



**Céline Besse**

**Kontakt**

Céline Besse

## Publikationen (2)

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, Gallert 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.

## Projekte (0)

Keine Resultate gefunden.

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