

S6 Table. Polymorphisms associated with gastric cancer susceptibility with p-value $<5 \times 10^{-5}$ in discovery phase and results of the validation phase and meta-analysis in female

Character					Phase I (GWAS phase)				Phase II (Validation phase)				Meta-analysis		
SNP	CHR	BP	MA	No.	MAF in cases	MAF in controls	Odds ratio	p-value	No.	MAF in cases	MAF in controls	Odds ratio	p-value	Odds ratio	p-value
rs2976394	8	143763622	C	743	0.38	0.53	0.53 (0.4-0.69)	3.36E-06	2,305	0.43	0.50	0.77 (0.64-0.92)	0.0043	0.68 (0.58-0.79)	6.71E-07
rs10216533	8	143763690	G	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,312	0.43	0.50	0.77 (0.64-0.92)	0.0048	0.68 (0.59-0.8)	9.49E-07
rs1045547	8	143763757	T	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,312	0.43	0.50	0.77 (0.64-0.92)	0.0048	0.68 (0.59-0.8)	9.49E-07
rs1045574	8	143763958	G	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,313	0.43	0.50	0.77 (0.64-0.93)	0.0060	0.69 (0.59-0.8)	1.29E-06
rs1045605	8	143764101	C	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,312	0.43	0.50	0.77 (0.64-0.92)	0.0048	0.68 (0.59-0.8)	9.49E-07
rs2294008	8	143761931	C	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,298	0.43	0.50	0.76 (0.64-0.92)	0.0043	0.68 (0.59-0.79)	8E-07
rs2294010	8	143762430	A	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,312	0.43	0.50	0.77 (0.64-0.92)	0.0048	0.68 (0.59-0.8)	9.49E-07
rs2920279	8	143762135	A	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,303	0.43	0.50	0.77 (0.64-0.93)	0.0053	0.69 (0.59-0.8)	1.1E-06
rs2920296	8	143763109	A	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,312	0.43	0.50	0.77 (0.64-0.92)	0.0048	0.68 (0.59-0.8)	9.49E-07
rs2920297	8	143763083	A	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,312	0.43	0.50	0.77 (0.64-0.92)	0.0048	0.68 (0.59-0.8)	9.49E-07
rs2920298	8	143763043	A	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,312	0.43	0.50	0.77 (0.64-0.92)	0.0048	0.68 (0.59-0.8)	9.49E-07
rs2976392	8	143762932	G	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,312	0.43	0.50	0.77 (0.64-0.92)	0.0048	0.68 (0.59-0.8)	9.49E-07
rs2976393	8	143763618	C	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,312	0.43	0.50	0.77 (0.64-0.92)	0.0048	0.68 (0.59-0.8)	9.49E-07
rs2976395	8	143763750	G	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,312	0.43	0.50	0.77 (0.64-0.92)	0.0048	0.68 (0.59-0.8)	9.49E-07
rs2976396	8	143764001	G	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,313	0.43	0.50	0.77 (0.64-0.93)	0.0057	0.69 (0.59-0.8)	1.22E-06
rs2978982	8	143763490	T	748	0.38	0.53	0.53 (0.41-0.7)	5.1E-06	2,312	0.43	0.50	0.77 (0.64-0.92)	0.0048	0.68 (0.59-0.8)	9.49E-07
rs1045531	8	143763547	C	745	0.38	0.53	0.53 (0.41-0.7)	5.41E-06	2,308	0.43	0.50	0.77 (0.64-0.92)	0.0052	0.69 (0.59-0.8)	1.12E-06
rs34635647	8	143756890	G	716	0.61	0.46	1.89 (1.44-2.5)	5.5E-06	2,236	0.44	0.50	0.77 (0.64-0.93)	0.0056	1.46 (1.25-1.71)	1.33E-06
rs2920282	8	143757763	C	743	0.62	0.47	1.86 (1.42-2.44)	6.1E-06	2,274	0.43	0.50	0.76 (0.63-0.92)	0.0040	1.47 (1.26-1.71)	8.08E-07
rs2920283	8	143757037	C	743	0.62	0.47	1.86 (1.42-2.44)	6.1E-06	2,274	0.43	0.50	0.76 (0.63-0.92)	0.0041	1.47 (1.26-1.71)	8.29E-07
rs2976386	8	143757409	A	743	0.62	0.47	1.86 (1.42-2.44)	6.1E-06	2,273	0.43	0.50	0.76 (0.63-0.92)	0.0041	1.47 (1.26-1.71)	8.33E-07
rs2978979	8	143757286	A	743	0.62	0.47	1.86 (1.42-2.44)	6.1E-06	2,273	0.43	0.50	0.76 (0.63-0.92)	0.0041	1.47 (1.26-1.71)	8.33E-07
rs2978980	8	143757708	G	743	0.62	0.47	1.86 (1.42-2.44)	6.1E-06	2,274	0.43	0.50	0.76 (0.63-0.92)	0.0040	1.47 (1.26-1.71)	8.08E-07
rs2978981	8	143759137	T	743	0.62	0.47	1.86 (1.42-2.44)	6.1E-06	2,275	0.43	0.50	0.76 (0.63-0.92)	0.0040	1.47 (1.26-1.71)	8.04E-07
rs2978977	8	143755720	A	714	0.50	0.35	1.86 (1.42-2.45)	6.78E-06	2,255	0.46	0.38	1.35 (1.12-1.63)	0.0014	1.5 (1.29-1.74)	2.31E-07
rs13249440	8	143756892	A	722	0.61	0.46	1.88 (1.43-2.47)	6.93E-06	2,252	0.43	0.50	0.77 (0.64-0.92)	0.0054	1.46 (1.25-1.7)	1.39E-06
rs13256647	8	143756895	T	722	0.61	0.46	1.88 (1.43-2.47)	6.93E-06	2,252	0.43	0.50	0.77 (0.64-0.92)	0.0054	1.46 (1.25-1.7)	1.39E-06
rs2976389	8	143760421	C	736	0.61	0.47	1.86 (1.42-2.44)	7.04E-06	2,269	0.43	0.50	0.76 (0.63-0.92)	0.0039	1.47 (1.26-1.71)	8.75E-07
rs2920292	8	143765699	G	732	0.61	0.47	1.86 (1.41-2.44)	8.82E-06	2,298	0.43	0.50	0.77 (0.64-0.93)	0.0054	1.45 (1.25-1.69)	1.69E-06
rs2920294	8	143765326	G	732	0.61	0.47	1.86 (1.41-2.44)	8.82E-06	2,298	0.43	0.50	0.77 (0.64-0.93)	0.0054	1.45 (1.25-1.69)	1.69E-06
rs2920295	8	143764937	G	732	0.61	0.47	1.86 (1.41-2.44)	8.82E-06	2,298	0.43	0.50	0.77 (0.64-0.93)	0.0054	1.45 (1.25-1.69)	1.69E-06
rs2976398	8	143764879	C	732	0.61	0.47	1.86 (1.41-2.44)	8.82E-06	2,298	0.43	0.50	0.77 (0.64-0.93)	0.0054	1.45 (1.25-1.69)	1.69E-06

rs2585179	8	143774193	A	738	0.61	0.47	1.84 (1.4-2.42)	1.08E-05	2,298	0.43	0.50	0.78 (0.65-0.94)	0.0075	1.44 (1.24-1.67)	2.9E-06
rs2920281	8	143760444	T	733	0.50	0.36	1.81 (1.39-2.36)	1.2E-05	2,271	0.46	0.38	1.36 (1.13-1.63)	0.0012	1.49 (1.28-1.73)	2.5E-07
rs2920280	8	143761144	C	745	0.51	0.37	1.8 (1.38-2.35)	1.21E-05	2,287	0.46	0.38	1.36 (1.13-1.63)	0.0012	1.49 (1.28-1.73)	2.44E-07
rs2976388	8	143760256	A	745	0.51	0.37	1.8 (1.38-2.35)	1.21E-05	2,284	0.46	0.38	1.35 (1.13-1.63)	0.0013	1.49 (1.28-1.73)	2.59E-07
rs2976397	8	143764613	T	745	0.51	0.37	1.8 (1.38-2.35)	1.21E-05	2,311	0.46	0.38	1.35 (1.13-1.62)	0.0012	1.48 (1.28-1.72)	2.54E-07
rs2920284	8	143756919	G	729	0.61	0.47	1.84 (1.4-2.42)	1.22E-05	2,258	0.43	0.50	0.77 (0.64-0.92)	0.0050	1.45 (1.25-1.7)	1.75E-06
rs2920285	8	143756218	C	729	0.61	0.47	1.84 (1.4-2.42)	1.22E-05	2,257	0.43	0.50	0.77 (0.64-0.92)	0.0049	1.46 (1.25-1.7)	1.69E-06
rs2920286	8	143754728	A	729	0.61	0.47	1.84 (1.4-2.42)	1.22E-05	2,257	0.43	0.50	0.76 (0.64-0.92)	0.0047	1.46 (1.25-1.7)	1.62E-06
rs2976385	8	143755426	C	729	0.61	0.47	1.84 (1.4-2.42)	1.22E-05	2,257	0.43	0.50	0.77 (0.64-0.92)	0.0049	1.46 (1.25-1.7)	1.69E-06
rs2978978	8	143756530	C	729	0.61	0.47	1.84 (1.4-2.42)	1.22E-05	2,258	0.43	0.50	0.77 (0.64-0.92)	0.0050	1.45 (1.25-1.7)	1.75E-06
rs2976387	8	143759364	A	740	0.50	0.36	1.8 (1.38-2.35)	0.000014	2,281	0.46	0.38	1.35 (1.13-1.63)	0.0012	1.49 (1.28-1.73)	2.93E-07
rs2585183	8	143766059	G	724	0.50	0.36	1.8 (1.38-2.36)	1.6E-05	2,288	0.46	0.38	1.34 (1.12-1.61)	0.0016	1.47 (1.27-1.71)	4.89E-07
rs2572910	8	143770135	G	722	0.60	0.46	1.83 (1.39-2.41)	1.9E-05	2,289	0.43	0.50	0.78 (0.65-0.94)	0.0074	1.43 (1.23-1.67)	4.26E-06
rs2717562	8	143776668	C	722	0.60	0.46	1.83 (1.39-2.41)	1.9E-05	2,284	0.43	0.50	0.78 (0.65-0.94)	0.0088	1.43 (1.22-1.66)	5.33E-06
rs2976384	8	143752994	C	727	0.61	0.47	1.82 (1.38-2.39)	1.93E-05	2,262	0.44	0.50	0.78 (0.64-0.93)	0.0075	1.44 (1.23-1.67)	4.01E-06
rs2585181	8	143771714	A	724	0.61	0.47	1.82 (1.38-2.39)	1.93E-05	2,286	0.44	0.50	0.78 (0.65-0.94)	0.0090	1.43 (1.22-1.66)	5.32E-06
rs2717608	8	143771712	C	724	0.61	0.47	1.82 (1.38-2.39)	1.93E-05	2,286	0.44	0.50	0.78 (0.65-0.94)	0.0090	1.43 (1.22-1.66)	5.32E-06
rs12233126	2	100012200	A	734	0.38	0.27	1.77 (1.34-2.33)	4.97E-05	2,312	0.28	0.28	1.01 (0.82-1.24)	0.9352	1.23 (1.04-1.45)	0.01347
rs28382949	2	100028112	T	734	0.38	0.27	1.77 (1.34-2.33)	4.97E-05	2,313	0.28	0.28	1.01 (0.82-1.24)	0.9380	1.23 (1.04-1.45)	0.01347
rs72956387	2	100028890	G	734	0.38	0.27	1.77 (1.34-2.33)	4.97E-05	2,313	0.28	0.28	1.01 (0.82-1.24)	0.9380	1.23 (1.04-1.45)	0.01347

MA, minor allele; MAF, minor allele frequency.