Study	TE	seTE		HR	95%–CI	Weight (fixed)	Weight (random)
BLCA	−0.04	0.0252		0.96	[0.91; 1.01]	9.6%	9.6%
BRCA	−0.05	0.0308		0.95	[0.89; 1.01]	6.5%	6.5%
CESC	−0.03	0.0404		0.97	[0.90; 1.05]	3.8%	3.8%
CHOL	−0.00	0.0894		1.00	[0.84; 1.19]	0.8%	0.8%
COAD	−0.00	0.0443		1.00	[0.92; 1.09]	3.1%	3.1%
ESCA	−0.05	0.0737		0.95	[0.82; 1.10]	1.1%	1.1%
GBM	−0.01	0.0191		0.99	[0.95; 1.02]	16.9%	16.9%
HNSC	−0.04	0.0249		0.96	[0.91; 1.01]	9.9%	9.9%
KICH	−0.04	0.1627		0.96	[0.70; 1.33]	0.2%	0.2%
KIRC	−0.07	0.0431		0.93	[0.86; 1.02]	3.3%	3.3%
KIRP	0.05	0.0668		1.05	[0.92; 1.20]	1.4%	1.4%
LIHC	−0.05	0.0261		0.95	[0.91; 1.00]	9.0%	9.0%
LUAD	−0.07	0.0386		0.93	[0.87; 1.01]	4.1%	4.1%
LUSC	−0.05	0.0357		0.95	[0.89; 1.02]	4.8%	4.8%
PAAD	0.09	0.0935		1.10	[0.91; 1.32]	0.7%	0.7%
PCPG	−0.12	0.1536		0.88	[0.65; 1.19]	0.3%	0.3%
PRAD	−0.10	0.2127		0.91	[0.60; 1.37]	0.1%	0.1%
READ	−0.17	0.1125		0.85	[0.68; 1.05]	0.5%	0.5%
SARC	−0.03	0.0196		0.97	[0.94; 1.01]	16.0%	16.0%
STAD	0.09	0.0533		1.09	[0.98; 1.21]	2.2%	2.2%
THCA	−0.05	0.0769		0.95	[0.81; 1.10]	1.0%	1.0%
THYM	0.01	0.0636		1.01	[0.89; 1.14]	1.5%	1.5%
UCEC	0.05	0.0441		1.06	[0.97; 1.15]	3.1%	3.1%
Fixed effect model				**0.97**	**[0.96; 0.99]**	**100.0%**	**––**
Overall effect				**0.97**	**[0.96; 0.99]**	**––**	**100.0%**

Heterogeneity: I^2 = 0% [0%; 37%], p = 0.65

0.2   0.5   1   2   3