

Marcio Augusto Afonso de Almeida, Ph.D.

Assistant Professor of Human Genetics
South Texas Diabetes and Obesity Institute
University of Texas Rio Grande Valley
School of Medicine

Contact Information

BROBL Room 1.117
One West University Avenue
Brownsville, TX 78520-4933
Phone: 956-882-7519
Email: Marcio.Almeida@utrgv.edu

Education & Training

B.S. Immunology 2004 Pontifícia Universidade Católica de Campinas, Campinas, Brazil
Ph.D. Bioinformatics 2010 University of São Paulo, Cidade Universitária São Paulo, Brazil
Ph.D. Thesis title: Application of computational and statistical methods for the study of Cis gene regulation

Research Focus

I am a bioinformatics specialist born in Brazil where I started my studies on genetics and molecular biology. My scientific career started on the Heart Institute in Brazil where I started working on human genome data and especially with genomic annotation using public repositories. Using those tools, I developed my Ph.D. thesis in Brazil where I published my first 3 scientific manuscripts. I move to USA to complete my studies in 2011 and I was hired by the Texas Biomedical Research Institute, one of world leaders in methodological advances to study large human families with projects focused on diabetes and cardiovascular disease. Originally, my PhD training was focused on genomic annotation with the aid of public resources and the computational prediction of functional DNA elements by the application of a trained Hidden Markov Model (HMM) algorithm. During the past seven years, I was one of the main analysts working with the San Antonio Family Studies (SAFS) project whole genome sequencing data. I started to work at UTRGV in 2015 where we successfully continued our scientific investigations. In 2018, I was appointed as an assistant professor in the Department of Human Genetics at the UTRGV's school of medicine. During my scientific career, I have published a total of 49 scientific manuscripts.

Publications

1. Kim D., Justice A.E., Chittoor G., Blanco E., Burrows R., Graff M., Howard A.G., Wang Y., Rohde R., Buchanan V.L., Voruganti V.S., **Almeida M.**, Peralta J., Lehman D.M., Curran J.E., Comuzzie A.G., Duggirala R., Blangero J., Albala C., Santos J.L., Angel B., Lozoff B., Gahagan S., North K.E.. **Genetic determinants of metabolic biomarkers and their associations with cardiometabolic traits in Hispanic/Latino adolescents.** *Pediatric Research*, 2021 Oct 13. doi: 10.1038/s41390-021-01729-7.
2. Goyal S., Tanigawa Y., Zhang W., Chai J., **Almeida M.**, Sim X., Lerner M., Chainakul J., Ramiu J.G., Seraphin C., Apple B., Vaughan A., Muniu J., Peralta J., Lehman D.M., Ralhan S., Wander G.S., Singh J.R., Mehra N.K., Sidorov E., Peyton M.D., Blackett P.R., Curran J.E., Tai E.S., Dam R.V., Cheng C., Duggirala R., Blangero J., Chambers J.C., Sabanayagam C., Kooner J.S., Rivas M.A., Aston C.E., Sanghera D.K.. **APOC3 genetic variation, serum triglycerides, and risk of coronary artery disease in Asian Indians, Europeans, and other ethnic groups.** *Lipids in Health and Disease*, 2021 Sep 21;20(1):113. doi: 10.1186/s12944-021-01531-8.
3. Arya R., Lopez-Alvarenga J.C., **Almeida M.**, Kumar S., Peralta J., Diaz-Badillo A., Diego V.P., Resendez R.G., Fowler S.P., Jenkinson C.P., Lehman D., Curran J., Lynch J.L., Hale D.E., DeFronzo R.A., Mummidis S., Blangero J., Duggirala R.. **Exome-chip-wide Association Study of Biomarkers of Liver Function and Metabolic Dysfunction-Associated Fatty Liver Disease (MAFLD) in Mexican Americans.** *Frontiers in Medicine*, 2022 (Accepted for publication).
4. Grasby KL, Jahanshad N, Painter JN, Colodro-Conde L, Bralten J, Hibar DP, Lind PA, Pizzagalli F, Ching CRK, McMahon MAB, Shatokhina N, Zsembik LCP, Thomopoulos SI, Zhu AH, Strike LT, Agartz I, Alhusaini S, **Almeida MAA**, Alnæs D, Amlien IK, Andersson M, Ard T, Armstrong NJ, Ashley-Koch A, Atkins JR, Bernard M, Brouwer RM, Buimer EEL, Bülow R, Bürger C, Cannon DM, Chakravarty M, Chen Q, Cheung JW, Couvy-Duchesne B, Dale AM, Dalvie S, de Araujo TK, de Zubicaray GI, de Zwarte SMC, den Braber A, Doan NT, Dohm K, Ehrlich S, Engelbrecht HR, Erk S, Fan CC, Fedko IO, Foley SF, Ford JM, Fukunaga M, Garrett ME, Ge T, Giddaluru S, Goldman AL, Green MJ, Groenewold NA, Grotegerd D, Gurholt TP, Gutman BA, Hansell NK, Harris MA, Harrison MB, Haswell CC, Hauser M, Herms S, Heslenfeld DJ, Ho NF, Hoehn D, Hoffmann P, Holleran L, Hoogman M, Hottenga JJ, Ikeda M, Janowitz D, Jansen IE, Jia T, Jockwitz C, Kanai R, Karama S, Kasperaviciute D, Kaufmann T, Kelly S, Kikuchi M, Klein M, Knapp M, Knodt AR, Krämer B, Lam M, Lancaster TM, Lee PH, Lett TA, Lewis LB, Lopes-Cendes I, Luciano M, Macciardi F, Marquand AF, Mathias SR, Melzer TR, Milaneschi Y, Mirza-Schreiber N, Moreira JCV, Mühlleisen TW, Müller-Myhsok B, Najt P, Nakahara S, Nho K, Olde Loohuis LM, Orfanos DP, Pearson JF, Pitcher TL, Pütz B, Quidé Y, Ragothaman A, Rashid FM, Reay WR, Redlich R, Reinbold CS, Repple J, Richard G, Riedel BC, Risacher SL, Rocha CS, Mota NR, Salminen L, Saremi A, Saykin AJ, Schlag F, Schmaal L, Schofield PR, Secolin R, Shapland CY, Shen L, Shin J, Shumskaya E, Sønderby IE, Sprooten E, Tansey KE, Teumer A, Thalamuthu A, Tordesillas-Gutiérrez D, Turner JA, Uhlmann A, Vallerga CL, van der Meer D, van Donkelaar MMJ, van Eijk L, van Erp TGM, van Haren NEM, van Rooij D, van Tol MJ, Veldink JH, Verhoef E, Walton E, Wang M, Wang Y, Wardlaw JM, Wen W, Westlye LT, Whelan CD, Witt SH, Wittfeld K, Wolf C, Wolfers T, Wu JQ, Yasuda CL, Zaremba D, Zhang Z, Zwiers MP, Artiges E, Assareh AA, Ayesa-Arriola R, Belger A, Brandt CL, Brown GG, Cichon S, Curran JE, Davies GE, Degenhardt F, Dennis MF, Dietsche B, Djurovic S, Doherty CP, Espiritu R, Garijo D, Gil Y, Gowland PA, Green RC, Häusler AN, Heindel W, Ho BC, Hoffmann WU, Holsboer F, Homuth G, Hosten N, Jack CR Jr, Jang M, Jansen A, Kimbrel NA, Kolskår K, Koops S,

Krug A, Lim KO, Luykx JJ, Mathalon DH, Mather KA, Mattay VS, Matthews S, Mayoral Van Son J, McEwen SC, Melle I, Morris DW, Mueller BA, Nauck M, Nordvik JE, Nöthen MM, O'Leary DS, Opel N, Martinot MP, Pike GB, Preda A, Quinlan EB, Rasser PE, Ratnakar V, Reppermund S, Steen VM, Tooney PA, Torres FR, Veltman DJ, Voyvodic JT, Whelan R, White T, Yamamori H, Adams HHH, Bis JC, Debette S, Decarli C, Fornage M, Gudnason V, Hofer E, Ikram MA, Launer L, Longstreth WT, Lopez OL, Mazoyer B, Mosley TH, Roschupkin GV, Satizabal CL, Schmidt R, Seshadri S, Yang Q; Alzheimer's Disease Neuroimaging Initiative; CHARGE Consortium; EPIGEN Consortium; IMAGEN Consortium; SYS Consortium; Parkinson's Progression Markers Initiative, Alvim MKM, Ames D, Anderson TJ, Andreassen OA, Arias-Vasquez A, Bastin ME, Baune BT, Beckham JC, Blangero J, Boomsma DI, Brodaty H, Brunner HG, Buckner RL, Buitelaar JK, Bustillo JR, Cahn W, Cairns MJ, Calhoun V, Carr VJ, Caseras X, Caspers S, Cavalleri GL, Cendes F, Corvin A, Crespo-Facorro B, Dalrymple-Alford JC, Dannlowski U, de Geus EJC, Deary IJ, Delanty N, Depondt C, Desrivières S, Donohoe G, Espeseth T, Fernández G, Fisher SE, Flor H, Forstner AJ, Francks C, Franke B, Glahn DC, Gollub RL, Grabe HJ, Gruber O, Håberg AK, Hariri AR, Hartman CA, Hashimoto R, Heinz A, Henskens FA, Hillegers MHJ, Hoekstra PJ, Holmes AJ, Hong LE, Hopkins WD, Hulshoff Pol HE, Jernigan TL, Jönsson EG, Kahn RS, Kennedy MA, Kircher TTJ, Kochunov P, Kwok JBJ, Le Hellard S, Loughland CM, Martin NG, Martinot JL, McDonald C, McMahon KL, Meyer-Lindenberg A, Michie PT, Morey RA, Mowry B, Nyberg L, Oosterlaan J, Ophoff RA, Pantelis C, Paus T, Pausova Z, Penninx BWJH, Polderman TJC, Posthuma D, Rietschel M, Roffman JL, Rowland LM, Sachdev PS, Sämann PG, Schall U, Schumann G, Scott RJ, Sim K, Sisodiya SM, Smoller JW, Sommer IE, St Pourcain B, Stein DJ, Toga AW, Trollor JN, Van der Wee NJA, van 't Ent D, Völzke H, Walter H, Weber B, Weinberger DR, Wright MJ, Zhou J, Stein JL#, Thompson PM#, Medland SE#; Enhancing NeuroImaging Genetics through Meta-Analysis Consortium (ENIGMA)—Genetics working group. **The genetic architecture of the human cerebral cortex.** *Science*. 2020 Mar 20;367(6484). pii: eaay6690. doi: 10.1126/science.aay6690

5. Diego VP, Luu BW, Hofmann M, Dinh LV, **Almeida M**, Powell JS, Rajalingam R, Peralta JM, Kumar S, Curran JE, Sauna ZE, Kellerman R, Park Y, Key NS, Escobar MA, Huynh H, Verhagen AM, Williams-Blangero S, Lehmann PV, Maraskovsky E, Blangero J, Howard TE. **Quantitative HLA-class-II/factor VIII (FVIII) peptidomic variation in dendritic cells correlates with the immunogenic potential of therapeutic FVIII proteins in hemophilia A.** *Journal of Thrombosis Hemostasis*. 2020 Jan;18(1):201-216. doi: 10.1111/jth.14647
6. Blackburn N.B., Michael L.F., Meikle P.J., Peralta J.M., Mosior M., McAhren S., Bui H.H., Bellinger M.A., Giles C., Kumar S., Leandro A.C., **Almeida M.**, Weir J.M., Mahaney M.C., Dyer T.D., Almasy L., VandeBerg J.L., Williams-Blangero S., Glahn D.C., Duggirala R., Kowala M., Blangero J., Curran J.E. **Rare DEGS1 variant significantly alters de novo ceramide synthesis pathway.** *Journal of Lipid Research*. 2019 Jun 21. pii: jlr.P094433. doi: 10.1194/jlr.P094433
7. Tom E. Howard, Vincent P. Diego, Marco Hofmann, **Marcio Almeida**, Bernadette W. Luu, Long V. Dinh, Raja Rajalingam, Miguel Escobar, Joanne Curran, Sarah Williams-Blangero, Jerry Powell, John Blangero, Eugene Maraskovsky, Nigel S. Key, Zuben E. Sauna. **P059 Analysis of HLAII peptidomes presented by dendritic cells (DCs) from healthy donors and hemophilia-A (HA) patients with or without factor VIII (FVIII) inhibitors after ex vivo administration of different therapeutic FVIII proteins (tFVIII).** *Human Immunology*. 2018 Volume 79, Supplement, October, Pages 103-104. <https://doi.org/10.1016/j.humimm.2018.07.117>

8. Haakon E. Nustad, **Marcio Almeida**, Angelo J. Canty, Marissa LeBlanc, Christian M. Page, and Phillip E. Melton. **Epigenetics, heritability and longitudinal analysis.** *BMC Genetics.* 2018 Sep 17;19(Suppl 1):77. doi: 10.1186/s12863-018-0648-1
9. **Marcio Almeida**, Juan Peralta, Jose Garcia, Vincent Diego, Harald Goring, Sarah Williams-Blangero, and John Blangero. **Modeling methylation data as an additional genetic variance component.** *BMC Proceedings.* 2018 Sep 17;12(Suppl 9):29. doi: 10.1186/s12919-018-0128-7
10. Arya R, Escalante A, Farook VS, Restrepo JF, Battafarano DF, **Almeida M**, Kos MZ, Fourcaudot MJ, Mummidis S, Kumar S, Curran JE, Jenkinson CP, Blangero J, Duggirala R, Del Rincon I. **Data on genetic associations of carotid atherosclerosis markers in Mexican American and European American rheumatoid arthritis subjects.** *Data Briefing.* 2018 Feb 8; 17:820-829. doi: 10.1016/j.dib.2018.02.006. eCollection 2018 Apr.
11. Arya R, Escalante A, Farook VS, Restrepo JF, Battafarano DF, **Almeida M**, Kos MZ, Fourcaudot MJ, Mummidis S, Kumar S, Curran JE, Jenkinson CP, Blangero J, Duggirala R, Del Rincon I. **A genetic association study of carotid intima-media thickness (CIMT) and plaque in Mexican Americans and European Americans with rheumatoid arthritis.** *Atherosclerosis.* 2018 Apr; 271:92-101. doi: 10.1016/j.atherosclerosis.2017.11.024. Epub 2017 Nov 26.
12. Jun G, Manning A, **Almeida M**, Zawistowski M, Wood AR, Teslovich TM, Fuchsberger C, Feng S, Cingolani P, Gaulton KJ, Dyer T, Blackwell TW, Chen H, Chines PS, Choi S, Churchhouse C, Fontanillas P, King R, Lee S, Lincoln SE, Trubetskoy V, DePristo M, Fingerlin T, Grossman R, Grundstad J, Heath A, Kim J, Kim YJ, Laramie J, Lee J, Li H, Liu X, Livne O, Locke AE, Maller J, Mazur A, Morris AP, Pollin TI, Ragona D, Reich D, Rivas MA, Scott LJ, Sim X, Tearle RG, Teo YY, Williams AL, Zöllner S, Curran JE, Peralta J, Akolkar B, Bell GI, Burtt NP, Cox NJ, Florez JC, Hanis CL, McKeon C, Mohlke KL, Seielstad M, Wilson JG, Atzmon G, Below JE, Dupuis J, Nicolae DL, Lehman D, Park T, Won S, Sladek R, Altshuler D, McCarthy MI, Duggirala R, Boehnke M, Frayling TM, Abecasis GR, Blangero J. **Evaluating the contribution of rare variants to type 2 diabetes and related traits using pedigrees.** *PNAS (Proceeding National Academy of Science)- USA.* 2018 Jan 9;115(2):379-384. doi: 10.1073/pnas.1705859115. Epub 2017 Dec 26.
13. Cadby G, Melton PE, McCarthy NS, **Almeida M**, Williams-Blangero S, Curran JE, VandeBerg JL, Hui J, Beilby J, Musk AW, James AL, Hung J, Blangero J, Moses EK. **Pleiotropy of cardiometabolic syndrome with obesity-related anthropometric traits determined using empirically derived kinships from the Busselton Health Study.** *Human Genetics.* 2017 Nov 27. doi: 10.1007/s00439-017-1856-x. [Epub ahead of print]
14. Lake NJ, Taylor RL, Trahair H, Harikrishnan KN, Curran JE, **Almeida M**, Kulkarni H, Mukhamedova N, Hoang A, Low H, Murphy AJ, Johnson MP, Dyer TD, Mahaney MC, Göring HHH, Moses EK, Sviridov D, Blangero J, Jowett JBM, Bozaoglu K. **TRAK2, a novel regulator of ABCA1 expression, cholesterol efflux and HDL biogenesis.** *European Heart Journal.* 2017 Jun 26. doi: 10.1093/eurheartj/ehx315. [Epub ahead of print]
15. Kos MZ, Carless MA, Peralta J, Curran JE, Quillen EE, **Almeida M**, Blackburn A, Blondell L, Roalf DR, Pogue-Geile MF, Gur RC, Göring HHH, Nimagaonkar VL, Gur RE, Almasy L. **Exome sequences of multiplex, multigenerational families reveal schizophrenia risk loci with potential implications for neurocognitive performance.** *American Journal Medical Genetics B Neuropsychiatric Genetics.* 2017 Dec;174(8):817-827. doi: 10.1002/ajmg.b.32597. Epub 2017 Sep 13

16. Arya R, Escalante A, Farook VS, Restrepo JF, Battafarano DF, **Almeida M**, Kos MZ, Fourcaudot MJ, Mummidi S, Kumar S, Curran JE, Jenkinson, Blangero J, Duggirala R, Rincon ID. **A genetic association study of carotid intima-media thickness (CIMT) and plaque in Mexican Americans and European Americans with rheumatoid arthritis.** *Journal of atherosclerosis.* In press: DOI: <http://dx.doi.org/10.1016/j.atherosclerosis.2017.11.024>
17. Nicholson AM, Finch NA, **Almeida M**, Perkerson RB, van Blitterswijk M, Wojtas A, Cenik B, Rotondo S, Inskeep V, Almasy L, Dyer T, Peralta J, Jun G, Wood AR, Frayling TM, Fuchsberger C, Fowler S, Teslovich TM, Manning AK, Kumar S, Curran J, Lehman D, Abecasis G, Duggirala R, Pottier C, Zahir HA, Crook JE, Karydas A, Mitic L, Sun Y, Dickson DW, Bu G, Herz J, Yu G, Miller BL, Ferguson S, Petersen RC, Graff-Radford N, Blangero J and Rademakers R. **Prosaposin is a regulator of progranulin levels and oligomerization.** *Nature Communication.* 2016 Jun 30;7:11992. doi: 10.1038/ncomms11992
18. Peralta JM, **Almeida M**, Abraham LJ, Moses E and Blangero J. **Finding potential cis-regulatory loci using allele-specific chromatin accessibility as weights in a kernel-based variance component test.** *BMC Proceedings.* 2016 Oct 18;10(Suppl 7):103-108
19. **Almeida M**, Blondell L, Peralta J, Kent JW, Jun G, Teslovich TM, Fuchsberger C, Wood AR, Manning AK, Frayling TM, Cingolani PE, Sladek R, Dyer TS, Abecasis G, Duggirala R and Blangero J. **Independent test assessment using the extreme value distribution theory.** *BMC Proceedings.* 2016 Oct 18;10(Suppl 7):245-249
20. Blangero J, Teslovich TM, Sim X, **Almeida MA**, Peralta J, Jun G, Dyer TD and Almasy L. **Omics-squared: human genomic, transcriptomic and phenotypic data for Genetic Analysis Workshop 19.** *BMC Proceedings.* 2015;9 Suppl 8:S2 (In press)
21. Fiedrichs S, Malzahn D, Pugh EW, **Almeida M**, Liu XQ and Bailey JN. **Filtering genetic variants and placing informative prior based on putative biological function.** *BMC Genetics.* Feb 3:17 Suppl 2:8, 2016
22. Chittoor G, Kent JW Jr, **Almeida M**, Puppala S, Farook VS, Cole SA, Haack K, Göring HH, MacCluer JW, Curran JE, Carless MA, Johnson MP, Moses EK, Almasy L, Mahaney MC, Lehman DM, Duggirala R, Comuzzie AG, Blangero J and Voruganti VS. **GWAS and transcriptional analysis prioritize ITPR1 and CNTN4 for a serum uric acid 3p26 QTL in Mexican Americans.** *BMC Genomics.* 2016 Apr 2;17(1):276.
23. Kulkarni H, Mamani M, Peralta J, **Almeida M**, Dyer TD, Goring HH, Johnson MP, Duggirala R, Mahaney MC, Olvera RL, Almasy L, Glahn DC, Williams-Blangero S, Curran JE and Blangero J. **Soluble forms of intercellular and vascular cell adhesion molecules independently predict progression to type 2 diabetes in Mexican American families.** *PLoS One.* 2016 Mar 23;11(3):e0151177
24. Knowles EE, Kent JW Jr, McKay DR, Sprooten E, Mathias SR, Curran JE, Carless MA, **de Almeida MA**, Harald HH, Dyer TD, Olvera RL, Fox PT, Duggirala R, Almasy L, Blangero J and Glahn DC. **Genome-wide linkage on chromosome 10q26 for a dimensional scale of major depression.** *J Affect Disord.* 2016 Feb;191:123-31.
25. Kim YJ, Lee J, Kim BJ, Abecasis G, **Almeida M**, Altshuler D, Asimit JL, Atzmon G, Barber M, Barzilai N, Beer NL, Bell GI, Below J, Blackwell T, Blangero J, Boehnke M, Bowden DW, Burtt N, Chambers J, Chen H, Chen P, Chines PS, Choi S, Churchhouse C, Cingolani P, Cornes BK, Cox N, Day-Williams AG, Duggirala R, Dupuis J, Dyer T, Feng S, Fernandez-Tajes J, Ferreira T, Fingerlin TE, Flannick J, Florez J, Fontanillas P, Frayling TM, Fuchsberger C, Gamazon ER, Gaulton K, Ghosh S, Glaser B, Gloyn A, Grossman

- RL, Grundstad J, Hanis C, Heath A, Highland H, Horikoshi M, Huh IS, Huyghe JR, Ikram K, Jablonski KA, Jun G, Kato N, Kim J, Kim YJ, Kim BJ, Lee J, King CR, Kooner J, Kwon MS, Im HK, Laakso M, Lam KK, Lee J, Lee S, Lee S, Lehman DM, Li H, Lindgren CM, Liu X, Livne OE, Locke AE, Mahajan A, Maller JB, Manning AK, Maxwell TJ, Mazoure A, McCarthy MI, Meigs JB, Min B, Mohlke KL, Morris AP, Musani S, Nagai Y, Ng MC, Nicolae D, Oh S, Palmer N, Park T, Pollin TI, Prokopenko I, Reich D, Rivas MA, Scott LJ, Seielstad M, Cho YS, Sim X, Sladek R, Smith P, Tachmazidou I, Tai ES, Teo YY, Teslovich TM, Torres J, Trubetskoy V, Willems SM, Williams AL, Wilson JG, Wiltshire S, Won S, Wood AR, Xu W, Yoon J, Zawistowski M, Zeggini E, Zhang W, Zöllner S and Park T. **A new strategy for enhancing imputation quality of rare variants from next-generation sequencing data via combining SNP and exome chip data.** *BMC Genomics*. 2015 Dec 29;16:1109. doi: 10.1186/s12864-015-2192-y
26. Arya R, Del Rincon I, Farook VS, Restrepo JF, Winnier DA, Fourcaudot MJ, Battafarano DF, **de Almeida M**, Kumar S, Curran JE, Jenkinson CP, Blangero J, Duggirala Rand and Escalante A. **Genetic variants influencing joint damage in Mexican Americans and European Americans with rheumatoid arthritis.** *Genet Epidemiol*. 2015 Dec;39(8):678-88.
 27. Mathias SR, Knowles EE, Kent JW Jr, McKay DR, Curran JE, **de Almeida MA**, Dyer TD, Göring HH, Olvera RL, Duggirala R, Fox PT, Almasy L, Blangero J and Glahn DC. **Recurrent major depression and right hippocampal volume: A bivariate linkage and association study.** *Hum Brain Mapp*. 2016 Jan;37(1):191-202.
 28. Sprooten E, Gupta CN, Knowles EE, McKay DR, Mathias SR, Curran JE, Kent JW Jr, Carless MA, **Almeida MA**, Dyer TD4, Göring HH, Olvera RL, Kochunov P, Fox PT, Duggirala R, Almasy L, Calhoun VD, Blangero J, Turner JA and Glahn DC. **Genome-wide significant linkage of schizophrenia-related neuroanatomical trait to 12q24.** *Am J Med Genet B Neuropsychiatr Genet*. 2015 Dec;168(8):678-86.
 29. Kos MZ, Carless MA, Peralta J, Blackburn A, **Almeida M**, Roalf D, Pogue-Geile MF, Prasad K, Gur RC, Nimagaonkar V, Curran JE, Duggirala R, Glahn DC, Blangero J, Gur RE and Almasy L. **Exome sequence data from multigenerational families implicate AMPA Receptor trafficking in neurocognitive impairment and schizophrenia risk.** *Schizophr Bull*. 2015 Sep 24.
 30. Blackburn A, **Almeida M**, Dean A, Curran JE, Johnson MP, Moses EK, Abraham LJ, Carless MA, Dyer TD, Kumar S, Almasy L, Mahaney MC, Comuzzie A, Williams-Blangero S, Blangero J, Lehman DM and Göring HH. **Effects of copy number variable regions on local gene expression in white blood cells of Mexican Americans.** *European journal of human genetics*: EJHG 01/2015; DOI:10.1038/ejhg.2014.280
 31. Toledo RA, Hatakanaka R, Lourenço DM Jr, Lindsey SC, Camacho CP, **Almeida M**, Lima JV Jr, Sekiya T, Garralda E, Naslavsky MS, Yamamoto GL, Lazar M, Meirelles O, Sobreira TJ, Lebrao ML, Duarte YA, Blangero J, Zatz M, Cerutti JM, Maciel RM and Toledo SP. **Comprehensive assessment of the disputed RET Y791F shows no association with MTC susceptibility.** *Endocrine Related Cancer* 11/2014; DOI:10.1530/ERC-14-0491
 32. Wood AR, Tuke MA, Nalls M, Hernandez D, Gibbs JR, Lin H, Xu CS, Li Q, Shen J, Jun G, **Almeida M**, Tanaka T, Perry JR, Gaulton K, Rivas M, Pearson R, Curran JE, Johnson MP, Göring HH, Duggirala R, Blangero J, McCarthy MI, Bandinelli S, Murray A, Weedon MN, Singleton A, Melzer D, Ferrucci L and Frayling TM. **Whole-genome sequencing to understand the genetic architecture of common gene expression and biomarker phenotypes.** *Human Molecular Genetics* 11/2014; 24(5). DOI:10.1093/hmg/ddu560

33. Bellis C, Kulkarni H, Mamtani M, Kent JW Jr, Wong G, Weir JM, Barlow CK, Diego V, **Almeida M**, Dyer TD, Göring HH, Almasy L, Mahaney MC, Comuzzie AG, Williams-Blangero S, Meikle PJ, Blangero J and Curran JE. **Human plasma lipidome is pleiotropically associated with cardiovascular risk factors and death.** *Circulation Cardiovascular Genetics* 11/2014; 7(6). DOI:10.1161/CIRGENETICS.114.000600
34. Knowles EE, McKay DR, Kent JW Jr, Sprooten E, Carless MA, Curran JE, **de Almeida MA**, Dyer TD, Göring HH, Olvera RL, Duggirala R, Fox PT, Almasy L, Blangero J, Glahn DC. **Pleiotropic locus for emotion recognition and amygdala volume identified using univariate and bivariate linkage.** *American Journal of Psychiatry* 10/2014; 172(2). DOI:10.1176/appi.ajp.2014.14030311.
35. Aslibekyan S, **Almeida M** and Tintle N. **Pathway analysis approaches for rare and common variants: Insights from Genetic Analysis Workshop 18.** *Genetic Epidemiology* 09/2014; 38 Suppl 1:S86-91. DOI:10.1002/gepi.21831.
36. Almasy L, Dyer TD, Peralta JM, Jun G, Wood AR, Fuchsberger C, **Almeida MA**, Kent JW Jr, Fowler S, Blackwell TW, Puppala S, Kumar S, Curran JE, Lehman D, Abecasis G, Duggirala R, Blangero J; **T2D-GENES Consortium.** **Data for Genetic Analysis Workshop 18: Human whole genome sequence, blood pressure, and simulated phenotypes in extended pedigrees.** *BMC Proceedings* 06/2014; 8(Suppl 1):S2. DOI:10.1186/1753-6561-8-S1-S2
37. Quillen EE, Voruganti VS, Chittoor G, Rubicz R, Peralta JM, **Almeida MA**, Kent JW Jr, Diego VP, Dyer TD, Comuzzie AG, Göring HH, Duggirala R, Almasy L and Blangero J. **Evaluation of estimated genetic values and their application to genome-wide investigation of systolic blood pressure.** *BMC proceedings* 06/2014; 8(Suppl 1):S66. DOI:10.1186/1753-6561-8-S1-S66.
38. Peralta JM, **Almeida M**, Kent JW Jr and Blangero J. **A variance component-based gene burden test.** *BMC proceedings* 06/2014; 8 (Suppl 1 Genetic Analysis Workshop 18):S49. DOI:10.1186/1753-6561-8-S1-S49.
39. **Almeida M**, Peralta JM, Farook V, Puppala S, Kent JW Jr, Duggirala R and Blangero J. **Pedigree-based random effect tests to screen gene pathways.** *BMC proceedings* 06/2014; 8(Suppl 1 Genetic Analysis Workshop):S100. DOI:10.1186/1753-6561-8-S1-S100.
40. Kulkarni H, Meikle PJ, Mamtani M, Weir JM, **Almeida M**, Diego V, Peralta JM, Barlow CK, Bellis C, Dyer TD, Almasy L, Mahaney MC, Comuzzie AG, Göring HH, Curran JE, Blangero J. **Plasma lipidome is independently associated with variability in metabolic syndrome in Mexican American families.** *The Journal of Lipid Research*. 03/2014; 55(5). DOI:10.1194/jlr.M044065
41. Knowles EE, Carless MA, **de Almeida MA**, Curran JE, McKay DR, Sprooten E, Dyer TD, Göring HH, Olvera R, Fox P, Almasy L, Duggirala R, Kent JW Jr, Blangero J, Glahn DC. **Genome-wide significant localization for working and spatial memory: Identifying genes for psychosis using models of cognition.** *American Journal of Medical Genetics Part B Neuropsychiatric Genetics* 01/2014; 165(1). DOI:10.1002/ajmg.b.32211.
42. Valentim CL, Cioli D, Chevalier FD, Cao X, Taylor AB, Holloway SP, Pica-Mattoccia L, Guidi A, Basso A, Tsai IJ, Berriman M, Carvalho-Queiroz C, **Almeida M**, Aguilar H, Frantz DE, Hart PJ, LoVerde PT, Anderson TJ. **Genetic and Molecular Basis of Drug Resistance and Species-Specific Drug Action in Schistosome Parasites.** *Science* 11/2013; 342(6164). DOI:10.1126/science.1243106 .
43. Havill LM, Coan HB, Dyer TD, Kent JW, Choudary A, Nicolella DP, Carless MA, Kumar S,

- Almeida MA**, Duggirala R, Glahn DC, Mahaney MC, Blangero J and Curran JE. **GWAS of self-reported osteoarthritis in Mexican Americans from the San Antonio Family Study**. *Osteoarthritis and Cartilage* 04/2013; 21:S173. DOI:10.1016/j.joca.2013.02.373
44. Coan HB, Curran JE, Dyer TD, Kent JW, Choudary A, Nicolella DP, Carless MA, Kumar S, **Almeida MA**, Duggirala R, Glahn DC, Mahaney MC, Blangero J and Havill LM. **Variation in osteoarthritis biomarker serum comp levels in Mexican Americans is associated with SNPs in a region of chromosome 22q encompassing MICAL3, BCL2L13, and BID**. *Osteoarthritis and Cartilage*, 04/2013; 21:S172. DOI:10.1016/j.joca.2013.02.371.
45. Blangero J, Diego VP, Dyer TD, **Almeida M**, Peralta J, Kent JW Jr, Williams JT, Almasy L, Göring HH. **A Kernel of Truth. Statistical Advances in Polygenic Variance Component Models for Complex Human Pedigrees**. *Advances in genetics*, 01/2013; 81:1-31. DOI:10.1016/B978-0-12-407677-8.00001-4.
46. Stein JL, Medland SE, Vasquez AA, Hibar DP, Senstad RE, Winkler AM, Toro R, Appel K, Bartecik R, Bergmann Ø, Bernard M, Brown AA, Cannon DM, Chakravarty MM, Christoforou A, Domin M, Grimm O, Hollinshead M, Holmes AJ, Homuth G, Hottenga JJ, Langan C, Lopez LM, Hansell NK, Hwang KS, Kim S, Laje G, Lee PH, Liu X, Loth E, Lourdusamy A, Mattingtsdal M, Mohnke S, Maniega SM, Nho K, Nugent AC, O'Brien C, Papmeyer M, Pütz B, Ramasamy A, Rasmussen J, Rijpkema M, Risacher SL, Roddey JC, Rose EJ, Ryten M, Shen L, Sprooten E, Strengman E, Teumer A, Trabzuni D, Turner J, van Eijk K, van Erp TG, van Tol MJ, Wittfeld K, Wolf C, Woudstra S, Aleman A, Alhusaini S, Almasy L, Binder EB, Brohawn DG, Cantor RM, Carless MA, Corvin A, Czisch M, Curran JE, Davies G, **de Almeida MA**, Delanty N, Depondt C, Duggirala R, Dyer TD, Erk S, Fagerness J, Fox PT, Freimer NB, Gill M, Göring HH, Hagler DJ, Hoehn D, Holsboer F, Hoogman M, Hosten N, Jahanshad N, Johnson MP, Kasperaviciute D, Kent JW Jr, Kochunov P, Lancaster JL, Lawrie SM, Liewald DC, Mandl R, Matarin M, Mattheisen M, Meisenzahl E, Melle I, Moses EK, Mühlleisen TW, Nauck M, Nöthen MM, Olvera RL, Pandolfo M, Pike GB, Puls R, Reinvang I, Rentería ME, Rietschel M, Roffman JL, Royle NA, Rujescu D, Savitz J, Schnack HG, Schnell K, Seiferth N, Smith C, Steen VM, Valdés Hernández MC, Van den Heuvel M, van der Wee NJ, Van Haren NE, Veltman JA, Völzke H, Walker R, Westlye LT, Whelan CD, Agartz I, Boomsma DI, Cavalleri GL, Dale AM, Djurovic S, Drevets WC, Hagoort P, Hall J, Heinz A, Jack CR Jr, Foroud TM, Le Hellard S, Maciardi F, Montgomery GW, Poline JB, Porteous DJ, Sisodiya SM, Starr JM, Sussmann J, Toga AW, Veltman DJ, Walter H, Weiner MW; Alzheimer's Disease Neuroimaging Initiative; EPIGEN Consortium; IMAGEN Consortium; Saguenay Youth Study Group, Bis JC, Ikram MA, Smith AV, Gudnason V, Tzourio C, Vernooij MW, Launer LJ, DeCarli C, Seshadri S; Cohorts for Heart and Aging Research in Genomic Epidemiology Consortium, Andreassen OA, Apostolova LG, Bastin ME, Blangero J, Brunner HG, Buckner RL, Cichon S, Coppola G, de Zubicaray GI, Deary IJ, Donohoe G, de Geus EJ, Espeseth T, Fernández G, Glahn DC, Grabe HJ, Hardy J, Hulshoff Pol HE, Jenkinson M, Kahn RS, McDonald C, McIntosh AM, McMahon FJ, McMahon KL, Meyer-Lindenberg A, Morris DW, Müller-Myhsok B, Nichols TE, Ophoff RA, Paus T, Pausova Z, Penninx BW, Potkin SG, Sämann PG, Saykin AJ, Schumann G, Smoller JW, Wardlaw JM, Weale ME, Martin NG, Franke B, Wright MJ, Thompson PM; **Enhancing Neuro Imaging Genetics through Meta-Analysis Consortium. Identification of common variants associated with human hippocampal and intracranial volumes**. *Nature Genetics* 04/2012; 44(5):552-61. DOI:10.1038/ng.2250.
47. Afonso de **Almeida MA**, Vançan Russo Horimoto AR, Lopes de Oliveira PS, Krieger JE, da Costa Pereira A. **Different approaches for dealing with rare variants in family-based genetic studies: An application of a Genetic Analysis Workshop 17 problem**.

- BMC proceedings*, 11/2011; 5 Suppl 9(Suppl 9):S78. DOI:10.1186/1753-6561-5-S9-S78.
- 48. Almeida MA, Oliveira PS, Pereira TV, Krieger JE, Pereira AC. **An empirical evaluation of imputation accuracy for association statistics reveals increased type-I error rates in genome-wide associations.** *BMC Genetics* 01/2011; 12:10. DOI:10.1186/1471-2156-12-10
 - 49. Castillo HA, Cravo RM, Azambuja AP, Simões-Costa MS, Sura-Trueba S, Gonzalez J, Slonimsky E, Almeida K, Abreu JG, **de Almeida MA**, Sobreira TP, de Oliveira SH, de Oliveira PS, Signore IA, Colombo A, Concha ML, Spengler TS, Bronner-Fraser M, Nobrega M, Rosenthal N, Xavier-Neto J. **Insights into the organization of dorsal spinal cord pathways from an evolutionarily conserved raldh2 intronic enhancer.** *Development* 02/2010; 137(3):507-18. DOI:10.1242/dev.043257