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Publications (3)

Deloukas P, Escher S, Dedoussis G, Blakemore A, Waldenberger M, Tsafantakis E, Tomaszewski M, Strauch K, Stanton A, Silveira A, Shields D, Sever P, Sennblad B, Sabater-Lleal M, Kooner J, McCarthy M, Palmer C, Samani N, Franks P, Munroe P, Wareham N, Chambers J, Gieger C, Zeggini E, Jarvelin M, Tobin M, Frayling T, Caulfield M, Hamsten A, Rolandsson O, Renström F, An Hashim N, Lataniotis L, Strawbridge R, Couto Alves A, Müller-Nurasyid M, Yaghootkar H, Zhang W, Southam L, Scott R, Warren H, Varga T, Stirrups K, Masca N, Besse C, Boland A, Braund P, Rayner N, Poulter N, Peters A, Pasko D, Matchan A, Keinänen-Kiukaanniemi S, Karaleftheri M, Jansson J, Grallert H, Franks S, Farmaki A, Dominiczak A, Connell J, Kanoni S. Analysis with the exome array identifies multiple new independent variants in lipid loci. *Hum Mol Genet* 2016; 25:4094–4106.

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.

Patin E, Stewart G, Booth D, George J, Casanova J, Bréchet C, Rice C, Talal A, Jacobson I, Bourlière M, Theodorou I, Poynard T, Negro F, Pol S, Bochud P, Abel L, Swiss Hepatitis C Cohort Study Group, International Hepatitis C Genetics Consortium, Suppiah V, Martinetti G, Hirsch H, Kutalik Z, Guergnon J, Bibert S, Nalpas B, Jouanguy E, Munteanu M, Bousquet L, Argiro L, Halfon P, Boland A, Mullhaupt B, Semela D, Dufour J, Heim M, Moradpour D, Cerny A, Malinverni R, French ANRS HC EP 26 Genoscan Study Group. Genome-wide association study identifies variants associated with progression of liver fibrosis from HCV infection. *Gastroenterology* 2012; 143:1244–52.e1-12.

Projects (0)

No results found.

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