Study	TE	seTE		HR	95%–CI	Weight (fixed)	Weight (random)
BLCA	0.07	0.1277		1.07	[0.83; 1.38]	9.3%	8.7%
BRCA	−0.04	0.1345		0.96	[0.73; 1.24]	8.4%	8.1%
CESC	0.12	0.2061		1.13	[0.75; 1.69]	3.6%	4.0%
CHOL	−0.07	0.5671		0.93	[0.31; 2.82]	0.5%	0.6%
COAD	−0.11	0.1705		0.89	[0.64; 1.25]	5.2%	5.5%
ESCA	0.14	0.2966		1.15	[0.64; 2.06]	1.7%	2.1%
GBM	−0.37	0.1760		0.69	[0.49; 0.97]	4.9%	5.3%
HNSC	−0.01	0.1481		0.99	[0.74; 1.33]	6.9%	7.0%
KICH	−1.02	0.9844		0.36	[0.05; 2.47]	0.2%	0.2%
KIRC	−0.45	0.1790		0.64	[0.45; 0.91]	4.7%	5.1%
KIRP	−0.59	0.3698		0.55	[0.27; 1.15]	1.1%	1.4%
LIHC	0.04	0.1346		1.04	[0.80; 1.35]	8.4%	8.1%
LUAD	−0.12	0.1224		0.89	[0.70; 1.13]	10.1%	9.3%
LUSC	0.03	0.1269		1.03	[0.80; 1.32]	9.4%	8.8%
PAAD	−0.70	0.2614		0.50	[0.30; 0.83]	2.2%	2.6%
PCPG	0.10	1.1270		1.11	[0.12; 10.09]	0.1%	0.2%
PRAD	0.67	0.7366		1.95	[0.46; 8.27]	0.3%	0.4%
READ	0.67	0.4893		1.96	[0.75; 5.10]	0.6%	0.8%
SARC	0.07	0.1367		1.07	[0.82; 1.40]	8.1%	7.9%
STAD	0.04	0.1289		1.04	[0.81; 1.34]	9.1%	8.6%
THCA	−0.52	0.9908		0.60	[0.09; 4.16]	0.2%	0.2%
THYM	−0.34	0.7843		0.71	[0.15; 3.30]	0.2%	0.3%
UCEC	−0.07	0.1773		0.94	[0.66; 1.33]	4.8%	5.2%
Fixed effect model				**0.95**	**[0.88; 1.02]**	**100.0%**	**--**
Overall effect				**0.94**	**[0.86; 1.03]**	**--**	**100.0%**

Heterogeneity: $I^2 = 14\%$ [0%; 48%], $p = 0.27$