



**Pedro Marques-Vidal**

## Contact

Pedro Marques-Vidal

## Publications (8)

Quteineh L, Mohacsi P, Dufour J, Soccal P, Katalik Z, Marques-Vidal P, Vollenweider P, Recher M, Hess C, Pascual M, Eap C, Steiger J, Van Delden C, Wójtowicz A, Bochud P, Crettol S, Vandenberghe F, Venetz J, Manuel O, Golshayan D, Lehmann R, Mueller N, Binet F, Swiss Transplant Cohort Study. Genetic immune and inflammatory markers associated with diabetes in solid organ transplant recipients. *Am J Transplant* 2018; 19:238–246.

Zhao W, Stott D, Staessen J, Snieder H, Slagboom P, Silbernagel G, Siemelink M, Sennblad B, Scott R, Schurmann C, Scholtens S, Schlessinger D, Savonen K, Sattar N, Sarzynski M, Sanna S, Rudan I, Bien S, Swertz M, Swift A, Taylor K, Wright A, Wong A, Wolffenbuttel B, Willemsen G, Wild S, Westendorp R, Waldenberger M, Waeber G, Vonk J, Völzke H, Vohl M, Vandenput L, Uitterlinden A, Tuomilehto J, Thuillier D, Thorand B, Tayo B, Rose L, Ridker P, Rice T, McKenzie C, Martin N, Marques-Vidal P, Männikkö R, Madden P, Lupoli S, Ludolph-Donislawski A, Lubke G, Luben R, Lorentzon M, Loh M, Lobbens S, Linneberg A, Lindgren C, Lind L, Lee N, Leander K, McKnight B, Mellström D, Menni C, Rawal R, Rasmussen-Torvik L, Rao D, Rankinen T, Raitakari O, Porteous D, Pisinger C, Peyser P, Padmanabhan S, Ong K, Olden M, Oldehinkel A, Nolte I, Nauck M, Narisu N, Musk A, Montgomery G, Launer L, Cupples L, Whitfield J, Weir D, Wareham N, Vollenweider P, Vestergaard H, van der Harst P, Tremoli E, Tiemeier H, Strauch K, Sørensen T, Smith B, Rauramaa R, Qi L, Psaty B, Polasek O, Peters U, Peters A, Wilson J, Tyrrell J, Frayling T, North K, Borecki I, Liu C, Kilpeläinen T, Loos R, Marchini J, Mohlke K, Heid I, van Duijn C, Strachan D, Spector T, Hunter D, Hirschhorn J, Fox C, Deloukas P, Boehnke M, Barroso I, Pedersen O, Pasterkamp G, Palmer L, Froguel P, Franks P, Ferrucci L, Faire U, Cusi D, Cucca F, Cooper R, Chines P, Chen Y, Chasman D, Bruinenberg M, Bouchard C, Boomsma D, Böger C, Bandinelli S, Balkau B, Baldassarre D, Gordon-Larsen P, Grabe H, Gudnason V, Ohlsson C, Morris A, Metspalu A, McCarthy M, März W, Marchand L, Lehtimäki T, Laakso M, Kuh D, Kooner J, Kivimäki M, Kardia S, Wouter Jukema J, Johnson A, Hveem K, Hayward C, Haiman C, Zillikens M, Langenberg C, Zhang W, Pérusse L, Navarro P, Müller-Nurasyid M, Middelberg R, Marten J, Manichaikul A, Mangino M, Lyytikäinen L, Luan J, Lim U, Kumari M, Komulainen P, Kleber M, Kinnunen L, Kanoni S, Justesen J, Jackson A, Pervjakova N, Sarti C, Smith A, Yengo L, Wu Y, Vitart V, Vink J, Verweij N, Vedantam S, van Vliet-Ostaptchouk J, van der Most P, van der Laan S, Trompet S, Teumer A, Tanaka T, Sung Y, Stringham H, Strawbridge R, Stančáková A, Smith J, Huffman J, Huang T, Horikoshi M, Katalik Z, Eicher J, Perez J, Lim E, Heard-Costa N, Chu A, Ahluwalia T, Ngwa J, Hadley D, Czajkowski J, Deng X, Barata L, Young K, Fisher V, Graff M, Feitosa M, Winkler T, Xue L, Mahajan A, Renström F, Neergaard Harder M, Gorski M, Fischer K, Eriksson J, Direk N, Corre T, Coggeshall S, Chittani M, Cadby G, Bragg J, Bonnefond A, Bielak L, Amin N, Alfred T, Ahmad S, Qi Q, Wu J, Justice A, Lakka T, Homuth G, Holmen O, Hollensted M, Hocking L, Hindorff L, Hernandez D, Heath A, Hastie N, Hassinen M, Hartman C, Harris T, Hansen T, Hamsten A, Hallmans G, Haitjema S, Grarup N, Grammer T, Jan Hottenga J, Huang J, Hung J, Kvaløy K, Kuusisto J, Krämer B, Kooperberg C, Kolovou G, Kolcic I, Koistinen H, Karlsson M, Kähönen M, Juonala M, Jørgensen M, Jhun M, Jarvelin M, Jansson J, James A, Ingelsson E, Hutri-Kähönen N, Grallert H, Gong J, Glorioso N, Braga D, Bottinger E, Bonnycastle L, Boerwinkle E, Blangero J, Biffar R, Bergmann S, Bergman R, Beilby J, Bartz T, Bakker S, Afzal U, Afaq S, Adair L, Abecasis G, Zubair N, Zimmermann M, Buckley B, Buyske S, Campbell H, Gigante B, Gertow K, Forrester T, Ford I, Faul J, Esko T, Eriksson A, Eiriksdottir G, den Ruijter H, Delgado G, Dedoussis G, de Geus E, de Craen A, de Borst G, Curran J, Collins F, Chambers J, Hua Zhao J. Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. *Nat Commun* 2017; 8:14977.

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, Amuzi 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, Katalik Z, Grarup N, Kilpeläinen T, Männistö S, Kraft 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, Demirkiran 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.

Bochud M, Simone E, Andrea F, Frei M, Claudius F, Jakob F, Ali G, Matthias G, Denis H, Marcel H, Walter H, Simon H, Felix H, Paul H, Eva K, Kaplan V, Peter D, Corinne C, Jean Michel G, Christoph H, Thomas L, Pedro M, Paccaud F, von Känel R, Aidacic V, Jürg B, Markus B, Heinz B, Martin B, Hans-Ulrich B, Ivo B, Reto C, Isabelle C, Daniel K, Stephan K, Beat K, Georg S, Pietro S, Jacques S, Alfred S, Alois S, Claudia S, Othmar S, Phuoc T, Marco V, Alessandro V, René V, Hans W, Fritz W, Johanna W, Joseph W, Frank S, Hana S, Benedict K, Heidi K, Vesna L, Giovanni L, Werner L, Phillippe L, Severin L, Christoph M, Jürgen M, Damian M, Werner M, Titus M, Valentina N, Jakob R, Rosemann T, Marco Z, Hasse B, CoLaus Cohort, FIRE and the Swiss HIV Cohort Study, Aubert V, Barth J, Battegay M, Bernasconi E, Böni J, Bucher H, Burton-Jeangros C, Calmy A, Cavassini M, Egger M, Elzi L, Fehr J, Fellay J, Furrer H, Ledergerber B, Vollenweider P, Tarr P, Marques-Vidal P, Waeber G, Preisig M, Mooser V, Valeri F, Djalali S, Rauch A, Bernasconi E, Calmy A, Cavassini M, Vernazza P, Battegay M, Weber R, Senn O, Fux C, Gorgievski M, Günthard H, Rauch A, Regenass S, Rickenbach M, Rudin C, Schöni-Affolter F, Schmid P, Schultze D, Schüpbach J, Speck R, Staehelin C, Tarr P, Telenti A, Trkola A, Weber R, Yerly S, Pantaleo G, Nadal D, Haerry D, Hasse B, Hirsch H, Hösl I, Kahler C, Kaiser L, Keiser O, Klimkait T, Kouyos R, Kovari H, Ledergerber B, Martinetti G, Martinez de Tejada B, Metzner K, Müller N, Jean-Michel A. Strong Impact of Smoking on Multimorbidity and Cardiovascular Risk Among Human Immunodeficiency Virus-Infected Individuals in Comparison With the General Population. *Open Forum Infect Dis* 2015; 2:ofv108.

Mukamal K, Liu Y, Danesh J, Rasheed A, Mason M, Zonderman A, Franke L, Kristal B, Karjalainen J, Reed D, Westra H, Evans M, Saleheen D, Harris T, Dedoussis G, Curhan G, Stumvoll M, Beilby J, Houston D, Lorentzon M, Psaty B, Döring A, Heath A, Montgomery G, Dahmen N, Carithers T, Tucker K, Ferrucci L, Boyd H, Melbye M, Treur J, Mellström D, Hottenga J, Prokopenko I, Tönjes A, Deloukas P, Kanoni S, Pasquale L, Feenstra B, Bandinelli S, Caporaso N, Grabe H, Neuhausen M, Wolffenbuttel B, Hu F, Hyppönen E, Jarvelin M, Cupples L, Franks P, Ridker P, van Duijn C, Heiss G, Metspalu A, North K, Ingelsson E, Nettleton J, van Dam R, Vollenweider P, Wareham N, Ordovas J, Chan A, Peters U, Ohlsson C, Gieger C, Martin N, Waldenberger M, Siscovick D, Raitakari O, Eriksson J, Mitchell P, Hunter D, Kraft P, Rimm E, Boomsma D, Borecki I, Loos R, Chasman D, Mozaffarian D, Wojczynski M, Vink J, Zhao J, Burlutsky G, Lahti J, Mikkilä V, Lemaitre R, Eriksson J, Musani S, Tanaka T, Geller F, Luan J, Hui J, Mägi R, Dimitriou M, Garcia M, Ho W, Manichaikul A, Rawal R, Cornelis M, Byrne E, Esko T, Nalls M, Ganna A, Paynter N, Monda K, Amin N, Fischer K, Renström F, Ngwa J, Huikari V, Cavadino A, Nolte I, Teumer A, Yu K, Marques-Vidal P, Wright M, Rose L, Magnusson P, Bastardot F, Kutalik Z, Jacobs D, Forouhi N, Mihailov E, Lind L, Lindgren C, Michaëlsson K, Morris A, Jensen M, Khaw K, Luben R, Wang J, Männistö S, Perälä M, Kähönen M, Lehtimäki T, Gu F, Biffar R, Pedersen N, Couper D, Oostra B, Hofman A, Ikram M, Tiemeier H, Uitterlinden A, van Rooij F, Barroso I, Johansson I, Xue L, Kaakinen M, Milani L, Power C, Snieder H, Stolk R, Baumeister S, Viikari J. Genome-wide meta-analysis identifies six novel loci associated with habitual coffee consumption. *Mol Psychiatry* 2014; 20:647–656.

Hansen T, Mozaffarian D, North K, Pedersen O, Raitakari O, Rissanen H, Tuomilehto J, Van der Schouw Y, Uitterlinden A, Zillikens M, Franco O, Shyong Tai E, Mikkilä V, Long J, Lohman K, Christiansen L, Hofman A, Johansson I, Jørgensen T, Karasawa S, Khaw K, Kim M, Kristiansson K, Li H, Lin X, Liu Y, Ou Shu X, Siscovick D, Uusitupa M, Cupples L, Rankinen T, Orho-Melander M, Wang T, Chasman D, Franks P, Sørensen T, Hu F, Loos R, Nettleton J, Mohlke K, Hunter D, Pasquale L, Toft U, Verschuren W, Vollenweider P, Wareham N, Witteman J, Zheng W, Ridker P, Kang J, Liang L, Jensen M, Curhan G, Qi L, Forouhi N, Qi Q, Xu M, Ahluwalia T, Boer J, Chen P, Daimon M, Eriksson J, Perola M, Friedlander Y, Gao Y, Heppe D, Holloway J, Asif Ali M, Li Y, Liu Z, Kilpeläinen T, Downer M, Tanaka T, Smith C, Sluijs I, Sonestedt E, Chu A, Renström F, Lin X, Ångquist L, Huang J, Houston D, Kanoni S, van Rooij F, Xiang Y, Wen W, Wojczynski M, Zhu J, Borecki I, Bouchard C, Cai Q, Cooper C, Dedoussis G, Deloukas P, Pérusse L, Ngwa J, Monda K, Kim Y, Laaksonen M, Jääskeläinen T, Lee N, Lehtimäki T, Lemaitre R, Lu W, Luben R, Manichaikul A, Männistö S, Marques-Vidal P, Ferrucci L. FTO genetic variants, dietary intake and body mass index: insights from 177,330 individuals. *Hum Mol Genet* 2014; 23:6961–72.

Faeh D, Marques-Vidal P, Brändle M, Braun J, Rohrmann S. Diabetes risk scores and death: predictability and practicability in two different populations. *Eur J Public Health* 2014; 25:26–8.

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, Sturm voll 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.

## 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](mailto:support.forschung@kssg.ch)