

Table S2. Linkage disequilibrium test results in the whole population.

Chr	Position (base pair)	SNP1	Gene	Chr	Position (base pair)	SNP2	Gene	R ²	D'
12	47844974	rs731236	VDR	12	47845054	rs7975232	VDR	0.52241	0.971043
12	47844974	rs731236	VDR	12	47846052	rs1544410	VDR	0.67492	0.898375
12	47845054	rs7975232	VDR	12	47846052	rs1544410	VDR	0.438847	0.838338
12	57764205	rs4646536	CYP27B1	12	57768115	rs3782130	CYP27B1	0.410515	0.853081
12	57764205	rs4646536	CYP27B1	12	57768302	rs10877012	CYP27B1	0.731762	0.905063
12	57764205	rs4646536	CYP27B1	12	57768956	rs703842	CYP27B1	0.747667	0.89186
12	57768115	rs3782130	CYP27B1	12	57768302	rs10877012	CYP27B1	0.386295	0.854472
12	57768115	rs3782130	CYP27B1	12	57768956	rs703842	CYP27B1	0.37292	0.836158
12	57768302	rs10877012	CYP27B1	12	57768956	rs703842	CYP27B1	0.758471	0.904393
Chr: Chromosome; SNP: Single Nucleotide Polymorphism.									