Kantonsspital St.Gallen

Yoichiro Kamatani

Kontakt

Yoichiro Kamatani

Publikationen (6)

Knol M, Poot R, Evans T, Satizabal C, Mishra A, Sargurupremraj M, van der Auwera S, Duperron M, Jian X, Hostettler I, van Dam-Nolen D, Lamballais S, Pawlak M, Lewis C, Carrion-Castillo A, van Erp T, Reinbold C, Shin J, Scholz M, Håberg A, Kämpe A, Li G, Avinun R, Atkins J, Hsu F, Amod A, Lam M, Tsuchida A, Teunissen M, Aygün N, Patel Y, Liang D, Beiser A, Beyer F, Bis J, Bos D, Bryan R, Bülow R, Caspers S, Catheline G, Cecil C, Dalvie S, Dartigues J, DeCarli C, Enlund-Cerullo M, Ford J, Franke B, Freedman B, Friedrich N, Green M, Haworth S, Helmer C, Hoffmann P, Homuth G, Ikram M, Jack C, Jahanshad N, Jockwitz C, Kamatani Y, Knodt A, Li S, Lim K, Longstreth W, Macciardi F, Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium, Enhancing NeuroImaging Genetics through Meta-Analysis (ENIGMA) Consortium, Mäkitie O, Mazoyer B, Medland S, Miyamoto S, Moebus S, Mosley T, Muetzel R, Mühleisen T, Nagata M, Nakahara S, Palmer N, Pausova Z, Preda A, Quidé Y, Reay W, Roshchupkin G, Schmidt R, Schreiner P, Setoh K, Shapland C, Sidney S, St Pourcain B, Stein J, Tabara Y, Teumer A, Uhlmann A, van der Lugt A, Vernooij M, Werring D, Windham B, Witte A, Wittfeld K, Yang Q, Yoshida K, Brunner H, Le Grand Q, Sim K, Stein D, Bowden D, Cairns M, Hariri A, Cheung C, Andersson S, Villringer A, Paus T, Cichon S, Calhoun V, Crivello F, Launer L, White T, Koudstaal P, Houlden H, Fornage M, Matsuda F, Grabe H, Debette S, Thompson P, Seshadri S, Adams H. Genetic variants for head size share genes and pathways with cancer. Cell Rep Med 2024:101529.

Rahmioglu N, Mortlock S, Ghiasi M, Møller P, Stefansdottir L, Galarneau G, Turman C, Danning R, Law M, Sapkota Y, Christofidou P, Skarp S, Giri A, Banasik K, Krassowski M, Lepamets M, Marciniak B, Nõukas M, Perro D, Sliz E, Sobalska-Kwapis M, Thorleifsson G, Topbas-Selcuki N, Vitonis A, Westergaard D, Arnadottir R, Burgdorf K, Campbell A, Cheuk C, Clementi C, Cook J, De Vivo I, DiVasta A, Dorien O, Donoghue J, Edwards T, Fontanillas P, Fung J, Geirsson R, Girling J, Harkki P, Harris H, Healey M, Heikinheimo O, Holdsworth-Carson S, Hostettler I, Houlden H, Houshdaran S, Irwin J, Jarvelin M, Kamatani Y, Kennedy S, Kepka E, Kettunen J, Kubo M, Kulig B, Kurra V, Laivuori H, Laufer M, Lindgren C, MacGregor S, Mangino M, Martin N, Matalliotaki C, Matalliotakis M, Murray A, Ndungu A, Nezhat C, Olsen C, Opoku-Anane J, Padmanabhan S, Paranjpe M, Peters M, Polak G, Porteous D, Rabban J, Rexrode K, Romanowicz H, Saare M, Saavalainen L, Schork A, Sen S, Shafrir A, Siewierska-Górska A, Słomka M, Smith B, Smolarz B, Szaflik T, Szyłło K, Takahashi A, Terry K, Tomassetti C, Treloar S, Vanhie A, Vincent K, Vo K, Werring D, Zeggini E, Zervou M, DBDS Genomic Consortium, FinnGen Study, FinnGen Endometriosis Taskforce, Celmatix Research Team, 23andMe Research Team, Adachi S, Buring J, Ridker P, D'Hooghe T, Goulielmos G, Hapangama D, Hayward C, Horne A, Low S, Martikainen H, Chasman D, Rogers P, Saunders P, Sirota M, Spector T, Strapagiel D, Tung J, Whiteman D, Giudice L, Velez Edwards D, Uimari O, Kraft P, Salumets A, Nyholt D, Mägi R, Stefansson K, Becker C, Yurttas-Beim P, Steinthorsdottir V, Nyegaard M, Missmer S, Montgomery G, Morris A, Zondervan K. The genetic basis of endometriosis and comorbidity with other pain and inflammatory conditions. Nat Genet 2023; 55:423-436.

Bakker M, Kanning J, Abraham G, Martinsen A, Winsvold B, Zwart J, Bourcier R, Sawada T, Koido M, Kamatani Y, Morel S, Amouyel P, Debette S, Bijlenga P, Berrandou T, Ganesh S, Bouatia-Naji N, Jones G, Bown M, Rinkel G, Veldink J, Ruigrok Y, Hostettler I. Genetic Risk Score for Intracranial Aneurysms: Prediction of Subarachnoid Hemorrhage and Role in Clinical Heterogeneity. Stroke 2023

Ruigrok Y, Sargurupremraj M, Dichgans M, Malik R, Klijn C, Zaroff J, Breen G, Coleman J, Kim H, Ko N, Bown M, Jones G, Martin O, Dauvillier J, Schilling S, Hirsch S, Tatlisumak T, Amouyel P, Debette S, Veldink J, Kamatani Y, Bijlenga P, Redon R, Woo D, Werring D, Broderick J, Lindgren A, von Und Zu Fraunberg M, Jaaskelainen J, Niemelä M, Gaal-Paavola E, Slowik A, Pera J, Worrall B, Rinkel G, Friedrich C, Verschuren W, Zwart J, Chen Z, Millwood I, Liang L, Lin K, Walters R, Matsuda K, Terao C, Akiyama M, Koido M, van Eijk K, Alg V, Hostettler I, Bourcier R, Morel S, van Rheenen W, van der Spek R, Rouleau G, Zhou S, Rannikmäe K, Hveem K, Willer C, Sandvei M, Brumpton B, Johnsen M, Børte S, Winsvold B, Desal H, Eugène F, Shotar E, Gentric J, Naggara O, Dina C, van den Berg L, Houlden H, Sudlow C, Bakker M. Author Correction: Genome-wide association study of intracranial aneurysms identifies 17 risk loci and genetic overlap with clinical risk factors. Nat Genet 2021; 53:254.

Ruigrok Y, Sargurupremraj M, Dichgans M, Malik R, Klijn C, Zaroff J, Breen G, Coleman J, Kim H, Ko N, Bown M, Jones G, Martin O, Dauvillier J, Schilling S, Hirsch S, Tatlisumak T, Amouyel P, Debette S, Veldink J, Kamatani Y, Bijlenga P, Redon R, Woo D, Werring D, Broderick J, Lindgren A, von Und Zu Fraunberg M, Jaaskelainen J, Niemelä M, Gaal-Paavola E, Slowik A, Pera J, Worrall B, Rinkel G, Friedrich C, Verschuren W, Zwart J, Chen Z, Millwood I, Liang L, Lin K, Walters R, Matsuda K, Terao C, Akiyama M, Koido M, van Eijk K, Alg V, Hostettler I, Bourcier R, Morel S, van Rheenen W, van der Spek R, Rouleau G, Zhou S, Rannikmäe K, Hveem K, Willer C, Sandvei M, Brumpton B, Johnsen M, Børte S, Winsvold B, Desal H, Eugène F, Shotar E, Gentric J, Naggara O, Dina C, van den Berg L, Houlden H, Sudlow C, Bakker M. Genome-wide association study of intracranial aneurysms identifies 17 risk loci and genetic overlap with clinical risk factors. Nat Genet 2020; 52:1303-1313.

Bowden D, Snieder H, Smith J, Sitlani C, Sever P, Seshadri S, Scott W, Schreiner P, Schmidt C, Sandow K, Salako B, Sabanayagam C, Rudan I, Rose L, Robinson J, Robino A, Ridker P, Starr J, Strauch K, Tang H, Boehnke M, Becker D, Zonderman A, Yuan J, Yao J, Wojczynski M, Wilson G, Williams C, Wei W, Wang Y, Wang L, Waldenberger M, Uitterlinden A, Tham Y, Teo Y, Taylor K, Rice T, Renström F, Raitakari O, Metspalu A, Meitinger T, Mahajan A, Mägi R, Louie T, Long J, Lohman K, Loh M, Liu Y, Liu K, Liu J, Liu J, Liu C, Lin S, Lim S, Lifelines Cohort Study, Milani L, Momozawa Y, Morris A, Polasek O, Peyser P, Peters A, Pedersen N, Pankow J, Palmer N, Palmas W, Padmanabhan S, Ogunniyi A, North K, Norris J, Nasri U, Nalls M, Murray A, Munson P, Mosley T, Li Y, Chasman D, Hayward C, Fox E, Kelly T, Mook-Kanamori D, Arnett D, Sims M, van Dam R, Psaty B, O'Connell J, Levy D, Kritchevsky S, Kardia S, Gudnason V, Evans M, Cooper R, Bouchard C, Fornage M, Rotimi C, Province M, Rao D, Cupples L, Morrison A, Munroe P, Rice K, Elliott P, Caulfield M, Gauderman W, Bierut L, Zhu X, Rotter J, Reiner A, Loos R, Wong T, Tai E, van Duijn C, Laurie C, Kamatani Y, Zheng W, Kooner J, Kato N, Jonas J, Hung Y, Horta B, Gieger C, Gasparini P, Froguel P, Freedman B, Franks P, Forrester T, Farrall M, Esko T, Deary I, de Faire U, Chen Y, Laakso M, Lehtimäki T, Liang K, Wu T, Wickremasinghe A, Weir D, Watkins H, Wareham N, Wagenknecht L, van der Harst P, Shu X, Scott J, Samani N, Rettig R, Redline S, Pereira A, Oldehinkel A, Newman A, Magnusson P, Chambers J, Lewis C, Yanek L, Leander K, Kühnel B, Kasturiratne A, Kähönen M, Jackson A, Hsu F, Horimoto A, Hartwig F, Harris S, Goel A, Giulianini F, Gao C, Gandin I, Divers J, Chen X, Chai J, Lee W, Lin K, 'an Luan J, Wen W, Weiss S, Ware E, Wang Y, Wang H, Varga T, van der Most P, Takeuchi F, Stančáková A, Sheu W, Scott R, Schupf N, Rauramaa R, Nelson C, He M, McKenzie C, Boissel M, Amini M, Alver M, Li C, Musani S, Marten J, Vojinovic D, Sim X, Cheng C, Lu Y, Franceschini N, Guo X, Ntalla I, Schwander K, Kraja A, Brown M, Bentley A, de Las Fuentes L, Winkler T, Feitosa M, Kilpeläinen T, Richard M, Sofer T, Matoba N, Zhou Y, Zhao W, Warren H, Tayo B, Tajuddin S, Smith A, Rankinen T, Manning A, Liu Y, Dorajoo R, Bartz T, Aschard H, Aslibekyan S, Noordam R, Sung Y, Lehne B, Howard B, Hofman A, Hirata M, Heng C, Heikkinen S, He J, Harris T, Hagenaars S, Gupta P, Gu D, Gu C, Graff M, Gigante B, GIANT Consortium, Gao H, Friedlander Y, Hunt S, Irvin M, Jia Y, Launer L, Langenberg C, Langefeld C, Kuusisto J, Kubo M, Krieger J, Kooperberg C, Komulainen P, Koistinen H, Koh W, Khor C, Kerrison N, Kaufman J, Katsuya T, Justice A, Joehanes R, Franco O, Forouhi N, Fisher V, Caizheng Y, Cade B, Cabrera C, Broeckel U, Brody J, Braund P, Bottinger E, Boerwinkle E, Bielak L, Barr R, Aung T, Arking D, Amin N, Alfred T, Afaq S, Zhao J, Campbell A, Canouil M, Chakravarti A, Faul J, Evangelou E, Ehret G, Eaton C, Duan Q, Dörr M, Debette S, de Silva H, de Mutsert R, Connell J, Collins F, COGENT-Kidney Consortium, Cocca M, Christensen K, Chauhan G, CHARGE Neurology Working Group, Zhang W. A Large-Scale Multi-ancestry Genome-wide Study Accounting for Smoking Behavior Identifies Multiple Significant Loci for Blood Pressure. Am J Hum Genet 2018; 102:375-400.

Projekte (0)

Keine Resultate gefunden.

Kantonsspital St.Gallen Rorschacher Strasse 95 CH-9007 St.Gallen T: +41 71 494 11 11 support.forschung@kssg.ch