

Supplemental Table 1. Genetic instrument for chronic kidney disease stage ≥ 3 phenotype

Chromosome	Position	SNP	Gene	Minor allele	Other allele	Beta	standard error	p-value
1	243464416	rs2636321	<i>SDCCAG8</i>	A	G	0.090773	0.016491	3.70E-08
2	73691144	rs9678093	<i>ALMS1</i>	C	A	-0.10081	0.017078	3.57E-09
2	73800881	rs758884626	<i>ALMS1</i>	G	GT	-0.0944	0.017193	4.01E-08
2	73834035	rs4852308	<i>ALMS1</i>	T	G	-0.09382	0.017201	4.92E-08
2	73934095	rs560203661	-	G	GA	-0.10024	0.01761	1.25E-08
4	77373079	rs2870238	<i>SHROOM3</i>	C	T	-0.10615	0.016497	1.24E-10
5	697380	rs56139727	<i>TPPP</i>	G	T	0.117045	0.019558	2.17E-09
5	176794191	rs12654812	<i>RGS14</i>	A	G	0.101139	0.017068	3.11E-09
7	1281064	rs12702509	<i>UNCX</i>	G	A	-0.10745	0.019558	3.93E-08
7	151415746	7:151415746_GT_G	<i>WDR86-AS1</i>	G	GT	0.10696	0.017587	1.19E-09
11	30757133	11:30757133_CAAAT_C	C	CAAAT	0.090371	0.016505	4.37E-08	
11	30777014	rs34830202		GA	G	-0.09653	0.016649	6.71E-09
11	65493191	rs4930159	<i>RNASEH2C</i>	T	G	0.095528	0.016506	7.15E-09
15	45650360	15:45650360_CT_C	<i>SQOR</i>	C	CT	0.105908	0.018382	8.32E-09
15	45682277	rs1145078	<i>GATM</i>	T	C	0.101627	0.018447	3.60E-08
16	20408529	rs66891425	<i>PDILT</i>	G	A	-0.12304	0.016973	4.19E-13
17	19474875	rs111653425	<i>SLC47A1</i>	T	C	0.356842	0.06479	3.63E-08