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Profile

Statement and Interests

Publications



Select Publications

Ye Z, Atkinson E, Fridley B, De Andrade M. Comparison of variable and model selection methods for genetic association studies using the GAW15 simulated data. BMC Proceedings. 2007;1(Suppl 1):S34.

Hebbring S, Slager S, Epperla N, Mazza JJ, Ye Z, Zhou Z, Achenbach S, Vasco DA, Call T, Rabe K, Kay NE, Caporaso NE, Lanasa M, Camp N, Strom S, Goldin L, Cerhan J, Brilliant MH, Schrodi SJ. Genetic evidence of PTPN22 effects on chronic lymphocytic leukemia. BLOOD. 2013 Jan;121(1):237-8. PubMed ID: 23287625 (http://www.ncbi.nlm.nih.gov/pubmed/23287625)

Hebbring S, Schrodi SJ, Ye Z, Zhou Z, Page D, Brilliant MH. A PheWAS approach in studying HLA-DRB1*1501 GENES AND IMMUNITY. 2013 Apr;14(3):187-91.

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Acharya A, Hernandez P, Thyvalikakath TP, Ye Z, Song M, Schleyer T. Development and initial validation of a content taxonomy for patient records in general dentistry. INTERNATIONAL JOURNAL OF MEDICAL INFORMATICS. 2013 Dec;82(12):1171-1182.

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PubMed ID: 23989841 (http://www.ncbi.nlm.nih.gov/pubmed/23989841)

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Ye Z, Vasco DA, Carter TC, Brilliant MH, Schrodi SJ, Shukla SK. Genome wide association study of SNP-, gene-, and pathway-based approaches to identify genes influencing susceptibility to Staphylococcus aureus infections. Front Genet. 2014 May;5:125.

PubMed ID: <u>24847357</u> (http://www.ncbi.nlm.nih.gov/pubmed/24847357)

Finamore JD, Ray W, Kadolph C, Rastegar-Mojarad M, Ye Z, Bohne J, Tachinardi U, Mendonca E, Finnegan BJ, Bartkowiak BA, Weichelt B, Lin S. D1-3: Marshfield Dictionary of Clinical and Translational Science (MD-CTS): An Online Reference for Clinical and Translational Science Terminology Clinical Medicine & Research. 2014 Sep;12(1-2):d1-3.

Anderson ER, Ye Z, Caldwell MD, Burmester JK. SNPs Previously Associated with Dupuytren's Disease Replicated in a North American Cohort. Clinical Medicine & Research. 2014 Dec;12(3-4):133-7.

PubMed ID: <u>24573701</u> (http://www.ncbi.nlm.nih.gov/pubmed/24573701)

McPherson EW, Zaleski CA, Ye Z, Lin S. Rodriguez syndrome with SF3B4 mutation: A severe form of Nager syndrome? American Journal of Medical Genetics Part A. 2014 Jul;164(7):1841-5. PubMed ID: 24715698 (http://www.ncbi.nlm.nih.gov/pubmed/24715698)

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He M, Person TN, Hebbring S, Heinzen E, Ye Z, Schrodi SJ, McPherson EW, Lin S, Peissig PL, Brilliant MH, O'Rawe J, Robinson RJ, Lyon GJ, Wang K. SeqHBase: a big data toolset for family based sequencing data analysis. JOURNAL OF MEDICAL GENETICS. 2015 Apr;52(4):282-8. PubMed ID: 25587064 (http://www.ncbi.nlm.nih.gov/pubmed/25587064)

Ye Z, Kadolph C, Strenn R, Wall D, McPherson EW, Lin S. WHATIF: An open-source desktop application for extraction and management of the incidental findings from next-generation sequencing variant data. Comput Biol Med. 2016 Jan;68:165-9..