

Universitätsmedizin Göttingen
Publikationen und Hochschulschriften 2014
Genetische Epidemiologie

Journalbeiträge

1. Albrecht C, Malzahn D, Brameier M, Hermes M, Ansari AA, Walter L (2014) Progression to AIDS in SIV-Infected Rhesus Macaques is Associated with Distinct KIR and MHC class I Polymorphisms and NK Cell Dysfunction. *Front Immunol* 5: 600, doi: 10.3389/fimmu.2014.00600
2. Badowska DM, Brzózka MM, Chowdhury A, Malzahn D, Rossner MJ (2014) Data calibration and reduction allows to visualize behavioural profiles of psychosocial influences in mice towards clinical domains. *EUR ARCH PSY CLIN N* E-Pub: E-Pub, doi: 10.1007/s00406-014-0532-6
3. Bickeböller H, Bailey JN, Beyene J, Cantor RM, Cordell HJ, Culverhouse RC, Engelma CD, Fardo DW, Ghosh S, König IR, Lorenzo Bermejo J, Melton PE, Santorico SA, Satten GA, Sun L, Tintle NL, Ziegler A, MacCluer JW, Almasy L (2014) Genetic Analysis Workshop 18: Methods and strategies for analyzing human sequence and phenotype data in members of extended pedigrees. *BMC Proc* 8(Suppl 1): S1, doi: 10.1186/1753-6561-8-S1-S1
4. Chen H, Malzahn D, Balliu B, Li C, Bailey JN (2014) Testing genetic association with rare and common variants in family data. *GENET EPIDEMIOL* 38 Suppl 1: S37-43, doi: 10.1002/gepi.21823
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9. Malzahn D, Friedrichs S, Rosenberger A, Bickeböller H (2014) Kernel score statistic for dependent data. *BMC Proc* 8(Suppl 1): S41, doi: 10.1186/1753-6561-8-S1-S41
10. Malzahn D, Müller-Nurasyid M, Heid IM, Wichmann HE, KORA study group, Bickeböller H (2014) Controversial association results for INSIG2 on body mass index may be explained by interactions with age and with MC4R. *EUR J HUM GENET* 22(10): 1217-24, doi: 10.1038/ejhg.2014.3
11. Wang Y, McKay JD, Rafnar T, Wang Z, Timofeeva MN, Broderick P, Zong X, Laplana M, Wei Y, Han Y, Lloyd A, Delahaye-Sourdeix M, Chubb D, Gaborieau V, Wheeler W, Chatterjee N, Thorleifsson G, Sulem P, Liu G, Kaaks R, Henrion M, Kinnersley B, Vallée M, LeCalvez-Kelm F, Stevens VL, Gapstur SM, Chen WV, Zaridze D, Szeszenia-Dabrowska N, Lissowska J, Rudnai P, Fabianova E, Mates D, Bencko V, Foretova L, Janout V, Krokan HE, Gabrielsen ME, Skorpen F, Vatten L, Njølstad I, Chen C, Goodman G, Benhamou S, Vooder T, Välk K, Nelis M, Metspalu A, Lener M, Lubiński J, Johansson M, Vineis P, Agudo A, Clavel-Chapelon F, Bueno-de-Mesquita HB, Trichopoulos D, Khaw KT, Johansson M, Weiderpass E, Tjønneland A, Riboli E, Lathrop M, Scelo G, Albanes D, Caporaso NE, Ye Y, Gu J, Wu X, Spitz MR, Dienemann H, Rosenberger A, Su L, Matakidou A, Eisen T, Stefansson K, Risch A, Chanock SJ, Christiani DC, Hung RJ, Brennan P, Landi MT, Houlston RS, Amos CI (2014) Rare variants of large effect in BRCA2 and CHEK2 affect risk of lung cancer. *NAT GENET* 46(7): 736-41, doi: 10.1038/ng.3002
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