Study	TE	seTE		HR	95%–CI	Weight (fixed)	Weight (random)
BLCA	−0.42	0.1537		0.66	[0.49; 0.89]	6.8%	6.3%
BRCA	−0.25	0.1322		0.78	[0.60; 1.01]	9.2%	6.9%
CESC	−0.99	0.2622		0.37	[0.22; 0.62]	2.3%	4.0%
CHOL	−0.32	0.3894		0.72	[0.34; 1.55]	1.1%	2.4%
COAD	−0.21	0.1941		0.81	[0.56; 1.19]	4.2%	5.3%
ESCA	−0.05	0.2147		0.95	[0.62; 1.45]	3.5%	4.9%
GBM	−0.23	0.1910		0.79	[0.55; 1.15]	4.4%	5.4%
HNSC	−0.55	0.1301		0.58	[0.45; 0.75]	9.5%	6.9%
KICH	1.99	1.3510		7.34	[0.52; 103.66]	0.1%	0.3%
KIRC	0.14	0.1528		1.15	[0.85; 1.55]	6.9%	6.3%
KIRP	0.39	0.3575		1.48	[0.73; 2.98]	1.3%	2.7%
LIHC	−0.17	0.1816		0.84	[0.59; 1.20]	4.9%	5.6%
LUAD	−0.43	0.1294		0.65	[0.51; 0.84]	9.6%	6.9%
LUSC	0.04	0.1134		1.04	[0.84; 1.30]	12.4%	7.3%
PAAD	−0.38	0.2062		0.68	[0.46; 1.02]	3.8%	5.1%
PCPG	2.70	1.2842		14.91	[1.20; 184.75]	0.1%	0.3%
PRAD	1.18	0.7238		3.25	[0.79; 13.44]	0.3%	0.8%
READ	−0.21	0.5286		0.81	[0.29; 2.28]	0.6%	1.5%
SARC	−0.27	0.1854		0.76	[0.53; 1.10]	4.7%	5.5%
STAD	−0.44	0.1387		0.64	[0.49; 0.84]	8.3%	6.7%
THCA	−0.35	0.7416		0.70	[0.16; 3.02]	0.3%	0.8%
THYM	−1.03	0.3741		0.36	[0.17; 0.75]	1.1%	2.5%
UCEC	−0.18	0.1798		0.83	[0.59; 1.19]	5.0%	5.7%
Fixed effect model				**0.77**	**[0.71; 0.84]**	**100.0%**	**--**
Overall effect				**0.78**	**[0.68; 0.89]**	**--**	**100.0%**

Heterogeneity: I^2 = 58% [34%; 74%], $p < 0.01$

0.2   0.5   1   2   3