvania. Nov. 11, 2008.
 68. Liu C. Variants of Human brain expressed miRNAs and disease association. XVI World Congress of Psychiatric Genetics. Osaka, Japan. Oct. 11, 2008.
 69. Liu C, Cheng L, Badner JA, Craig DW, Josephson M, Christian SL, and Gershon ES. Genomic mapping of gene expression regulation in human frontal cortex. 58th Annual Meeting of the American Society of Human Genetics. Philadelphia, Pennsylvania. Nov. 11, 2008.
 70. Liu C, Cheng L, Badner JA, Craig DW, Josephson M, Christian SL, and Gershon ES. Genomic mapping of gene expression regulation in human frontal cortex. The 5th Symposium for Chinese Neuroscientists Worldwide. Changsha, China. Aug. 1, 2008.
 71. Reid J, Liu C, Cheng L, Alvi O, Lucas AS, Lewis L, Hall O, Nazareth L, Wheeler D, Muzny D, Gershon ES, and Gibbs RA. microRNA Variation and Schizophrenia. The Biology of Genomes 2008. Cold Spring Harbor, NY. May 6, 2008.
 72. Reid J, Liu C, Nagaraja A, Creighton C, Cheng L, Alvi O, Lucas AS, Lewis L, Hall O, Nazareth L, Wheeler D, Anderson M, Matzuk M, Muzny D, Gunaratne P, Gershon ES, and Gibbs RA. microRNAs and deep sequencing of the short RNA-ome. Advances in Genome Biology and Technology 2008. Marco Island, FL. Feb. 6, 2008.
 73. Tang J, Cheng L, Craig DW, Josephson M, Christian SL, Gershon ES, Chen X, and Liu C. DNA copy number variations in major psychosis. 58th Annual Meeting of the American Society of Human Genetics. Philadelphia, Pennsylvania. Nov. 11, 2008.

74. Zhang D, Cheng L, and Liu C. Telomere length in brain and psychiatric disorders. 58th Annual Meeting of the American Society of Human Genetics. Philadelphia, Pennsylvania. Nov. 11, 2008.
75. Cheng L, Tang Y, Gershon ES, and Liu C. G72/G30 transgenic mice and brain expression profiling. The Society of Biological Psychiatry's 62nd Annual Scientific Program and Convention. San Diego, CA. May 17, 2007.
76. Liu C and Armoskus C. Missing parts of the whole genome association studies. The 1st Annual Midwest Symposium on Computational Biology and Bioinformatics. Evanston, IL. Oct. 6, 2007.
77. Shi J, Wittke-Thompson JK, Badner JA, Hattori E, Gershon ES, and Liu C. Clock genes may influence Bipolar disorder susceptibility and dysfunctional circadian rhythm. 57th Annual Meeting of the American Society of Human Genetics. San Diego. Oct. 23, 2007.
78. Shi J, Maheshwari M, Skol A, Badner JA, Gershon ES, Gibbs RA, and Liu C. Association of the DAOA gene and mental illness: a meta-analysis and resequencing. 62nd Annual Scientific Convention & Program. San Diego, CA. May 17, 2007.
79. Cheng L, Hattori E, Nakajima A, Tang Y, Gershon ES, and Liu C. G72/G30 transgenic mice and gene expression profiling. 56th Annual Meeting of the American Society of Human Genetics. New Orleans, Louisiana. Oct. 9, 2006.
80. Detera-Wadleigh SD, Liu C, Cardona I, Corona W, Akula N, Steele CJM, Kundu M, Bonner TI, Badner JA, Maheshwari M, Gibbs RA, Gershon ES, and McMahon FJ. DOCK9 variants are associated with several bipolar disorder phenotypes and interact with G72. XIV World Congress of Psychiatric Genetics. Cagliari, Sardinia, Italy. Oct. 28, 2006.
81. Liu C. Evaluation of sequencing analysis software for complex disease studies. The 8th International Meeting on Human Genome Variation and Complex Genome Analysis. Hong Kong. Sept. 14, 2006.
82. Liu C. Evaluation of sequencing Analysis software for complex disease studies. 56th Annual Meeting of the American Society of Human Genetics. New Orleans, Louisiana. Oct. 9, 2006.
83. Liu C, Cheng L, Hattori E, Nakajima A, Tang Y, and Gershon ES. G72/G30 transgenic mice and gene expression profiling. XIV World Congress of Psychiatric Genetics. Cagliari, Sardinia, Italy. Oct. 28, 2006.
84. Shi J, Badner JA, Qian Y, Gershon ES, and Liu C. An Association analysis of Cannabinoid Receptor 1 gene with Bipolar disorder. 56th Annual Meeting of the American Society of Human Genetics. New Orleans, Louisiana. Oct. 9, 2006.
85. Detera-Wadleigh SD, Liu C, Badner JA, Corona W, Akula N, Cardona I, Kundu M, Steele CJM, Maheshwari M, Bonner T, Gershon ES, and McMahon FJ. DOCK9, a candidate bipolar disorder susceptibility gene on chromosome 13q32.3. XIII World Congress of Psychiatric Genetics. Boston, Massachusetts. Oct. 14, 2005.
86. Detera-Wadleigh SD, Liu C, Badner JA, Corona W, Akula N, Cardona I, Kundu M, Steele CJM, Maheshwari M, Bonner T, Gershon ES, and McMahon FJ. DOCK9, a candidate bipolar disorder susceptibility gene on chromosome 13q32.3. 55th Annual Meeting of the American Society of Human Genetics. Salt Lake City, Utah. Oct. 25, 2005.
87. Gershon ES and Liu C. Linkage and association findings in Bipolar disorder and Schizophrenia. American College of Neuropsychopharmacology. Waikoloa, Hawaii. Dec. 11, 2005.
88. Liu C, Badner JA, Zou H, Christian SL, and Gershon ES. Association of DISC1 with bipolar disorder in two sample sets. XIII World Congress of Psychiatric Genetics. Boston, Massachusetts. Oct. 14, 2005.
89. Liu C, Shi J, Zou H, Badner JA, Christian SL, and Gershon ES. Association study of trace Amine receptors in Bipolar disorder. 55th Annual Meeting of the American Society of Human Genetics. Salt Lake City, Utah. Oct. 25, 2005.
90. Maheshwari M, Liu C, Badner JA, Smith CK, Christian SL, Zou H, Detera-Wadleigh SD, Bonner TI, McMahon FJ, Gershon ES, and Gibbs RA. Resequencing & fine map association study of G72/G30 region in bipolar disorder. 55th Annual Meeting of the American Society of Human Genetics. Salt Lake City, Utah. Oct. 25, 2005.
91. Shi J, Hattori E, Zou H, Badner JA, Christian SL, Gershon ES, and Liu C. Cholinergic system genes and Bipolar disorder. 55th Annual Meeting of the American Society of Human Genetics. Salt Lake City, Utah. Oct. 25, 2005.
92. Liu C, Yao F, Zhang R, Zhu Z, Xia J, and Gershon ES. MutScreen: a web-based tool of primer design for mutation screening by PCR-direct sequencing. 54th Annual Meeting of the American Society of Human Genetics. Toronto, Canada. Oct. 27, 2004.
93. Liu C, Nguyen T, Zhang R, Yao F, Zhu Z, and Gershon ES. SNP information mining pipeline (SIMP) for

- complex disease studies. 53rd Annual Meeting of the American Society of Human Genetics. Los Angeles, CA. Nov. 4, 2003.
94. Hattori E, Liu C, Badner JA, Christian SL, Maheshwari M, Detera-Wadleigh SD, Bonner TI, Gibbs RA, and Gershon ES. Polymorphisms of LG72 (on 13q33) associated with Bipolar disorder by TDT and partition of linkage evidence. 52nd Annual Meeting of the American Society of Human Genetics. Baltimore MD. Oct. 15, 2002.
 95. Liu C, Bonner TI, Liu Y, Lyons JL, and Gershon ES. DNannotator: a web-based sequence annotation tool kit and its application to analyses of the sequence of human chromosome 13q32-33. 52nd Annual Meeting of the American Society of Human Genetics. Baltimore MD. Oct. 15, 2002.
 96. Liu C, Zhang D, Zou H, Christian SL, Badner JA, and Gershon ES. Mutation screening of 3 candidate genes on 13q32 for bipolar disorder. Society of biological psychiatry. New Orleans, Louisiana. May 3, 2001.
 97. Liu C and Gershon ES. SNP-Cruncher: A Perl script toolbox for large-scale SNP data mining and its application on 13q32. 51th Annual Meeting of the American Society of Human Genetics. San Diego, California. Oct. 12, 2001.
 98. Liu C and Gershon ES. SNP Cruncher: A Perl script toolbox for large-scale SNP data mining. The IXth World Congress on Psychiatric Genetics. St. Louis, Missouri. Oct. 6, 2001.
 99. Christian SL, Vlamakis V, Liu C, and Gershon ES. An ~10 Mb Integrated transcript map of chromosome 13q32. Eighth World Congress on Psychiatric Genetics. Versailles, France. Aug. 27, 2000.
 100. Christian SL, Vivian V, Liu C, Badner JA, and Gershon ES. An integrated transcript map of the chromosome 13q32 region linked to bipolar disorder and schizophrenia. 50th Annual Meeting of the American Society of Human Genetics. Philadelphia, Pennsylvania. Oct. 3, 2000.
 101. Liu C, Badner JA, Christian SL, Guroff JJ, Detera-Wadleigh SD, and Gershon ES. Further linkage evidence for a bipolar disorder susceptibility locus on 13q32. 50th Annual Meeting of the American Society of Human Genetics. Philadelphia, Pennsylvania. Oct. 3, 2000.
 102. Liu C, Badner JA, Christian SL, Guroff JJ, Detera-Wadleigh SD, and Gershon ES. Fine mapping provides further linkage evidence for a bipolar disorder susceptibility locus on 13q32. Eighth World Congress on Psychiatric Genetics. Versailles, France. Aug. 27, 2000.
 103. Liu C, Xia J, Tang D, and Pan Q. Molecular cloning of complete coding sequence of human M6ba. 48th Annual Meeting of the American Society of Human Genetics. Denver, Colorado. Oct. 27, 1998.
 104. Xia J, Liu C, Yu K, Pan Q, Zheng D, and Dai H. Molecular cloning of the human Dimethylglycine Dehydrogenase-like gene (DMGDHL1) from the sarcosinemia critical region at 9q34. 18th International Congress of Genetics. Beijing, China. June 20, 1998.
 105. Xia J, Liu C, Wang D, Pan Q, Liao X, Dai H, and Deng H. Molecular cloning and localization of human atrophin-1-like gene. 18th International Congress of Genetics. Beijing, China. June 20, 1998.
 106. Xia J, Liu C, Huang L, Pan Q, and Dai H. Molecular cloning and mapping of human Cx31.1 gene. 48th Annual Meeting of the American Society of Human Genetics. Denver, Colorado. Oct. 27, 1998.
 107. Xia J, Wang D, Liu C, Ruan Q, Pan Q, Dai H, and Deng H. Molecular cloning and Localization of human ataxin-2 like gene. 18th International Congress of Genetics. Beijing, China. June 20, 1998.
 108. Tang D, Xia J, Li L, Dai H, Jiang S, Liu C, and Long Z. Repository of Human Chromosomal Anomalies in China. 47th Annual meeting of American Society of Human Genetics. Baltimore, Maryland. Oct. 28, 1997.
 109. Xia J, Ruan Q, Xu L, Tang D, Liu C, and Deng H. Construction and Application of Chromosome Band-specific Probe Pools and pUC19 Libraries. 47th Annual meeting of American Society of Human Genetics. Baltimore, Maryland. Oct. 28, 1997.
 110. Xu L, Xia J, Ruan Q, Liu C, and Deng H. Mapping of human MAD2 gene to chromosome 5q23.3. Human Genome Meeting. Toronto, Canada. Mar. 6, 1997.
 111. Zhang C, Chen Y, Liu C, and Gui M. Hereditary and variation of domesticated silkworm generated by introduction of eri silkworm DNA. Proceedings of the 3rd National Symposium on Transgenic Animals. Beijing, China. Nov. 1, 1996.
 112. Zhang C, Chen Y, Gui M, and Liu C. RAPD analysis of hereditary and variation of domesticated silkworm generated by introduction of eri silkworm DNA. The First International Conference on Transgenic Animals. Beijing, China. Nov. 1, 1996.
 113. Liu C, Zhang C, Chen Y, and Gui M. RAPD analysis of the offspring generated by cross or DNA introduction between domesticated silkworm and eri silkworm. The 5th Congress of Genetics Society of

China. Beijing, China. Oct., 1995.