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Kenyan Bacteraemia Study Group, ; Wellcome Trust Case Control Consortium 2 (WTCCC2), ; Rautanen, A; Pirinen, M; Mills, TC; Rockett, KA; Strange, A; Ndungu, AW; Naranbhai, V; Gilchrist, JJ; Bellenguez, C; Freeman, C; Band, G; Bumpstead, SJ; Edkins, S; Giannoulatou, E; Gray, E; Dronov, S; Hunt, SE; Langford, C; Pearson, RD; Su, Z; Vukcevic, D; Macharia, AW; Uyoga, S; Ndila, C; Mturi, N; Njuguna, P; Mohammed, S; Berkley, JA; Mwangi, I; Mwarumba, S; Kitsao, BS; Lowe, BS; Morpeth, SC; Khandwalla, I; Kilifi Bacteraemia Surveillance Group, ; Blackwell, JM; Bramon, E; Brown, MA; Casas, JP; Corvin, A; Duncanson, A; Jankowski, J; Markus, HS; Mathew, CG; Palmer, CN; Plomin, R; Sawcer, SJ; Trembath, RC; Viswanathan, AC; Wood, NW; Deloukas, P; Peltonen, L; Williams, TN; Scott, JA; Chapman, SJ; Donnelly, P; Hill, AV; Spencer, CC (2016) Polymorphism in a lincRNA Associates with a Doubled Risk of Pneumococcal Bacteremia in Kenyan Children. *American journal of human genetics*, 98 (6). pp. 1092-100. ISSN 0002-9297 DOI: <https://doi.org/10.1016/j.ajhg.2016.03.025>

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