



**Chris Wallace**

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## Publications (3)

Lyons P, Moosig F, Neumann T, Ohlsson S, Quickert S, Ramirez G, Rewerska B, Schett G, Sinico R, Szczeklik W, Tesar V, Vukcevic D, European Vasculitis Genetics Consortium, Terrier B, Watts R, Vaglio A, Holle J, Wallace C, Martorana D, Little M, Peters J, Alberici F, Liley J, Coulson R, Astle W, Baldini C, Francesco B, Cid M, Elding H, Emmi G, Epplen J, Guillevin L, Jayne D, Jiang T, Gunnarsson I, Lamprecht P, Leslie S, Smith K. Genome-wide association study of eosinophilic granulomatosis with polyangiitis reveals genomic loci stratified by ANCA status. *Nat Commun* 2019; 10:5120.

Neumann T, Martorana D, Moosig F, Ohlsson S, Quickert S, Ramirez S, Rewerska B, Schett G, Sinico R, Szczeklik W, Tesar V, Vukcevic D, Terrier B, Watts R, Vaglio A, Holle J, Wallace C, Little M, Leslie S, Lamprecht P, Lyons P, Peters J, Alberici F, Liley J, Coulson R, Astle W, Baldini C, Francesco B, Cid M, Elding H, Emmi G, Epplen J, Guillevin L, Jayne D, Jiang T, Gunnarsson I, Smith K. Genetically distinct clinical subsets, and associations with asthma and eosinophil abundance, within Eosinophilic Granulomatosis with Polyangiitis. *bioRxiv (The Preprint Server for Biology)* 2018

Morris A, Stringham H, Swift A, Tuomi T, Uda M, Vollenweider P, Waeber G, Wallace C, Walters G, Weedon M, Wittman 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, Ruukonen 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.

## Projects (0)

No results found.

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