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Impact of common genetic determinants of Hemoglobin A1c on type 2 diabetes risk and diagnosis in ancestrally diverse populations: A transethnic genome-wide meta-analysis

DOI: 10.1371/journal.pmed.1002383
PLoS Medicine, 2017, 14, e1002383.

Source: <https://exaly.com/paper-pdf/68470479/citation-report.pdf>

Version: 2024-04-28

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#	Paper	IF	Citations
298	Genetic aetiology of glycaemic traits: approaches and insights. 2017 , 26, R172-R184		8
297	Genome-wide meta-analysis in Japanese populations identifies novel variants at the TMC6-TMC8 and SIX3-SIX2 loci associated with HbA. <i>Scientific Reports</i> , 2017 , 7, 16147	4.9	25
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