**Supplementary Table S4** Candidate causal variants selected for validation study through SNAP (high LD with rs1413299).

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| SNP | Chr. | Positiona | Alleleb | r2 c |
| rs7027650 | chr9 | 101741969 | T/A | 0.792 |
| rs10988451 | chr9 | 101741666 | G/A | 0.789 |

a Position is GRCh37.

b Risk allele/other allele.

c r2 of linkage disequilibrium between variants with rs1413299.