



Albert V Smith

Contact

Albert V Smith

Publications (5)

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.

Packard C, Pers T, Person T, Peters A, Petersen E, Peyser P, Pirie A, Polasek O, Polderman T, Puolijoki H, Raitakari O, Perry J, Perry J, Perola M, Padmanabhan S, Palmer C, Palmer N, Pasterkamp G, Patel A, Pattie A, Pedersen O, Peissig P, Peloso G, Pennell C, Rasheed A, Rauramaa R, Reilly D, Samani N, Sapkota Y, Sattar N, Schoen R, Schreiner P, Schulze M, Scott R, Segura-Lepe M, Shah S, Sheu W, Salomaa V, Saleheen D, Ruth K, Renström F, Rheinberger M, Ridker P, Rioux J, Rivas M, Roberts D, Robertson N, Robino A, Rolandsson O, Rudan I, Sim X, Lin K, Lubitz S, Lyytikäinen L, Männistö S, Marenne G, Mazul A, McCarthy M, McKean-Cowdin R, Medland S, Meidtner K, Milani L, Luan J, Loukola A, Lotery A, Lin L, Lin X, Lind L, Lindström J, Linneberg A, Liu C, Liu D, Liu Y, Lo K, Lophatananon A, Mistry V, Mitchell P, Mohlke K, Neville M, Nielsen S, Nikus K, Njølstad P, Nordestgaard B, Nyholt D, O'Connell J, O'Donoghue M, Olde Loohuis L, Ophoff R, Nelson C, Narisu N, Nalls M, Moilanen L, Moitry M, Montgomery G, Mook-Kanamori D, Moore C, Mori T, Morris A, Müller-Nurasyid M, Munroe P, Owen K, Slater A, Walker M, Witte D, Wood A, Wu Y, Yaghoobkar H, Yao J, Yao P, Yerges-Armstrong L, Young R, Zeggini E, Zhan X, Wilson J, Willer C, White H, Wallentin L, Wang F, Wang C, Wang S, Wang Y, Ware E, Wareham N, Warren H, Waterworth D, Wessel J, Zhang W, Zhao J, Zhao W, CHD Exome+ Consortium, EPIC-CVD Consortium, ExomeBP Consortium, Global Lipids Genetic Consortium, GoT2D Genes Consortium, EPIC InterAct Consortium, INTERVAL Study, ReproGen Consortium, T2D-Genes Consortium, MAGIC Investigators, Loos R, Hirschhorn J, Lindgren C, Zhou W, Zondervan K, Rotter J, Pospisilik J, Rivadeneira F, Borecki I, Deloukas P, Frayling T, Lettre G, North K, Understanding Society Scientific Group, Small K, Swift A, Tada H, Tansey K, Tardif J, Taylor K, Teumer A, Thompson D, Thorleifsson G, Thorsteinsdottir U, Thuesen B, Surendran P, Sun L, Stumvoll M, Smith A, Southam L, Spector T, Speliotes E, Starr J, Stefansson K, Steinthorsdottir V, Stirrups K, Strauch K, Stringham H, Tönjes A, Tromp G, Trompet S, Varga T, Varma R, Velez Edwards D, Vermeulen S, Veronesi G, Vestergaard H, Vitart V, Vogt T, Völker U, Vuckovic D, Varbo A, Vanhala M, van Setten J, Tsfantakis E, Tuomilehto J, Tybjaerg-Hansen A, Tyrer J, Uher R, Uitterlinden A, Uusitupa M, Laan S, Duijn C, Leeuwen N, Wagenknecht L, Lin H, Bots M, Caulfield M, Chambers J, Chasman D, Chen Y, Chowdhury R, Christensen C, Chu A, Cocca M, Collins F, Cook J, Catamo E, Carey D, Cappellani S, Bottinger E, Bowden D, Brandslund I, Breen G, Brilliant M, Broer L, Brumat M, Burt A, Butterworth A, Campbell P, Corley J, Corominas Galbany J, Cox A, Ruijter H, Dennis J, Denny J, Di Angelantonio E, Drenos F, Du M, Dubé L, Dunning A, Easton D, Edwards T, Hollander A, Heijer M, Demerath E, Crosslin D, Cuellar-Partida G, D'Eustacchio A, Danesh J, Davies G, Bakker P, Groot M, Mutsert R, Deary I, Dedoussis G, Ellinghaus D, Turcot V, Locke A, Mahajan A, Marouli E, Sivapalaratnam S, Young K, Alfred T, Feitosa M, Masca N, Manning A, Medina-Gomez C, Lempradl A, Karaderi T, Hendricks A, Lu Y, Highland H, Schurmann C, Justice A, Fine R, Bradfield J, Esko T, Giri A, Graff M, Guo X, Mudgal P, Ng M, Reiner A, Barroso I, Bastarache L, Benn M, Bergmann S, Bielak L, Blüher M, Boehnke M, Boeing H, Boerwinkle E, Böger C, Bang L, Balkau B, Auer P, Vedantam S, Willems S, Winkler T, Abecasis G, Aben K, Alam D, Alharthi S, Allison M, Amouyel P, Asselbergs F, Bork-Jensen J, Ellnor P, Howson J, Jukema J, Kahali B, Kahn R, Kähönen M, Kamstrup P, Kanoni S, Kaprio J, Karaleftheri M, Kardina S, Karpe F, Jørgensen T, Jørgensen M, Johansson S, Hu Y, Huang P, Huffman J, Ikram M, Ingelsson E, Jackson A, Jansson J, Jarvik G, Jensen G, Jia Y, Kathiresan S, Kee F, Kiemeny L, Lamparter D, Lange E, Lange L, Langenberg C, Larson E, Lee N, Lehtimäki T, Lewis C, Li H, Li J, Lakka T, Laakso M, Kuusisto J, Kim E, Kitajima H, Komulainen P, Kooner J, Kooperberg C, Korhonen T, Kovacs P, Kuivaniemi H, Kutalik Z, Kuulasmaa K, Li-Gao R, Elliott P, Franks P, Friedrich N, Frikke-Schmidt R, Galesloot T, Gan W, Gandin I, Gasparini P, Gibson J, Giedraitis V, Gjesing A, Franke A, Franco O, Fornage M, Evangelou E, Farmaki A, Farooqi I, Faul J, Fauser S, Feng S, Ferrannini E, Ferrieres J, Florez J, Ford I, Gordon-Larsen P, Gorski M, Grabe H, Have C, Hayward C, He L, Heard-Costa N, Heath A, Heid I, Helgeland Ø, Hernessniemi J, Hewitt A, Holmen O, Hattersley A, Harris T, Harris K, Grant S, Grarup N, Griffiths H, Grove M, Gudnason V, Gustafsson S, Haessler J, Hakonarson H, Hammerschlag A, Hansen T, Hovingh G. Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. *Nat Genet* 2017; 50:26-41.

Sim X, Ridker P, Rich S, Renström F, Reiner A, Reilly D, Rauramaa R, Rasheed A, Rader D, Psaty B, Poulter N, Polasek O, Pistis G, Rioux J, Robertson N, Shaffer C, Sevilla R, Sever P, Scott R, Schmidt E, Sattar N, Sanna S, Samani N, Salomaa V, Rudan I, Rotter J, Roden D, Pisinger C, Peters A, Perola M, Neville M, Munroe P, Müller-Nurasyid M, Mulas A, Morrison A, Morris A, Metspalu A, Melander O, Meigs J, Maschio A, Masca N, Marouli E, Nielsen J, Nielsen S, Pedersen O, Patel A, Pasko D, Palmer C, Padmanabhan S, Muntendam P, Molony C, Orho-Melander M, O'Donnell C, Mehran R, Ordovas J, Nordestgaard B, Mäntyselkä P, Kathiresan S, Zheng N, Zhang H, Zeggini E, Young R, Yaghoobkar H, Xu M, Wilson P, Wilson J, Wessel J, Weinstock J, Weeke P, Warren H, Zhang W, Zhang Y, Willer C, Musunuru K, Deloukas P, Abecasis G, Cowan C, McCarthy M, Danesh J, Howson J, Zoledziewska M, Zhou Y, Zhou W, Wareham N, Wang N, Waldenberger M, Stringham H, Strauch K, Stitzel N, Stirrups K, Starr J, Speliotes E, Spector T, Southam L, Somayajula S, Smith B, Smith A, Small K, Surendran P, Tada H, Virtamo J, Varga T, Varbo A, van Zuydam N, Tybjaerg-Hansen A, Tuomilehto J, Tsao P, Trompet S, Taylor K, Tardif J, Tang H, Tall A, Sivapalaratnam S, Frikke-Schmidt R, Dedoussis G, Deary I, Davies G, Damrauer S, Cupples L, Cucca F, Connell J, Chu A, Christensen C, Chowdhury R, Chen Y, Chen Y, Denny J, Dominiczak A, Frayling T, Franks P, Fornage M, Ford I, Ferrieres J, Ferrario M, Feitosa M, Farmaki A, Esko T, Eiriksdottir G, Ebeling T, Dubé M, Chasman D, Chambers J, Caulfield M, Arveiler D, Di Angelantonio E, Amouyel P, Alves A, Alam D, Emdin C, Saleheen D, Mahajan A, Wang X, Butterworth A, Yu H, Peloso G, Assimes T, Auer P, Busonero F, Brown M, Brandslund I, Bottinger E, Bork-Jensen J, Boerwinkle E, Boehnke M, Bis J, Benn M, Bang L, Ballantyne C, Baber U, Liu D, Manning A, Laakso M, Kuusisto J, Kuulasmaa K, Kooperberg C, Kooner J, Koistinen H, Klarin D, Khera A, Kee F, Karpe F, Kanoni S, Kamstrup P, Lakka T, Langenberg C, Manichaikul A, Malarstig A, Mägi R, Lu X, Lu Y, Loos R, Linneberg A, Lin L, Liewald D, Lauritzen T, Launer L, Langsted A, Justesen J, Jukema J, Jørgensen M, Grove M, Groop L, Grarup N, Grallert H, Goodarzi M, Giulianini F, Gieger C, Garcia M, Gao W, Ganesh S, Fuster V, Frossard P, Gudnason V, Hansen T, Jensen G, Jarvelin M, Jakobsdottir J, Jackson A, Jabeen S, Hveem K, Huo Y, Huffman J, Holmen O, Hirschhorn J, Hayward C, Harris T, Fritsche L. Exome-wide association study of plasma lipids in >300,000 individuals. *Nat Genet* 2017; 49:1758-1766.

Tremoli E, Sebert S, Schunkert H, Saramies J, Sanna S, Samani N, Rossouw J, Rose L, Roberts R, Rice K, Rettig R, Renström F, Rayner N, Rasheed A, Rao D, Quertermous T, Psaty B, Pramstaller P, Sheu W, Shin Y, Sim X, Tobin M, Thorsteinsdottir U, Teumer A, Tarasov K, Tanaka T, Tai E, Syvänen A, Swift A, Sundstrom J, Stringham H, Stirrups K, Stanton A, Stančáková A, Spector T, Sosa M, Smith A, Smit J, Poulter N, Peters A, Meneton P, McKenzie C, McCarthy M, Marouli E, Markus H, Mach F, Müller-Nurasyid M, Müller G, Männistö S, Lobbens S, Lindström J, Lin L, Lin H, Liang K, Li X, Levy D, Lee W, Menni C, Metspalu A, Mijatovic V, Perola M, Penninx B, Pedersen N, Parsa A, Palmer C, Paccard F, Ong K, O'Reilly P, O'Donnell C, Nikus K, Narisu N, Nagaraja R, Mulas A, Morrison A, Morris A, Montasser M, Moilanen L, Lee I, Uitterlinden A, Munroe P, Palmas W, Hveem K, Lehtimäki T, Ingelsson E, Tuomilehto J, Rauramaa R, Saltevo J, Laakso M, Melander O, Caulfield M, Snieder H, Langenberg C, Schwarz P, Njølstad I, Assimes T, Lind L, Wong A, März W, Kumari M, Salomaa V, Newton-Cheh C, Morris A, Deloukas P, Chakravarti A, Loos R, Ridker P, Stefansson K, Palmer C, Wichmann H, Hamsten A, Franks P, Kuulasmaa K, Lakatta E, Jarvelin M, Froguel P, Rotter J, Chen Y, Sever P, Dedoussis G, Wilks R, Westra H, Weder A, Watkins H, Wareham N, Wain L, Wagner A, Vollenweider P, Voight B, Vitart V, Virtamo J, Verwoert G, Vasan R, van Iperen E, van Duijn C, Vaidya D, Vaez A, Willsgaard T, Wilson J, Wong T, Taylor K, Hovingh G, Franke L, Willer C, Gierman H, Elliott P, Lee J, Saleheen D, Mohlke K, Cooper R, Bovet P, Zhu X, Zhao J, Zhang W, Yengo L, Yao J, Yang T, Uusitupa M, de Faire U, Bevan S, Barroso I, Barnes M, Bandinelli S, Balkau B, Baldassarre D, Axelsson T, Arking D, Amin N, Adair L, Absher D, Bochud M, Tayo B, Wu Y, Zhao W, Kim Y, Bis J, Björnsdottir G, Boehnke M, Danesh J, Dallongeville J, Döring A, Connell J, Collins F, Chung R, Chines P, Cheng C, Chang I, Chambers J, Cabrera C, Burnier M, Brown M, Bornstein S, Boomsma D, Bonnycastle L, Boerwinkle E, Warren H, Johnson A, Bouatia-Naji N, Kaakinen M, Meirelles O, Hughes M, Shungin D, Strawbridge R, Pihur V, Petersen A, Kanoni S, Donnelly L, Luan J, Thorleifsson G, Johnson T, Schmidt E, Jackson A, Chasman D, Ferreira T, Kristiansson K, Shah S, Kleber M, Jansen R, Lataniotis L, Kim S, Joehanes R, Pers T, Witkowska K, Folkersen L, Smith A, Theusch E, Rallidis L, Salfati E, Magnusson P, Nolte I, Eriksson N, Fava C, Lyytikäinen L, Guo X, Ehret G, Delgado G, Lee N, Iribarren C, Illig T, Ikram M, Hunt S, Hsiung C, Howard P, Hottenga J, Holmen O, Holmen J, Hofman A, Hirschhorn J, Hingorani A, Hicks A, Herzig K, Hercberg S, Hayward C, Havulinna A, Jensen R, Kähönen M, Kang H, Lakka T, Kvaløy K, Kuusisto J, Kutalik Z, Kuh D, Krauss R, Kosova G, Kooner J, Komulainen P, Kolovou G, Klopp N, Kivimäki M, Kim E, Kim Y, Khaw K, Keating B, Kathiresan S, Hassinen M, Hartikainen A, Fornage M, Ferrucci L, Ferrieres J, Felix J, Farrall M, Fall T, Evans A, Evangelou E, Esko T, Eriksson P, Erdmann J, Enroth S, Elosua R, Eicher J, Edkins S, Drenos F, Doney A, Forrester T, Franceschini N, Duran O, Hallmans G, Gyllenstein U, Gudnason V, Groves C, Gräßler J, Gorski M, Goodarzi M, Goodall A, Goel A, Giulianini F, Gigante B, Gianfagna F, Gertow K, Gao H, Ganesh S, Fraser R, Franco-Cereceda A, Dominiczak A. The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. *Nat Genet* 2016; 48:1171-1184.

Franco O, Lorenzo C, Karter A, Ingelsson E, Hansen T, Cupples L, Brown J, Bis J, Becker D, Zengini E, Yanek L, Mathias R, Norris J, Peloso G, Ferrannini E, Deloukas P, Dedoussis G, Bottinger E, Boeing H, Wagenknecht L, Varma R, Vaidya D, Toniolo D, Sheu W, Javad S, Tsafantakis E, Traglia M, Rayner N, Peter A, Pasko D, Palmer N, Ntalla I, Muzny D, Mohlke K, Metcalf G, McLeod O, McKean-Cowdin R, Renström F, Rice K, Sala C, Torres M, Thanopoulou A, Tentolouris N, Stirrups K, Stahl E, Speliotes E, Soranzo N, Smith J, Serafetinidis I, Sennblad B, Matchan A, Goodarzi M, van Duijn C, Tai E, Psaty B, Pedersen O, Chasman D, Borecki I, Laakso M, Zeggini E, Wong T, Wareham N, Waterworth D, Boerwinkle E, Scott R, Meigs J, Rotter J, Dupuis J, Siscovick D, Frayling T, Wilson J, Loos R, Florez J, Kao W, Watkins H, Walker M, Uitterlinden A, Launer L, Langenberg C, Jansson J, Hofman A, Hayward C, Hattersley A, Harris T, Hamsten A, Gudnason V, Gibbs R, Levy D, Oostra B, O'Donnell C, Smith B, Schulze M, Rudan I, Ridker P, Rich S, Province M, Polasek O, Pankow J, Padmanabhan S, O'Rahilly S, Franks P, Maruthur N, Amin N, Meidtner K, Hua Zhao J, Li M, Layton J, Lange L, Jakobsdottir J, Isaacs A, Hara K, Guo X, Garcia M, Morrison A, Nalls M, Peters M, Allin K, Varga T, Taylor K, Strawbridge R, Stoiber M, Southam L, Smith A, Silveira A, Schurmann C, Sabater-Lleal M, Freitag D, Fornage M, Bork-Jensen J, Hidalgo B, Lipovich L, Raghavan S, Hivert M, Dauriz M, Brody J, Yaghootkar H, Wang S, Willems S, Chu A, Fox K, Huffman J, An P, Boland A, Besse C, Abrol R, Stančáková A, Baldrige A, Li L, Ehm M, Grarup N, Rasmussen-Torvik L, Lu Y, Wessel J, Marouli E, Kirkpatrick A, Khor C, Karaleftheri M, Jørgensen T, Jørgensen M, Jensen R, Ikram M, Hoffmann P, Heo J, Hallmans G, Kraja A, Kuusisto J, Lange E, Mamakou V, Malerba G, Linneberg A, Lindgren C, Liu Y, Liu C, Liao J, Leong A, Lee W, Lee I, Hai Y, Gustafsson S, Grove M, Cheng C, Chen Y, Chen Y, Burns S, Bowden D, Bombieri C, Boehnke M, Bihlmeyer N, Barbieri C, Aung T, Correa A, Czajkowski J, Dehghan A, Gottesman O, Goel A, Goddard W, Giulianini F, Gambaro G, Frånberg M, Farmaki A, Escher S, Eiriksdottir G, Ehret G, Aponte J. Low-frequency and rare exome chip variants associate with fasting glucose and type 2 diabetes susceptibility. *Nat Commun* 2015; 6:5897.

Projects (0)

No results found.

Kantonsspital St.Gallen

Rorschacher Strasse 95

CH-9007 St.Gallen

T: +41 71 494 11 11

support.forschung@kssg.ch