



## Mohammad Arfan Ikram

### Kontakt

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## Publikationen (2)

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.

Mukamal K, Liu Y, Danesh J, Rasheed A, Mason M, Zonderman A, Franke L, Kristal B, Karjalainen J, Reed D, Westra H, Evans M, Saleheen D, Harris T, Dedoussis G, Curhan G, Stumvoll M, Beilby J, Houston D, Lorentzon M, Psaty B, Döring A, Heath A, Montgomery G, Dahmen N, Carithers T, Tucker K, Ferrucci L, Boyd H, Melbye M, Treur J, Mellström D, Hottenga J, Prokopenko I, Tönjes A, Deloukas P, Kanoni S, Pasquale L, Feenstra B, Bandinelli S, Caporaso N, Grabe H, Neuhauser M, Wolffenbuttel B, Hu F, Hyppönen E, Jarvelin M, Cupples L, Franks P, Ridker P, van Duijn C, Heiss G, Metspalu A, North K, Ingelsson E, Nettleton J, van Dam R, Vollenweider P, Wareham N, Ordovas J, Chan A, Peters U, Ohlsson C, Gieger C, Martin N, Waldenberger M, Siscovick D, Raitakari O, Eriksson J, Mitchell P, Hunter D, Kraft P, Rimm E, Boomsma D, Borecki I, Loos R, Chasman D, Mozaffarian D, Wojczynski M, Vink J, Zhao J, Burlutsky G, Lahti J, Mikkilä V, Lemaitre R, Eriksson J, Musani S, Tanaka T, Geller F, Luan J, Hui J, Mägi R, Dimitriou M, Garcia M, Ho W, Manichaikul A, Rawal R, Cornelis M, Byrne E, Esko T, Nalls M, Ganna A, Paynter N, Monda K, Amin N, Fischer K, Renström F, Ngwa J, Huikari V, Cavadino A, Nolte I, Teumer A, Yu K, Marques-Vidal P, Wright M, Rose L, Magnusson P, Bastardot F, Kutalik Z, Jacobs D, Forouhi N, Mihailov E, Lind L, Lindgren C, Michaëlsson K, Morris A, Jensen M, Khaw K, Luben R, Wang J, Männistö S, Perälä M, Kähönen M, Lehtimäki T, Gu F, Biffar R, Pedersen N, Couper D, Oostra B, Hofman A, Ikram M, Tiemeier H, Uitterlinden A, van Rooij F, Barroso I, Johansson I, Xue L, Kaakinen M, Milani L, Power C, Snieder H, Stolck R, Baumeister S, Viikari J. Genome-wide meta-analysis identifies six novel loci associated with habitual coffee consumption. *Mol Psychiatry* 2014; 20:647–656.

## Projekte (0)

Keine Resultate gefunden.

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