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Publications (2)

Kovacs P, Pankow J, Payne F, Prokopenko I, Psaty B, Renström F, Rice K, Rotter J, Rybin D, Sandholt C, Sayer A, Shrader P, North K, Müller G, Kumari M, Kuusisto J, Laakso M, Lecoeur C, Lévy-Marchal C, Li G, Loos R, Lyssenko V, Marmot M, Marques-Vidal P, Morken M, Schwarz P, Siscovich D, Stančáková A, Vollenweider P, Walker M, Watanabe R, Pedersen O, Meigs J, Ingelsson E, Barroso I, Florez J, Franks P, Dupuis J, Wareham N, Kao W, Groop L, Sturmöller M, Teslovich T, Waeber G, Williams G, Witte D, Wood A, Xie W, Boehnke M, Cooper C, Ferrucci L, Froguel P, Langenberg C, Kivimäki M, Scott R, Marek D, Rasmussen-Torvik L, Stringham H, Tanaka T, Aadahl M, Arking D, Bergmann S, Boerwinkle E, Bonnycastle L, Bornstein S, Brunner E, Lagou V, Kutalik Z, Chu A, Grarup N, Manning A, Hivert M, Shungin D, Tönjes A, Yesupriya A, Barnes D, Bouatia-Naji N, Glazer N, Jackson A, Bumpstead S, Brage S, Carlson O, Gräßler J, Gustafsson S, Hallmans G, Hansen T, Hingorani A, Holloway J, Hu F, Isomaa B, Jameson K, Johansson I, Jonsson A, Goodarzi M, Fox C, Chen H, Chen Y, Chines P, Collins F, Couper D, Dennison E, Dowling N, Egan J, Ekelund U, Erdos M, Forouhi N, Jørgensen T. No interactions between previously associated 2-hour glucose gene variants and physical activity or BMI on 2-hour glucose levels. *Diabetes* 2012; 61:1291-6.

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, Ruokonen 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, Kooper 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.

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