

Biobank - 9160

Albrecht E, Waldenberger M, Krumsiek J, Evans AM, Jeratsch U, Breier M, Adamski J, Koenig W, Zeilinger S, Fuchs C, Klopp N, Theis FJ, Wichmann HE, Suhre K, Illig T, Strauch K, Peters A, Gieger C, Kastenmüller G, Doering A, Meisinger C. Metabolite profiling reveals new insights into the regulation of serum urate in humans. *Metabolomics* 2014;10:141-151

Originalpublikation - IF 3,965

Volltext: <http://dx.doi.org/10.1007/s11306-013-0565-2>

Biobank - 9160

Altmaier E, Fobo G, Heier M, Thorand B, Meisinger C, Römisch-Margl W, Waldenberger M, Gieger C, Illig T, Adamski J, Suhre K, Kastenmüller G. Metabolomics approach reveals effects of antihypertensives and lipid-lowering drugs on the human metabolism. *Eur J Epidemiol* 2014;29(5):325-336

Originalpublikation - IF 5,147

Volltext: <http://dx.doi.org/10.1007/s10654-014-9910-7>

Klinik für Päd. Nieren-, Leber und Stoffwechselerkrankungen - 6720 ; Institut für Diagnostische und Interventionelle Neuroradiologie - 8210

Bauer A, Loos S, Wehrmann C, Horstmann D, Donnerstag F, Lemke J, Hillebrand G, Lobel U, Pape L, Haffner D, Bindt C, Ahlenstiel T, Melk A, Lehnhardt A, Kemper MJ, Oh J, Hartmann H. Neurological involvement in children with *E. coli* O104:H4-induced hemolytic uremic syndrome. *Pediatr Nephrol* 2014;29(9):1607-1615

Originalpublikation - IF 2,881

Volltext: <http://dx.doi.org/10.1007/s00467-014-2803-x>

Biobank - 9160

Baumert J, Huang J, McKnight B, Sabater-Lleal M, Steri M, Chu AY, Trompet S, Lopez LM, Fornage M, Teumer A, Tang W, Rudnicka AR, Malarstig A, Hottenga JJ, Kavousi M, Lahti J, Tanaka T, Hayward C, Huffman JE, Morange PE, Rose LM, Basu S, Rumley A, Stott DJ, Buckley BM, de Craen AJ, Sanna S, Masala M, Biffar R, Homuth G, Silveira A, Sennblad B, Goel A, Watkins H, Müller-Nurasyid M, Ruckerl R, Taylor K, Chen MH, de Geus EJ, Hofman A, Witteman JC, de Maat MP, Palotie A, Davies G, Siscovick DS, Kolcic I, Wild SH, Song J, McArdle WL, Ford I, Sattar N, Schlessinger D, Grotevendt A, Franzosi MG, Illig T, Waldenberger M, Lumley T, Tofler GH, Willemsemen G, Uitterlinden AG, Rivadeneira F, Raikkonen K, Chasman DI, Folsom AR, Lowe GD, Westendorp RG, Slagboom PE, Cucca F, Wallachofski H, Strawbridge RJ, Seedorf U, Koenig W, Bis JC, Mukamal KJ, van Dongen J, Widen E, Franco OH, Starr JM, Liu K, Ferrucci L, Polasek O, Wilson JF, Oudot-Mellakh T, Campbell H, Navarro P, Bandinelli S, Eriksson J, Boomsma DI, Dehghan A, Clarke R, Hamsten A, Boerwinkle E, Jukema JW, Naitza S, Ridker PM, Volzke H, Deary IJ, Reiner AP, Tregouet DA, O'Donnell CJ, Strachan DP, Peters A, Smith NL. No evidence for genome-wide interactions on plasma fibrinogen by smoking, alcohol consumption and body mass index: results from meta-analyses of 80,607 subjects. *PLoS One* 2014;9(12):e111156

Originalpublikation - IF 3,534

Volltext: <http://dx.doi.org/10.1371/journal.pone.0111156>

Biobank - 9160

Beilina A, Rudenko IN, Kaganovich A, Civiero L, Chau H, Kalia SK, Kalia LV, Lobbstaedt E, Chia R, Ndukwue K, Ding J, Nalls MA, International Parkinson's Disease Genomics Consortium, North American Brain Expression Consortium, Olszewski M, Hauser DN, Kumaran R, Lozano AM, Baekelandt V, Greene LE, Taymans JM, Greggio E, Cookson MR. Unbiased screen for interactors of leucine-rich repeat kinase 2 supports a common pathway for sporadic and familial Parkinson disease. *Proc Natl Acad Sci U S A* 2014;111(7):2626-2631

Originalpublikation - IF 9,809

Volltext: <http://dx.doi.org/10.1073/pnas.1318306111>

Biobank - 9160

Claussnitzer M, Dankel SN, Klocke B, Grallert H, Glunk V, Berulava T, Lee H, Oskolkov N, Fadista J, Ehlers K, Wahl S, Hoffmann C, Qian K, Rönn T, Riess H, Müller-Nurasyid M, Bretschneider N, Schroeder T, Skurk T, Horsthemke B, DIAGRAM+ Consortium, Spieler D, Klingenspor M, Seifert M, Kern MJ, Mejhert N, Dahlman I, Hansson O, Hauck SM, Blüher M, Arner P, Groop L, Illig T, Suhre K, Hsu YH, Mellgren G, Hauner H, Laumen H. Leveraging cross-species transcription factor binding site patterns: from diabetes risk loci to disease mechanisms. *Cell* 2014;156(1-2):343-358

Originalpublikation - IF 33,116

Volltext: <http://dx.doi.org/10.1016/j.cell.2013.10.058>

Biobank - 9160

DIAbetes Genetics Replication And Meta-analysis (DIAGRAM) Consortium, Asian Genetic Epidemiology Network Type 2 Diabetes (AGEN-T2D) Consortium, South Asian Type 2 Diabetes (SAT2D) Consortium, Mexican American Type 2 Diabetes (MAT2D) Consortium, Type 2 Diabetes Genetic Exploration by Next-generation sequencing in multi-ethnic Samples (T2D-GENES) Consortium, Mahajan A, Go MJ, Zhang W, Below JE, Gaulton KJ, Ferreira T, Horikoshi M, Johnson AD, Ng MC, Prokopenko I, Saleheen D, Wang X, Zeggini E, Abecasis GR, Adair LS, Almgren P, Atalay M, Aung T, Baldassarre D, Balkau B, Bao Y, Barnett AH, Barroso I, Basit A, Been LF, Beilby J, Bell GI, Benediktsson R, Bergman RN, Boehm BO, Boerwinkle E, Bonycastle LL, Burtt N, Cai Q, Campbell H, Carey J, Cauchi S, Caulfield M, Chan JC, Chang LC, Chang TJ, Chang YC, Charpentier G, Chen CH, Chen H, Chen YT, Chia KS, Chidambaram M, Chines PS, Cho NH, Cho YM, Chuang LM, Collins FS, Cornelis MC, Couper DJ, Crenshaw AT, van Dam RM, Danesh J, Das D, de Faire U, Dedousis G, Deloukas P, Dimas AS, Dina C, Doney AS, Donnelly PJ, Dorkhan M, van Duijn C, Dupuis J, Edkins S, Elliott P, Emilsson V, Erbel R, Eriksson JG, Escobedo J, Esko T, Eury E, Florez JC, Fontanillas P, Forouhi NG, Forsen T, Fox C, Fraser RM, Frayling TM, Froguel P, Frossard P, Gao Y, Gertow K, Gieger C, Gigante B, Grallert H, Grant GB, Grrop LC, Groves CJ, Grundberg E, Guiducci C, Hamsten A, Han BG, Hara K, Hassanali N, Hattersley AT, Hayward C, Hedman AK, Herder C, Hofman A, Holmen OL, Hovingh K, Hreidarsson AB, Hu C, Hu FB, Hui J, Humphries SE, Hunt SE, Hunter DJ, Hveem K, Hydrie ZI, Ikegami H, Illig T, Ingelsson E, Islam M, Isomaa B, Jackson AU, Jafar T, James A, Jia W, Jockel KH, Jonsson A, Jowett JB, Kadokawa T, Kang HM, Kanoni S, Kao WH, Kathiresan S, Kato N, Kutalanda P, Keinanen-Kiukaanniemi KM, Kelly AM, Khan H, Khaw KT, Khor CC, Kim HL, Kim S, Kim YJ, Kinnunen L, Klopp N, Kong A, Korpi-Hyovalti E, Kowlessur S, Kraft P, Kravic J, Kristensen MM, Krishika S, Kumar A, Kumate J, Kuusisto J, Kwak SH, Laakso M, Lagou V, Lakka TA, Langenberg C, Langford C, Lawrence R, Leander K, Lee JM, Lee NR, Li M, Li X, Li Y, Liang J, Liju S, Lim WY, Lind L, Lindgren CM, Lindholm E, Liu CT, Liu JJ, Lobbens S, Long J, Loos RJ, Lu W, Luan J, Lyssenko V, Ma RC, Maeda S, Magi R, Mannisto S, Matthews DR, Meigs JB, Melander O, Metspalu A, Meyer J, Mirza G, Mihailov E, Moebus S, Mohan V, Mohlke KL, Morris AD, Muhleisen TW, Müller-Nurasyid M, Musk B, Nakamura J, Nakashima E, Navarro P, Ng PK, Nica AC, Nilsson PM, Njolstad I, Nothen MM, Ohnaka K, Ong TH, Owen KR, Palmer CN, Pankow JS, Park KS, Parkin M, Pechlivanis S, Pedersen NL, Peltonen L, Perry JR, Peters A, Pinidiyapathirage JM, Platou CG, Potter S, Price JF, Qi L, Radha V, Rallidis L, Rasheed A, Rathman W, Rauramaa R, Raychaudhuri S, Rayner NW, Rees SD, Rehnberg E, Ripatti S, Robertson N, Roden M, Rossin EJ, Rudan I, Rybin D, Saaristo TE, Salomaa V, Saltevo J, Samuel M, Sanghera DK, Saramies J, Scott J, Scott LJ, Scott RA, Segre AV, Sehmi J, Sennblad B, Shah N, Shah S, Shera AS, Shu XO, Shuldiner AR, Sigurdsson G, Sijbrands E, Silveira A, Sim X, Sivapalaratnam S, Small KS, So WY, Stancakova A, Stefansson K, Steinbach G, Steinhorsdottir V, Stirrups K, Strawbridge

RJ, Stringham HM, Sun Q, Suo C, Syvanen AC, Takayanagi R, Takeuchi F, Tay WT, Teslovich TM, Thorand B, Thorleifsson G, Thorsteinsdottir U, Tikkainen E, Trakalo J, Tremoli E, Trip MD, Tsai FJ, Tuomi T, Tuomilehto J, Uitterlinden AG, Valladares-Salgado A, Vedantam S, Veglia F, Voight BF, Wang C, Wareham NJ, Wennauer R, Wickremasinghe AR, Wilsgaard T, Wilson JF, Wiltshire S, Winckler W, Wong TY, Wood AR, Wu JY, Wu Y, Yamamoto K, Yamauchi T, Yang M, Yengo L, Yokota M, Young R, Zabaneh D, Zhang F, Zhang R, Zheng W, Zimmet PZ, Altshuler D, Bowden DW, Cho YS, Cox NJ, Cruz M, Hanis CL, Kooner J, Lee JY, Seielstad M, Teo YY, Boehnke M, Parra EJ, Chambers JC, Tai ES, McCarthy MI, Morris AP. Genome-wide trans-ancestry meta-analysis provides insight into the genetic architecture of type 2 diabetes susceptibility. *Nat Genet* 2014;46(3):234-244

Originalpublikation - IF 29,648

Volltext: <http://dx.doi.org/10.1038/ng.2897>

Biobank - 9160

Dong J, Gao J, Nalls M, Gao X, Huang X, Han J, Singleton AB, Chen H, International Parkinson's Disease Genomics Consortium (IPDGC). Susceptibility loci for pigmentation and melanoma in relation to Parkinson's disease. *Neurobiol Aging* 2014;35(6):1512.e5-1512.10

Originalpublikation - IF 4,853

Volltext: <http://dx.doi.org/10.1016/j.neurobiolaging.2013.12.020>

Klinik für Psychiatrie, Sozialpsychiatrie und Psychotherapie - 7110

Emrich HM. Trauma and fiction - trauma and concreteness in film: represented in the fotofilm fiasco (2010). In: Elm M, Kabalek K, Kühne JB [Hrsg.]: *The horrors of trauma in cinema : violence void visualization*. 1. Publ. Newcastle upon Tyne: Cambridge Scholars Publ., 2014. S. 177-191

Buchbeitrag, Monografie

Biobank - 9160

Fan YM, Hernesniemi J, Oksala N, Levula M, Raitoharju E, Collings A, Hutri-Kähönen N, Juonala M, Marniemi J, Lyytikäinen LP, Seppälä I, Mennander A, Tarkka M, Kangas AJ, Soininen P, Salenius JP, Klopp N, Illig T, Laitinen T, Ala-Korpela M, Laaksonen R, Viikari J, Kähönen M, Raitakari OT, Lehtimäki T. Upstream Transcription Factor 1 (USF1) allelic variants regulate lipoprotein metabolism in women and USF1 expression in atherosclerotic plaque. *Sci Rep* 2014;4:4650

Originalpublikation - IF 5,078

Volltext: <http://dx.doi.org/10.1038/srep04650>

Biobank - 9160

Ganesh SK, Chasman DI, Larson MG, Guo X, Verwoert G, Bis JC, Gu X, Smith AV, Yang ML, Zhang Y, Ehret G, Rose LM, Hwang SJ, Papanicolaou GJ, Sijbrands EJ, Rice K, Eiriksdottir G, Pihur V, Ridker PM, Vasan RS, Newton-Cheh C, Global Blood Pressure Genetics Consortium, Raffel LJ, Amin N, Rotter JI, Liu K, Launer LJ, Xu M, Caulfield M, Morrison AC, Johnson AD, Vaidya D, Dehghan A, Li G, Bouchard C, Harris TB, Zhang H, Boerwinkle E, Siscovick DS, Gao W, Uitterlinden AG, Rivadeneira F, Hofman A, Willer CJ, Franco OH, Huo Y, Witteman JC, Munroe PB, Gudnason V, Palmas W, van Duijn C, Fornage M, Levy D, Psaty BM, Chakravarti A. Effects of long-term averaging of quantitative blood pressure traits on the detection of genetic associations. *Am J Hum Genet* 2014;95(1):49-65

Originalpublikation - IF 10,987

Volltext: <http://dx.doi.org/10.1016/j.ajhg.2014.06.002>

Biobank - 9160

Hara K, Fujita H, Johnson TA, Yamauchi T, Yasuda K, Horikoshi M, Peng C, Hu C, Ma RC, Imamura M, Iwata M, Tsunoda T, Morizono T, Shojima N, So WY, Leung TF, Kwan P, Zhang R, Wang J, Yu W, Maegawa H, Hirose H, DIAGRAM consortium, Kaku K, Ito C, Watada H, Tanaka Y, Tobe K, Kashiwagi A, Kawamori R, Jia W, Chan JC, Teo YY, Shyong TE, Kamatani N, Kubo M, Maeda S, Kadokawa T. Genome-wide association study identifies three novel loci for type 2 diabetes. *Hum Mol Genet* 2014;23(1):239-246

Originalpublikation - IF 6,677

Volltext: <http://dx.doi.org/10.1093/hmg/ddt399>

Biobank - 9160

Herder C, Illig T. Neue Biomarker und Gene in der Prädiktion des Typ-2-Diabetes. *Diabetologe* 2014;10(7):566-571

Originalpublikation - IF 0,062

Volltext: <http://dx.doi.org/10.1007/s11428-014-1211-y>

Biobank - 9160

Hoggart CJ, Venturini G, Mangino M, Gomez F, Aszari G, Zhao JH, Teumer A, Winkler TW, Tsernikova N, Luan J, Mihailov E, Ehret GB, Zhang W, Lamparter D, Esko T, Mace A, Rueger S, Bochud PY, Barcella M, Dauvilliers Y, Benyamin B, Evans DM, Hayward C, Lopez MF, Franke L, Russo A, Heid IM, Salvi E, Vendantam S, Arking DE, Boerwinkle E, Chambers JC, Fiorito G, Grallert H, Guerrera S, Homuth G, Huffman JE, Porteous D, Generation Scotland Consortium, LifeLines Cohort study, GIANT Consortium, Moradpour D, Iranzo A, Hebebrand J, Kemp JP, Lammers GJ, Aubert V, Heim MH, Martin NG, Montgomery GW, Peraita-Adrados R, Santamaría J, Negro F, Schmidt CO, Scott RA, Spector TD, Strauch K, Volzke H, Wareham NJ, Yuan W, Bell JT, Chakravarti A, Kooner JS, Peters A, Matullo G, Wallaschofski H, Whitfield JB, Paccaud F, Vollenweider P, Bergmann S, Beckmann JS, Tafti M, Hastie ND, Cusi D, Bochud M, Frayling TM, Metspalu A, Jarvelin MR, Scherag A, Smith GD, Borecki IB, Rousson V, Hirschhorn JN, Rivolta C, Loos RJ, Kutalik Z. Novel approach identifies SNPs in SLC2A10 and KCNK9 with evidence for parent-of-origin effect on body mass index. *PLoS Genet* 2014;10(7):e1004508

Originalpublikation - IF 8,167

Volltext: <http://dx.doi.org/10.1371/journal.pgen.1004508>

Biobank - 9160

Holmes MV, Exeter HJ, Folkersen L, Nelson CP, Guardiola M, Cooper JA, Sofat R, Boekholdt SM, Khaw KT, Li KW, Smith AJ, Van't Hooft F, Eriksson P, Franco-Cereceda A, Asselbergs FW, Boer JM, Onland-Moret NC, Hofker M, Erdmann J, Kivimaki M, Kumari M, Reiner AP, Keating BJ, Humphries SE, Hingorani AD, Mallat Z, Samani NJ, Talmud PJ, CARDIoGRAM Consortium. Novel genetic approach to investigate the role of plasma secretory phospholipase A2 (sPLA2)-V isoenzyme in coronary heart disease: modified Mendelian randomization analysis using PLA2G5 expression levels. *Circ Cardiovasc Genet* 2014;7(2):144-150

Originalpublikation - IF 5,337

Volltext: <http://dx.doi.org/10.1161/CIRCGENETICS.113.000271>

Biobank - 9160

Hornhardt S, Rößler U, Sauter W, Rosenberger A, Illig T, Bickeböller H, Wichmann HE, Gomolka M. Genetic factors in individual radiation sensitivity. *DNA Repair (Amst)* 2014;16:54-65

Originalpublikation - IF 3,362

Volltext: <http://dx.doi.org/10.1016/j.dnarep.2014.02.001>

Biobank - 9160

Jourdan C, Linseisen J, Meisinger C, Petersen AK, Gieger C, Rawal R, Illig T, Heier M, Peters A, Wallaschofski H, Nauck M, Kastenmüller G, Suhre K, Prehn C, Adamski J, Koenig W, Roden M, Wichmann HE, Völzke H. Associations between thyroid hormones and serum metabolite profiles in an euthyroid population. *Metabolomics* 2014;10:152-164

Originalpublikation - IF 3,965

Volltext: <http://dx.doi.org/10.1007/s11306-013-0563-4>

Biobank - 9160

Knoll N, Jarick I, Volckmar AL, Klingenspor M, Illig T, Grallert H, Gieger C, Wichmann HE, Peters A, Wiegand S, Biebermann H, Fischer-Pozovszky P, Wabitsch M, Völzke H, Nauck M, Teumer A, Rosskopf D, Rimmbach C, Schreiber S, Jacobs G, Lieb W, Franke A, Hebebrand J, Hinney A. Mitochondrial DNA variants in obesity. *PLoS One* 2014;9(5):e94882

Originalpublikation - IF 3,534

Volltext: <http://dx.doi.org/10.1371/journal.pone.0094882>

Biobank - 9160

Kretschmer A, Möller G, Lee H, Laumen H, von Toerne C, Schramm K, Prokisch H, Eyerich S, Wahl S, Baurecht H, Franke A, Claussnitzer M, Eyerich K, Teumer A, Milani L, Klopp N, Hauck SM, Illig T, Peters A, Waldenberger M, Adamski J, Reischl E, Weidinger S. A common atopy-associated variant in the Th2 cytokine locus control region impacts transcriptional regulation and alters SMAD3 and SP1 binding. *Allergy* 2014;69(5):632-642

Originalpublikation - IF 5,995

Volltext: <http://dx.doi.org/10.1111/all.12394>

Hannover Unified Biobank ; Biobank - 9160 ; TwinCore - Experimentelle Infektionsforschung - 9651

Kühn A, Nieters A, Köttgen A, Goek ON, Michels K, Nöthlings U, Jacobs G, Meisinger C, Pessler F, Akmatov MF, Kuhnisch J, Moebus S, Glocker E, Naus S, Keimling M, Leitzmann M, Linseisen J, Sarioglu H, von Toerne C, Hauck SM, Wallaschofski H, Wichmann HE, Illig T. Machbarkeitsprüfung und Qualitätsentwicklung zur Gewinnung von Biomaterialien in den Pretest-Studien der Deutschen Nationalen Kohorte. *Bundesgesundheitsblatt Gesundheitsforschung Gesundheitsschutz* 2014;57(11):1255-1263

Originalpublikation - IF 1,006

Volltext: <http://dx.doi.org/10.1007/s00103-014-2048-7>

Biobank - 9160

Liu CT, Buchkovich ML, Winkler TW, Heid IM, African Ancestry Anthropometry Genetics Consortium, GIANT Consortium, Borecki IB, Fox CS, Mohlke KL, North KE, Adrienne Cupples L. Multi-ethnic fine-mapping of 14 central adiposity loci. *Hum Mol Genet* 2014;23(17):4738-4744

Originalpublikation - IF 6,677

Volltext: <http://dx.doi.org/10.1093/hmg/ddu183>

Biobank - 9160

Lüneburg N, Lieb W, Zeller T, Chen MH, Maas R, Carter AM, Xanthakis V, Glazer NL, Schwedhelm E, Seshadri S, Ikram MA, Longstreth WT,Jr, Fornage M, Konig IR, Loley C, Ojeda FM, Schillert A, Wang TJ, Sticht H, Kittel A, Konig J, Benjamin EJ, Sullivan LM, Bernges I, Anderssohn M, Ziegler A, Gieger C, Illig T, Meisinger C, Wichmann HE, Wild PS, Schunkert H, Psaty BM, Wiggins KL, Heckbert SR, Smith N, Lackner K, Lunetta KL, Blankenberg S, Erdmann J, Munzel T, Grant PJ, Vasan RS, Boger RH. Genome-Wide Association Study of L-Arginine and Dimethylarginines Reveals Novel Metabolic Pathway for Symmetric Dimethylarginine. *Circ Cardiovasc Genet* 2014;7(6):864-872

Originalpublikation - IF 5,337

Volltext: <http://dx.doi.org/10.1161/CIRGENETICS.113.000264>

Arbeitsbereich Klinische Psychologie in der Klinik für Psychiatrie, Sozialpsychiatrie und Psychotherapie - 7180

Magid M, Reichenberg JS, Poth PE, Robertson HT, LaViolette AK, Kruger TH, Wollmer MA. Treatment of major depressive disorder using botulinum toxin A: a 24-week randomized, double-blind, placebo-controlled study. *J Clin Psychiatry* 2014;75(8):837-844

Originalpublikation - IF 5,139

Volltext: <http://dx.doi.org/10.4088/JCP.13m08845>

Biobank - 9160

Majcher-Peszynska J, Loebermann M, Klammt S, Frimmel S, Mundkowski RG, Welte T, Reisinger EC, Drewelow B, CAPNETZ Study Group. Ampicillin/sulbactam in elderly patients with community-acquired pneumonia. *Infection* 2014;42(1):79-87

Originalpublikation - IF 2,864

Volltext: <http://dx.doi.org/10.1007/s15010-013-0518-8>

Biobank - 9160 ; Institut für Biometrie - 8410 ; Institut für Allgemeinmedizin - 5440

Mand P, Roth K, Biertz F, Kersting M, Kruschinski C, Schmiemann G, Hummers-Pradier E. Drug-disease interaction in elderly patients in family practice. *Int J Clin Pharmacol Ther* 2014;52(5):337-345

Originalpublikation - IF 1,044

Volltext: <http://dx.doi.org/10.5414/CP202003>

Biobank - 9160

Missios P, Zhou Y, Guachalla LM, von Figura G, Wegner A, Chakkarappan SR, Binz T, Gompf A, Hartleben G, Burkhalter MD, Wulff V, Gunes C, Sattler RW, Song Z, Illig T, Klaus S, Bohm BO, Wenz T, Hiller K, Rudolph KL. Glucose substitution prolongs maintenance of energy homeostasis and lifespan of telomere dysfunctional mice. *Nat Commun* 2014;5:4924

Originalpublikation - IF 10,742

Volltext: <http://dx.doi.org/10.1038/ncomms5924>

Biobank - 9160

Mook-Kanamori DO, Róisch-Margl W, Kastenmüller G, Prehn C, Petersen AK, Illig T, Gieger C, Wang-Sattler R, Meisinger C, Peters A, Adamski J, Suhre K. Increased amino acids levels and the risk of developing of hypertriglyceridemia in a 7-year follow-up. *J Endocrinol Invest* 2014;37(4):369-374

Originalpublikation - IF 1,552

Volltext: <http://dx.doi.org/10.1007/s40618-013-0044-7>

Biobank - 9160

Nalls MA, Saad M, Noyce AJ, Keller MF, Schrag A, Bestwick JP, Traynor BJ, Gibbs JR, Hernandez DG, Cookson MR, Morris HR, Williams N, Gasser T, Heutink P, Wood N, Hardy J, Martinez M, Singleton AB, International Parkinson's Disease Genomics Consortium (IPDGC), Wellcome Trust Case Control Consortium 2 (WTCCC2), North American Brain Expression

Consortium (NABEC), United Kingdom Brain Expression Consortium (UKBEC). Genetic comorbidities in Parkinson's disease. *Hum Mol Genet* 2014; 23(3):831-841

Originalpublikation - IF 6,677

Volltext: <http://dx.doi.org/10.1093/hmg/ddt465>

Biobank - 9160

Ng MC, Shriner D, Chen BH, Li J, Chen WM, Guo X, Liu J, Bielinski SJ, Yanek LR, Nalls MA, Comeau ME, Rasmussen-Torvik LJ, Jensen RA, Evans DS, Sun YV, An P, Patel SR, Lu Y, Long J, Armstrong LL, Wagenknecht L, Yang L, Snively BM, Palmer ND, Mudgal P, Langefeld CD, Keene KL, Freedman BI, Mychaleckyj JC, Nayak U, Raffel LJ, Goodarzi MO, Chen YD, Taylor HA,Jr, Correa A, Sims M, Couper D, Pankow JS, Boerwinkle E, Adeyemo A, Doumatey A, Chen G, Mathias RA, Vaidya D, Singleton AB, Zonderman AB, Igo RP,Jr, Sedor JR, FIND Consortium, Kabagambe EK, Siscovick DS, McKnight B, Rice K, Liu Y, Hsueh WC, Zhao W, Bielak LF, Kraja A, Province MA, Bottinger EP, Gottesman O, Cai Q, Zheng W, Blot WJ, Lowe WL, Pacheco JA, Crawford DC, eMERGE Consortium, DIAGRAM Consortium, Grundberg E, MuTHER Consortium, Rich SS, Hayes MG, Shu XO, Loos RJ, Borecki IB, Peyser PA, Cummings SR, Psaty BM, Fornage M, Iyengar SK, Evans MK, Becker DM, Kao WH, Wilson JG, Rotter JI, Sale MM, Liu S, Rotimi CN, Bowden DW, META-analysis of type 2 Diabetes in African Americans Consortium. Meta-analysis of genome-wide association studies in African Americans provides insights into the genetic architecture of type 2 diabetes. *PLoS Genet* 2014; 10(8):e1004517

Originalpublikation - IF 8,167

Volltext: <http://dx.doi.org/10.1371/journal.pgen.1004517>

Biobank - 9160

Obayashi M, Stevanin G, Synofzik M, Monin ML, Duyckaerts C, Sato N, Streichenberger N, Viguetto A, Desestret V, Tesson C, Wichmann HE, Illig T, Huttenlocher J, Kita Y, Izumi Y, Mizusawa H, Schols L, Klopstock T, Brice A, Ishikawa K, Durr A. Spinocerebellar ataxia type 36 exists in diverse populations and can be caused by a short hexanucleotide GGCCTG repeat expansion. *J Neurol Neurosurg Psychiatry* 2015; 86(9):986-995

Originalpublikation - IF 5,580

Volltext: <http://dx.doi.org/10.1136/jnnp-2014-309153>

Zentralapotheke - 8500

Picksak G, Kantelhardt P. Medikationsfehler - Liegezeitverlängerung durch Falschlagerung einer Infusion? *KHP* 2014; 35(5):159-160

Originalpublikation

Biobank - 9160

Polonikov AV, Ivanov VP, Bogomazov AD, Freidin MB, Illig T, Solodilova MA. Antioxidant Defense Enzyme Genes and Asthma Susceptibility: Gender-Specific Effects and Heterogeneity in Gene-Gene Interactions between Pathogenetic Variants of the Disease. *Biomed Res Int* 2014; 2014: 708903

Originalpublikation - IF 2,706

Volltext: <http://dx.doi.org/10.1155/2014/708903>

Biobank - 9160

Raedler D, Ballenberger N, Klucker E, Bock A, Otto R, Prazeres da Costa O, Holst O, Illig T, Buch T, von Mutius E, Schaub B. Identification of novel immune phenotypes for allergic and nonallergic childhood asthma. *J Allergy Clin Immunol* 2015; 135(1):81-91

Originalpublikation - IF 11,248

Volltext: <http://dx.doi.org/10.1016/j.jaci.2014.07.046>

Biobank - 9160

Rahmioglu N, Macgregor S, Drong AW, Hedman AK, Harris HR, Randall JC, Prokopenko I, The International Endogene Consortium (IEC), The GIANT Consortium, Nyholt DR, Morris AP, Montgomery GW, Missmer SA, Lindgren CM, Zondervan KT, The International Endogene Consortium IEC The GIANT Consortium. Genome-wide enrichment analysis between endometriosis and obesity-related traits reveals novel susceptibility loci. *Hum Mol Genet* 2015; 24(4):1185-1199

Originalpublikation - IF 6,677

Volltext: <http://dx.doi.org/10.1093/hmg/ddu516>

Biobank - 9160

Raitoharju E, Seppälä I, Oksala N, Lyytikäinen LP, Raitakari O, Viikari J, Ala-Korpela M, Soininen P, Kangas AJ, Waldenberger M, Klopp N, Illig T, Leiviskä J, Loo BM, Hutili-Kähönen N, Kähönen M, Laaksonen R, Lehtimäki T. Blood microRNA profile associates with the levels of serum lipids and metabolites associated with glucose metabolism and insulin resistance and pinpoints pathways underlying metabolic syndrome: The cardiovascular risk in Young Finns Study. *Mol Cell Endocrinol* 2014; 391(1-2):41-49

Originalpublikation - IF 4,241

Volltext: <http://dx.doi.org/10.1016/j.mce.2014.04.013>

Biobank - 9160

Reinehr T, Wolters B, Knop C, Lass N, Hellmuth C, Harder U, Peissner W, Wahl S, Grallert H, Adamski J, Illig T, Prehn C, Yu Z, Wang-Sattler R, Koletzko B. Changes in the serum metabolite profile in obese children with weight loss. *Eur J Nutr* 2015; 54(2):173-181

Originalpublikation - IF 3,840

Volltext: <http://dx.doi.org/10.1007/s00394-014-0698-8>

Biobank - 9160

Ried JS, Shin SY, Krumsiek J, Illig T, Theis FJ, Spector TD, Adamski J, Wichmann HE, Strauch K, Soranzo N, Suhre K, Gieger C. Novel Genetic Associations with Serum Level Metabolites Identified by Phenotype Set Enrichment Analyses. *Hum Mol Genet* 2014; 23(21):5847-5857

Originalpublikation - IF 6,677

Volltext: <http://dx.doi.org/10.1093/hmg/ddu301>

Biobank - 9160

Rodriguez E, Baurecht H, Wahn AF, Kretschmer A, Hotze M, Zeilinger S, Klopp N, Illig T, Schramm K, Prokisch H, Kühnel B, Gieger C, Harder J, Cifuentes L, Novak N, Weidinger S. An integrated epigenetic and transcriptomic analysis reveals distinct tissue-specific patterns of DNA methylation associated with atopic dermatitis. *J Invest Dermatol* 2014; 134(7):1873-1883

Originalpublikation - IF 6,372

Volltext: <http://dx.doi.org/10.1038/jid.2014.87>

Biobank - 9160

Schäfer N, Yu Z, Wagener A, Millrose MK, Reissmann M, Bortfeldt R, Dieterich C, Adamski J, Wang-Sattler R, Illig T, Brockmann GA. Changes in metabolite profiles caused by genetically determined obesity in mice. *Metabolomics* 2014;10: 461-472

Originalpublikation - IF 3,965

Volltext: <http://dx.doi.org/10.1007/s11306-013-0590-1>

Institut für Klinische Pharmakologie - 5350

Schneider JY, Pham VV, Fröhlich JC, Tsikas D. DDAH activity is not associated with oxidative stress in elderly patients with peripheral arterial occlusive disease. *Exp Gerontol* 2014;55: 159-160

Letter ; Originalpublikation - IF 3,529

Volltext: <http://dx.doi.org/10.1016/j.exger.2014.03.012>

Biobank - 9160

Schramm K, Marzi C, Schurmann C, Carstensen M, Reinmaa E, Biffar R, Eckstein G, Gieger C, Grabe HJ, Homuth G, Kastenmüller G, Mägi R, Metspalu A, Mihailov E, Peters A, Petersmann A, Roden M, Strauch K, Suhre K, Teumer A, Völker U, Völzke H, Wang-Sattler R, Waldenberger M, Meitinger T, Illig T, Herder C, Grallert H, Prokisch H. Mapping the genetic architecture of gene regulation in whole blood. *PLoS One* 2014;9(4):e93844

Originalpublikation - IF 3,534

Volltext: <http://dx.doi.org/10.1371/journal.pone.0093844>

Biobank - 9160

Schulte EC, Schramm K, Schurmann C, Lichtner P, Herder C, Roden M, Gieger C, Peters A, Trenkwalder C, Högl B, Frauscher B, Berger K, Fietze I, Gross N, Stiasny-Kolster K, Oertel W, Bachmann CG, Paulus W, Zimprich A, Völzke H, Schminke U, Nauck M, Illig T, Meitinger T, Müller-Myhsok B, Prokisch H, Winkelmann J. Blood cis-eQTL Analysis Fails to Identify Novel Association Signals among Sub-Threshold Candidates from Genome-Wide Association Studies in Restless Legs Syndrome. *PLoS One* 2014;9(5):e98092

Originalpublikation - IF 3,534

Volltext: <http://dx.doi.org/10.1371/journal.pone.0098092>

TwinCore: Institut für Molekulare Bakteriologie - 9641

Schulz S, Häussler S. Chromatin immunoprecipitation for ChIP-chip and ChIP-seq. *Methods Mol Biol* 2014;1149: 591-605

Buchbeitrag, Monografie

Volltext: http://dx.doi.org/10.1007/978-1-4939-0473-0_45

DZL Deutsches Zentrum für Lungenforschung ; Klinik für Päd. Pneumologie, Allergologie und Neonatologie - 6710 ; Biobank - 9160

Sharma V, Michel S, Gaertner V, Franke A, Vogelberg C, von Berg A, Bufe A, Heinzmann A, Laub O, Rietschel E, Simma B, Frischer T, Genuneit J, Zeilinger S, Illig T, Schedel M, Potaczek DP, Kabesch M. Fine-mapping of IgE-associated loci 1q23, 5q31, and 12q13 using 1000 Genomes Project data. *Allergy* 2014;69(8):1077-1084

Originalpublikation - IF 5,995

Volltext: <http://dx.doi.org/10.1111/all.12431>

Biobank - 9160

Shin SY, Petersen AK, Wahl S, Zhai G, Römischi-Margl W, Small KS, Döring A, Kato BS, Peters A, Grundberg E, Prehn C, Wang-Sattler R, Wichmann HE, Hrabe de Angelis M, Illig T, Adamski J, Deloukas P, Spector TD, Suhre K, Gieger C, Soranzo N. Interrogating causal pathways linking genetic variants, small molecule metabolites and circulating lipids. *Genome Med* 2014;6(3):25

Originalpublikation - IF 4,942

Volltext: <http://dx.doi.org/gm542>

Biobank - 9160

Stratmann B, Xu T, Meisinger C, Menart B, Roden M, Herder C, Grallert H, Peters A, Koenig W, Illig T, Wichmann HE, Wang-Sattler R, Rathmann W, Tschoepe D. PLA1A2 platelet polymorphism predicts mortality in prediabetic subjects of the population based KORA S4-Cohort. *Cardiovasc Diabetol* 2014;13: 90-2840-13-90

Originalpublikation - IF 3,706

Volltext: <http://dx.doi.org/10.1186/1475-2840-13-90>

Biobank - 9160

Tragante V, Barnes MR, Ganesh SK, Lanktree MB, Guo W, Franceschini N, Smith EN, Johnson T, Holmes MV, Padmanabhan S, Karczewski KJ, Almoguera B, Barnard J, Baumert J, Chang YP, Elbers CC, Farrall M, Fischer ME, Gaunt TR, Gho JM, Gieger C, Goel A, Gong Y, Isaacs A, Kleber ME, Mateo Leach I, McDonough CW, Meijis MF, Melander O, Nelson CP, Nolte IM, Pankratz N, Price TS, Shaffer J, Shah S, Tomaszewski M, van der Most PJ, Van Iperen EP, Vonk JM, Witkowska K, Wong CO, Zhang L, Beitelshes AL, Berenson GS, Bhatt DL, Brown M, Burt A, Cooper-DeHoff RM, Connell JM, Cruickshanks KJ, Curtis SP, Davey-Smith G, Delles C, Gansevoort RT, Guo X, Haiqing S, Hastie CE, Hofker MH, Hovingh GK, Kim DS, Kirkland SA, Klein BE, Klein R, Li YR, Maiwald S, Newton-Cheh C, O'Brien ET, Onland-Moret NC, Palmas W, Parsa A, Penninx BW, Pettinger M, Vasan RS, Ranchalis JE, M Ridker P, Rose LM, Sever P, Shimbo D, Steele L, Stolk RP, Thorand B, Trip MD, van Duijn CM, Verschuren WM, Wijmenga C, Wyatt S, Young JH, Zwinderian AH, Bezzina CR, Boerwinkle E, Casas JP, Caulfield MJ, Chakravarti A, Chasman DI, Davidson KW, Doevedans PA, Dominiczak AF, FitzGerald GA, Gums JG, Fornage M, Hakonarson H, Halder I, Hillege HL, Illig T, Jarvik GP, Johnson JA, Kastelein JJ, Koenig W, Kumari M, Marz W, Murray SS, O'Connell JR, Oldehinkel AJ, Pankow JS, Rader DJ, Redline S, Reilly MP, Schadt EE, Kottke-Marchant K, Snieder H, Snyder M, Stanton AV, Tobin MD, Uitterlinden AG, van der Harst P, van der Schouw YT, Samani NJ, Watkins H, Johnson AD, Reiner AP, Zhu X, de Bakker PI, Levy D, Asselbergs FW, Munroe PB, Keating BJ. Gene-centric meta-analysis in 87,736 individuals of European ancestry identifies multiple blood-pressure-related loci. *Am J Hum Genet* 2014;94(3):349-360

Originalpublikation - IF 10,987

Volltext: <http://dx.doi.org/10.1016/j.ajhg.2013.12.016>

Biobank - 9160

Walsh KM, Codd V, Smirnov IV, Rice T, Decker PA, Hansen HM, Kollmeyer T, Kosel ML, Molinaro AM, McCoy LS, Bracci PM, Cabriga BS, Pekmezci M, Zheng S, Wiemels JL, Pico AR, Tihan T, Berger MS, Chang SM, Prados MD, Lachance DH, O'Neill BP, Sicotte H, Eckel-Passow JE, ENGAGE Consortium Telomere Group, van der Harst P, Wiencke JK, Samani NJ, Jenkins RB, Wrensch MR. Variants near TERT and TERC influencing telomere length are associated with high-grade glioma risk. *Nat Genet* 2014;46(7):731-735

Originalpublikation - IF 29,648

Volltext: <http://dx.doi.org/10.1038/ng.3004>

Biobank - 9160

Winkler TW, Day FR, Croteau-Chonka DC, Wood AR, Locke AE, Magi R, Ferreira T, Fall T, Graff M, Justice AE, Luan J, Gustafsson S, Randall JC, Vedantam S, Workalemahu T, Kilpelainen TO, Scherag A, Esko T, Kutalik Z, Heid IM, Loos RJ, Genetic Investigation of Anthropometric Traits (GIANT) Consortium. Quality control and conduct of genome-wide association meta-analyses. *Nat Protoc* 2014;9(5):1192-1212

Originalpublikation - IF 7,782

Volltext: <http://dx.doi.org/10.1038/nprot.2014.071>

Biobank - 9160

Wood AR, Esko T, Yang J, Vedantam S, Pers TH, Gustafsson S, Chu AY, Estrada K, Luan J, Kutalik Z, Amin N, Buchkovich ML, Croteau-Chonka DC, Day FR, Duan Y, Fall T, Fehrmann R, Ferreira T, Jackson AU, Karjalainen J, Lo KS, Locke AE, Magi R, Mihailov E, Porcu E, Randall JC, Scherag A, Vinkhuyzen AA, Westra HJ, Winkler TW, Workalemahu T, Zhao JH, Absher D, Albrecht E, Anderson D, Baron J, Beekman M, Demirkiran A, Ehret GB, Feenstra B, Feitosa MF, Fischer K, Fraser RM, Goel A, Gong J, Justice AE, Kanoni S, Kleber ME, Kristiansson K, Lim U, Lotay V, Lui JC, Mangino M, Mateo Leach I, Medina-Gomez C, Nalls MA, Nyholt DR, Palmer CD, Pasko D, Peclivanis S, Prokopenko I, Ried JS, Ripke S, Shungin D, Stancakova A, Strawbridge RJ, Sung YJ, Tanaka T, Teumer A, Trompet S, van der Laan SW, van Setten J, Van Vliet-Ostaptchouk JV, Wang Z, Yengo L, Zhang W, Afzal U, Arnlov J, Arscott GM, Bandinelli S, Barrett A, Bellis C, Bennett AJ, Berne C, Bluher M, Bolton JL, Bottcher Y, Boyd HA, Bruinenberg M, Buckley BM, Buyske S, Caspersen IH, Chines PS, Clarke R, Claudi-Boehm S, Cooper M, Daw EW, De Jong PA, Deelen J, Delgado G, Denny JC, Dhomukshe-Rutten R, Dimitriou M, Doney AS, Dorr M, Eklund N, Eury E, FolkerSEN L, Garcia ME, Geller F, Giedraitis V, Go AS, Grallert H, Grammer TB, Grassler J, Gronberg H, de Groot LC, Groves CJ, Haessler J, Hall P, Haller T, Hallmans G, Hannemann A, Hartman CA, Hassinen M, Hayward C, Heard-Costa NL, Helmer Q, Hemani G, Henders AK, Hillege HL, Hlatky MA, Hoffmann W, Hoffmann P, Holmen O, Houwing-Duistermaat JJ, Illig T, Isaacs A, James AL, Jeff J, Johansen B, Johansson A, Jolley J, Juliusdottir T, Junntila J, Kho AN, Kinnunen L, Klopp N, Kocher T, Kratzer W, Lichtner P, Lind L, Lindstrom J, Lobbens S, Lorentzon M, Lu Y, Lysenko V, Magnusson PK, Mahajan A, Maillard M, McArdle WL, McKenzie CA, McLachlan S, McLaren PJ, Menni C, Merger S, Milani L, Moayyeri A, Monda KL, Morken MA, Muller G, Muller-Nurasyid M, Musk AW, Narisu N, Nauck M, Nolte IM, Nothen MM, Oozageer L, Pilz S, Rayner NW, Renstrom F, Robertson NR, Rose LM, Roussel R, Sanna S, Scharnagl H, Scholtens S, Schumacher FR, Schunkert H, Scott RA, Sehmi J, Seufferlein T, Shi J, Silventoinen K, Smit JH, Smith AV, Smolonska J, Stanton AV, Stirrups K, Stott DJ, Stringham HM, Sundstrom J, Swertz MA, Syvanen AC, Tayo BO, Thorleifsson G, Tyrer JP, van Dijk S, van Schoor NM, van der Velde N, van Heemst D, van Oort FV, Vermeulen SH, Verweij N, Vonk JM, Waite LL, Waldenberger M, Wennauer R, Wilkens LR, Willenborg C, Wilsgaard T, Wojczynski MK, Wong A, Wright AF, Zhang Q, Arveiler D, Bakker SJ, Beilby J, Bergman RN, Bergmann S, Biffar R, Blangero J, Boomsma DI, Bornstein SR, Bovet P, Brambillia P, Brown MJ, Campbell H, Caulfield MJ, Chakravarti A, Collins R, Collins FS, Crawford DC, Cupples LA, Danesh J, de Faire U, den Ruijter HM, Erbel R, Erdmann J, Eriksson JG, Farrall M, Ferrannini E, Ferrieres J, Ford I, Forouhi NG, Forrester T, Gansevoort RT, Gejman PV, Gieger C, Golay A, Gottesman O, Gudnason V, Gyllensten U, Haas DW, Hall AS, Harris TB, Hattersley AT, Heath AC, Hengstenberg C, Hicks AA, Hindorff LA, Hingorani AD, Hofman A, Hovingh GK, Humphries SE, Hunt SC, Hypponen E, Jacobs KB, Jarvelin MR, Jousilahti P, Jula AM, Kaprio J, Kastelein JJ, Kayser M, Kee F, Keinanen-Kiukaanniemi SM, Kiemeneij LA, Kooner JS, Kooperberg C, Koskinen S, Kovacs P, Kraja AT, Kumari M, Kuusisto J, Lakka TA, Langenberg C, Le Marchand L, Lehtimaki T, Lupoli S, Madden PA, Mannisto S, Manunta P, Marette A, Matise TC, McKnight B, Meitinger T, Moll FL, Montgomery GW, Morris AD, Morris AP, Murray JC, Nelis M, Ohlsson C, Oldehinkel AJ, Ong KK, Ouwehand WH, Pasterkamp G, Peters A, Pramstaller PP, Price JF, Qi L, Raitakari OT, Rankinen T, Rao DC, Rice TK, Ritchie M, Rudan I, Salomaa V, Samani NJ, Saramies J, Sarzynski MA, Schwarz PE, Sebert S, Sever P, Shuldiner AR, Sinisalo J, Steinthorsdottir V, Stolk RP, Tardif JC, Tonjes A, Tremblay A, Tremoli E, Virtamo J, Vohl MC, Electronic Medical Records and Genomics (eMERGE) Consortium, MIGen Consortium, PAGEGE Consortium, LifeLines Cohort Study, Amouyel P, Asselbergs FW, Assimes TL, Bochud M, Boehm BO, Boerwinkle E, Bottinger EP, Bouchard C, Cauchi S, Chambers JC, Chanock SJ, Cooper RS, de Bakker PI, Dedoussis G, Ferrucci L, Franks PW, Froguel P, Groop LC, Haiman CA, Hamsten A, Hayes MG, Hui J, Hunter DJ, Hveem K, Jukema JW, Kaplan RC, Kivimaki M, Kuh D, Laakso M, Liu Y, Martin NG, Marz W, Melbye M, Moebus S, Munroe PB, Njolstad I, Oostra BA, Palmer CN, Pedersen NL, Perola M, Perusse L, Peters U, Powell JE, Power C, Quertermous T, Rauramaa R, Reinmaa E, Ridker PM, Rivadeneira F, Rotter JI, Saaristo TE, Saleheen D, Schlessinger D, Slagboom PE, Snieder H, Spector TD, Strauch K, Stumvoll M, Tuomilehto J, Uusitupa M, van der Harst P, Volzke H, Walker M, Wareham NJ, Watkins H, Wichmann HE, Wilson JF, Zanen P, Deloukas P, Heid IM, Lindgren CM, Mohlke KL, Speliotes EK, Thorsteinsdottir U, Barroso I, Fox CS, North KE, Strachan DP, Beckmann JS, Berndt SL, Boehnke M, Borecki IB, McCarthy MI, Metspalu A, Stefansson K, Uitterlinden AG, van Duijn CM, Franke L, Willer CJ, Price AL, Lettre G, Loos RJ, Weedon MN, Ingelsson E, O'Connell JR, Abecasis GR, Chasman DI, Goddard ME, Visscher PM, Hirschhorn JN, Frayling TM. Defining the role of common variation in the genomic and biological architecture of adult human height. *Nat Genet* 2014;46(11):1173-1186

Originalpublikation - IF 29,648

Volltext: <http://dx.doi.org/10.1038/ng.3097>

Biobank - 9160

Yousri NA, Kastenmuller G, Gieger C, Shin SY, Erte I, Menni C, Peters A, Meisinger C, Mohney RP, Illig T, Adamski J, Soranzo N, Spector TD, Suhre K. Long term conservation of human metabolic phenotypes and link to heritability. *Metabolomics* 2014;10(5):1005-1017

Originalpublikation - IF 3,965

Volltext: <http://dx.doi.org/10.1007/s11306-014-0629-y>