

Figure S4. Power differentials (a) between MLC and MinP-M, and (b) between MLC and PC80, each plotted against power differentials between MLC and SSBw. The latter is on the x axis. Each point corresponds to one gene with power estimated in 1000 replicated datasets (nominal $\alpha=0.05$). Data were generated under simulation models 1-5, as defined in Table 2, and were analysed with (typed) and without (untyped) the causal variant(s) in the regression model.

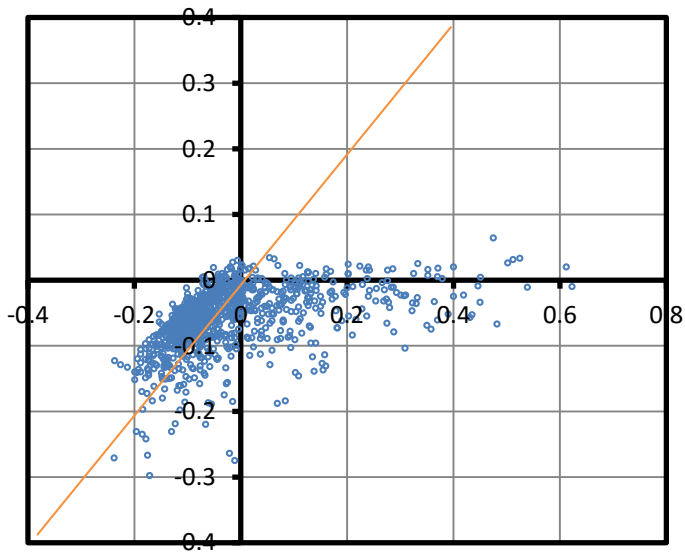
In the upper panels: MLC power is greater than MinP-M power for genes above the x axis, MLC power is greater than SSBw power for genes on the right of the y axis, and the upper right quadrant includes all genes for which MLC is more powerful than both MinP-M and SSBw. Genes close to the diagonal line have MinP-M power \approx SSBw power, while those below the line have MinP-M power $>$ SSBw power.

In the lower panels: MLC power is greater than PC80 power for genes above the x axis, MLC power is greater than SSBw power for genes on the right of the y axis, and the upper right quadrant includes all genes for which MLC is more powerful than both PC80 and SSBw. Genes close to the diagonal line have PC80 power \approx SSBw power, while those below the line have PC80 power $>$ SSBw power.

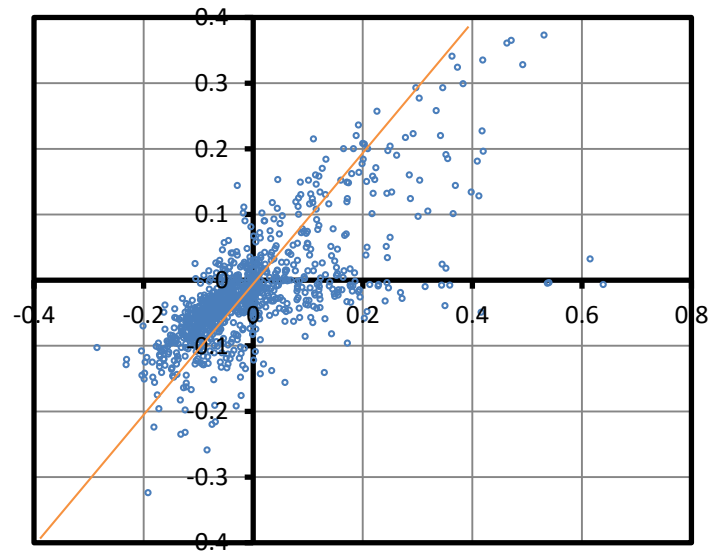
The table summarizes the number of genes according to power inequalities among the methods.

Model	# of genes	Analysis	Power Differential				
			MLC>MinPM	MLC>SSBw	MLC>PC80	MLC>Min,SSB	MLC>all 3
1	1000	Typed	80	369	260	53	32
		Untyped	258	348	349	193	132
2	993	Typed	62	51	142	27	9
		Untyped	223	279	249	170	97
3	935	Typed	372	295	278	220	136
		Untyped	312	275	296	199	109
4	993	Typed	119	167	269	89	54
		Untyped	286	354	344	236	157
5	935	Typed	359	315	244	234	130
		Untyped	313	302	267	220	108

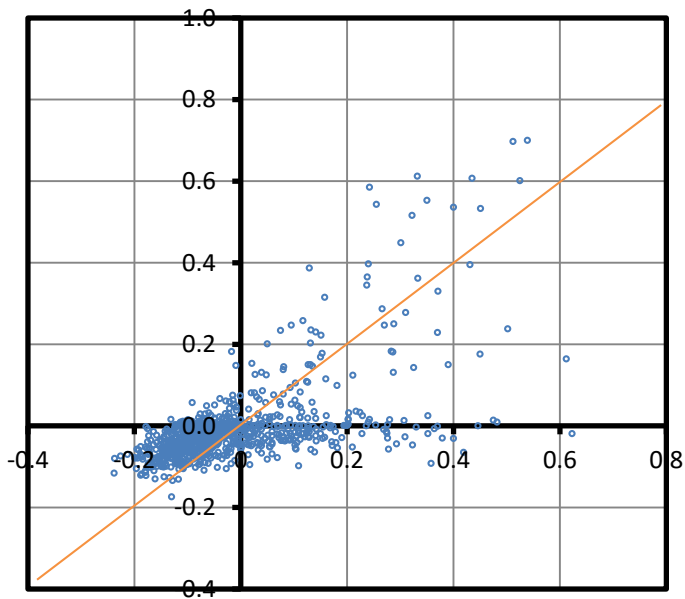
(a) Model 1: MLC-minP vs MLC-SSBw (typed)



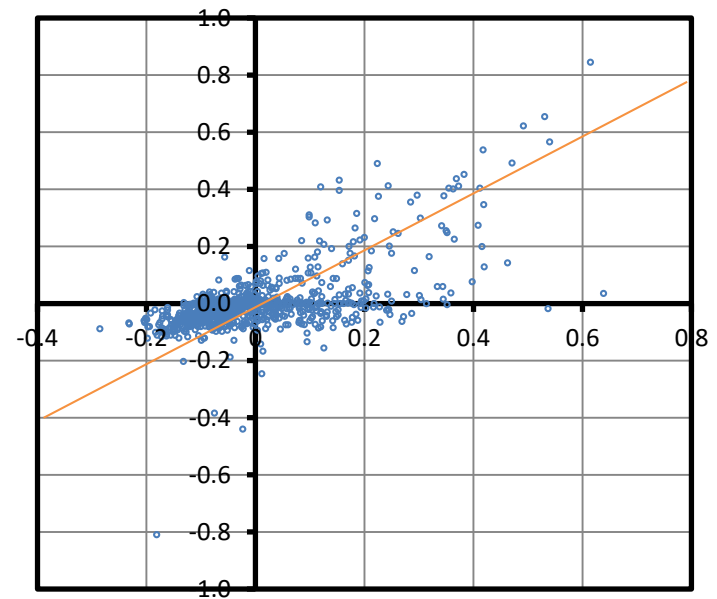
Model 1: MLC-minP vs MLC-SSBw (untyped)



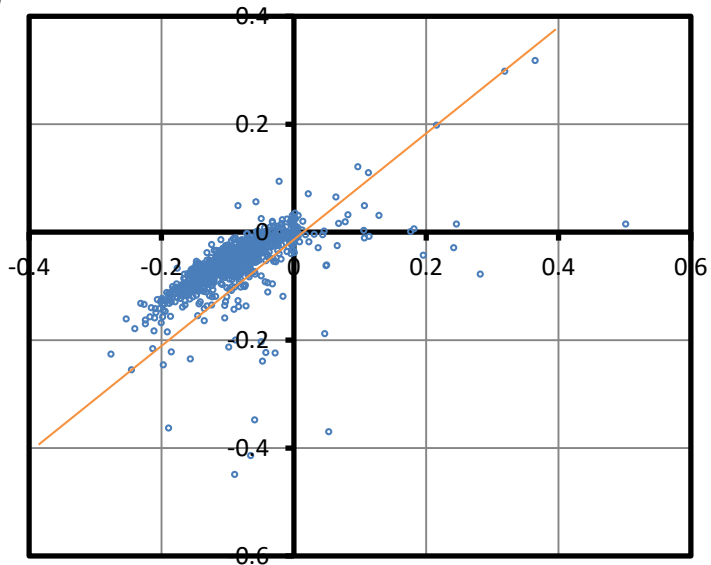
(b) Model 1: MLC-PC80 vs MLC-SSBw (typed)



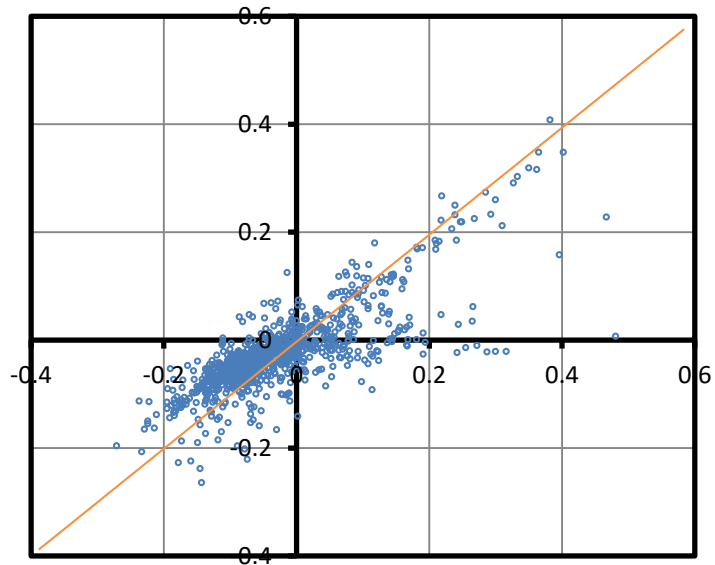
Model 1: MLC-PC80 vs MLC-SSBw (untyped)



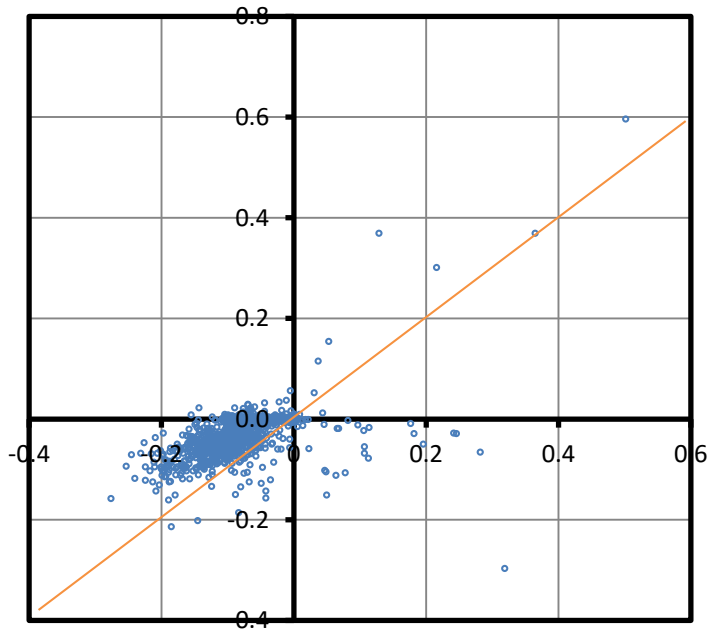
(a) **Model 2: MLC-minP vs MLC-SSBw (typed)**



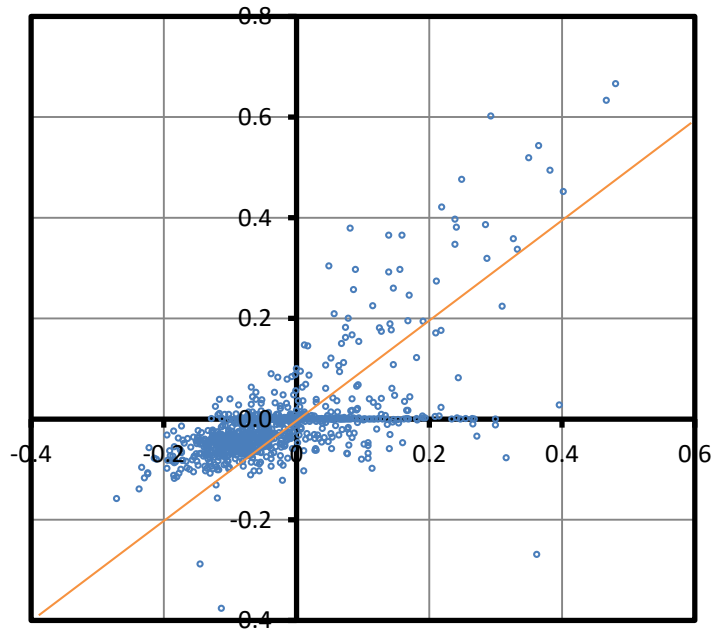
Model 2: MLC-minP vs MLC-SSBw (untyped)



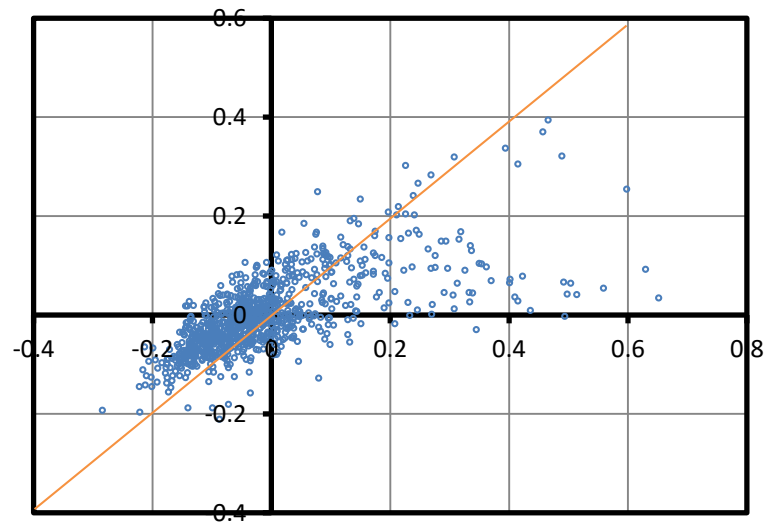
(b) **Model 2: MLC-PC80 vs MLC-SSBw (typed)**



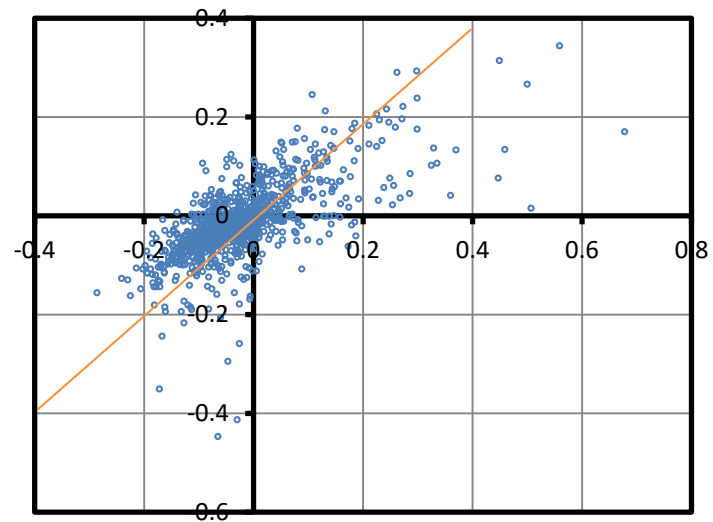
Model 2: MLC-PC80 vs MLC-SSBw (untyped)



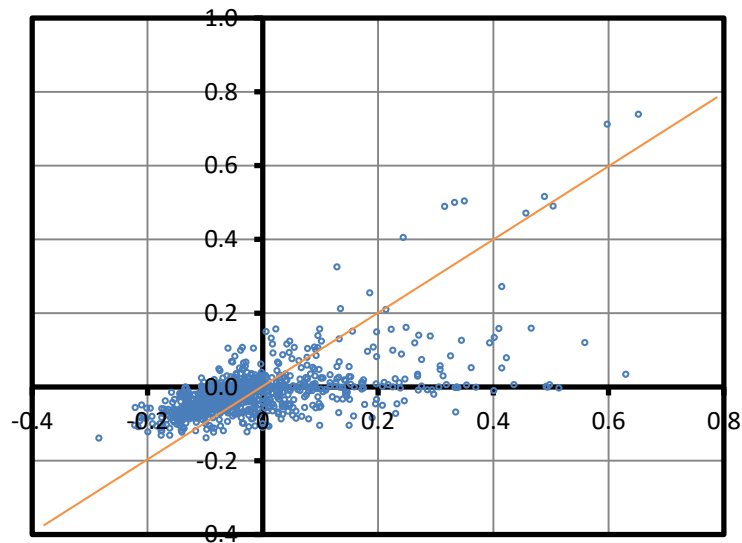
(a) Model 3: MLC-minP vs MLC-SSBw (typed)



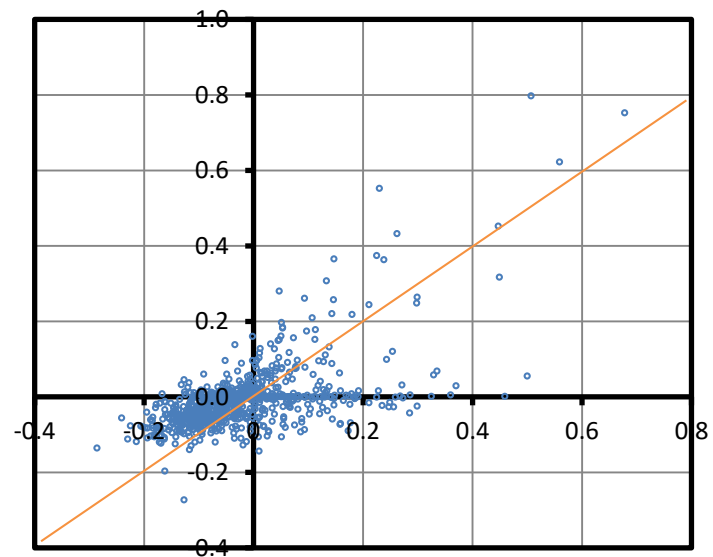
Model 3: MLC-minP vs MLC-SSBw (untyped)



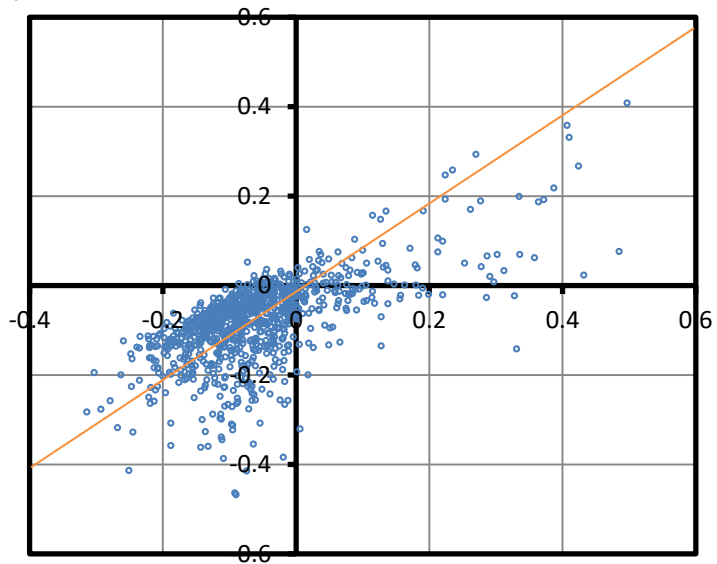
(b) Model 3: MLC-PC80 vs MLC-SSBw (typed)



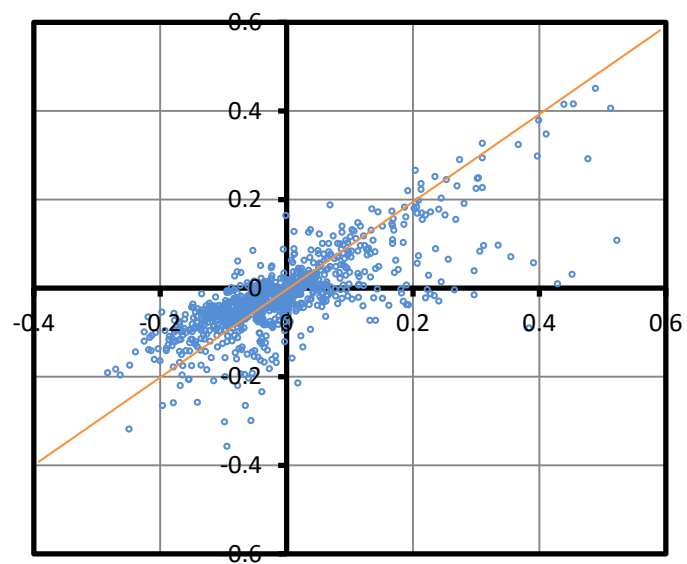
Model 3: MLC-PC80 vs MLC-SSBw (untyped)



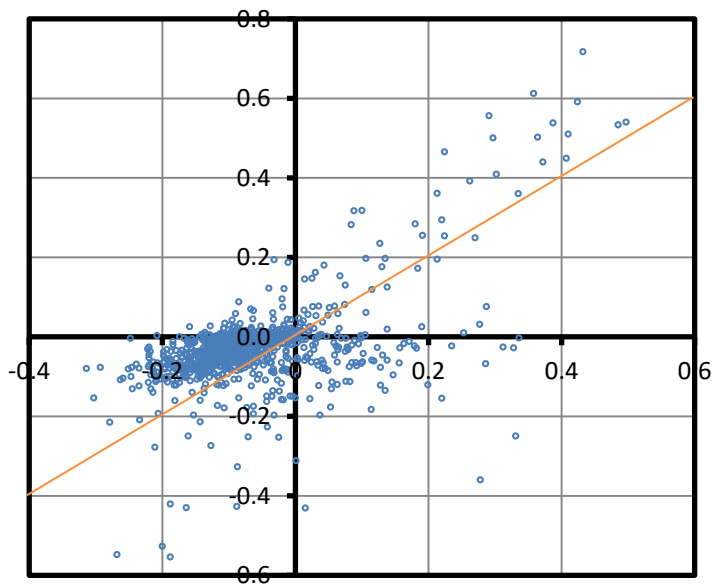
(a) Model 4: MLC-minP vs SSBw (typed)



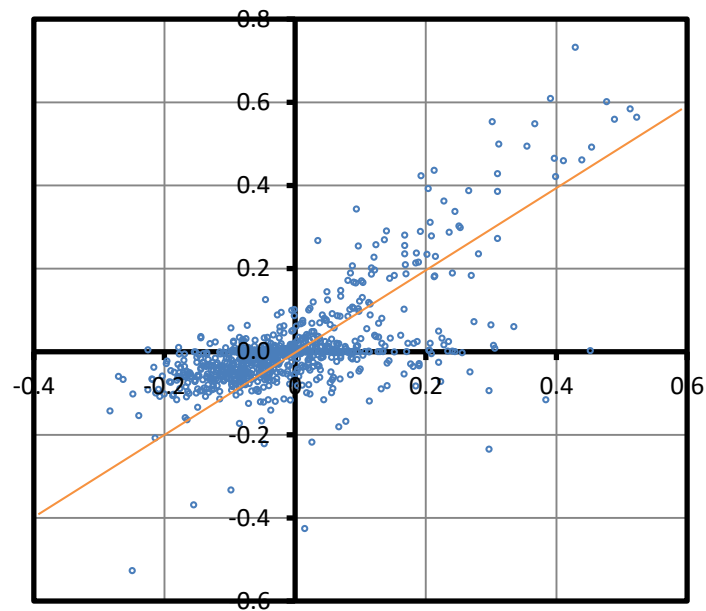
Model 4: MLC-minP vs SSBw (untyped)



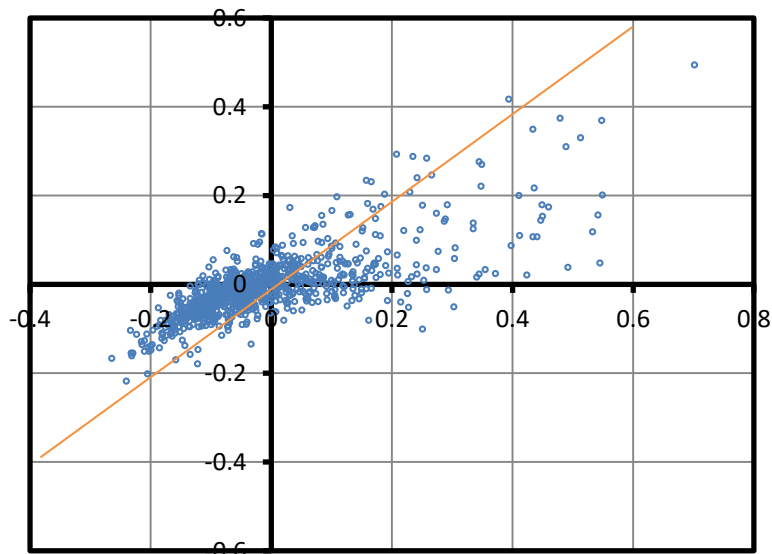
(b) Model 4: MLC-PC80 vs MLC-SSBw (typed)



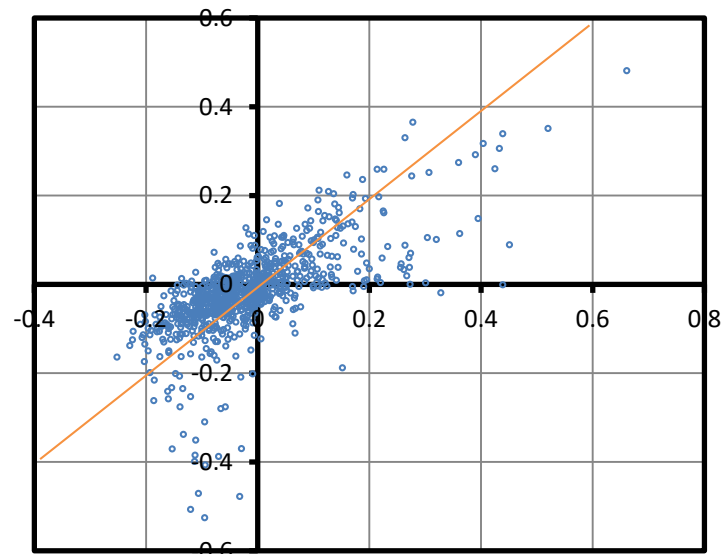
Model 4: MLC-PC80 vs MLC-SSBw (untyped)



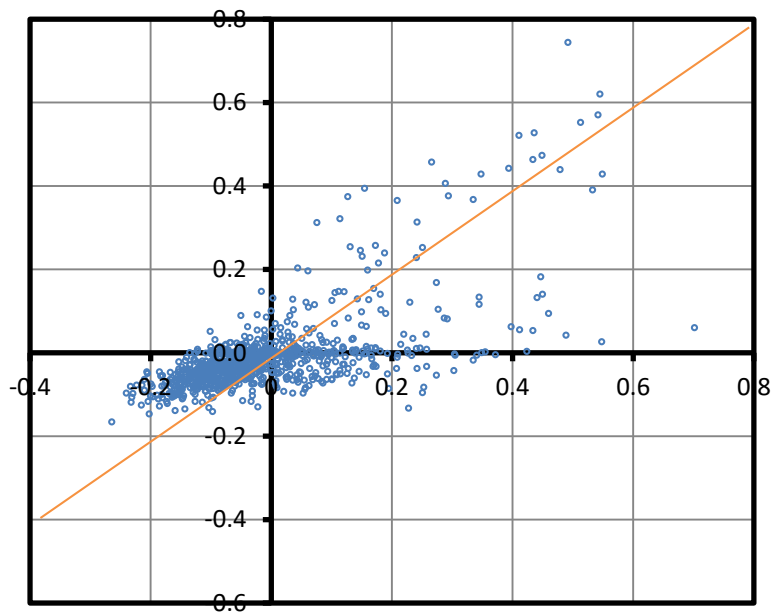
(a) Model 5: MLC-minP vs MLC-SSBw (typed)



Model 5: MLC-minP vs SSBw (untyped)



(b) Model 5: MLC-PC80 vs MLC-SSBw (typed)



Model 5: MLC-PC80 vs MLC-SSBw (untyped)

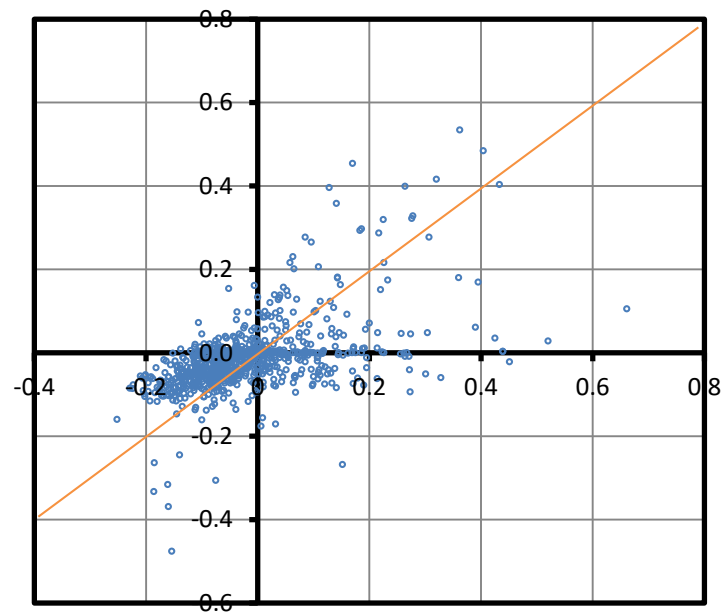


Figure S5. Power differentials (a) between MLC and MinP-M, and (b) between MLC and SKAT, each plotted against power differentials between MLC and SKATO. The latter is on the x axis. Each point corresponds to one gene with power estimated in 1000 replicated datasets (nominal alpha=0.05). Data were generated under simulation models 1-5, as defined in Table 2, and were analysed with (typed) and without (untyped) the causal variant(s) in the regression model.

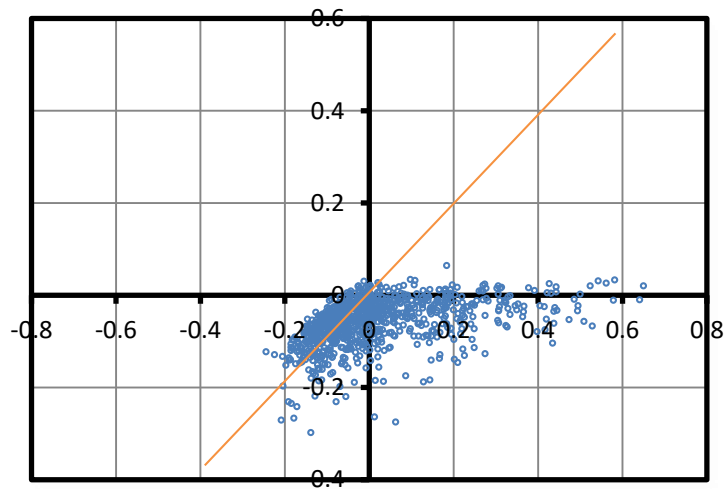
In the upper panels: MLC power is greater than MinP-M power for genes above the x axis, MLC power is greater than SKATO power for genes on the right of the y axis, and the upper right quadrant includes all genes for which MLC is more powerful than both MinP-M and SKATO. Genes close to the diagonal line have MinP-M power \approx SKATO power, while those below the line have MinP-M power $>$ SKATO power.

In the lower panels: MLC power is greater than SKAT power for genes above the x axis, MLC power is greater than SKATO power for genes on the right of the y axis, and the upper right quadrant includes all genes for which MLC is more powerful than both SKAT and SKATO). Genes close to the diagonal line have SKAT power \approx SKATO power, while those below the line have SKAT power $>$ SKATO power..

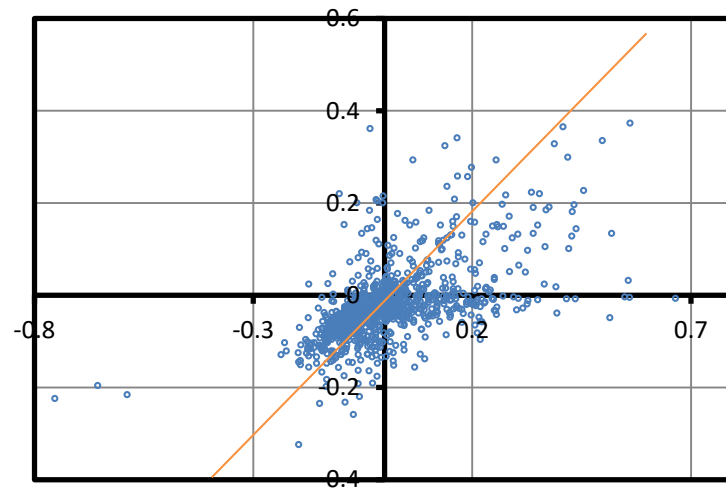
The table summarizes the number of genes according to power inequalities among the methods.

Model	# of genes	Analysis	Power Differential				
			MLC>MinPM	MLC>SKATO	MLC>SKAT	MLC>Min,SKATO	MLC>all 3
1	1000	Typed	80	380	296	62	53
		Untyped	258	438	390	195	173
2	993	Typed	62	106	67	37	25
		Untyped	223	371	314	142	125
3	935	Typed	372	284	326	193	178
		Untyped	312	370	317	198	165
4	993	Typed	119	238	179	98	88
		Untyped	286	452	395	221	203
5	935	Typed	359	458	334	294	238
		Untyped	313	416	351	230	202

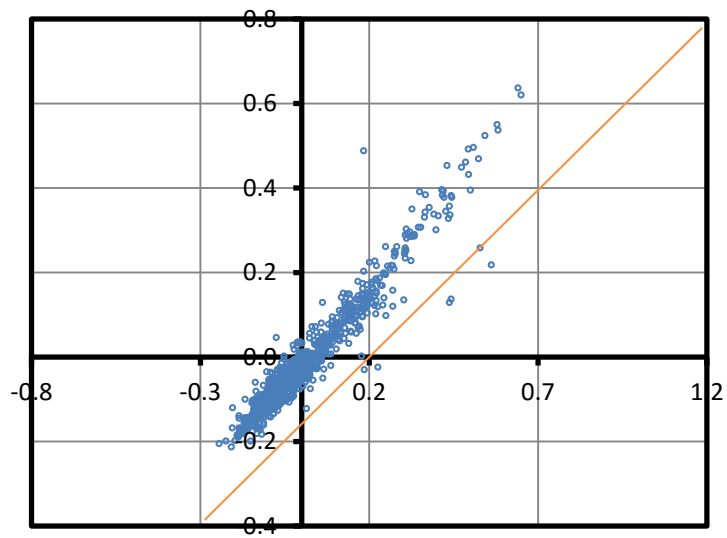
(a) Model 1: MLC-minP vs MLC-SKATO (typed)



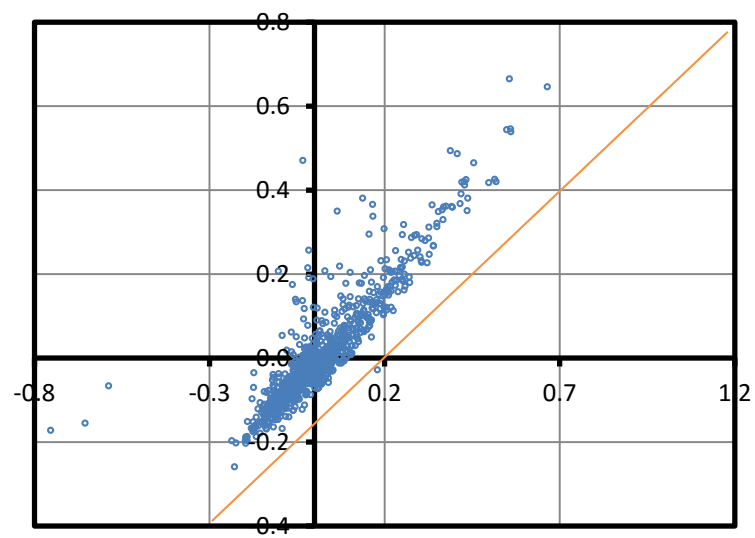
Model 1: MLC-minP vs MLC-SKATO (untyped)



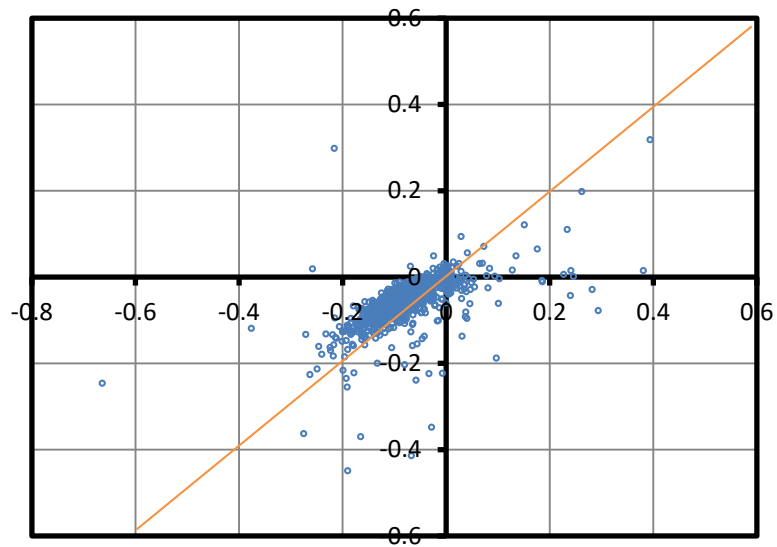
(b) Model 1: MLC-SKAT vs MLC-SKATO (typed)



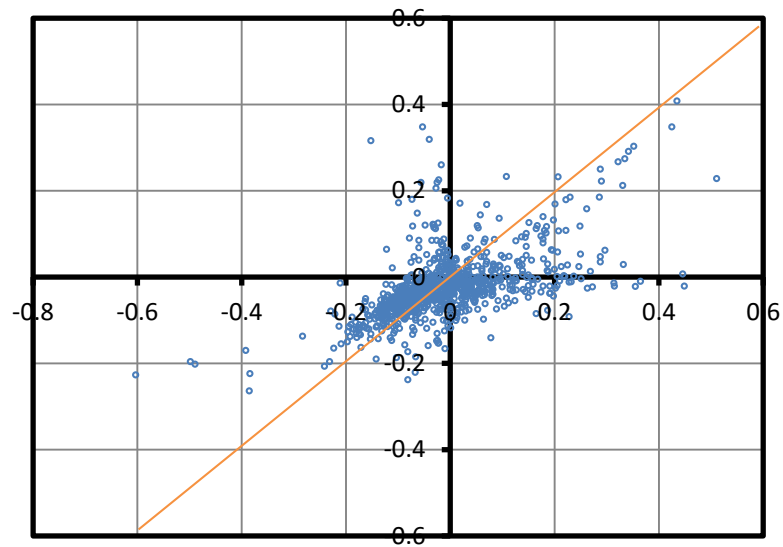
Model 1: MLC-SKAT vs MLC-SKATO (untyped)



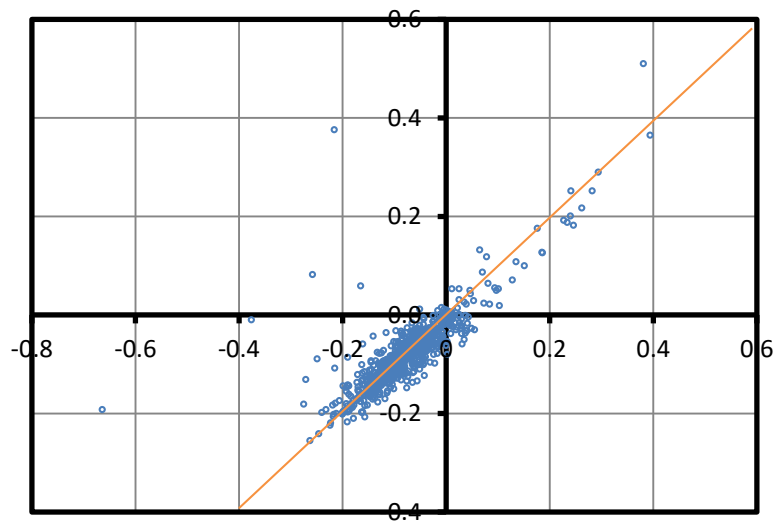
(a) Model 2: MLC-minP vs MLC-SKATO (typed)



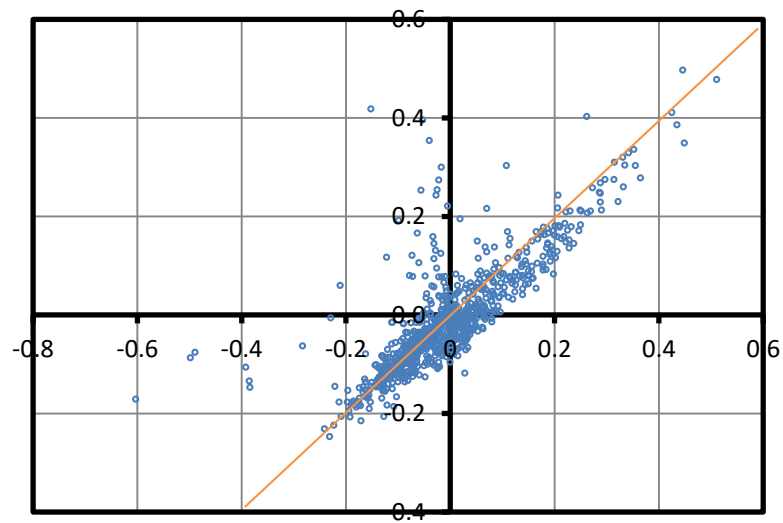
Model 2: MLC-minP vs MLC-SKATO (untyped)



(b) Model 2: MLC-SKAT vs MLC-SKATO (typed)

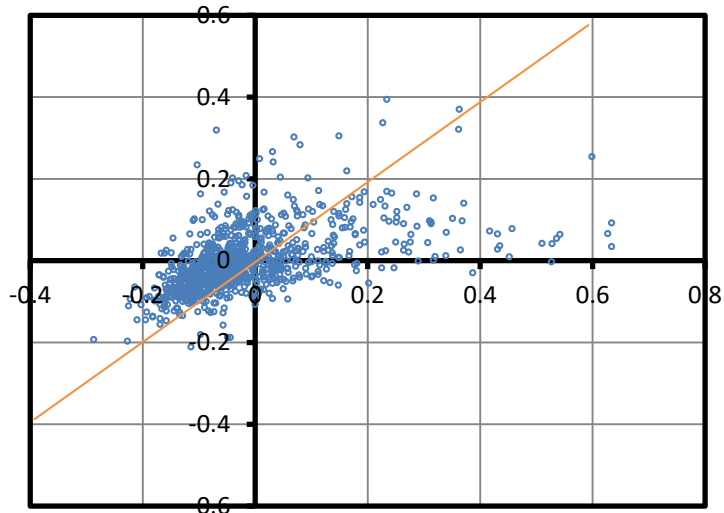


Model 2: MLC-SKAT vs MLC-SKATO (untyped)

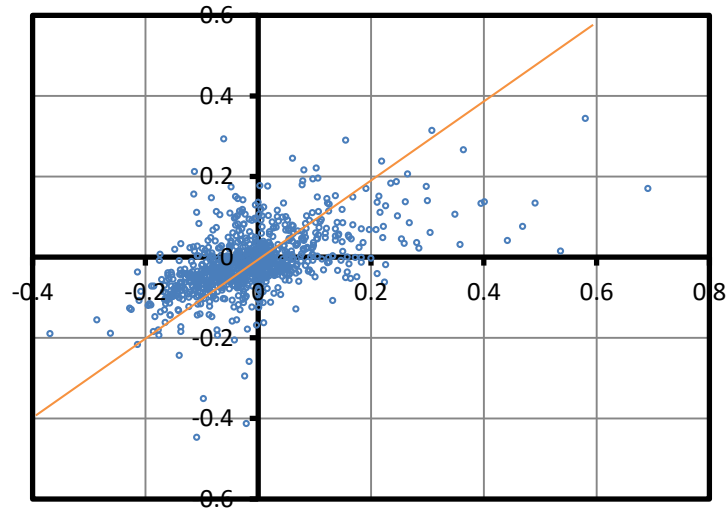


Model 3: MLC-minP vs MLC-SKATO (typed)

(a)

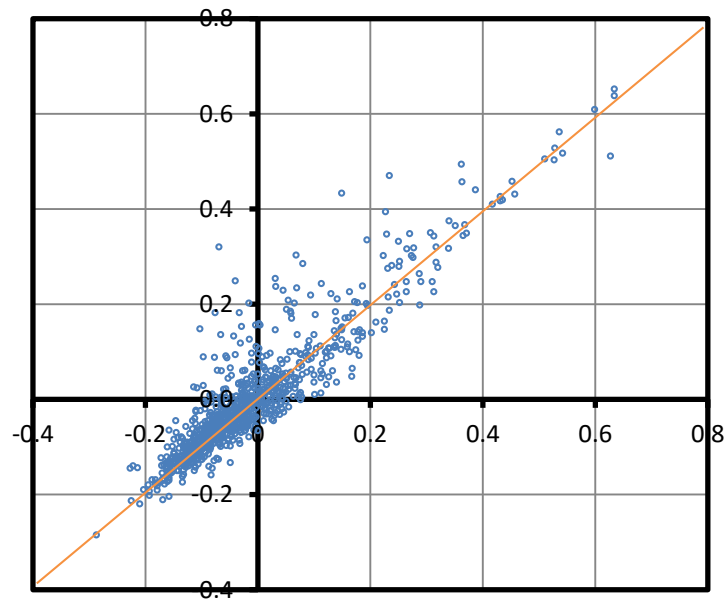


Model 3: MLC-minP vs MLC-SKATO (untyped)

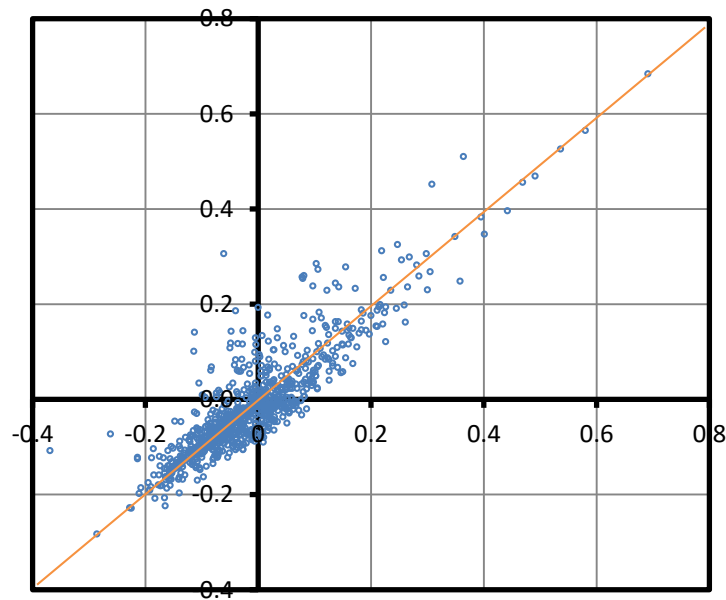


Model 3: MLC-SKAT vs MLC-SKATO (typed)

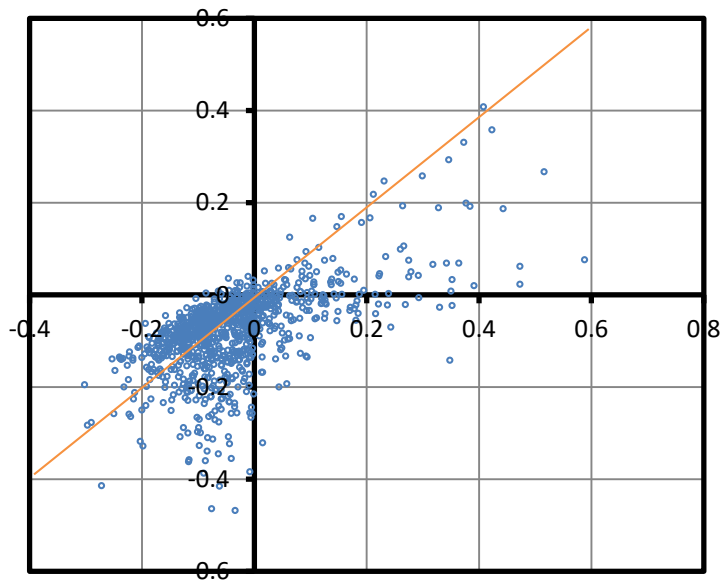
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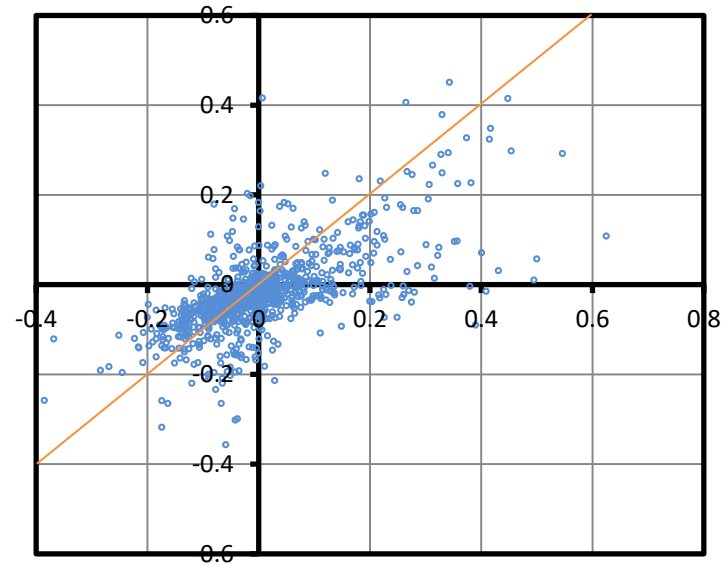
Model 3: MLC-SKAT vs MLC-SKATO (untyped)



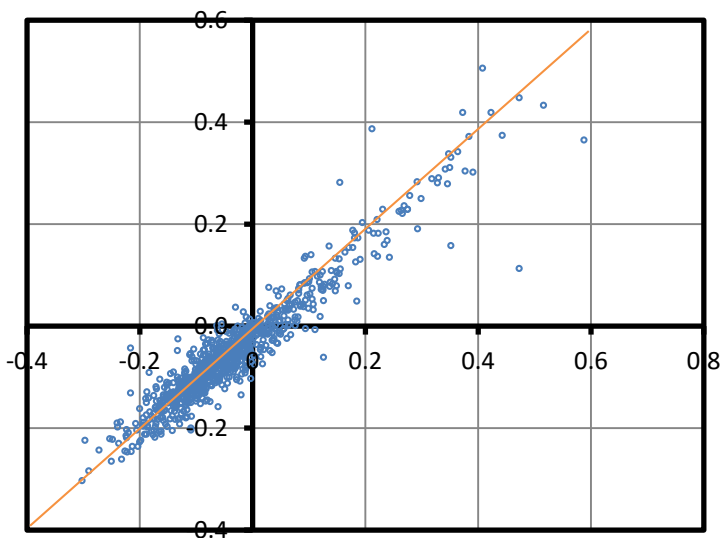
(a) Model 4: MLC-minP vs MLC-SKATO (typed)



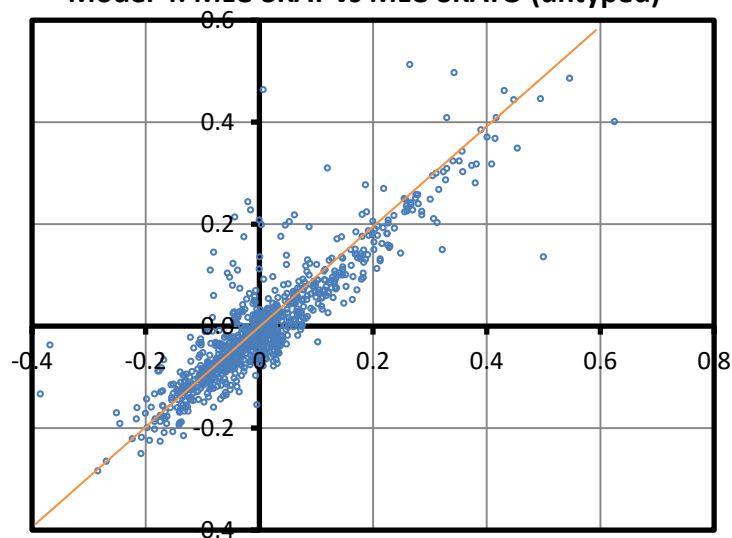
Model 4: MLC-minP vs MLC-SKATO (untyped)



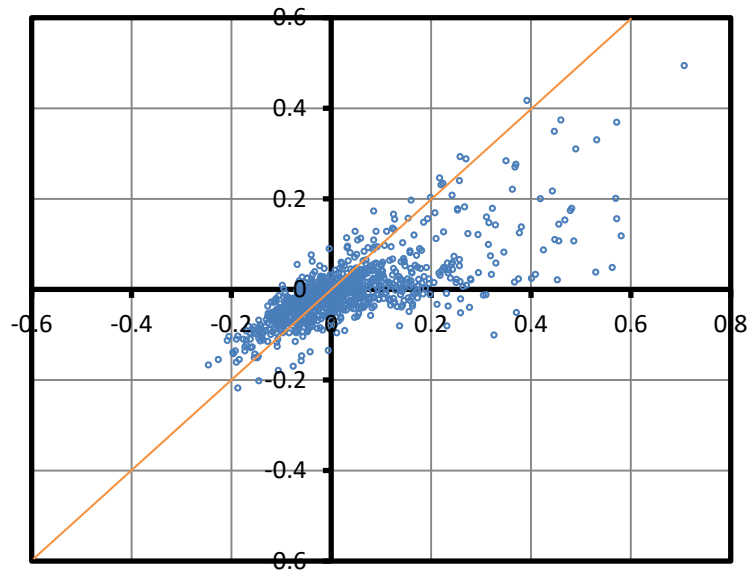
(b) Model 4: MLC-SKAT vs MLC-SKATO (typed)



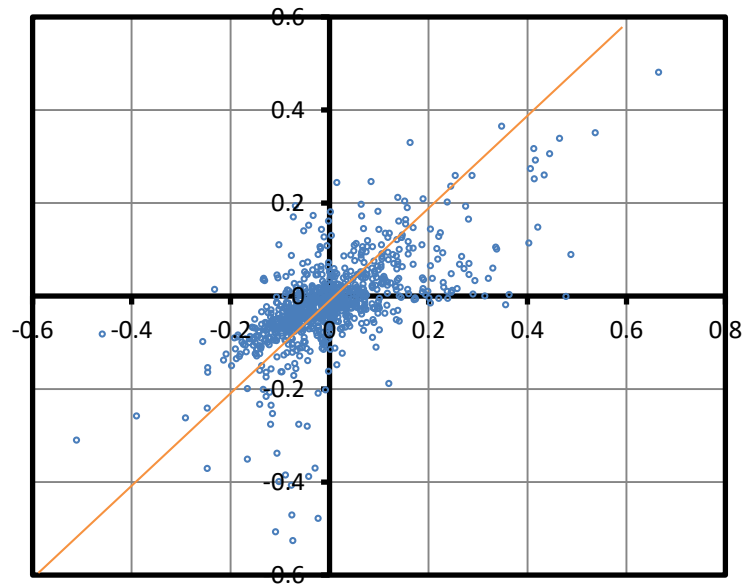
Model 4: MLC-SKAT vs MLC-SKATO (untyped)



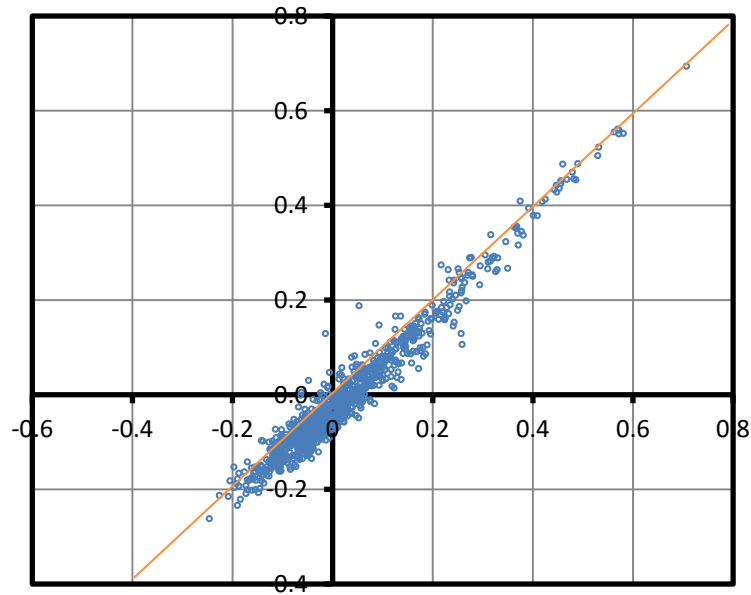
(a) Model 5: MLC-minP vs MLC-SKATO (typed)



Model 5: MLC-minP vs SKATO (untyped)



(b) Model 5: MLC-SKAT vs MLC-SKATO (typed)



Model 5: MLC-SKAT vs MLC-SKATO (untyped)

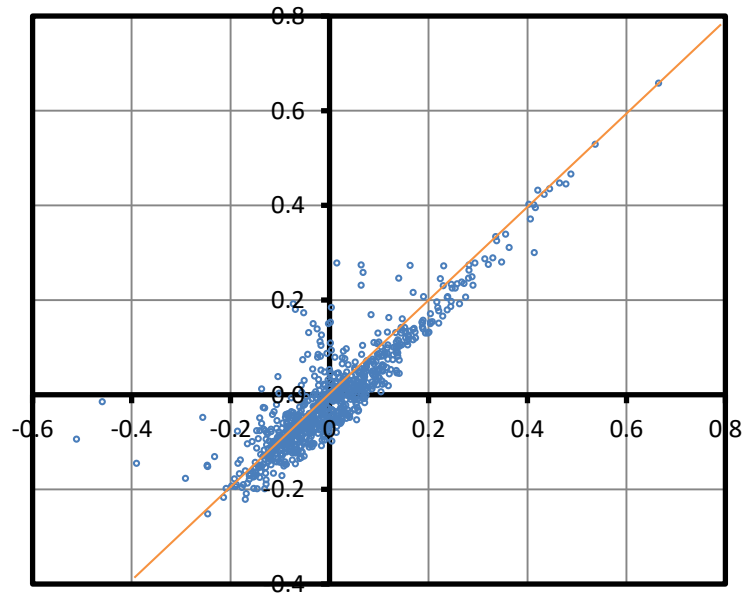


Figure S6. Power differentials (a) between MLC and MinP-M, and (b) between MLC and Wald, each plotted against power differentials between MLC and LC. The latter is on the x axis. Each point corresponds to one gene with power estimated in 1000 replicated datasets (nominal $\alpha=0.05$). Data were generated under simulation models 1-5, as defined in Table 2, and were analysed with (typed) and without (untyped) the causal variant(s) in the regression model.

In the upper panels: MLC power is greater than MinP-M power for genes above the x axis, MLC power is greater than LC power for genes on the right of the y axis, and the upper right quadrant includes all genes for which MLC is more powerful than both MinP-M and LC. Genes close to the diagonal line have MinP-M power \approx LC power, while those below the line have MinP-M power $>$ LC power.

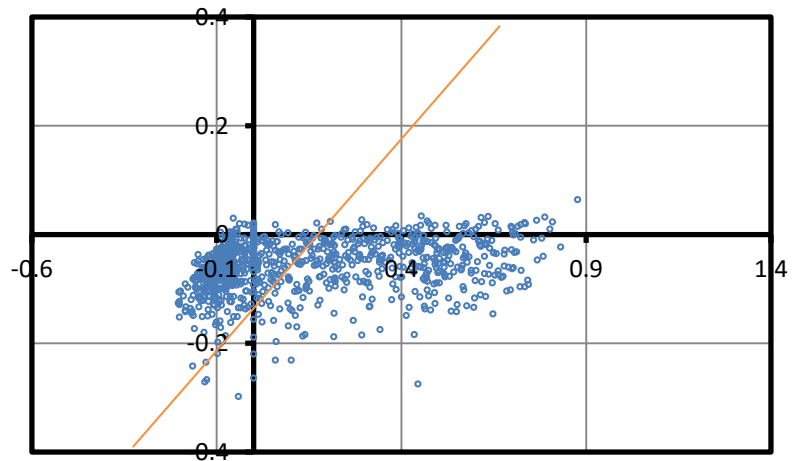
In the lower panels: MLC power is greater than Wald power for genes above the x axis, MLC power is greater than LC power for genes on the right of the y axis, and the upper right quadrant includes all genes for which MLC is more powerful than both Wald and LC). Genes close to the diagonal line have Wald power \approx LC power, while those below the line have Wald power $>$ LC power.

The table summarizes the number of genes according to power inequalities among the methods.

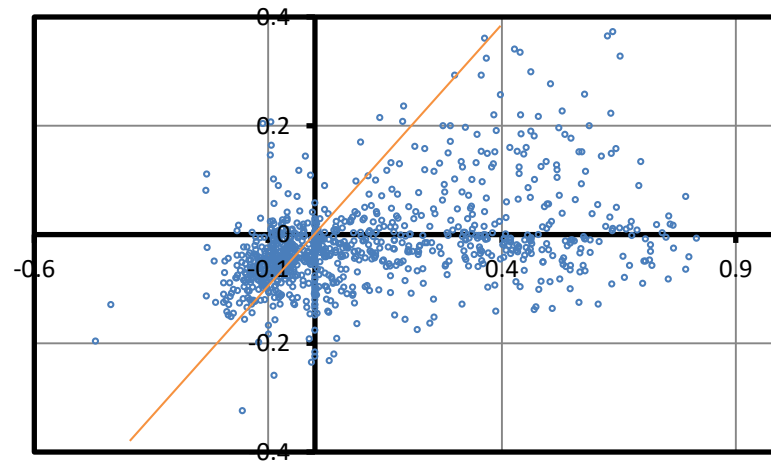
Model	# of genes	Analysis	Power Differential				
			MLC>MinPM	MLC>LC	MLC>Wald	MLC>Min,LC	MLC>all 3
1	1000	Typed	80	502	964	60	60
		Untyped	258	489	866	199	171
2	993	Typed	62	267	947	31	29
		Untyped	223	397	792	146	107
3	935	Typed	372	510	893	298	283
		Untyped	312	388	764	221	182
4	993	Typed	119	427	516	100	64
		Untyped	286	490	603	218	141
5	935	Typed	359	585	905	319	310
		Untyped	313	498	741	264	218

(a)

Model 1: MLC-minP vs MLC-LC (typed)

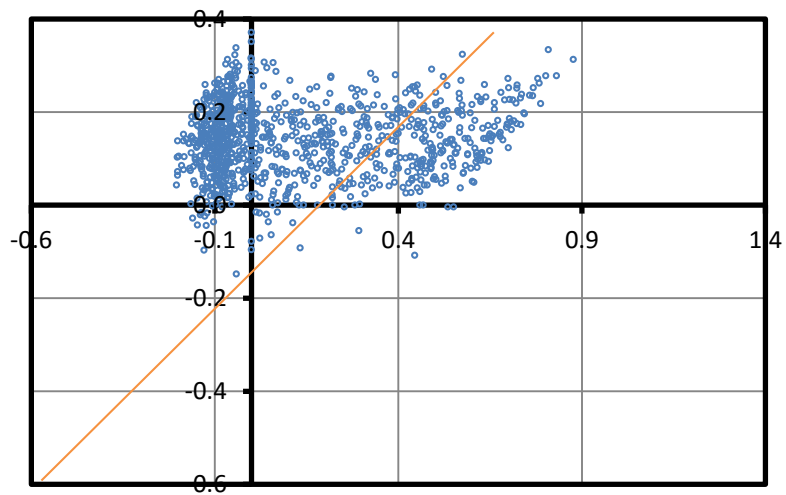


Model 1: MLC-minP vs MLC-LC (untyped)

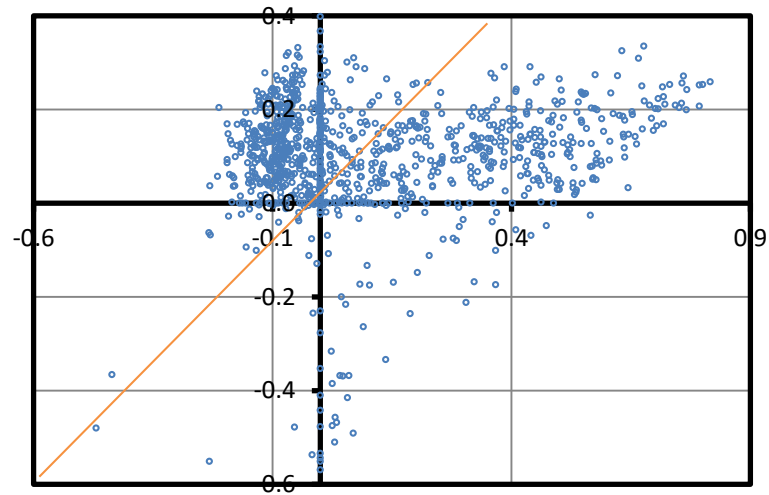


(b)

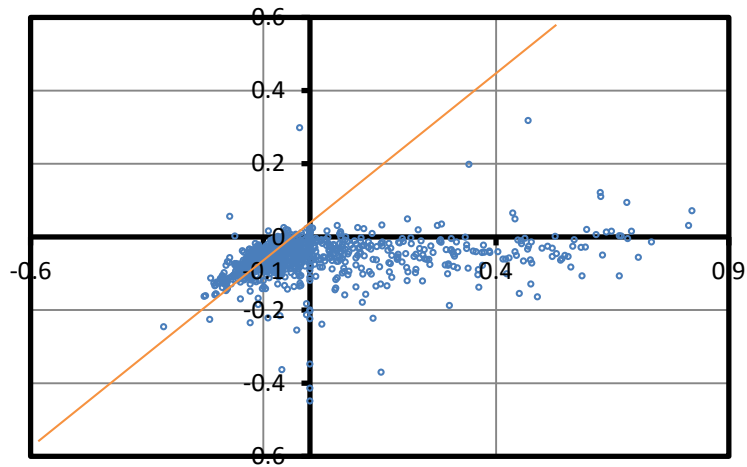
Model 1: MLC-Wald vs MLC-LC (typed)



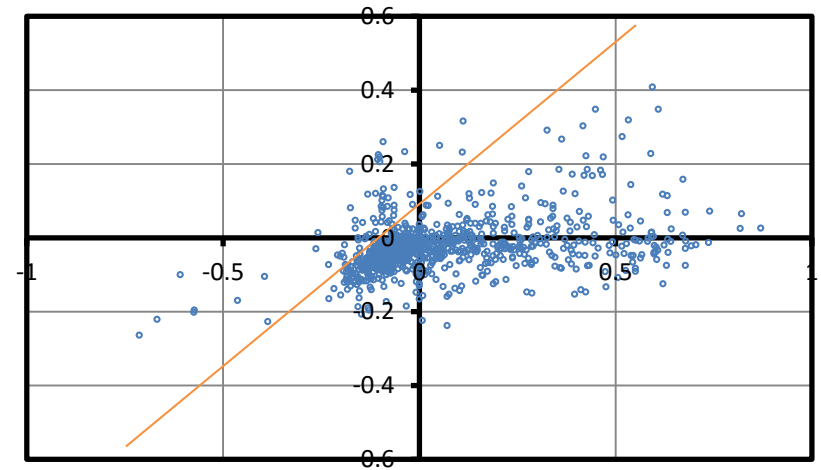
Model 1: MLC-Wald vs MLC-LC (untyped)



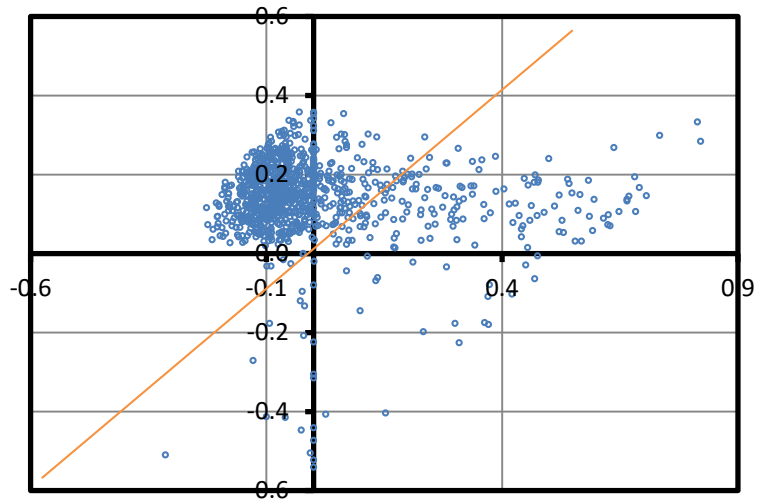
(a) Model 2: MLC-minP vs MLC-LC (typed)



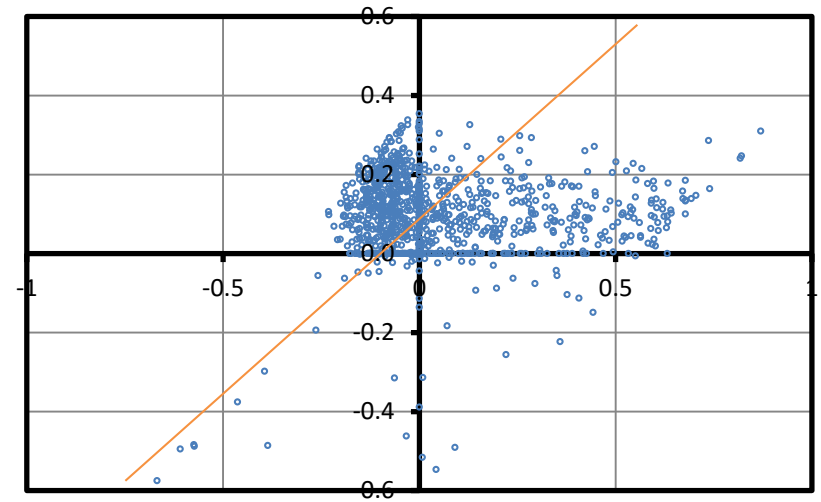
Model 2: MLC-minP vs MLC-LC (untyped)



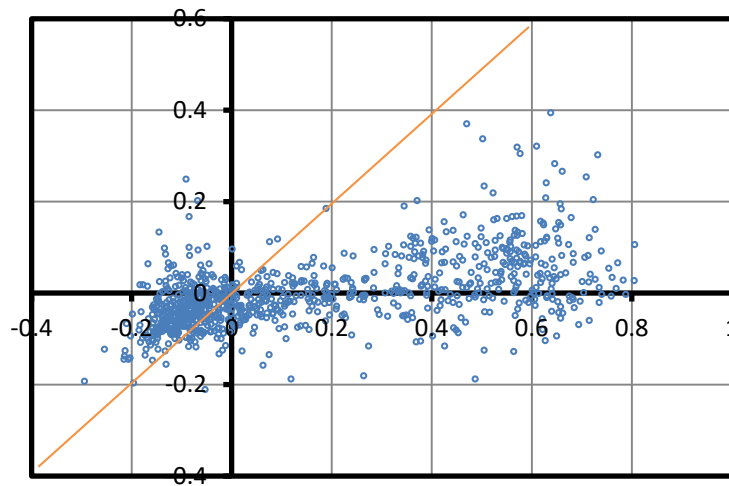
(b) Model 2: MLC-Wald vs MLC-LC (typed)



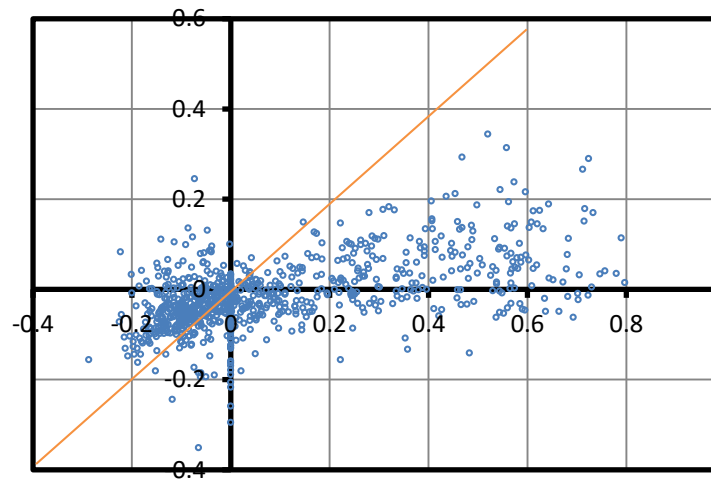
Model 2: MLC-Wald vs MLC-LC (untyped)



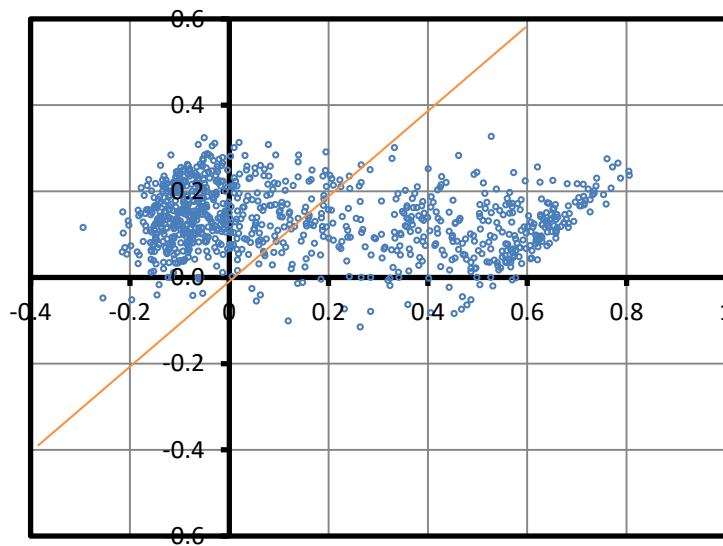
(a) Model 3: MLC-minP vs MLC-LC (typed)



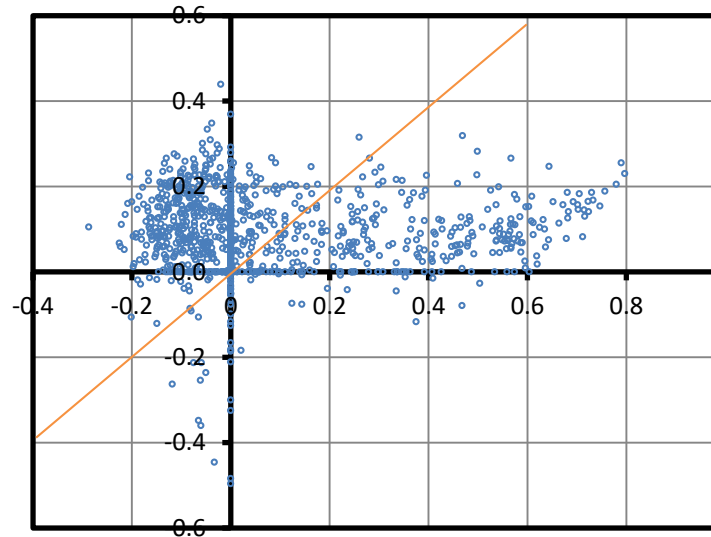
Model 3: MLC-minP vs MLC-LC (untyped)



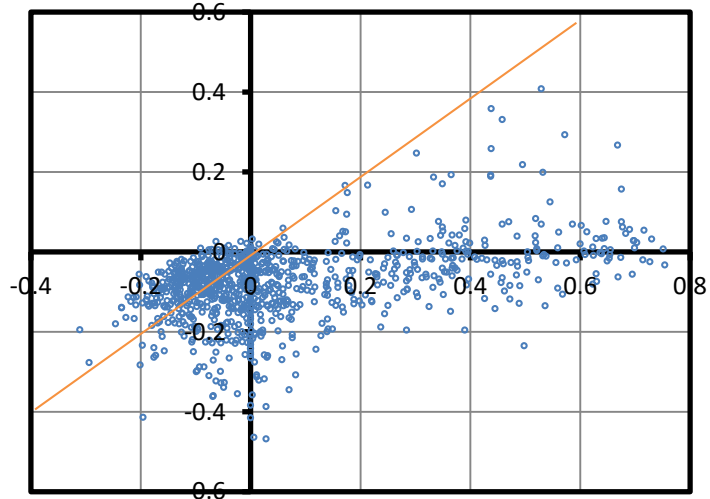
(b) Model 3: MLC-Wald vs MLC-LC (typed)



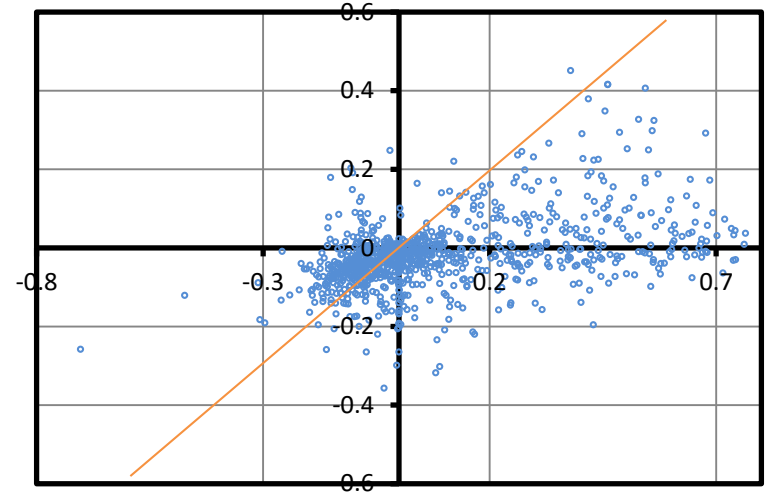
Model 3: MLC-Wald vs MLC-LC (untyped)



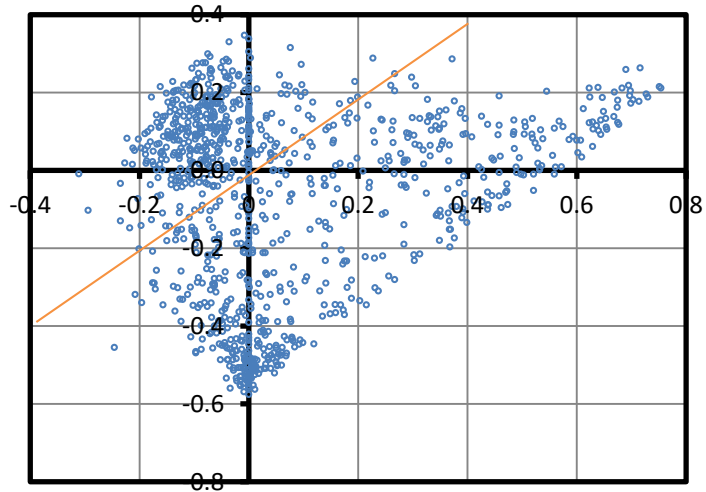
(a) Model 4: MLC-minP vs MLC-LC (typed)



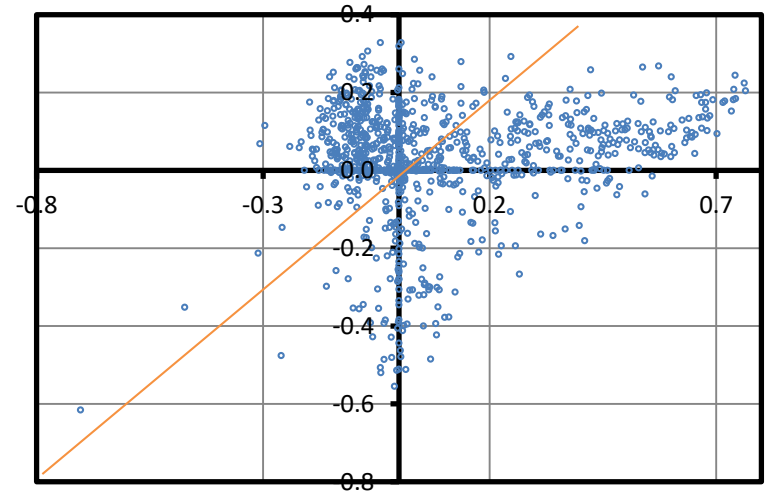
Model 4: MLC-minP vs MLC-LC (untyped)



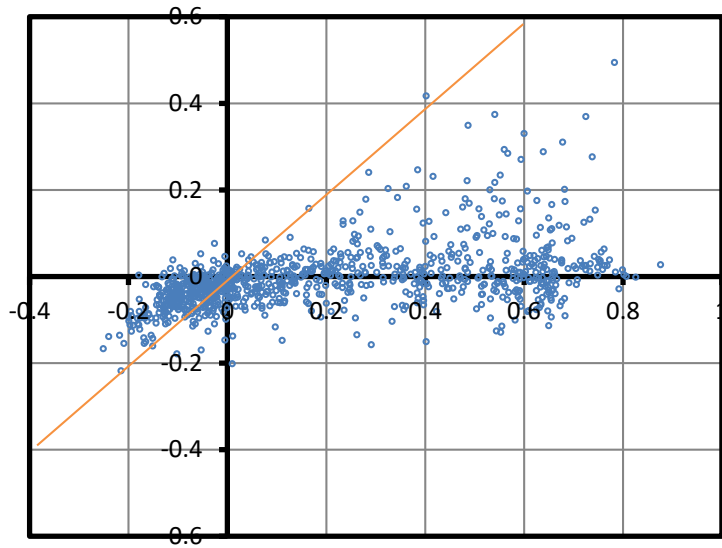
(b) Model 4: MLC-Wald vs MLC-LC (typed)



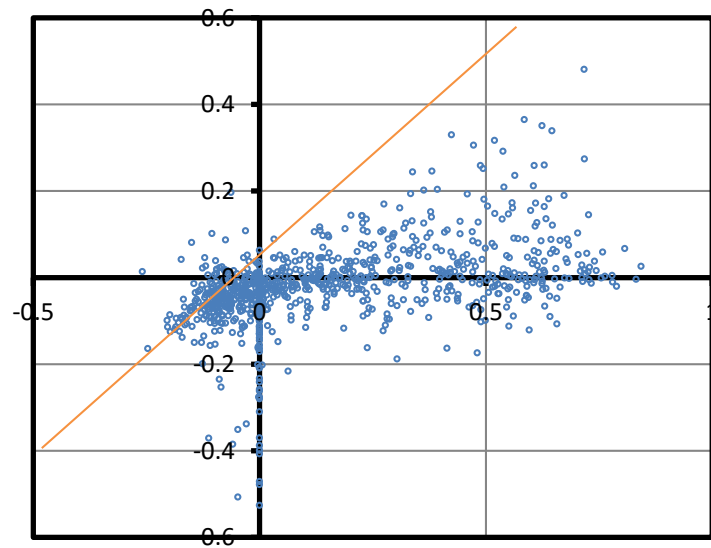
Model 4: MLC-Wald vs MLC-LC (untyped)



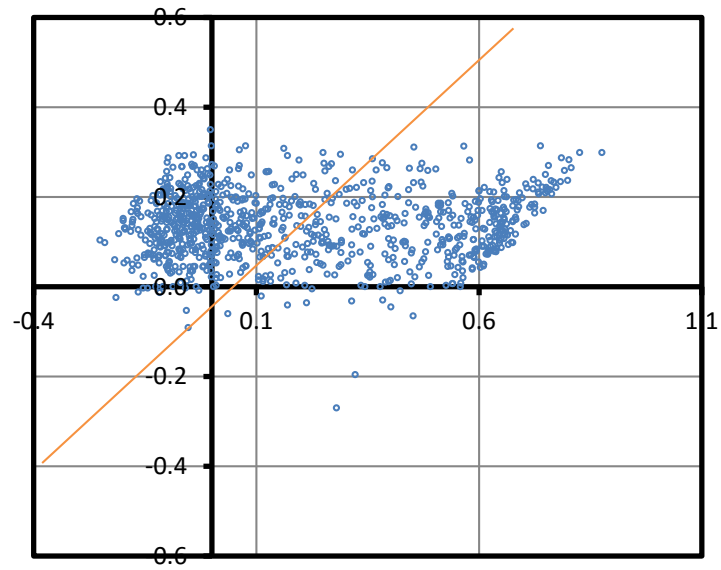
(a) Model 5: MLC-minP vs MLC-LC (typed)



Model 5: MLC-minP vs LC (untyped)



(b) Model 5: MLC-Wald vs MLC-LC (typed)



Model 5: MLC-Wald vs MLC-LC (untyped)

