

**Table S4.** Results of TaqMan validation for GWAS and replication set.

	Concordance rate at each CNP locus (%)				
	rs8192723 (41,356,054*)	rs10418304 (41,356,751*)	rs10422346 (41,362,173*)	rs8102683 (41,363,765*)	rs8105704 (41,363,898*)
GWAS set <sup>†</sup> (N=445)	96.6	98.9	98.2	99.6	99.3
Replication set <sup>‡</sup> (N=759)	96.7	97.8	97.4	99.2	98.8

We calculated the concordance rates by comparing the result of PlatinumCNV (maximum a posteriori copy number dosages) and the TaqMan validation result.

\* NCBI Build. 37; hg19.

<sup>†</sup> qPCR assay position: 41,355,978-41,356,078 (Build 37; hg19)

<sup>‡</sup> qPCR assay position: 41,363,935-41,364,006 (Build 37; hg19)