



**Dawn M Waterworth**

**Kontakt**

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## Publikationen (4)

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, 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, Tsafantakis 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 M, 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, Ellinor P, Howson J, Jukema J, Kahali B, Kahn R, Kähönen M, Kamstrup P, Kanoni S, Kaprio J, Karaleftheri M, Kardia 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, Hernesniemi 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.

Perola M, Vartiainen E, Jousilahti P, Rao D, Sarzynski M, Sørensen T, Astrup A, Kloppenburg M, Meulenbelt I, Zhang Y, Hofman A, Rivadeneira F, Jula A, Koskinen S, Knekt P, Heliövaara M, Eriksson J, Palotie A, Osmond C, Kajantie E, Uitterlinden A, Zeller T, Yao J, Sattar N, Ridker P, Brent Richards J, Rice T, Rathmann W, Rasmussen-Torvik L, Rankinen T, Pradhan A, Peters A, Savage D, Söderberg S, Wood A, Widén E, Wichmann H, Walker M, Vohl M, Uh H, van Heemst D, Vandenput L, Timpson N, Paternoster L, Loos R, Hansen T, Vestergaard H, Vollenweider P, Waeber G, Bergmann S, Waterworth D, Borecki I, Pramstaller P, Frayling T, Pedersen O, Hu F, Leibel R, Lindgren C, Franks P, Schadt E, Klein R, Jukema J, Spector T, Grallert H, Eline Slagboom P, Ferrucci L, Wareham N, März W, Bandinelli S, Casas J, Lorentzon M, Mellström D, Raitakari O, Lehtimäki T, Kähönen M, Viikari J, Isaacs A, van Dijk K, Langenberg C, Scott R, Lind L, Ohlsson C, Chasman D, Allison M, Bouchard C, Harris T, van Duijn C, Huupponen R, Nalls M, Buckley B, Liu Y, Guo X, Rose L, Mahajan A, Willems S, Renström F, Pasko D, Del Greco M F, Luan J, Kleber M, Pérusse L, Blangero J, Bellis C, Barroso I, Amuzu A, Amin N, Ramos Y, Ju Sung Y, Ahluwalia T, Gaunt T, Tanaka T, Lahti J, Zhao J, Hayes J, Drong A, Hedman Å, Feitosa M, Kriebel J, Sun Q, Skowronski A, Carli J, Pers T, Schick U, Henneman P, Eriksson J, Lyytikäinen L, Beekman M, Kristiansson K, Mangino M, Trompet S, Kutalik Z, Grarup N, Kilpeläinen T, Männistö S, Kraff P, Koenig W, Karlsson M, Jørgensen T, Jørgensen M, Jenny N, Jansson J, Ioan-Facsinay A, Ingelsson E, Kwekkeboom J, Laatikainen T, Myers R, Morris A, Menni C, Meisinger C, Marques-Vidal P, Lu Y, Lowe G, LeDuc C, Ladwig K, Illig T, Hunter D, Demirkan A, Delgado G, Deelen J, Day F, Dastani Z, Dale C, Crosslin D, de Craen A, I Chen Y, Finucane F, Ford I, Hicks A, Hernandez D, Herder C, Havulinna A, Hankinson S, Hallmans G, Gustafsson S, Gieger C, Garcia M, Böhringer S. Genome-wide meta-analysis uncovers novel loci influencing circulating leptin levels. *Nat Commun* 2016; 7:10494.

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, Yaghoobkar 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, Wittteman 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, 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.

## Projekte (0)

Keine Resultate gefunden.

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