Study	TE	seTE		HR	95%−CI	Weight (fixed)	Weight (random)
BLCA	0.37	0.1797		1.45	[1.02; 2.06]	9.2%	6.8%
BRCA	−0.69	0.2456		0.50	[0.31; 0.81]	4.9%	5.6%
CESC	−0.21	0.3773		0.81	[0.39; 1.70]	2.1%	3.6%
CHOL	−1.10	0.6374		0.33	[0.10; 1.16]	0.7%	1.7%
COAD	0.10	0.2405		1.11	[0.69; 1.78]	5.1%	5.6%
ESCA	−0.44	0.3458		0.64	[0.33; 1.26]	2.5%	4.0%
GBM	0.15	0.1995		1.16	[0.79; 1.72]	7.5%	6.4%
HNSC	−0.02	0.1869		0.98	[0.68; 1.41]	8.5%	6.6%
KICH	0.43	0.7076		1.53	[0.38; 6.13]	0.6%	1.5%
KIRC	−0.50	0.1873		0.61	[0.42; 0.87]	8.5%	6.6%
KIRP	−0.45	0.4394		0.64	[0.27; 1.51]	1.5%	3.0%
LIHC	−0.24	0.2161		0.78	[0.51; 1.20]	6.4%	6.1%
LUAD	−0.44	0.1973		0.64	[0.44; 0.94]	7.6%	6.4%
LUSC	−0.03	0.1648		0.97	[0.70; 1.33]	10.9%	7.1%
PAAD	−0.52	0.2442		0.60	[0.37; 0.96]	5.0%	5.6%
PCPG	−19.64	11929.0318		0.00	[0.00; Inf]	0.0%	0.0%
PRAD	−1.30	1.0595		0.27	[0.03; 2.18]	0.3%	0.7%
READ	1.08	0.4017		2.94	[1.34; 6.46]	1.8%	3.4%
SARC	−0.41	0.2985		0.66	[0.37; 1.19]	3.3%	4.7%
STAD	0.26	0.2053		1.30	[0.87; 1.94]	7.0%	6.3%
THCA	−1.04	0.6414		0.35	[0.10; 1.24]	0.7%	1.7%
THYM	−0.80	1.0698		0.45	[0.06; 3.66]	0.3%	0.7%
UCEC	0.39	0.2296		1.48	[0.94; 2.32]	5.6%	5.8%
Fixed effect model				**0.90**	**[0.81; 1.01]**	**100.0%**	−−
Overall effect				**0.87**	**[0.73; 1.05]**	−−	**100.0%**

Heterogeneity: $I^2 = 58\%$ [33%; 73%], $p < 0.01$