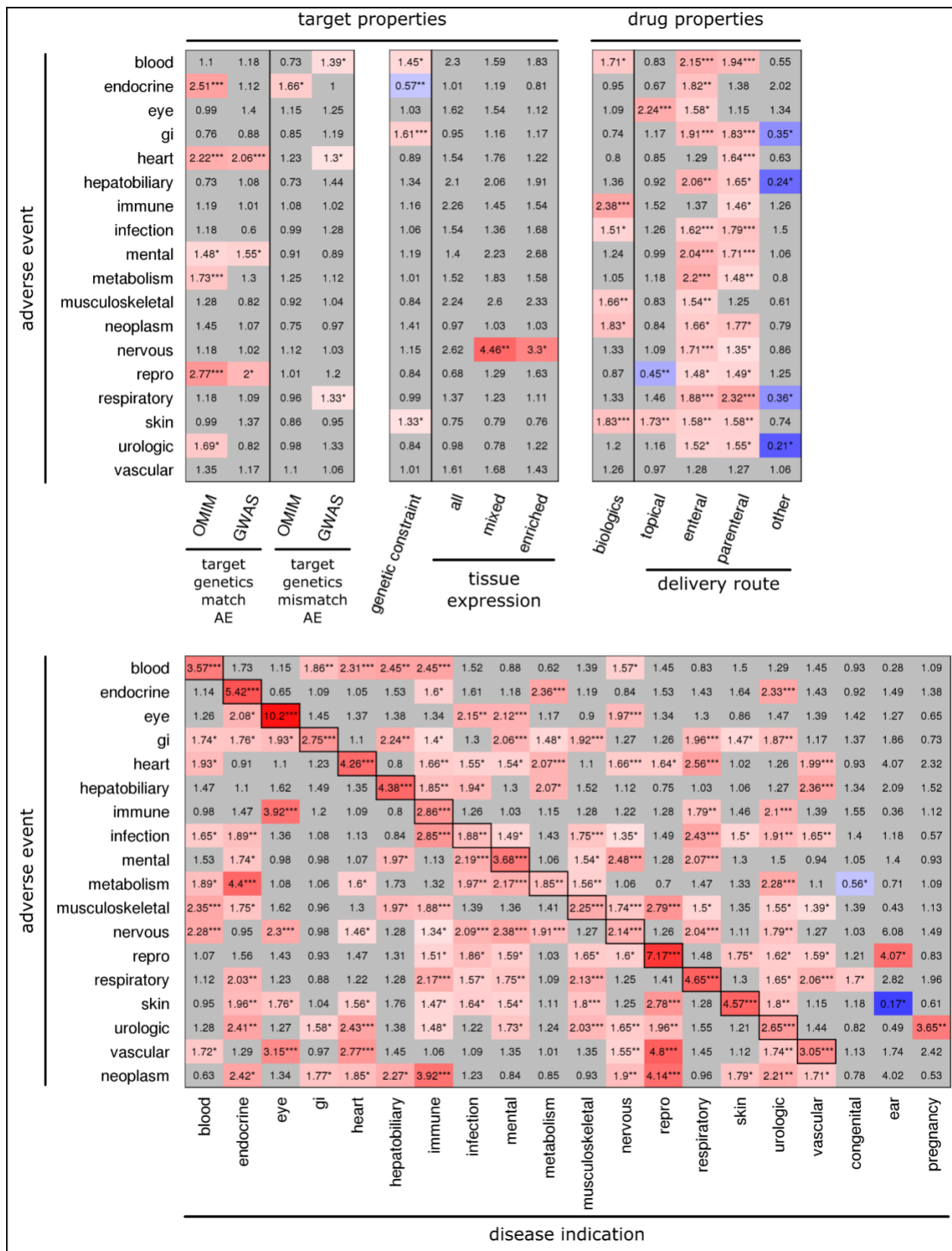


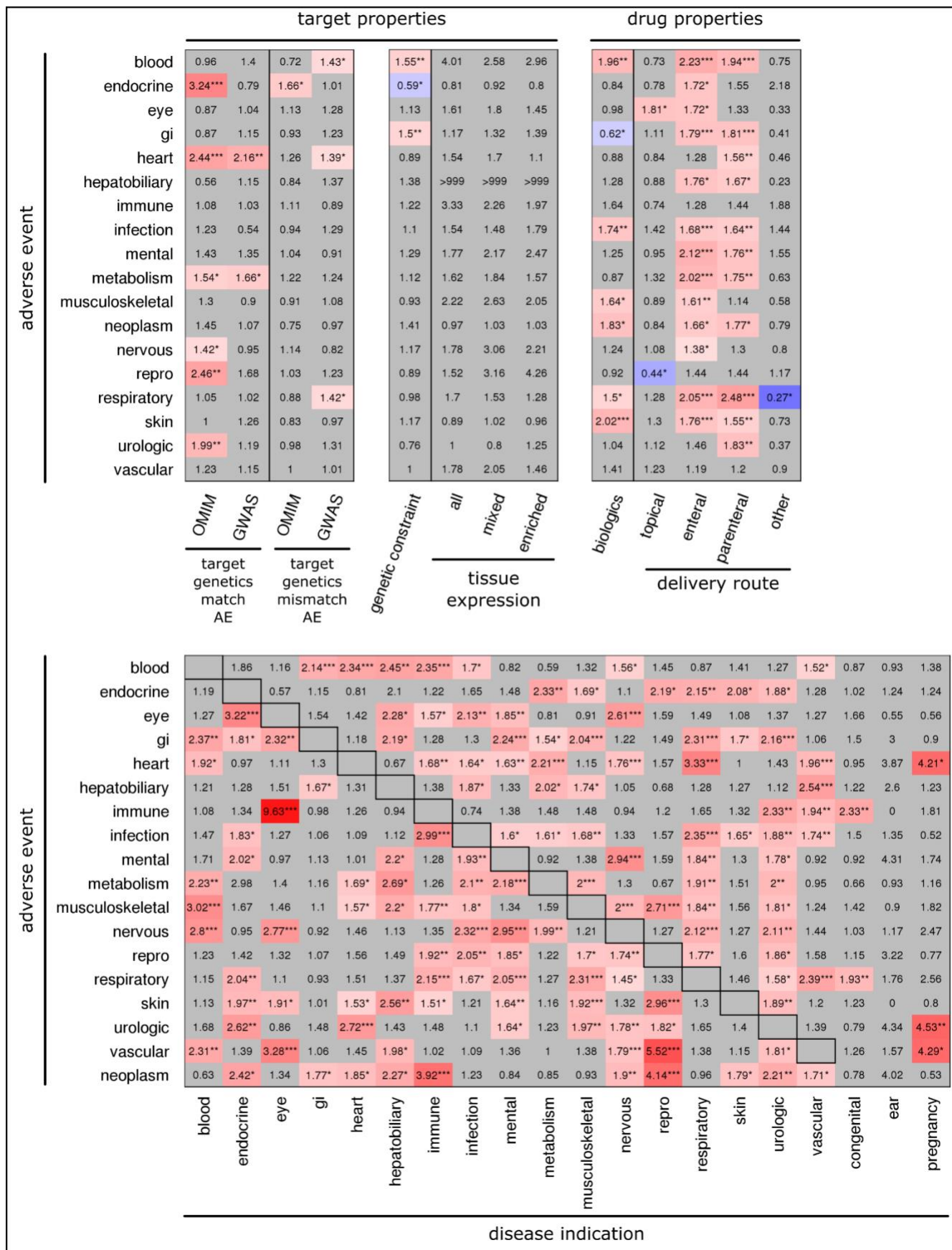
Supplementary Information for

Nguyen et al., “Phenotypes associated with genes encoding drug targets are predictive of clinical trial side effects”

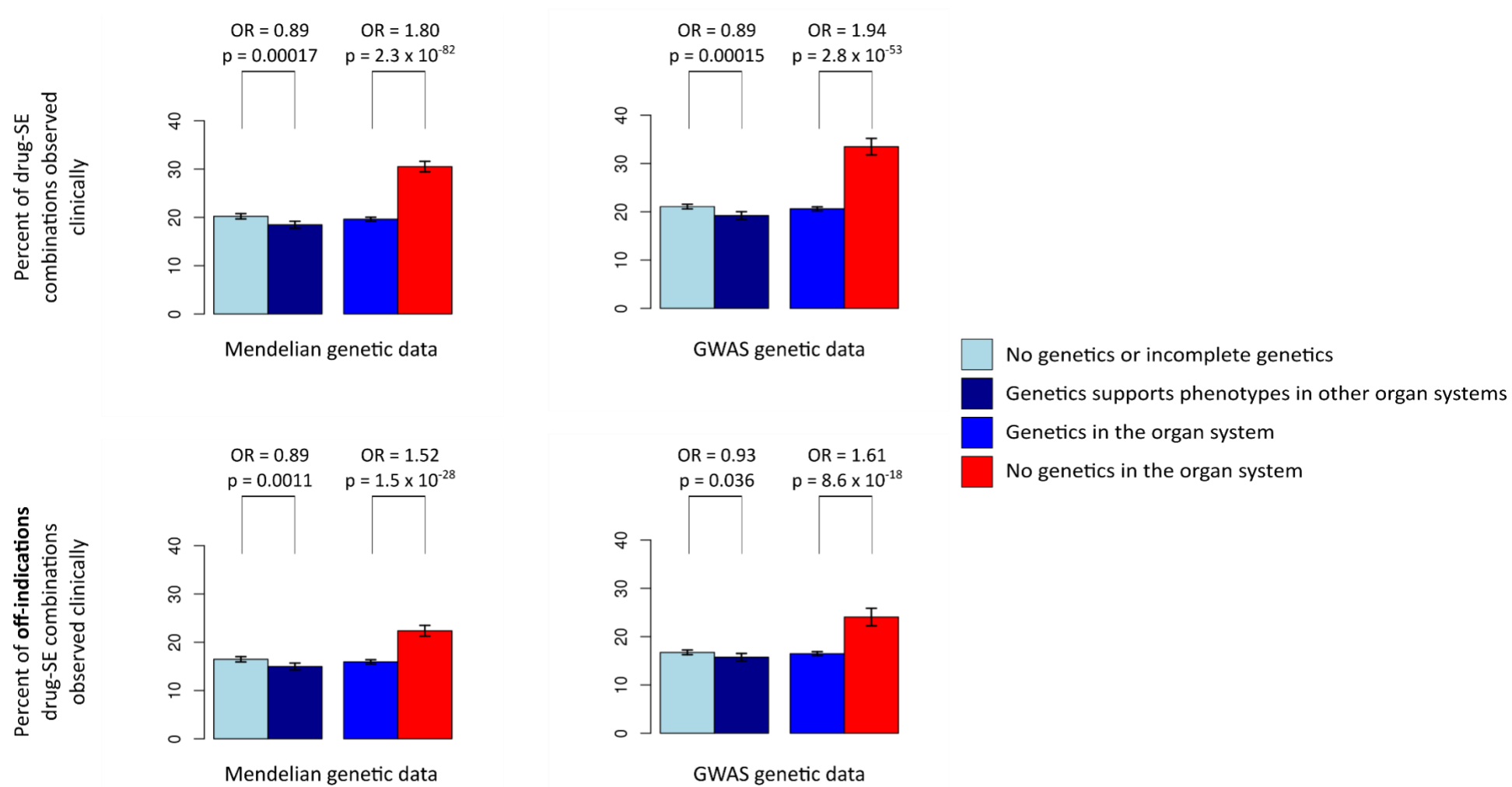
gen_gwas_blood_no (798)	0.29***	1.08	0.79	1.24	0.44**	1.79**	2.08	0.9	1.03	0.39**	0.65*	0.77	1.20*	0.38*	0.88	1.44**	1.11	0.33**	0.86	1.60**	1.11	0.73*	1.15	0.76	2.12***	0.65*	0.63*	1.67**	1.34*
gen_gwas_blood_eyes (194)	0.77***	1.82*	0.42**	0.96	1.47	1.06	2.44	1.8*	0.67	0.31**	0.56	0.37*	0.23**	0.55	0.37*	1.26	0.26	0.73	0.41**	0.75	0.29	0.19	0.12	0.16	0.35	1.28	0.59	2.41***	
gen_gwas_congenital_no (916)	0.21***	1.14	0.66**	1.20*	0.47**	1.9**	1.43	1	0.87	0.61*	0.28	0.59	1.16	0.58**	0.72*	1.03**	1.04	0.52**	0.86	1.07**	1.11	0.71**	1.13	0.76	2.12***	0.66**	0.62**	1.16	1.79**
gen_gwas_congenital_eyes (76)	2.44***	2.21*	0.41*	0.79	0.34**	0.77	1.95	1.7*	1.22	0.32**	0.52**	1.54	0.46	1.73	1.2	1.7	1.57	1.79*	0.98	0.22**	0.44*	1.18	0.89	1.54	1.11	1.4	2.6**	2.43**	1.8*
gen_gwas_ear_no (991)	0.24***	1.31	0.61**	1.22*	0.59**	1.62**	1.69	1.1	0.9	0.84	0.99	0.65	1.03	0.65*	0.79*	1.85**	1.08	0.57**	0.84	1.42**	0.99	0.73*	1.1	0.73	2.15***	0.69*	0.71	1.43*	2.05**
gen_gwas_ear_eyes (1)	inf	inf	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
gen_gwas_endocrine_no (841)	1.96**	1.29	0.64**	1.19	0.91**	1.53**	1	1.07	0.88	1.12	1.13	0.9	0.58**	0.72	1	1.58*	1.11	0.95**	0.96	1.43**	0.68**	0.81	1.09	0.79	1.05	0.85	0.82	1.11	1.68**
gen_gwas_endocrine_eyes (151)	2.37**	1.94	0.78	1.19	0.97*	1.79*	2.95*	1.1	1.07	0.57	0	0	0.34**	0.98	0.29**	1.48	1.06	0.67*	0.67	1.01	2.70**	0.89	1.04	1.15	4.20**	0.44*	0.56	1.91*	1.64**
gen_gwas_eye_no (903)	1.84**	1.13	0.71**	1.21*	0.64**	1.29*	2.08	1.09	0.84	1.34	0.97	0.48	0.87	0.42	0.38	0.79	1.69**	1.04	0.88	1.06	1.48**	0.98	1.14	0.98	0.99	0.96	0.79	0.67	1.94*
gen_gwas_eye_eyes (89)	0.63**	2.01*	0.24**	1.02	0.8	1.31	4.65**	1.05	1.43	1.28	1.05	2.42	1.43	3.46**	0.96	1.51	1.31	0.87	1.1	0.86	1.12	0.81	1.71*	0.82	0.44	0.62	1.28	1.82	1.34
gen_gwas_gi_no (860)	2.16**	0.98	0.8	1.27*	0.51**	1.73**	0.77	1.05	0.67*	0.92	1.05	0.67	1.1	0.42**	0.71*	1.77**	0.91	0.53**	0.8	1.69**	1.06	0.74*	1.25*	0.66	2.62**	0.57**	0.66**	1.4*	1.79**
gen_gwas_gi_eyes (132)	1.83**	2.64**	0.16**	0.86	0.84	1.21	4.68**	1.18	2.28**	0.69	0.76	0.85	0.81	2.67**	1.15	1.22	1.89	1.34	1.27	0.48**	0.78	0.97	0.61*	1.85	0.38	1.81*	1.92**	1.03	1.43
gen_gwas_heart_no (823)	1.36**	1.37**	0.58**	1.17	0.84	1.56**	1.04	1.13	0.76	0.86	1.06	0.55	1.21	0.72	0.86	1.69**	1.22	0.72**	0.86	1.27*	1.04	0.71**	0.96	0.94	1.25	0.7*	0.82	1.53**	1.94**
gen_gwas_heart_eyes (169)	6.29**	0.96	1.02	1.12	0.21**	1.58*	2.59*	0.92	1.56	0.93	0.79	1.4	0.6	0.69	0.5	1.19	0.78	0.45**	1.02	1.38	0.87	1.07	1.5*	0.65	2.98**	0.98	0.59	0.77	1.08
gen_gwas_hepatobiliary_no (924)	2.10**	1.08	0.71*	1.31*	0.66**	1.68**	1.76	1.13	0.9	0.82	1.04	0.58	0.67	0.71	1.72*	1.65**	1.19	0.59**	0.62	1.34*	0.91	0.72**	1.04	0.85	2.03**	0.72*	0.71*	1.4*	2.09**
gen_gwas_hepatobiliary_eyes (903)	0.54**	0.57**	0.29**	0.83	0.54**	0.97	1.74	0.67	0.8	1.18	0.62	0.71	0.87	0.42	0.38	0.79	1.69**	1.04	0.88	1.06	1.48**	0.98	1.14	0.98	0.99	0.96	0.79	0.67	1.94*
gen_gwas_immune_no (720)	2.04**	0.93	1.03	1.14	0.56**	1.65**	0.86	0.93	0.63*	1.14	1.15	0.69	1	0.41**	0.51**	0.99*	0.91	0.41**	0.69*	1.62**	1.02	0.66**	1.2	0.91	2.37**	0.57**	0.49**	1.25	2.23**
gen_gwas_immune_eyes (272)	1.97**	1.86**	0.18**	1.14	0.99	1.28	2.6*	1.35*	1.71**	0.47*	0.72	0.81	1.08	1.74*	1.69**	0.81	1.45	1.61**	1.42	0.9	0.95	1.16	0.86	0.81	0.66	1.33	1.64**	1.25	0.8
gen_gwas_infection_no (935)	2.54**	1.37**	0.82**	1.13	0.59**	1.77**	1.71	1.12	0.87	0.83	0.96	0.73	1.15	0.69	0.77*	2.0**	1.16	0.55**	0.71*	1.3*	1.1	0.72**	1.05	0.83	2.03**	0.75*	0.71*	1.4*	2.05**
gen_gwas_infection_eyes (57)	1.22	0.68	0.68	1.82*	0.89	1.23	0.81	0.86	1.27	1.12	1.25	0	0.3*	0.51	0.74	0.37	0.79	1.49	3.02**	1.92*	0.32**	1.17	1.47	1	1.54	0.55	1.03	1.11	0.85
gen_gwas_mental_no (744)	1.56**	1.37**	0.48**	1.28*	0.95	1.11	1.53	1.28*	1.24	1.04	1.15	0.87	1.4*	0.75	0.87	1.74**	1.61*	0.77*	0.96	0.66**	1.28*	0.83	0.81*	0.75	1.84**	0.93	0.96	1.48*	1.69**
gen_gwas_mental_eyes (278)	2.79**	0.92	1.37	0.94	0.25**	2.91**	1.18	0.74	0.42**	0.59	0.7	0.42	0.47**	0.7*	0.68	1.08	1.78	0.48**	0.78	3.63**	0.37**	0.74	1.63**	1.18	1.31	0.47**	0.44*	0.9	1.39*
gen_gwas_metabolism_no (674)	2.10**	0.96	0.15**	1.21*	0.65**	1.62**	1.12	1.05	0.76	0.83	1	0.53	1.28	0.6*	0.58**	0.23**	0.96	0.52**	0.81	1.35*	1.17	0.68**	1.03	1.04	1.48*	0.59**	0.66**	1.25	2.45**
gen_gwas_metabolism_eyes (318)	1.48**	2.1**	1.12	0.55**	0.88	1.31*	3.04**	1.09	0.82	0.81	1.18	0.31	2.03**	1.42	0.54**	2.28**	1.09	0.87	0.76	0.62**	2.7**	0.74	0.83	0.87	1.51	0.67	0.7	1.25	2.15**
gen_gwas_musculoskeletal_no (833)	2.16**	1.07	0.67*	1.32*	0.51**	1.74**	1.55	1.05	0.99	0.98	1.12	0.54	0.68*	0.7	0.81	2.01**	0.92	0.53**	0.8	1.60**	0.79*	0.79*	1.36**	0.92	1.18	0.74	0.82	1.16	2.21**
gen_gwas_musculoskeletal_eyes (159)	1.64*	1.77**	0.65	0.78	1.39	1.15	1.2	1.11	0.72	0.56	0.61	1.5	2.78**	0.74	0.74	0.68	1.71	1.21	0.85	0.59*	2.3**	0.89	0.48**	0.69	3.4**	0.81	0.58	1.68*	0.71
gen_gwas_neoplasm_no (835)	1.54**	1.02	0.77*	1.24*	0.65**	1.62**	1.12	1.05	0.76	0.83	1	0.53	1.28	0.6*	0.58**	0.23**	0.96	0.52**	0.81	1.35*	1.17	0.68**	1.03	1.04	1.48*	0.59**	0.66**	1.25	2.45**
gen_gwas_neoplasm_eyes (157)	4.7**	2.04**	0.32**	0.95	0.37	1.43	2.38	1.14	1.62	1.01	0.95	1.52	0.48**	1.2	1.88	0.34**	1.55	1.28	1.2	1.18	0.66**	1.34	1.24	3.24**	1.52	1.23	1.42	0.47**	
gen_gwas_nervous_no (756)	1.31**	1.1	0.8	1.12	0.61**	1.45**	0.98	1.05	0.84	1.01	1.11	0.47	1.24	0.47**	0.65**	1.44*	0.67	0.57**	0.79	1.51**	1.3*	0.58**	0.92	0.54	2.00**	0.57**	0.61**	0.75	1.57**
gen_gwas_nervous_eyes (236)	4.11**	1.48	0.97	1.21*	0.64**	1.29*	1.92	1.13	0.82	0.74	1.21	0.85	0.82	0.52	0.52	0.38	1.17	0.28	0.38	1.17	0.28	0.38	1.47	0.75	2.01**	0.63**	0.61**	1.13	2.09**
gen_gwas_pregnancy_no (957)	0.27**	1.43**	0.16**	1.06	0.64**	1.27*	1.82	1.08	0.86	0.91	1.01	0.7	1.08	0.47*	0.8	1.47**	1.17	0.62**	0.86	1.45**	1.05	0.79*	1.11	0.9	2.37**	0.74*	0.71*	1.47*	1.77**
gen_gwas_pregnancy_eyes (35)	0.82	0.14**	0.57**	1.79**	0.18**	2.47*	0	1.26	2.3*	0	0.64	0	0.51	1.35	1.16	3.32**	0.63	0.32*	0.58	0.84	0.43	0.56	0.93	0	0.37	0.44	0.95	0.56	3.91**
gen_gwas_repro_no (871)	1.9**	1.16	0.63**	1.28*	0.6**	1.73**	0.84	1.16	0.76	1	1.02	0.85	0.83	0.64*	0.78	2.35**	1.21	0.54**	0.85	1.51**	0.87	0.74*	1.31*	0.74	0.86	0.72*	0.67**	1.63*	2.21**
gen_gwas_repro_eyes (121)	3.86**	1.55*	0.77	0.8	0.91	1.22	4.46**	0.78	1.75*	0.36	0.84	0	0.12**	1.01	0.82	0.16**	0.73	1.22	1.07	0.79	1.6*	0.45**	1.46	7.13**	0.88	1.22	0.47	0.65	
gen_gwas_respiratory_no (932)	2.12**	1.08	0.61**	1.4**	0.58**	1.78**	1.72	1.12	0.8	0.8	1.02	0.57	1.1	0.62*	0.68**	1.85**	0.81	0.52**	0.77	1.47**	1.06	0.79*	1.14	0.84	2.21**	0.59**	0.6**	1.34	2.16**
gen_gwas_respiratory_eyes (60)	4.41**	0.79**	0.83	0.23**	1.11	1.28	0.77	0.87	2.0*	1.36	0.75	1.97	0.81	1.31	1.18	0.89	4.19**	1.98*	2	0.81	0.59	0.57	0.81	0.84	0.92	2.55*	2.57**	1.47	0.54
gen_gwas_skin_no (933)	2.10**	1.25	0.16**	1.16	0.37**	1.79**	0.67	1.07	0.88	1.01	0.97	0.57	1.06	0.62*	0.69**	1.92**	0.89	0.62**	0.73	1.07	1.37	0.73	0.73	2.01**	0.63**	0.61**	1.13	2.09**	
gen_gwas_skin_eyes (59)	0.98	1.48	0.39**	1.24	0.28	1.16	2.24	1.34	1.33	1.81	1.21	0.41	0.85	0.82	0.52	0.37	1.11	2.37**	0.95	0.29**	0.37*	0.35	0.47*	0.73	1.81	0.68	0.38	0.38	1.94**
gen_gwas_urologic_no (878)	2.07**	1.36**	0.44**	1.12	0.61**	1.78**	1.82	1.03	0.71**	0.91	1.01	0.7	1.1	0.63*	0.66**	1.87**	1.17	0.62**	0.88	1.39**	1.02	0.71**	1.18	0.7	2.24**	0.74*	0.68**	1.1	2.02**
gen_gwas_urologic_eyes (35)	0.82	0.58	0.25*	0.70**	0.62	1.35	0	2.06*	0.54	0	0.64	0	0.33	1.35	0.65**	0.19	0.63	0.32*	0.58	1.41	0.7	1.53	0.44	3.54	0.78	0.44	1.54	0.27	0.1*
gen_gwas_vascular_no (756)	1.6**	1.39**	0.68**	1.02	0.72**	1.47**	0.62	1.07	0.82	0.9	0.94	0.32	1.06	0.59**	0.83	1.09	1.1	0.74**	0.88	1.35*	1	0.82	1.01	0.63	1.91**	0.65**	0.67**	1.18	1.18
gen_gwas_vascular_eyes (236)	0.71**	0.88	0.7	1.49**	0.68**	1.63**	3.93**	1.07	1.19	0.85	1.1	2.26	0.92	1.13	0.76	2.44**	1.07	0.52**	0.94	1.14	0.8	0.78	1.21	1.56	1.23	1.13	1.09	1.44	2.66**
gen_omim_blood_no (884)	1.9**	0.79*	1.75**	0.87	0.95**	1.55**	0.9	0.84	1.29	0.29**	0.91	2.34	1.56**	0.81	0.68**	1	0.73	0.96**	0.76	2.05**	1.77**	1.19	1.34**	1.56	2.57**	1.09	0.79	1.29	0.97
gen_omim_blood_eyes (251)	1.94**	2.55**	0.99	0.41**	2.12**	0.74*	1.14	1.64**	0.75	0.79**	2.29**	0.41	0.41**	0.82	0.84*														



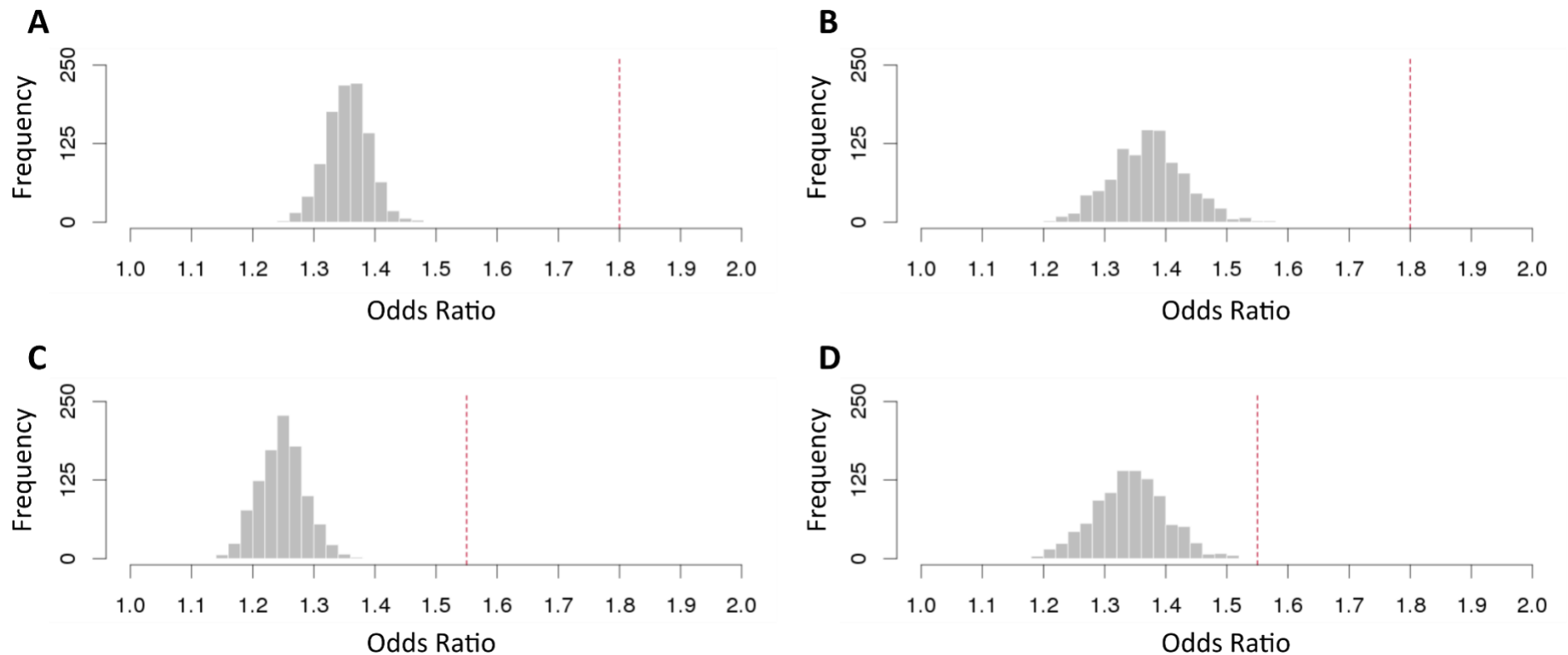
Supplementary Figure 2. All coefficients from each of the 18 side effect regression models, where “on-indication” side effects are retained in the data. Asterisks show P-value of the coefficient (*, $P < 0.05$; **, $P < 0.005$; ***, $P < 0.0005$).



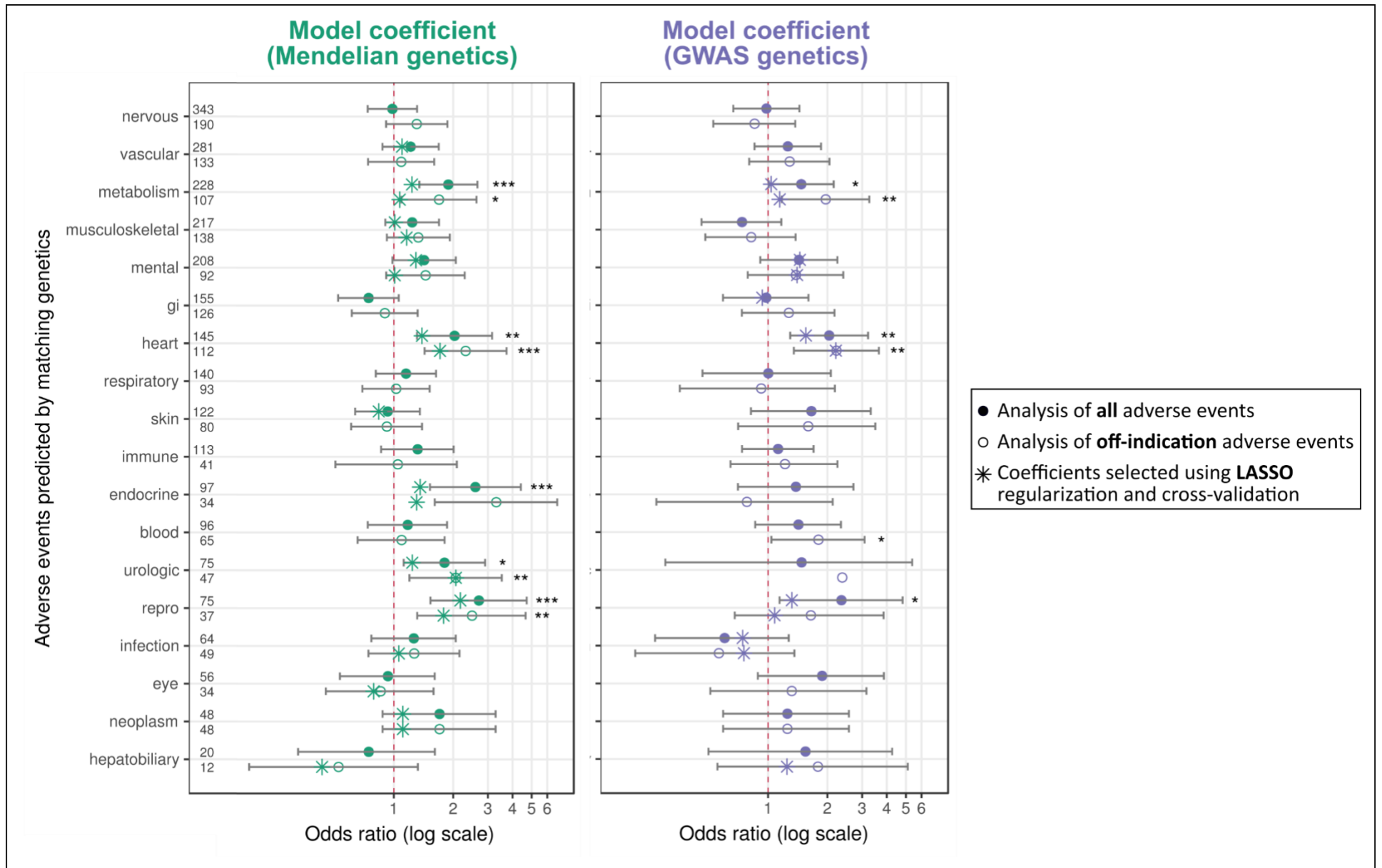
Supplementary Figure 3. All coefficients from each of the 18 side effect regression models, where only “off-indication” side effects are retained in the data. Asterisks show P-value of the coefficient (*, $P < 0.05$; **, $P < 0.005$; ***, $P < 0.0005$).



Supplementary Figure 4. Comparison of the rate at which all possible drug-SE (side effect) combinations are manifested as side effects in the clinical trial data set, between drug-SE pairs with or without genetic support. OR, odds ratio; P, P-value from Fisher's exact test (two-tailed). The top two panels show analyses based on all 38,199 drug-SE combinations; the bottom two panels show analyses based on 33,489 "off-indication" drug-SE combinations. The left two panels show analyses based on Mendelian genetic data; the right two panels show analyses based on GWAS genetic data. Error bars represent the 95% confidence interval of the reported proportions. Underlying data and confidence intervals for the OR values are shown in **Supplementary Table 1**.



Supplementary Figure 5. Result of permutation test for significance of genetics-SE enrichment relative to randomized SEs and randomized target genetics. In each analysis, gray histogram shows ORs resulting from 1,000 simulations and dashed vertical line shows the OR of the study data. Mean and standard deviation of the simulation ORs are provided in **Supplementary Table 2**. In each analysis, the simulation OR never exceeds the study data OR (Monte Carlo empirical $P < 0.001$). Analyses used: (A) all drug-SE pairs, randomized SEs; (B) all drug-SE pairs, randomized target genetics; (C) only off-indication drug-SE pairs, randomized SEs; (D) only off-indication drug-SE pairs, randomized target genetics.



Supplementary Figure 6. Summary of side effect modeling results with quantitative, tissue-specific target gene expression values included in the models. LEFT: coefficients of the Mendelian genetics predictors in each of the 18 side effect models (from **Supplementary Table 9**). RIGHT: coefficients of the GWAS genetics predictors in each of the 18 side effect models (from **Supplementary Table 9**). Coefficients from the regression models were exponentiated to obtain odds ratios. Along each row, circular points indicate odds ratios from the glm models; error bars are 95% confidence intervals from the glm models; large eight-pointed stars indicate odds ratios when the genetics predictors are selected as predictors in the glmnet models (which are built using feature selection, lasso regularization, and cross-validation). Small asterisks offset from the data indicate P-values of the the Fisher's exact test (LEFT) and P-values of the coefficients from the glm models (CENTER and RIGHT); (*, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.001$).

Data inclusion	Comparison	GS	GN	GS+GN	NS	NN	NS+NN	GS/GN	NS/NN	Odds Ratio	p	OR 95% CI	GS/GS+GN	NS/NS+NN	stderr(GS/GS+GN)	stderr(NS/NS+NN)
All drug-SE combinations, all genetics	Overall: Genetic support vs no genetic support	2512	5862	8374	5729	24096	29825	0.43	0.24	1.80	1.73E-94	1.71 1.90	30.00%	19.21%	0.50%	0.23%
	Among no genetic support: incomplete genetic info vs complete genetic info	2759	12165	14924	2970	11931	14901	0.23	0.25	0.91	0.00158	0.86 0.97	18.49%	19.93%	0.32%	0.33%
All drug-SE combinations, Mendelian genetics	Overall: Genetic support vs no genetic support	2092	4765	6857	6149	25193	31342	0.44	0.24	1.80	2.28E-82	1.70 1.91	30.51%	19.62%	0.56%	0.22%
	Among no genetic support: incomplete genetic info vs complete genetic info	2014	8893	10907	4135	16300	20435	0.23	0.25	0.89	0.00017	0.84 0.95	18.47%	20.23%	0.37%	0.28%
All drug-SE combinations, GWAS genetics	Overall: Genetic support vs no genetic support	962	1911	2873	7279	28047	35326	0.50	0.26	1.94	2.75E-53	1.79 2.11	33.48%	20.61%	0.88%	0.22%
	Among no genetic support: incomplete genetic info vs complete genetic info	1745	7334	9079	5534	20713	26247	0.24	0.27	0.89	0.00015	0.84 0.95	19.22%	21.08%	0.41%	0.25%
Off-indication drug-SE combinations, all genetics	Overall: Genetic support vs no genetic support	1458	5062	6520	4219	22750	26969	0.29	0.19	1.55	1.70E-36	1.45 1.66	22.36%	15.64%	0.52%	0.22%
	Among no genetic support: incomplete genetic info vs complete genetic info	2067	11595	13662	2152	11155	13307	0.18	0.19	0.92	0.01892	0.86 0.99	15.13%	16.17%	0.31%	0.32%
Off-indication drug-SE combinations, Mendelian genetics	Overall: Genetic support vs no genetic support	1180	4095	5275	4497	23717	28214	0.29	0.19	1.52	1.47E-28	1.41 1.63	22.37%	15.94%	0.57%	0.22%
	Among no genetic support: incomplete genetic info vs complete genetic info	1503	8528	10031	2994	15189	18183	0.18	0.20	0.89	0.00111	0.84 0.96	14.98%	16.47%	0.36%	0.28%
Off-indication drug-SE combinations, GWAS genetics	Overall: Genetic support vs no genetic support	506	1597	2103	5171	26215	31386	0.32	0.20	1.61	8.61E-18	1.44 1.78	24.06%	16.48%	0.93%	0.21%
	Among no genetic support: incomplete genetic info vs complete genetic info	1292	6919	8211	3879	19296	23175	0.19	0.20	0.93	0.0362	0.87 1.00	15.73%	16.74%	0.40%	0.25%

Supplementary Table 1. Comparison of the rate at which all possible drug-SE (side effect) combinations are manifested as side effects in the clinical trial data set, between drug-SE pairs with or without genetic support. OR, odds ratio; p, p-value from Fisher's exact test (two-tailed). GS = genetics + side effect; GN = genetics + no side effect; NS = no genetics + side effect; NN = no genetics + no side effect. For the overall comparison, "genetics" indicates a match between the side effect and genetic phenotype. For the "among no genetic support" comparison, "genetics" indicates that there is a genetic phenotype for the targets that mismatches; "no genetics" indicates there is no or incomplete genetic information for the target gene(s).

Phenotype category	Main analysis														Off-indication analysis																	
	(Drug x organ system) combinations	Drugs with genetics and side effect	Drugs with genetics and no side effect	Drugs with no genetics and side effect	Drugs with no genetics and no side effect	Rate of side effect when genetic association exists (precision) (PPV)	Rate of side effect when genetic association does not exist (FOR)	TPR (recall)	TNR	NPV	LR+	iLR-	OR	95% CI	p	(Drug x organ system) combinations	Drugs with genetics and side effect	Drugs with genetics and no side effect	Drugs with no genetics and side effect	Drugs with no genetics and no side effect	Rate of side effect when genetic association exists (precision) (PPV)	Rate of side effect when genetic association does not exist (FOR)	TPR (recall)	TNR	NPV	LR+	iLR-	OR	95% CI	p		
blood	96	269	234	1220	26.3%	16.1%	29.1%	81.9%	83.9%	1.61	1.16	1.86	1.40	2.46	1.34E-05	65	234	212	1193	21.7%	15.1%	23.5%	83.6%	84.9%	1.43	1.09	1.56	1.13	2.15	0.005724		
heart	145	176	451	1047	45.2%	30.1%	24.3%	85.6%	69.9%	1.69	1.13	1.91	1.48	2.46	3.92E-07	112	162	318	992	40.9%	24.3%	26.0%	86.0%	75.7%	1.86	1.16	2.16	1.63	2.85	5.90E-08		
congenital	17	572	22	1208	2.9%	1.8%	43.6%	67.9%	98.2%	1.36	1.20	1.63	0.81	3.25	0.165102913	9	507	11	1135	1.7%	1.0%	45.0%	69.1%	99.0%	1.46	1.26	1.83	0.67	4.89	0.222399		
ear	5	112	74	1628	4.3%	4.3%	6.3%	93.6%	95.7%	0.98	1.00	0.98	0.30	2.47	1	4	112	65	1622	3.4%	3.9%	5.8%	93.5%	96.1%	0.90	0.99	0.89	0.23	2.46	1		
endocrine	97	252	177	1293	27.8%	12.0%	35.4%	83.7%	88.0%	2.17	1.30	2.81	2.09	3.76	3.71E-12	34	207	96	1204	14.1%	7.4%	26.2%	85.3%	92.6%	1.78	1.16	2.06	1.31	3.17	0.001412		
eye	56	256	193	1314	17.9%	12.8%	22.5%	83.7%	87.2%	1.38	1.08	1.49	1.05	2.08	0.018630562	34	250	148	1268	12.0%	10.5%	18.7%	83.5%	89.5%	1.13	1.03	1.17	0.76	1.75	0.4617		
gl	155	276	497	891	36.0%	35.8%	23.8%	76.3%	64.2%	1.01	1.00	1.01	0.80	1.27	0.95420768	126	244	379	823	34.1%	31.5%	25.0%	77.1%	68.5%	1.09	1.03	1.12	0.87	1.44	0.373119		
hepatobiliary	20	153	160	1486	11.6%	9.7%	11.1%	90.7%	90.3%	1.19	1.02	1.21	0.70	2.01	0.42299002	12	150	134	1443	7.4%	8.5%	8.2%	90.6%	91.5%	0.87	0.99	0.86	0.42	1.60	0.766027		
immune	113	335	251	1120	25.2%	18.3%	31.0%	77.0%	81.7%	1.35	1.12	1.51	1.16	1.95	0.001757385	41	232	116	914	15.0%	11.3%	26.1%	79.8%	88.7%	1.29	1.08	1.39	0.92	2.07	0.09472		
infection	64	104	536	1115	38.1%	32.5%	10.7%	91.5%	67.5%	1.25	1.02	1.28	0.91	1.80	0.143708732	49	89	455	1056	35.5%	30.1%	9.7%	92.2%	69.9%	1.25	1.02	1.28	0.87	1.87	0.20933		
metabolism	228	400	280	911	36.3%	23.5%	44.9%	69.5%	76.5%	1.47	1.26	1.85	1.49	2.30	1.34E-08	107	292	185	820	26.8%	18.4%	36.6%	73.7%	81.6%	1.40	1.16	1.62	1.22	2.15	0.0006		
musculoskeletal	217	435	322	845	33.3%	27.6%	40.3%	66.0%	72.4%	1.18	1.11	1.31	1.06	1.62	0.01184709	138	373	221	749	27.0%	22.8%	38.4%	66.8%	77.2%	1.16	1.08	1.25	0.97	1.61	0.07435		
neoplasm	48	305	122	1344	13.6%	8.3%	28.2%	81.5%	91.7%	1.53	1.14	1.73	1.19	2.50	0.003098258	48	305	122	1344	13.6%	8.3%	28.2%	81.5%	91.7%	1.53	1.14	1.73	1.19	2.50	0.003098		
nervous	343	421	401	654	44.9%	38.0%	46.1%	60.8%	62.0%	1.18	1.13	1.33	1.09	1.61	0.003732245	190	312	266	558	37.8%	32.3%	41.7%	64.1%	67.7%	1.16	1.10	1.28	1.01	1.62	0.042694		
pregnancy	7	268	35	1509	2.5%	2.3%	16.7%	84.9%	97.7%	1.11	1.02	1.13	0.42	2.61	0.826676608	7	262	32	1486	2.6%	2.1%	17.9%	85.0%	97.9%	1.20	1.04	1.24	0.46	2.90	0.648859		
mental	208	350	255	1006	37.3%	20.2%	44.9%	74.2%	79.8%	1.74	1.35	2.34	1.87	2.94	5.16E-14	92	278	177	912	24.9%	16.3%	34.2%	76.6%	83.7%	1.46	1.16	1.71	1.27	2.29	0.000342		
urologic	75	245	220	1279	23.4%	14.7%	25.4%	83.9%	85.3%	1.58	1.13	1.78	1.30	2.41	0.000219979	47	215	162	1220	17.9%	11.7%	22.5%	85.0%	88.3%	1.50	1.10	1.65	1.13	2.37	0.008259		
repro	75	199	186	1359	27.4%	12.0%	28.7%	87.2%	88.0%	2.25	1.22	2.75	2.00	3.77	5.75E-10	37	185	148	1318	16.7%	10.1%	20.0%	87.7%	89.9%	1.62	1.10	1.78	1.17	2.66	0.005417		
respiratory	140	211	464	1004	39.9%	31.6%	23.2%	82.6%	68.4%	1.33	1.08	1.44	1.12	1.84	0.003676008	93	197	361	950	32.1%	27.5%	20.5%	82.8%	72.5%	1.19	1.04	1.24	0.93	1.65	0.130584		
skin	122	224	413	1060	35.3%	28.0%	22.8%	82.6%	72.0%	1.31	1.07	1.40	1.08	1.80	0.00874946	80	211	307	1004	27.5%	23.4%	20.7%	82.6%	76.6%	1.19	1.04	1.24	0.92	1.66	0.150424		
vascular	281	299	436	803	48.4%	35.2%	39.2%	72.9%	64.8%	1.44	1.20	1.73	1.41	2.13	8.31E-08	133	245	304	739	35.2%	29.1%	30.4%	75.1%	70.9%	1.22	1.08	1.32	1.02	1.71	0.031836		
TOTAL	38199	2512	5862	5729	24096	30.0%	19.2%	30.5%	80.4%	80.8%	1.56	1.16	1.80	1.71	1.90	1.73E-94	33489	1458	5062	4219	22750	22.4%	15.6%	25.7%	81.8%	84.4%	1.41	1.10	1.55	1.45	1.66	1.70E-36
Main analysis with randomized data categories																																
Random genetics (mean of 1000 randomizations)	38199	2623	7615	5618	22343	25.6%	20.1%	31.8%	74.6%	79.9%	1.25	1.09	1.37	1.30	1.45	1.99687E-11	33489	1762	6987	3915	20825	20.1%	15.8%	31.0%	74.9%	84.2%	1.24	1.09	1.34	1.26	1.43	2.78E-09
Standard deviation of random genetics	121	334	121	334	0.58%	0.22%	1.46%	1.12%	0.22%	0.04	0.02	0.06	0.06	0.06	6.11543E-10	85	317	85	317	0.5%	0.2%	1.5%	1.1%	0.2%	0.04	0.01	0.06	0.06	0.06	6.41E-08		
Random side effects (mean of 1000 randomizations)	38199	2528	5846	7213	22612	30.2%	24.2%	26.0%	79.5%	75.8%	1.26	1.07	1.36	1.28	1.43	3.06273E-18	33489	1705	4815	5964	21005	26.2%	22.1%	22.2%	81.4%	77.9%	1.19	1.05	1.25	1.17	1.33	2.53E-06
Standard deviation of random side effects	47	47	109	109	0.6%	0.4%	0.4%	0.1%	0.37%	0.02	0.01	0.03	0.03	0.04	8.94689E-17	37	37	100	100	0.6%	0.4%	0.4%	0.1%	0.4%	0.03	0.01	0.04	0.04	0.04	6.96E-05		
Main analysis using target annotations from single sources																																
DrugBank	15981	1402	2123	3788	8668	39.8%	30.4%	27.0%	80.3%	69.6%	1.37	1.10	1.51	1.40	1.63	4.28442E-25	13438	755	1861	2694	8128	28.9%	24.9%	21.9%	81.4%	75.1%	1.17	1.04	1.22	1.11	1.35	3.84E-05
Citeline Pharmaprojects	34944	1923	4480	5588	22953	30.0%	19.6%	25.6%	83.7%	80.4%	1.57	1.12	1.76	1.66	1.87	3.25909E-71	30661	1073	3797	4108	21683	22.0%	15.9%	20.7%	85.1%	84.1%	1.39	1.07	1.49	1.38	1.61	3.83E-24
Santos et al (2017)	14133	1395	2012	3535	7191	40.9%	33.0%	28.3%	78.1%	67.0%	1.29	1.09	1.41	1.30	1.53	2.96E-17	11770	765	1791	2481	6733	29.9%	26.9%	23.6%	79.0%	73.1%	1.12	1.03	1.16	1.05	1.28	0.002911
Replication analysis using OFFSIDES data																																
TOTAL	5523	1230	326	2940	1027	79.0%	74.1%	29.5%	75.9%	25.9%	1.22	1.08	1.32	1.14	1.52	0.000112548	4998	1003	317	2669	1009	76.0%	72.6%	27.3%	76.1%	27.4%	1.14	1.05	1.20	1.03	1.39	0.016452
Random genetics (mean of 1000 randomizations)	5523	1632	523	2538	830	75.7%	75.4%	39.1%	61.3%	24.6%	1.01	1.01	1.02	0.90	1.16	0.472879439	4998	1425	515	2247	811	73.5%	73.5%	38.8%	61.2%	26.5%	1.00	1.00	1.00	0.88	1.14	0.483419
Standard deviation of random genetics	85	35	85	35	0.8%	0.8%	0.5%	2.0%	2.6%	0.5%	0.04	0.03	0.07	0.06	0.08	0.301098261	79	34	79	34	0.9%	0.5%	2.1%	2.6%	0.5%	0.04	0.03	0.07	0.06	0.08	0.29747	
Analysis of placebo contribution to side effects using AACT data																																
Placebo-controlled side effects	4809	291	943	460	3115	23.6%	12.9%	38.7%	76.8%	87.1%	1.67	1.25	2.09	1.77	2.47	6.61954E-18	4275	198	798	379	2900	19.9%	11.6%	34.3%	78.4%	88.4%	1.59	1.19	1.90	1.56	2.30	1.11E-10
All placebo group side effects	4746	529	690	1151	2376	43.4%	32.6%	31.5%	77.5%	67.4%	1.40	1.13	1.58	1.38	1.81	2.32827E-11	4041	330	590	910	2211	35.9%	29.2%	26.6%	78.9%	70.8%	1.26	1.08	1.36	1.16	1.59	0.00013
All drug group side effects	4809	604	630	1232	2343	48.9%	34.5%	32.9%	78.8%	65.5%	1.55	1.17	1.82	1.60	2.08	4.12156E-19	4091	386	544	966	2195	41.5%	30.6%	28.6%	80.1%	69.4%	1.44	1.12	1.61	1.38	1.88	7.35E-10

Supplementary Table 2. Analysis of enrichment between genetic phenotypes and side effects in individual organ systems. P value is from Fisher's exact test (two-sided.) PPV, positive predictive value; FOR, false omission rate; TPR, true positive rate; TNR, true negative rate; SEN, sensitivity; SPE, specificity; PPV, positive predictive value; NPV, negative predictive value; LR+, likelihood ratio positive (SEN/1-SPE); iLR-, inverse likelihood ratio negative (SPE/1-SEN)

side effect modeled	AUC.real.genetics	AUC.no.genetics	median.AUC.random.genetics	CI95.AUC.random.genetics	p
heart	0.703	0.684	0.682	0.675-0.695	< 0.001
metabolism	0.706	0.701	0.691	0.687-0.7	< 0.001
repro	0.731	0.72	0.712	0.706-0.725	0.006
endocrine	0.704	0.678	0.671	0.658-0.695	0.007
urologic	0.739	0.732	0.723	0.716-0.736	0.007
mental	0.712	0.716	0.702	0.697-0.714	0.037

Supplementary Table 4. Results of cross validation analysis of six off-indication side effect models where genetics was significant in any of the regression models. Distribution of permutation values are shown in **Figure 4** and are reflected in 95% CI values. The p values are derived from the frequency of permutation runs that exceed the AUC of the true model.

Adverse Event	# drugs	% drugs	UMLS ID
headache	1446	54.9	C0018681
nausea	1209	45.9	C0027497
diarrhea	959	36.4	C0011991
emesis	708	26.9	C0042963
dizziness	617	23.4	C0012833,C0042571
fatigue	607	23.1	C0015672
pain	606	23.0	C0030193
rhinopharyngitis	531	20.2	C0027441
upper respiratory tract infection	498	18.9	C0041912
constipation	462	17.5	C0009806
vertigo	440	16.7	C0042571
skin rash	422	16.0	C0015230
abdominal pain	410	15.6	C0000737
gastrointestinal disease	388	14.7	C0017178
fever	386	14.7	C0015967
back pain	381	14.5	C0004604
pruritus	381	14.5	C0033774
drowsiness	368	14.0	C2830004
death	353	13.4	C1306577
insomnia	339	12.9	C0917801
cough	332	12.6	C0010200
dyspepsia	327	12.4	C0013395
arthralgia	322	12.2	C0003862
infectious disease	316	12.0	C0009450
urinary tract infection	307	11.7	C0042029
hypertension	304	11.5	C0020538
edema	285	10.8	C0013604

Supplementary Table 5. Common side effects removed from analysis.

enteral	parenteral	topical	other
buccal nasogastric oral oro-pharyngeal rectal sublingual	epidural infiltration intraarterial intraarterial (hepatic artery) intraarticular intracoronary intracutaneous intra-dermal intralymphatic intramuscular intramuscular (cremaster) intramuscular (deltoid) intramuscular (gluteal) intraspinal intrathecal intrathecal, lumbar intravenous intravenous (portal) parenteral subcutaneous subcutaneous (abdomen) subcutaneous (arm) subcutaneous (thigh) transarterial transdermal	dental inhaled intranasal intranasal spray mouth rinse ophthalmic percutaneous topical topical (buccal) topical (ophthalmic) topical (otic)	instillation, bladder instillation, intravesical intrabone intra-bronchial intracameral intracardiac intracavernosal intracerebral intracerebroventricular intradiscal intra-gastric intra-locus coeruleus intramammary intramyocardial intranodal intraocular intraotic intrapericardial intraperitoneal intrapleural intraprostatic intra-striatal intratendinous intratracheal intratumor intratumoral intratympanic intrauterine intravaginal intraventricular intravitreous percutaneous hepatic perfusion peritumoral subconjunctiva subgingivally subretinal subtenon urethral

Supplementary Table 6. Coding of delivery route.

Organ System Abbreviation	MedDRA System Organ Class (SOC) Term	UMLS id	Comment
blood	Blood and lymphatic system disorders	C0851353	
heart	Diseases, Heart	C0018799	
congenital	Congenital, familial and genetic disorders	C0851352	
ear	Ear and labyrinth disorders	C0851354	
endocrine	System Diseases, Endocrine	C0014130	
eye	Diseases, Eye	C0015397	
gi	Diseases, Gastrointestinal	C0017178	
general	General disorders and administration site conditions	C0851362	mapped to other SOC terms
hepatobiliary	Hepatobiliary disease NOS	C0267792	
immune	Diseases, Immune System	C0021053	
infection	Diseases, Communicable	C0009450	
injury	Injury, poisoning and procedural complications	C0947733	mapped to other SOC terms
investigation	Investigation NOS	C1261322	mapped to other SOC terms
metabolism	Metabolism and nutrition disorders	C0851358	
musculoskeletal	Musculoskeletal and connective tissue disorders	C0263660	
neoplasm	Neoplasm NOS	C0027651	
nervous	Diseases, Nervous System	C0027765	
pregnancy	Pregnancy, puerperium and perinatal conditions	C0851363	
mental	Mental disorder	C0004936	
urologic	Diseases, Urologic	C0042075	
repro	Reproductive system and breast disorders	C0851366	
respiratory	Respiratory, thoracic and mediastinal disorders	C0851355	
skin	Skin and subcutaneous tissue disorders	C0178298	
social	Social circumstances	C0851364	exclude from study
procedures	Surgical and medical procedures	C1948041	exclude from study
vascular	Diseases, Vascular	C0042373	

Supplementary Table 7. MedDRA System Organ Class (SOC) terms inclusion and remapping criteria.

		This study (human x 21 organ systems)	IQ Consortium (3 species x 12 organ systems)
<u>Translatability of positive findings</u>			
Number of species-phenotype combinations with:	LR+ \leq 1	1	7
	1 < LR+ < 10	20	22
	LR+ \geq 10	0	4
Sensitivity		30%	48%
PPV		30%	43%
<u>Translatability of negative findings</u>			
Number of species-phenotype combinations with:	iLR- \leq 1	1	10
	1 < iLR- < 10	20	25
	iLR- \geq 10	0	0
Specificity		80%	84%
NPV		81%	86%

Supplementary Table 8. Comparison of summary statistics between results of this study and the results of the IQ Consortium (Monticello *et al.*, 2017). PPV, positive predictive value. NPV, negative predictive value. LR+, positive likelihood ratio. iLR-, inverse negative likelihood ratio.

Model (Side effect predicted)	Variable	Model: glm (all data used)					Model: glm (only off-indication side effects used)					Model: glmnet (all data used)		Model: glmnet (only off-indication data used)					
		beta	95% CI	OR	95% CI	p	beta	95% CI	OR	95% CI	p	beta	OR	beta	OR				
blood	gen_omim_blood_yes	0.162092	-0.3047	0.621131	1.175968	0.737347	1.861032	0.491958	0.091146	-0.42316	0.59299	1.095429	0.654972	1.80939	0.724693	NA	NA	NA	NA
heart	gen_omim_heart_yes	0.709114	0.270634	1.147718	2.03219	1.310796	3.1493	0.001505	0.838638	0.359089	1.316276	2.313214	1.432024	3.729507	0.000586	0.287803	1.333495	0.325952305	1.385349293
congenital	gen_omim_congenital_yes	0.935395	-0.26439	2.229239	2.548219	0.767676	9.292789	0.13694	2.345718	0.124561	5.306846	10.44077	1.132651	201.7131	0.063	NA	NA	NA	NA
ear	gen_omim_ear_yes	0.184921	-1.08308	1.283147	1.203124	0.338553	3.607978	0.735659	-0.01987	-1.39101	1.135715	0.980325	0.248823	3.113398	0.974913	NA	NA	NA	NA
endocrine	gen_omim_endocrine_yes	0.952153	0.421856	1.482671	2.591283	1.524789	4.404696	0.000426	1.196117	0.478165	1.907841	3.307249	1.613112	6.738521	0.000998	0.217877	1.243434	0.154896338	1.167536926
eye	gen_omim_eye_yes	-0.06875	-0.63011	0.477691	0.933563	0.532535	1.612347	0.807445	-0.15367	-0.79461	0.464039	0.857556	0.451755	1.590485	0.612354	NA	NA	NA	NA
gi	gen_omim_gi_yes	-0.29415	-0.65022	0.057828	0.745167	0.521299	1.059533	0.10319	-0.10457	-0.4915	0.27835	0.900714	0.611708	1.320948	0.594114	NA	NA	NA	NA
hepatobiliary	gen_omim_hepatobiliary_yes	-0.293	-1.1182	0.479707	0.746022	0.326867	1.615602	0.470619	-0.64547	-1.68957	0.280241	0.524418	0.1846	1.323449	0.195131	NA	NA	NA	NA
immune	gen_omim_immune_yes	0.278185	-0.14869	0.698659	1.320731	0.861839	2.010873	0.197499	0.046765	-0.68232	0.73583	1.047875	0.505443	2.087213	0.896641	NA	NA	NA	NA
infection	gen_omim_infection_yes	0.233047	-0.26189	0.722432	1.262441	0.769594	2.059435	0.352636	0.238231	-0.29628	0.766419	1.269002	0.743579	2.152046	0.378714	NA	NA	NA	NA
metabolism	gen_omim_metabolism_yes	0.636062	0.298069	0.97566	1.889027	1.347254	2.652917	0.000231	0.527928	0.08995	0.965871	1.695416	1.09412	2.627075	0.018008	0.084695	1.088385	0.011376563	1.011441523
musculoskeletal	gen_omim_musculoskeletal_yes	0.214049	-0.09963	0.527754	1.238684	0.905175	1.69512	0.180807	0.2859	-0.08112	0.653439	1.33096	0.922084	1.92214	0.126746	NA	NA	0.039245178	1.040025443
neoplasm	gen_omim_neoplasm_yes	0.533812	-0.13093	1.188392	1.705421	0.87728	3.281801	0.111976	0.533812	-0.13093	1.188392	1.705421	0.87728	3.281801	0.111976	NA	NA	NA	NA
nervous	gen_omim_nervous_yes	-0.01654	-0.3053	0.271444	0.983596	0.736905	1.311857	0.910448	0.267309	-0.09063	0.625795	1.306444	0.913354	1.869732	0.14326	NA	NA	NA	NA
pregnancy	gen_omim_pregnancy_yes	0.311179	-0.01433	1.568157	1.365033	0.362644	4.797795	0.632798	0.559077	-0.48173	1.924256	1.749058	0.622443	6.850052	0.462027	NA	NA	NA	NA
mental	gen_omim_mental_yes	0.353576	-0.01715	0.723029	1.424151	0.982991	2.060665	0.609296	0.371056	-0.08904	0.82766	1.449264	0.914806	2.729959	0.112109	0.163284	1.177371	NA	NA
urologic	gen_omim_urologic_yes	0.591618	0.115107	1.06582	1.806909	1.121993	2.903218	0.014587	0.723119	0.181844	1.260413	2.060851	1.199427	3.526876	0.00846	0.11313	1.119777	0.296989895	1.3458017
repro	gen_omim_repro_yes	0.993491	0.428062	1.551227	2.700646	1.534282	4.717257	0.000513	0.913327	0.27293	1.538904	2.492601	1.313808	4.65948	0.004574	0.575129	1.777359	0.328021306	1.38821855
respiratory	gen_omim_respiratory_yes	0.142477	-0.21035	0.493275	1.153126	0.8103	1.637671	0.427019	0.027181	-0.36721	0.41789	1.027553	0.692664	1.518753	0.891956	NA	NA	NA	NA
skin	gen_omim_skin_yes	-0.07187	-0.45144	0.303625	0.930655	0.636711	1.359761	0.708848	-0.08137	-0.49811	0.329332	0.921856	0.607676	1.590039	0.699591	NA	NA	NA	NA
vascular	gen_omim_vascular_yes	0.196397	-0.13257	0.524479	1.21701	0.875839	1.689579	0.241024	0.086637	-0.3011	0.47102	1.090501	0.740006	1.601266	0.65975	0.016293	1.016426	NA	NA
blood	gen_omim_blood_no	-0.06826	-0.41805	0.281478	0.934015	0.658332	1.325087	0.701802	-0.07765	-0.44624	0.290561	0.925292	0.640033	1.337178	0.679215	NA	NA	NA	NA
heart	gen_omim_heart_no	0.18758	-0.08777	0.464246	1.206327	0.91597	1.590814	0.182585	0.217087	-0.08664	0.522831	1.242452	0.917011	1.686795	0.162311	NA	NA	NA	NA
congenital	gen_omim_congenital_no	0.556453	-0.75215	1.875506	1.744474	0.471353	6.524119	0.400224	-0.429	-3.69902	2.711292	0.651157	0.024748	15.04871	0.786016	NA	NA	NA	NA
ear	gen_omim_ear_no	0.283705	-0.36565	0.951975	1.328042	0.693746	2.59082	0.396474	0.290216	-0.3766	0.977593	1.336716	0.686187	2.658051	0.398499	NA	NA	NA	NA
endocrine	gen_omim_endocrine_no	0.435031	0.012475	0.863699	1.545012	1.012554	2.371919	0.044802	0.488916	-0.04701	1.03844	1.630547	0.954077	2.824806	0.076558	NA	NA	NA	NA
eye	gen_omim_eye_no	0.047614	-0.35236	0.449448	1.048766	0.703028	1.567447	0.815684	0.031798	-0.42396	0.489774	1.032309	0.654445	1.631947	0.891298	NA	NA	NA	NA
gi	gen_omim_gi_no	-0.19492	-0.47221	0.081635	0.822901	0.623622	1.08506	0.167483	-0.08399	-0.38774	0.219655	0.919442	0.678591	1.245646	0.587535	NA	NA	NA	NA
hepatobiliary	gen_omim_hepatobiliary_no	-0.15745	-0.59371	0.28023	0.854317	0.552273	1.323434	0.479237	-0.01814	-0.48015	0.448807	0.982025	0.618691	1.566442	0.938856	NA	NA	NA	NA
immune	gen_omim_immune_no	0.179531	-0.16292	0.523773	1.196655	0.849661	1.688386	0.305004	0.034616	-0.44613	0.522257	1.035222	0.6401	1.685828	0.888343	NA	NA	NA	NA
infection	gen_omim_infection_no	0.035065	-0.24121	0.312199	1.035687	0.785677	1.366426	0.803738	-0.02538	-0.31735	0.267362	0.974942	0.728073	1.306513	0.864824	NA	NA	NA	NA
metabolism	gen_omim_metabolism_no	0.197369	-0.14211	0.536591	1.218193	0.867528	1.701167	0.253936	0.22929	-0.1657	0.624887	1.257707	0.847297	1.868034	0.255031	NA	NA	NA	NA
musculoskeletal	gen_omim_musculoskeletal_no	-0.08626	-0.39685	0.222583	0.917359	0.672435	1.2493	0.584903	-0.08239	-0.44194	0.274764	0.92091	0.642789	1.31622	0.651931	NA	NA	NA	NA
neoplasm	gen_omim_neoplasm_no	-0.20252	-0.69404	0.287333	0.816667	0.499554	1.332861	0.417514	-0.20252	-0.69404	0.287333	0.816667	0.499554	1.332861	0.417514	NA	NA	NA	NA
nervous	gen_omim_nervous_no	0.123526	-0.17412	0.420969	1.131479	0.8402	1.523437	0.415581	0.209155	-0.13489	0.553539	1.232635	0.873812	1.739397	0.233317	NA	NA	NA	NA
pregnancy	gen_omim_pregnancy_no	-0.21143	-1.1365	0.706452	0.809426	0.32094	2.026786	0.650078	-0.24671	-1.24838	0.746712	0.781365	0.286968	2.11005	0.624435	NA	NA	NA	NA
mental	gen_omim_mental_no	0.076743	-0.27227	0.426049	1.079764	0.761648	1.531196	0.666353	0.2366	-0.15859	0.634078	1.266934	0.85335	1.885283	0.241378	NA	NA	NA	NA
urologic	gen_omim_urologic_no	-0.06371	-0.45245	0.325373	0.938276	0.636068	1.384547	0.747837	-0.03926	-0.47496	0.39765	0.961501	0.621908	1.488233	0.859769	NA	NA	NA	NA
repro	gen_omim_repro_no	-0.18886	-0.62197	0.243867	0.827903	0.536884	1.761174	0.391827	-0.15285	-0.62701	0.32164	0.858262	0.534189	1.379388	0.526901	NA	NA	NA	NA
respiratory	gen_omim_respiratory_no	0.017435	-0.28374	0.318731	1.017588	0.752966	1.375381	0.909625	-0.0855	-0.41672	0.245148	0.918052	0.659204	1.277811	0.612323	NA	NA	NA	NA
skin	gen_omim_skin_no	-0.24749	-0.54618	0.05007	0.78076	0.579156	1.051345	0.103498	-0.30631	-0.63671	0.022203	0.73616	0.529032	1.022451	0.068186	NA	NA	NA	NA
vascular	gen_omim_vascular_no	0.017479	-0.25792	0.292845	1.017632	0.772658	1.304235	0.900945	-0.05025	-0.3606	0.259874	0.950993	0.697255	1.296766	0.750731	NA	NA	NA	NA
blood	gen_gwas_blood_yes	0.354919	-0.14976	0.850343	1.426065	0.860912	2.34045	0.163569	0.588198	0.038603	1.127301	1.80074	1.039358	3.087131	0.033845	NA	NA	NA	NA
heart	gen_gwas_heart_yes	0.713666	0.258691	1.16798	2.041461	1.295323	3.215489	0.002073	0.798489	0.300843	1.293995	2.22218	1.350997	3.647328	0.001604	0.326082	1.38553	0.411220872	1.50865854
congenital	gen_gwas_congenital_yes	1.375561	-0.3206	2.978908	3.957297	0.725716	19.66634	0.096448	-18.9322	NA	132.4311	6.00E-09	NA	3.27E+57	0.994258	NA	NA	NA	NA
ear	gen_gwas_ear_yes	-14.9689	NA	93.37754	3.16E-07	NA	3.58E+40	0.977696	-14.911	NA	93.39451	3.34E-07	NA	3.64E+40	0.977782	NA	NA	NA	NA
endocrine	gen_gwas_endocrine_yes	0.325005	-0.35164	0.996775	1.384038	0.703535	2.709529	0.344111	-0.2471	-1.30438	0.754428	0.781066	0.271339	2.126396	0.6366	NA	NA	NA	NA
eye	gen_gwas_eye_yes	0.63102	-0.11954	1.352384	1.879528	0.887313	3.866634	0.202303	0.276686	-0.67362	1.148453	1.318752	0.509861	3.15331	0.549397	NA	NA	NA	NA
gi	gen																		