



Nicole Soranzo

Contact

Nicole Soranzo

Publications (3)

Palotie A, Linneberg A, Husemoen L, Jarvelin M, Franks S, Blakemore A, Kooner J, Chambers J, Oksa H, Korpi-Hyövähti E, Petersen E, Christensen C, Skaaby T, Thuesen B, Käräjämäki A, Groop L, Tuomi T, Willer C, Hveem K, Holmen O, Palmer C, Morris A, Doney A, Tuomilehto J, Karpe F, Brandslund I, Numans M, Jukema J, Connell J, Dominiczak A, Brown M, Wareham N, Langenberg C, Starr J, Deary I, Männistö S, Jousilahti P, Spector T, Sattar N, Ford I, de Bakker P, Lind L, Ingelsson E, Gambaro G, van der Harst P, van der Meer P, de Boer R, Metspalu A, Mägi R, Esko T, Packard C, Dedoussis G, Ripatti S, Munroe P, Tobin M, Newton-Cheh C, Frossard P, Stringham H, Boehnke M, Pedersen O, Hansen T, Mohlke K, Collins F, Scotland G, Hayward C, Nordestgaard B, Caulfield M, Howson J, Butterworth A, Wain L, Danesh J, Lindgren C, Asselbergs F, Saleheen D, Samani N, Tomaszewski M, Morris A, Mahajan A, Porteous D, Padmanabhan S, Blankenberg S, Arveiler D, Amouyel P, Sever P, Stanton A, Poulter N, McCarthy M, Chowdhury R, Di Angelantonio E, Shafi Majumder A, Alam D, Ferrieres J, Kee F, Laakso M, Kuusisto J, Melander O, Kathiresan S, Zeggini E, Elliott P, Deloukas P, Virtamo J, Veronesi G, Müller-Nurasyid M, Kuulasmaa K, Salomaa V, Franks P, Yiorkas A, Matchan A, Ohlsson T, Fava C, Stančáková A, Huyghe J, Marten J, Southam L, Swift A, Narisu N, Jackson A, Bonnycastle L, Stirrups K, Bork-Jensen J, Robertson N, Neville M, Rayner N, Groves C, Donnelly L, Zhang H, Havulinna A, Shaw-Hawkins S, Perola M, Kontto J, Gjesing A, Zhao W, Samuel M, Tragante V, Staley J, Witkowska K, Barnes D, Sim X, Grarup N, Manning A, Cook J, Warren H, Young R, Drenos F, Tukiainen T, Yaghootkar H, Rasheed A, Fallgaard Nielsen S, Kraja A, Liu C, Mihailov E, Harakalova M, Tinker A, Giannakopoulou O, Ferreira T, Freitag D, Masca N, Surendran P, Herzig K, Rolandsson O, Lin H, Vogt T, Hoek M, Reily D, Malarstig A, Uria-Nickelsen M, Felix J, Vasani R, Burgess S, Hassinen M, Lieb W, Traylor M, Rudan I, Polasek O, Rauramaa R, Lakka T, Komulainen P, Uusitupa M, Lindström J, Marouli E, Justice A, Highland H, Markus H, Huffman J, Renström F, Nelson C, Vergnaud A, Moayyeri A, Evangelou E, Verweij N, Trabetti E, Soranzo N, Malerba G, Lannfelt L, Willems S, Zhang W, Poveda A, Varga T, Hallmans G, Farmaki A, Menni C, Marioni R, Liewald D, Harris S, Scott R, Luan J, Trompet S, de Craen A, Caslake M, Kajantie E. Trans-ancestry meta-analyses identify rare and common variants associated with blood pressure and hypertension. *Nat Genet* 2016; 48:1151-1161.

Franco O, Lorenzo C, Karter A, Ingelsson E, Hansen T, Cupples L, Brown J, Bis J, Becker D, Zeggini 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, Tsfantakis 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.

Morris A, Stringham H, Swift A, Tuomi T, Uda M, Vollenweider P, Waeber G, Wallace C, Walters G, Weedon M, Witteman J, Zhang C, Zhang W, Caulfield M, Collins F, Davey Smith G, Yuan X, Song K, Munroe P, Narisu N, Nordström A, Nordström P, Oostra B, Palmer C, Payne F, Peden J, Prokopenko I, Renström F, Ruukonen A, Salomaa V, Sandhu M, Scott L, Scuteri A, Silander K, Day I, Franks P, Hugh Watkins, Waterworth D, Boehnke M, Deloukas P, Groop L, Hunter D, Thorsteinsdottir U, Schlessinger D, Wichmann H, Frayling T, Abecasis G, Hirschhorn J, Loos R, Stefansson K, Mohlke K, Barroso I, Wareham N, van Duijn C, Hattersley A, Hu F, Jarvelin M, Kong A, Kooner J, Laakso M, Lakatta E, Mooser V, Morris A, Peltonen L, Samani N, Spector T, Strachan D, Tanaka T, Tuomilehto J, Uitterlinden A, McCarthy M, Lindgren C, Lyon H, Rivadeneira F, Sanna S, Timpson N, Zillikens M, Zhao J, Almgren P, Bandinelli S, Bennett A, Bergman R, Bonnycastle L, Bumpstead S, Chanock S, Cherkas L, Chines P, Coin L, Luan J, Drong A, Heid I, Randall J, Lamina C, Steinthorsdottir V, Qi L, Speliotes E, Thorleifsson G, Willer C, Herrera B, Jackson A, Lim N, Scheet P, Soranzo N, Amin N, Aulchenko Y, Chambers J, Cooper C, Crawford G, Hofman A, Holle R, Holloway J, Illig T, Isomaa B, Jacobs L, Jameson K, Jousilahti P, Karpe F, Kuusisto J, Laitinen J, Lathrop G, Lawlor D, Mangino M, McArdle W, Meitinger T, Havulinna A, Hamsten A, Doering A, Dominiczak A, Doney A, Ebrahim S, Elliott P, Erdos M, Estrada K, Ferrucci L, Fischer G, Forouhi N, Gieger C, Grallert H, Groves C, Grundy S, Guiducci C, Hadley D, Morken M. Genome-wide association scan meta-analysis identifies three Loci influencing adiposity and fat distribution. *PLoS Genet* 2009; 5:e1000508.

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