



**Sudha Seshadri**

**Kontakt**

Sudha Seshadri

## Publikationen (3)

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, Debetto S, Thompson P, Seshadri S, Adams H. Genetic variants for head size share genes and pathways with cancer. *Cell Rep Med* 2024;101529.

MacKinnon A, Gottesman R, Schmidt R, Houlden H, Stott D, Koh J, Beekman M, Amin N, Vernooij M, Tozer D, Beiser A, Morris Z, Yang Q, Himali J, DeCarli C, Gudnason V, Markus H, Launer L, Seshadri S, Ikram M, Wardlaw J, Schmidt H, Werring D, Mosley T, Jukema J, Rost N, Wong T, Slagboom P, van Duijn C, Deary I, Kantarci K, Pirpamer L, van den Akker E, Hilal S, Giese A, Trompet S, Luciano M, Hostettler I, Liu J, Hofer E, Fornage M, Smith A, Romero J, Adams H, Traylor M, Lu D, Vojinovic D, Li S, Cheng C, Windham B, Bastin M, van der Lugt A, Saba Y, Chen C, van der Grond J, Liewald D, Satizabal C, Yilmaz P, Wilson D, Jack C, van der Lee S, Sigurdsson S, Knol M. Association of common genetic variants with brain microbleeds: A genome-wide association study. *Neurology* 2020; 95:e3331-e3343.

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, Joeanes 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, Debetto 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](mailto:support.forschung@kssg.ch)