

Charleston W.K. Chiang

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Curriculum Vitae

Academic Appointment

- 2019-present Assistant Professor, University of Southern California
Quantitative and Computational Biology Section, Department of Biological Sciences
Dornsife College of Letters, Arts, and Sciences
- 2018-present Assistant Professor, University of Southern California
Center for Genetic Epidemiology, Department of Preventive Medicine
Keck School of Medicine

Education

- 2011 Ph.D., Genetics, Harvard University
Division of Medical Sciences, Program in Biological and Biomedical Science
Leder Human Biology and Translational Medicine Program Scholar
- 2005 B.S., Microbiology, Immunology, and Molecular Genetics, UCLA
Mathematics (minor), Statistics (minor)
Summa Cum Laude, College Honors Program Scholar

Past Employment / Research Experience

parental leaves in 2013 and 2016.

- 2015-2017# Postdoctoral Research Fellow, UCLA
Center for Neurobehavioral Genetics, Semel Institute for Neuroscience
Advisor: Nelson B. Freimer, M.D.
- 2011-2015# Postdoctoral Research Fellow, UCLA
Department of Ecology and Evolutionary Biology
Advisor: John Novembre, Ph.D.
- 2005-2011 Graduate student, Harvard Medical School, Broad Institute
Department of Genetics
Thesis Advisor: Joel N. Hirschhorn, M.D. Ph.D.
- 2002-2005 Undergraduate Research Assistant, UCLA
Department of Human Genetics
Advisor: Eric Vilain, M.D. Ph.D.

Publications

11 (co-) first authorship (numerically bolded), **6** (co-) corresponding authorship (numerically bolded)

* equal contribution, † corresponding author, **bolded** name denotes group member

Through 2018, 14 out of 26 publications in journals with impact factor (IF) > 8.

- 36. Sun H***, **Lin M***, Russell EM, Minster RL, Chan TF, Naseri T, Reupena MS, Lum-Jones A, OLaGA Study Group, Cheng I, Wilkens LR, Le Marchand L, Haiman CA, **Chiang CWK**[†]. "The impact of global and local Polynesian genetic ancestry on complex traits in Native Hawaiians." *In preparation*
- 35.** Graff M, Justice AE, et al. "Discovery and fine-mapping of height loci using high density imputation of genome-wide association studies in individuals of African ancestry – the African Ancestry Anthropometry Genetics Consortium." *In preparation*

34. Chen MH, Raffield LM, Mousas A, et al. "Trans-ethnic and ancestry-specific blood-cell genetics in 746,667 individuals." *In revision at Cell*.
33. Pua CJ, et al. "Genetic studies of hypertrophic cardiomyopathy in Singaporeans identify variants in TNNI3 and TNNT2 that are common in Chinese patients." *In press at Cir Genom Precis Med*.
32. Lin M[†], Caberto C, Wan P, Li Y, Lum-Jones A, Tiirikainen M, Pooler L, Nakamura B, Sheng X, Porcel J, Lim U, Setiawan VW, Le Marchand L, Wilkens LR, Haiman CA, Cheng I, **Chiang CWK[†]**. "Population specific reference panels are crucial for the genetic analyses of Native Hawaiians: an example of the *CREBRF* locus." *Accepted at Hum Mol Genet*; bioRxiv 2019: <https://www.biorxiv.org/content/10.1101/789073v1>. doi: 10.1101/789073
31. **Chen M[†]**, Sidore C, Akiyama M, Ishigaki K, Kamatani Y, Schlessinger D, Cucca F, Okada Y, **Chiang CWK[†]**. "Evidence of polygenic adaptation at height-associated loci in mainland Europeans and Sardinians." *In revision at Am J Hum Genet*. bioRxiv 2019: <https://www.biorxiv.org/content/10.1101/776377v1>. doi: 10.1101/776377
30. Marcus JH, Posth C, Ringbauer H, Lai L, Skeates R, Sidore C, Beckett J, Furtwangler A, Olivieri A, **Chiang CWK**, Al-asadi H, Dey K, Joseph TA, Liu CC, Der Sarkissian C, Radzeviciute R, Michel M, Gradoli MG, Marongiu P, Rubino S, Mazzarello V, Rovina D, La Fragola A, Serra RM, Bandiera P, Bianucci R, Pompianu E, Murgia C, Guirguis M, Orquin RP, Tuross N, van Dommelen P, Haak W, Reich D, Schlessinger D, Cucca F, Krause J, Novembre J. "Genetic history from the Middle Neolithic to present on the Mediterranean island of Sardinia." **Nat Comm** 2020 Feb 24;11(1):939. doi: 10.1038/s41467-020-14523-6.
29. de Smith AJ, Walsh KM, Morimoto LM, Francis SS, Hansen HM, **Jeon S**, Gonseth S, **Chen M**, **Sun H**, Luna-Fineman S, Antillón F, Girón V, Kang AY, Smirnov I, Shao X, Whitehead TP, Barcellos LF, Jolly KW, Healy J, Laverdière C, Sinnett D, Taub JW, Birch JM, Thompson PD, Pombo-de-Oliveira MS, Spector LG, DeWan AT, Mueller BA, **Chiang CWK**, Metayer C, Ma X, Wiemels JL. "Heritable variation at the chromosome 21 gene *ERG* is associated with acute lymphoblastic leukemia risk in children with and without Down syndrome." **Leukemia** 2019 Nov;33(11):2746-2751. doi: 10.1038/s41375-019-0514-9.
28. Locke AE*, Steinberg KM*, **Chiang CWK***, Service S*, Havulinna A, Stell L, Pirinen M, Abel HJ, Chiang CC, Fulton RS, Jackson AU, Kang CJ, Kanchi KL, Koboldt DC, Larson DE, Nelson J, Nicholas TJ, Pietila A, Ramensky V, Ray D, Scott LJ, Stringham HM, Vangipurapu J, Welch R, Yajnik P, Yin X, Eriksson JG, Ala-Korpela M, Jarvelin MR, Manniko M, Laivouri H, FinnGen Project, Dutcher SK, Stitzel NO, Wilson RK, Hall IM, Sabatti C, Palotie A, Salomaa V, Laakso M, Ripatti S, Boehnke M, Freimer NB. "Exome sequencing of Finnish isolates enhances rare-variant association power." **Nature** 2019 Aug;572(7769):323-328. doi: 10.1038/s41586-019-1457-z.
27. Sohail M, Maier RM, Ganna A, Bloemendal A, Martin AR, Turchin MC, **Chiang CWK**, Hirschhorn JN, Daly MJ, Patterson N, Neale B, Mathieson I, Reich D, Sunyaev SR. "Polygenic adaptation on height is overestimated due to uncorrected stratification in genome-wide association studies." **eLife**. 2019 Mar 21;8. pii: e39702. doi: 10.7554/eLife.39702.
26. **Chiang CWK[†]**, Mangul S, Robles C, Sankararaman S. "A comprehensive map of genetic variation in the world's largest ethnic group – Han Chinese." **Mol Bio Evol** 2018 Nov 1;35(11):2736-2750. doi: 10.1093/molbev/msy170.
25. **Chiang CWK[†]**, Marcus J, Sidore C, Biddanda A, Al-asadi H, Zoledziwska M, Pitzalis M, Busonero F, Maschio A, Pistis G, Steri M, Angius A, Lohmueller KE, Abecasis G, Schlessinger D, Cucca F, Novembre J[†]. "Genomic history of the Sardinian population." **Nat Genet** 2018 Oct;50(10):1426-1434. doi: 10.1038/s41588-018-0215-8.

24. Ganna A, et al. "Quantifying the impact of rare and ultra-rare coding variation across the phenotypic spectrum." **Am J Hum Genet**. 2018 Jun 7;102(6):1204-1211. doi: 10.1016/j.ajhg.2018.05.002.
23. Kim DS, et al. "Novel association of TM6SF2 rs58542926 genotype with increased serum tyrosine levels and decreased apoB-100 particles in Finns." **J Lipid Res**. 2017 Jul;58(7):1471-1481. doi: 10.1194/jlr.P076034.
22. Ganel L, et al. "SVScore: an impact prediction tool for structural variation." **Bioinformatics**. 2017 Apr 1;33(7):1083-1085. doi: 10.1093/bioinformatics/btw789.
21. Reid JS, Jeff J, et al. "A principal component meta-analysis on multiple anthropometric traits identifies novel loci for body shape in >170,000 individuals of the GIANT Consortium." **Nat Commun**. 2016 Nov 23;7:13357. doi: 10.1038/ncomms13357.
20. Chen CH, Yang JH, **Chiang CWK**, Hsiung CN, Wu PE, Chang LC, Chang J, Song IW, Yang SL, Chen YT, Liu FT, Shen CY. "Population structure of Han Chinese in the modern Taiwanese population based on 10,000 participants in the Taiwan Biobank project." **Hum Mol Genet**. 2016 Dec 15;25(24):5321-5331. doi: 10.1093/hmg/ddw346.
19. **Chiang CWK**[†], Ralph P, Novembre J[†]. "Conflations of short identity-by-descent segments bias their inferred length distribution." **G3: Genes, Genomes, Genetics**. 2016 May 1; 6(5):1287-1296. doi: 10.1534/g3.116.027581.
18. Winkler TW, Justic AE, Graff M, Barata L, et al. "The influence of age and sex on genetic associations with body composition: a large-scale genome-wide interaction study." **PLoS Genet**. 2015 Oct 1;11(10):e1005378. doi: 10.1371/journal.pgen.1005378. eCollection 2015 Oct.
17. Zoledziwska M*, Sidore C*, **Chiang CWK***, Sanna S*, Mulas A, Steri M, Busonero F, Marcus JH, Marongiu M, Maschio A, Del Vecchio DO, Floris M, Meloni A, Delitala A, Concas MP, Murgia F, Biino G, Vaccargiu S, Nagaraja R, Lohmueller KE; UK10K Consortium, Timpson NJ, Soranzo N, Tachmazidou I, Dedoussis G, Zeggini E; Understanding Society Scientific Group, Uzzau S, Jones C, Lyons R, Angius A, Abecasis GR, Novembre J, Schlessinger D, Cucca F. "Height-reducing variants and selection for short stature in Sardinia." **Nat Genet**. 2015 Nov;47(11):1272-81. doi: 10.1038/ng.3368.
16. Sidore C, Busonero F, Maschio A, Porcu E, Naitza S, Zoledziwska M, Mulas A, Pistis G, Steri M, Danjou F, Kwong A, Ortega Del Vecchio VD, **Chiang CWK**, Bragg-Gresham J, Pitzalis M, Nagaraja R, Tarrier B, Brennan C, Uzzau S, Fuchsberger C, Atzeni R, Reinier F, Berutti R, Huang J, Timpson NJ, Toniolo D, Gasparini P, Malerba G, Dedoussis G, Zeggini E, Soranzo N, Jones C, Lyons R, Angius A, Kang HM, Novembre J, Sanna S, Schlessinger D, Cucca F, Abecasis GR. "Genome sequencing elucidates Sardinian genetic architecture and augments association analyses for lipid and blood inflammatory markers." **Nat Genet**. 2015 Nov;47(11):1352-6. doi: 10.1038/ng.3403.
15. Yang WY, Platt A, **Chiang CWK**, Eskin E, Novembre J, Pasaniuc B. "Spatial localization of recent ancestors for admixed individuals." **G3: Genes, Genomes, Genetics**. 2014 Nov 3; 4(12):2505-18. doi: 10.1534/g3.114.014274.
14. Wang SR, Agarwala V, Flannick J, **Chiang CWK**, Altshuler D, GoT2D Consortium, Hirschhorn JN. "Simulation of Finnish population history, guided by empirical genetic data, to assess power of rare variant tests in Finland." **Am J Hum Genet**. 2014 May 1;94(5):710-20. doi: 10.1016/j.ajhg.2014.03.019.
13. Olalde I, Sanchez-Quinto F, Datta D, Marigorta UM, **Chiang CWK**, Rodriguez JA, Fernandez-

- Callejo M, Gonzalez I, Montfort M, Matas-Lalueza L, Civit S, Luiselli D, Charlier P, Pettener D, Ramirez O, Navarro A, Himmelbauer H, Marques-Bonet T, Lalueza-Fox C. "Genomic analysis of the blood attributed to Louis XVI (1754-1793), king of France." **Sci Rep**. 2014 Apr 24;4:4666. doi: 10.1038/srep04666.
12. Olalde I, Allentoft ME, Sanchez-Quinto F, Santpere G, **Chiang CWK**, DeGiorgio M, Prado-Martinez J, Rodriguez JA, Quilez J, Rasmussen S, Ramirez O, Marigorta UM, Fernandez-Callejo M, Prada ME, Encinas JM, Nielsen R, Netea MG, Novembre J, Sturm RA, Sabeti P, Marques-Bonet T, Navarro A, Willerslev E, Lalueza-Fox C. "Derived immune and ancestral pigmentation alleles in a 7,000-year-old Mesolithic European." **Nature**. 2014 Mar 13;507(7491):225-8. doi: 10.1038/nature12960.
 11. Monda KL, Chen GK, Taylor KC, Palmer CD, et al. "A meta-analysis identifies new loci associated with body mass index in individuals of African ancestry." **Nat Genet**. 2013 Jun;45(6):690-6. doi: 10.1038/ng.2608.
 10. Turchin MC*, **Chiang CWK***, Palmer CD, Sankararaman S, Reich D, Genetic Investigation of ANthropometric Traits (GIANT) Consortium, Hirschhorn JN. "Evidence of widespread selection on standing variation in Europe at height-associated SNPs." **Nat Genet**. 2012 Sep;44(9):1015-9. doi: 10.1038/ng.2368.
 9. **Chiang CWK**, Liu CT, Lettre G, Lange LA, Jorgensen NW, Keating BJ, Vedantam S, Nock NL, Franceschini N, Reiner AP, Demerath EW, Boerwinkle E, Rotter JI, Wilson JG, North KE, Papanicolaou GJ, Cupples LA, Genetic Investigation of ANthropometric Traits (GIANT) consortium, Murabito JM, Hirschhorn JN. "Ultraconserved elements in the human genome: association and transmission analyses of highly constrained single-nucleotide polymorphisms." **Genetics**. 2012 Sep;192(1):253-66.
 8. Dauber A, Yu Y, Turchin MC, **Chiang CWK**, Meng Y, Demerath EW, Patel SR, Rich SS, Rotter JI, Schreiner PJ, Wilson JG, Shen Y, Wu BL, Hirschhorn JN. "Genome-wide association of copy number variation reveals an association between short stature and the presence of low frequency genomic deletions." **Am J Hum Genet**. 2011 Dec 9;89(6):751-9.
 7. **Chiang CWK**, Gajdos ZKZ, Korn JM, Butler JL, Hackett R, Guiducci C, Nguyen T, Wilks R, Forrester T, Henderson KD, Le Marchand L, Henderson BE, Haiman CA, Cooper RS, Lyon HN, Zhu X, McKenzie CA, Palmert MR, Hirschhorn JN. "The efficacy of detecting variants with small effects on the Affymetrix 6.0 platform using pooled DNA." **Hum Genet**. 2011 Nov;130(5):607-21.
 6. Zhu X, Young JH, Fox E, et al. "Combined admixture mapping and association analysis identifies a novel blood pressure genetic locus on 5p13: contributions from the CARE consortium." **Hum Mol Genet**. 2011 Jun 1;20(11):2285-95.
 5. Kang SJ*, **Chiang CWK***, Palmer CD*, Tayo BO, Lettre G, Butler JL, Hackett R, Adeyemo AA, Guiducci C, Berzins I, Nguyen TT, Feng T, Luke A, Shriner D, Ardlie K, Rotimi C, Wilks R, Forrester T, McKenzie CA, Lyon HN, Cooper RS, Zhu X, Hirschhorn JN. "Genome wide association of anthropometric traits in African and African derived populations." **Hum Mol Genet**. 2010 Jul 1;19(13):2725-38.
 4. **Chiang CWK**, Gajdos ZKZ, Korn JM, Kuruvilla FG, Butler JL, Hackett R, Guiducci C, Nguyen T, Wilks R, Forrester T, Haiman C, Henderson KD, Le Marchand L, Henderson BE, Palmert MR, McKenzie CA, Lyon HN, Cooper RS, Zhu X, Hirschhorn JN. "Rapid assessment of genetic ancestry in populations of unknown origin by genome-wide genotyping of pooled samples." **PLoS Genet**. 2010;6(3):e1000866.

3. **Chiang CWK***, Derti A*, Schwartz D, Chou MF, Hirschhorn JN, Wu C-t. "Ultraconserved elements: Analyses of dosage sensitivity, motifs, and boundaries." **Genetics**. 2008 Dec;180(4):2277-93.
2. Keating BJ, Tischfield S, Murray SS, *et al.* "Concept, Design and Implementation of a Cardiovascular Gene-centric 50K SNP Array for Large-scale Genomic Association Studies." **PLoS ONE**. 2008;3(10):e3583.
1. Dewing P, **Chiang CWK**, Sinchak K, Sim H, Fernagut P, Kelly S, Chesselet M, Micevych P, Albrecht KH, Harley VR, Vilain E. "Direct regulation of adult brain function by the male-specific factor SRY." **Curr Biol**. 2006, Feb 21;16(4): 415-420.

Contributed Abstracts: Talks

bolded name denotes group member

11. **Chen M**, Sidore C, Akiyama M, Ishigaki K, Kamatani Y, Schlessinger D, Cucca F, Okada Y, **Chiang CWK**. "Evidence of polygenic adaptation at height-associated loci in mainland Europeans and Sardinians." Bay Area Population Genetics Meeting (BAPG), Berkeley, California, Nov 2019.
10. **Chen M**, Sidore C, Akiyama M, Ishigaki K, Kamatani Y, Schlessinger D, Cucca F, Okada Y, **Chiang CWK**. "Evidence of polygenic adaptation at height-associated loci in mainland Europeans and Sardinians." Southern California Evolution Meeting (SCalE), Irvine, California, Oct 2019.
9. **Lin M**, Caberto C, Wan P, Lum-Jones A, Tiirikainen M, Le Marchand L, Wilkens L, Haiman C, Cheng I, **Chiang CWK**. "Oceanian reference panels are crucial for understanding the impact of a Polynesian-specific variant on adiposity in Native Hawaiians." 23andMe Genome Research Day, Mountain View, California, Jun 2019.
8. **Chiang CWK**, Mangul S, Robles C, Kretzschmar WW, Cai N, Kendler KS, Sankararaman S, Flint J. "A comprehensive map of genetic variation in the world's largest ethnic group – Han Chinese." Society for Molecular Biology & Evolution (SMBE), Yokohama, Japan, Jul 2018.
7. **Chiang CWK**, Locke AE, Steinberg KM, Service S, Abel H, Havulinna AS, Chiang C, Stell L, Stringham HM, Jackson AU, Pirinen M, Ray D, Larson DE, Koboldt DC, Scott LJ, Fulton RS, Nelson J, Nicholas TJ, Yajnik P, Ramensky V, Stitzel NO, Wilson RK, Hall IM, Sabatti C, Ripatti S, Salomaa V, Palotie A, Laakso M, Boehnke M, Freimer NB. "Insights to the population structure and genetic architecture of cardiometabolic traits in 20,029 Finnish exomes." American Society of Human Genetics (ASHG) Orlando, FL, Oct 2017.
6. **Chiang CWK**, Locke AE, Steinberg KM, Service S, Abel H, Havulinna AS, Chiang C, Stell L, Stringham HM, Jackson AU, Pirinen M, Ray D, Larson DE, Koboldt DC, Scott LJ, Fulton RS, Nelson J, Nicholas TJ, Yajnik P, Ramensky V, Stitzel NO, Wilson RK, Hall IM, Sabatti C, Ripatti S, Salomaa V, Palotie A, Laakso M, Boehnke M, Freimer NB. "Insights to the genetic architecture of cardiometabolic traits empowered by 20,000 exomes and the history of Finland." Gordon Research Conference (GRC), Human Genetics & Genomics, Stowe, VT, Jul 2017.
5. **Chiang CWK**, Sidore C, Zoledziwska M, Marcus JH, Al-asadi H, Abecasis GR, Schlessinger D, Cucca F, Novembre J. "Population genetic insight to the study of human height." International Society Evolution, Medicine & Public Health (ISEMPH), Durham, NC, Jun 2016.
4. Novembre J, **Chiang CWK**, Marcus J, Sidore C, Zoledziwska M, Steri M, Al-asadi H, Sanna S, Abecasis G, Schlessinger D, Cucca F. "The population structure and demographic history of Sardinia in relationship to neighboring populations." American Society of Human Genetics (ASHG), San Diego, CA, Oct 2014.

3. **Chiang CWK**, Ralph P, Novembre J. "Conflations of short IBD blocks can bias inferred length of IBD blocks." Society for Molecular Biology & Evolution (SMBE), San Juan, PR, Jun 2014.
2. Turchin MC, **Chiang CWK**, Palmer CD, Hirschhorn JN, GIANT consortium. "Intra-European allele frequency differences at height-associated SNPs suggest widespread selection on standing variation." Cold Spring Harbor Laboratory (CSHL), The Biology of Genomes, Cold Spring Harbor, NY, May 2011.
1. **Chiang CWK**, Dewing P, Vilain E. "The genetic effect of the male-determining gene Sry on brain sexual dimorphism and function." Southern California Conference of Undergraduate Research (SCCUR), Whittier College, Whittier, CA, Nov. 2004.

Contributed Abstracts: Posters (Last 5 Years, out of 25 total)

bolded name denotes group member

14. **Chiang CWK**. "The impact of Polynesian genetic ancestry on the architecture of complex traits in Native Hawaiians, and the challenges to study them." Society for Molecular Biology & Evolution (SMBE), Quebec City, Canada, July 2020. (Conference canceled due to COVID-19)
13. **Jeon S**, de Smith AJ, Musken IS, Metayer C, Ma X, Wiemels JL, **Chiang CWK**. "Multi-ethnic genome-wide association study of Acute Lymphoblastic Leukemia." International Genetic Epidemiology Society (IGES), Seoul, Korea, July 2020. (Conference canceled due to COVID-19)
12. **Fan C**, Mancuso N, **Chiang CWK**. "A genealogical estimate of genetic relationships." Biology of Genomes, CSHL, Cold Spring Harbor, NY, May 2020 (Online due to COVID-19).
11. **Lin M**, Caberto C, Wan P, Lum-Jones A, Tiirikainen M, Pooler L, Nakamura B, Sheng X, Le Marchand L, Wilkens L, Haiman C, Cheng I, **Chiang CWK**. "Oceanian reference panels are crucial for understanding the impact of a Polynesian-specific variant on adiposity in Native Hawaiians." American Society of Human Genetics (ASHG), Houston, TX, Oct 2019.
10. **Chen M**, Akiyama M, Sidore C, Ishigaki K, Okada Y, Kamatani Y, Schlessinger D, **Chiang CWK**. "Re-examining the evidence for polygenic adaptation at height-associated loci in mainland Europe and Sardinia." American Society of Human Genetics (ASHG), Houston, TX, Oct 2019. Reviewer's Choice Abstract
9. Lettre G, Chen MH, Raffield L, Mousas A, Jiang T, Akbari P, Sakaue S, Bao EL, Lareau CA, **Chen M**, **Chiang CWK**, Okada Y, Sankaran VG, Soranzo N, Reiner A, Johnson AD, Auer P, Blood-Cell Consortium. "The genetic architecture of hematological traits within and between populations." American Society of Human Genetics (ASHG), Houston, TX, Oct 2019.
8. **Chen M**, Akiyama M, Sidore C, Ishigaki K, Okada Y, Kamatani Y, Schlessinger D, **Chiang CWK**. "Re-examining the evidence for polygenic adaptation at height-associated loci in mainland Europe and Sardinia." Gordon Research Conference (GRC), Human Genetics & Genomics, Waterville, NH, Jul 2019.
7. **Sun H**, **Lin M**, Wilkens L, Le Marchand L, Haiman CA, **Chiang CWK**. "The impact of global and local Polynesian genetic ancestry on complex traits in Native Hawaiians." Biology of Genomes, CSHL, Cold Spring Harbor, NY, May 2019.
6. **Chen M**, Akiyama M, Sidore C, Ishigaki K, Okada Y, Kamatani Y, Schlessinger D, **Chiang CWK**. "Re-examining the evidence of polygenic adaptation at height-associated loci in mainland Europe and

Sardinia.” Biology of Genomes, CSHL, Cold Spring Harbor, NY, May 2019.

5. **Chiang CWK**, Mangul S, Robles C, Sankararaman S. “A comprehensive map of genetic variation in the world’s largest ethnic group – Han Chinese.” American Society of Human Genetics (ASHG), San Diego, CA, Oct 2018.
4. Noursome D, Jiang X, McKean-Cowdin R, **Chiang CWK**, Burkemper B, Torres M, Varma R. “Admixture and genome-wide association mapping of the retinal nerve fiber layer thickness in a multiethnic population.” Association for Research in Vision and Ophthalmology (ARVO), Honolulu, HI. Apr 2018.
3. **Chiang CWK**, Marcus J, Sidore C, Zoledziwska M, Steri M, Al-asadi H, Sanna S, Abecasis GR, Schlessinger D, Cucca F, Novembre J. “Using whole-genome sequencing to shed insight on the complex prehistory of Sardinia.” American Society of Human Genetics (ASHG), Vancouver, Canada, Oct 2016. *Reviewer’s Choice Abstract*
2. Marcus JH, **Chiang CWK**, Novembre J. “Testing directional selection on polygenic traits using ancient DNA.” American Society of Human Genetics (ASHG), Vancouver, Canada, Oct 2016.
1. **Chiang CWK**, Locke AE, Steinberg KM, Service S, Abel H, Ramensky V, Pirinen M, Stell L, Stringham HM, Jackson AU, Yajnik P, Ray D, Larson DE, Koboldt DC, Scott LJ, Fulton RS, Nelson J, Nicholas TJ, Stitzel NO, Hall IM, Sabatti C, Ripatti S, Salomaa V, Palotie A, Laakso M, Boehnke M, Wilson RK, Freimer NB. “Insights to the genetic architecture of cardiometabolic traits and population structure in 20,029 Finnish exomes.” UCLA Cardiovascular Symposium, Los Angeles, CA, Sep. 2016.

Invited Talks and Colloquia

2020.01.22	Department Seminar, Department of Genetics University of Georgia, Athens, GA.
2019.09.09	Institute Seminar, International Laboratory of Human Genome Research (LIIGH) National University of Mexico (UNAM), Queretero, Mexico.
2019.08.29	Department Seminar, Quantitative Computational Biology Section Department of Biological Sciences, USC, Los Angeles, CA.
2018.06.08	Informatics Center for Neurogenetics and Neurogenomics (ICNN) Symposium University of California, Los Angeles (UCLA), Los Angeles, CA.
2018.04.18	Asian Evo Conference, Symposium on “Human Evolution and Adaptation” China National GeneBank, Shenzhen, China.
2018.04.13	Institute Seminar, Institute of Biomedical Sciences (IBMS) Academia Sinica, Taipei, Taiwan.
2017.04.13	Department Seminar, Center for Genetic Epidemiology University of Southern California, Los Angeles, CA.
2017.02.23	Department Seminar, Virginia Institute for Psychiatric and Behavioral Genetics (VIPBG) Virginia Commonwealth University, Richmond, VA.
2017.01.26	Department Seminar, Department of Anthropology University of Utah, Salt Lake City, UT.
2015.05.06	Institute Seminar, CAS-MPG Partnered Institute of Computational Biology Chinese Academy of Science, Shanghai, China.
2015.04.30	Department Seminar, Center for Genomic Sciences University of Hong Kong, Hong Kong, SAR, China.
2015.04.09	Institute Seminar, Institute of Biomedical Sciences (IBMS)

Academic Awards and Recognitions

2020-2021	Undergraduate Research Associate Program Award, USC (To support undergraduate researcher stipend in academic year 2020-21)
2017	ASHG Charles J. Epstein Trainee Award for Excellence in Human Genetics Research (Semifinalist)
2013-2015	National Institute of Health NRSA Postdoctoral Fellowship, F32
2010	Summer Institute in Statistical Genetics Travel Scholarship
2007-2010	National Science Foundation Graduate Fellowship
2005	Director's Fellowship, Yale University (declined)
2005	Phi Beta Kappa Society
2004	Ira J. and Shirley Spoon Honors Collegium Scholarship, UCLA
2003	Golden Key International Honour Society
2001	CRC Press Freshmen Chemistry Achievement Award
2001	The National Society of Collegiate Scholar
2001	Alpha Lambda Delta Honor Society
2001	Phi Eta Sigma Honor Society

Mentored Student and Postdocs

Abbreviations: **PIBBS**, Programs in Biomedical and Biological Sciences PhD program; **CBB**, Computational Biology and Bioinformatics PhD program; **QB**, quantitative biology undergraduate program, USC.

Year	Name	Program	Role
2018-current	Minhui Chen, Ph.D.	Postdoctoral Fellow, USC	--
2018-current	Meng Lin, Ph.D.	Postdoctoral Fellow, USC	--
2018-current	Soyoung Jeon	Ph.D. student, PIBBS, USC	Co-PhD advisor
2019-current	Caoqi Fan	Ph.D. student, CBB, USC	PhD advisor
2019-current	Tsz Fung Chan	Ph.D. student, Epidemiology, USC	Co-PhD advisor
2019-current	Olzhas Iksan	Ph.D. student, Genetics, Kazakh National Univ.	External committee member
2019-current	Samuel Sommerer	B.S. student, QB, USC	--
2020-current	Sydney Rashid	B.S. student, QB, USC	--
2020-current	Jiemiao Chen	B.S. student, Mathematics, Shandong University	--
2018-2019	Hanxiao Sun	M.S. student, Biostatistics, USC	MS advisor
2020 Spring	Bryan Dinh	Rotation student, CBB, USC	--
2018 Fall	Audrey Nickle	Rotation student, PIBBS, USC	--
2018 Spring	Kuochang Tseng	Rotation student, PIBBS, USC	--
2018 Spring	Charles Bramlett	Rotation student, PIBBS, USC	--

Teaching

2019	Guest Lecturer, University of Southern California BISC 577A (Computational Biology Laboratory), Fall Term
2009	Teaching Assistant, Harvard University. LS1B (Genetics, Genomics, and Evolution), Spring Term
2008	Guest Lecturer, Harvard Medical School. Genetics 228 (Genetics in Medicine; "Using Genetics Databases"), Fall Term

2007 Teaching Assistant, Harvard Medical School.
Genetics 228 (Genetics in Medicine), Fall Term

Departmental and University Service

2019-2020 Epidemiology Ph.D. Admission Committee (chair, 2020)
2019-2020 Program in Biomedical and Biological Sciences (PIBBS) Ph.D. Admission Committee
2019 Quantitative Computational Biology Section Faculty Search Committee, Department
of Biological Sciences, USC
2018 Biostatistics Ph.D. Admission Committee

Professional Services and Scientific Communities

Editorship and Editorial Board: Review Editor (Frontier in Genetics, 2018-current)

Invited Manuscript Reviewer:

Nature	Nature Genetics	Nature Communications
Am J of Hum Genet	Science	Science Advances
PNAS	Molecular Biology and Evolution	Human Molecular Genetics
Genetics	G3: Genes, Genomes, Genetics	Bioinformatics
PLoS Genetics	PLoS ONE	Eur J of Hum Genet
Human Genetics	Frontiers in Genetics	Nutrition and Diabetes
Scientific Reports		

Invited Abstract Reviewer: American Society of Human Genetics (Evolutionary and Population Genetics, 2017)

Invited Grant Reviewer: Human Frontier Science Program (Mail-in Reviewer, 2018); Medical Research Council NIRG (Reviewer, 2019)

Symposia Organizer: 1st AsiaEvo Conference (2018)

Mentor: Trainee-Mentor Luncheon, American Society of Human Genetics (Evolution and Population Genetics, 2019)

Member: Society for Molecular Biology & Evolution (SMBE); American Society for Human Genetics (ASHG)

Media Appearances and Engagement

“Accuracy of Genotyping Chips Called into Question” *The Scientist*, Jul 24, 2019
<https://www.the-scientist.com/news-opinion/accuracy-of-genotyping-chips-called-into-question--66198>

“Hidden Stories of Chinese Migration and Culture Found in Giant Genetic Study” *The New York Times*, Oct 5, 2018
<https://www.nytimes.com/2018/10/05/science/china-genetics.html>

Outreach and Community Services

Hoyt Art Gallery, USC

Community outreach pairing local visual artist with biomedical researcher to collaborate for innovative artistic representation of ongoing research at Keck School of Medicine

2018 Partnered Biomedical Researcher

DNA Day Essay Contest, American Society of Human Genetics (ASHG)

2014-2015, 2017 Second-round Judge

“How to get a postdoc” workshop, Department of Ecology and Evolutionary Biology, UCLA

2015 Panelist

Science in the News, Boston, MA.

Science in the News is a free evening seminar series for the public aimed at explaining and discussing today's hottest scientific topics without the hype or exaggeration often introduced by media coverage.

2005-2006 Lecture Coordinator

“Stem Cells: Hope or Hype?” lecture series (2005)

“Obesity and Diabetes” lecture series (2006)

MATHCOUNTS, Suzanne Middle School, Walnut, CA

MATHCOUNTS is a nation-wide program for 6th to 8th grade students designed to elicit interest in mathematics through teamwork and competition.

2003-2005 Head Coach, Suzanne Middle School

1999-2003 Assistant Coach, Suzanne Middle School

2006, 2008, 2013 Volunteer, East San Gabriel Valley Chapter, CA