

S5 Table. The pQTL of eight proteins showed a moderate linkage disequilibrium (LD) with previously identified breast cancer variants

Protein	Breast cancer susceptibility variants	PMID	LD (r^2)
GP1BA	rs11065822	32424353	0.254788
IL2RB	rs11065822	32424353	0.321178
VCAM1	rs11065822	32424353	0.321178
ICAM2	rs11065822	32424353	0.321178
KDELC2	rs74911261	29058716	0.470168
LAG3	rs11065822	32424353	0.320864
MICB	rs71557345	29059683	0.628132
PDGFRL	rs6796502	29059683	0.521154