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Publications (2)

Pers T, Hirschhorn J, Fehrmann R, Raychaudhuri S, Boehnke M, Renström F, Speliotes E, Frayling T, Esko T, Gustafsson S, Vedantam S, Lui J, Yang J, Wood A, Westra H, Chan Y, Karjalainen J, Franke L. Biological interpretation of genome-wide association studies using predicted gene functions. *Nat Commun* 2015; 6:5890.

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Defining the role of common variation in the genomic and biological architecture of adult human height. *Nat Genet* 2014; 46:1173-86.

Projects (0)

No results found.

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