Study	TE	seTE		HR	95%−CI	Weight (fixed)	Weight (random)
BLCA	−0.05	0.0264		0.95	[0.90; 1.00]	10.6%	9.1%
BRCA	−0.08	0.0298		0.92	[0.87; 0.98]	8.3%	7.8%
CESC	−0.09	0.0377		0.91	[0.85; 0.98]	5.2%	5.5%
CHOL	0.01	0.0638		1.01	[0.89; 1.14]	1.8%	2.2%
COAD	−0.11	0.0474		0.89	[0.81; 0.98]	3.3%	3.8%
ESCA	−0.00	0.0667		1.00	[0.88; 1.14]	1.7%	2.1%
GBM	−0.01	0.0305		0.99	[0.93; 1.05]	7.9%	7.5%
HNSC	−0.08	0.0357		0.93	[0.86; 0.99]	5.8%	6.0%
KICH	0.02	0.1480		1.03	[0.77; 1.37]	0.3%	0.4%
KIRC	−0.11	0.0358		0.89	[0.83; 0.96]	5.7%	5.9%
KIRP	−0.04	0.0595		0.96	[0.86; 1.08]	2.1%	2.5%
LIHC	−0.07	0.0293		0.93	[0.88; 0.98]	8.6%	8.0%
LUAD	−0.13	0.0375		0.88	[0.81; 0.94]	5.2%	5.5%
LUSC	−0.01	0.0421		0.99	[0.91; 1.08]	4.2%	4.6%
PAAD	−0.03	0.0558		0.97	[0.87; 1.08]	2.4%	2.8%
PCPG	−0.01	0.1277		0.99	[0.77; 1.27]	0.5%	0.6%
PRAD	−0.01	0.1728		0.99	[0.70; 1.38]	0.2%	0.3%
READ	−0.03	0.0841		0.97	[0.82; 1.14]	1.0%	1.3%
SARC	−0.09	0.0259		0.91	[0.87; 0.96]	11.0%	9.4%
STAD	0.04	0.0429		1.04	[0.96; 1.13]	4.0%	4.5%
THCA	−0.02	0.1076		0.98	[0.79; 1.21]	0.6%	0.8%
THYM	−0.24	0.0879		0.78	[0.66; 0.93]	1.0%	1.2%
UCEC	−0.07	0.0292		0.93	[0.88; 0.99]	8.7%	8.0%
Fixed effect model				**0.94**	**[0.92; 0.95]**	**100.0%**	**−−**
Overall effect				**0.94**	**[0.92; 0.96]**	**−−**	**100.0%**

Heterogeneity: $I^2 = 19\%$ [0%; 51%], $p = 0.21$