

S4 Table. The association between infiltrate markers and *SYK(T)*, *SYK(S)* and *SYK(L)* for the total MATCH cohort, and the LNN and LNP subgroups of the MATCH cohort.

	PTPRC/CD45		VEGFA		% of infiltrating cells		% of invasive tumor cells	
	r _s	P value	r _s	P value	r _s	P value	r _s	P value
Total MATCH cohort								
<i>SYK(T)</i>	0.09	0.18	0.16	0.012	-0.04	0.56	-0.10	0.14
<i>SYK(S)</i>	0.12	0.07	0.16	0.015	-0.14	0.043	0.03	0.64
<i>SYK(L)</i>	0.08	0.21	0.12	0.06	0.04	0.56	-0.09	0.18
LNN cohort								
<i>SYK(T)</i>	0.02	0.81	0.16	0.05	-0.04	0.63	-0.10	0.19
<i>SYK(S)</i>	0.09	0.26	0.19	0.017	-0.24	0.007	0.04	0.59
<i>SYK(L)</i>	0.01	0.93	0.13	0.09	0.05	0.62	-0.14	0.07
LNP cohort								
<i>SYK(T)</i>	0.18	0.11	0.18	0.11	-0.03	0.80	-0.10	0.38
<i>SYK(S)</i>	0.10	0.38	0.09	0.42	0.02	0.87	-0.14	0.21
<i>SYK(L)</i>	0.08	0.48	0.10	0.39	0.07	0.57	-0.13	0.24