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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, Houlihan 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.

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, Houlihan 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, Houlihan 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.

Projekte (0)

Keine Resultate gefunden.

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