Study	TE	seTE		HR	95%−CI	Weight (fixed)	Weight (random)
BLCA	−0.02	0.0162		0.98	[0.95; 1.01]	10.7%	7.9%
BRCA	0.01	0.0207		1.01	[0.97; 1.05]	6.5%	6.4%
CESC	−0.01	0.0273		0.99	[0.93; 1.04]	3.8%	4.8%
CHOL	0.02	0.0494		1.02	[0.92; 1.12]	1.1%	2.0%
COAD	−0.02	0.0214		0.98	[0.94; 1.02]	6.1%	6.2%
ESCA	0.03	0.0381		1.03	[0.96; 1.11]	1.9%	3.0%
GBM	0.01	0.0209		1.01	[0.97; 1.06]	6.4%	6.4%
HNSC	0.01	0.0150		1.01	[0.98; 1.04]	12.4%	8.3%
KICH	−0.14	0.0670		0.87	[0.77; 1.00]	0.6%	1.2%
KIRC	0.03	0.0277		1.03	[0.97; 1.08]	3.6%	4.7%
KIRP	0.08	0.0425		1.09	[1.00; 1.18]	1.5%	2.5%
LIHC	−0.05	0.0236		0.95	[0.91; 1.00]	5.0%	5.6%
LUAD	−0.03	0.0176		0.97	[0.94; 1.01]	9.0%	7.4%
LUSC	0.02	0.0162		1.02	[0.99; 1.06]	10.7%	7.9%
PAAD	−0.10	0.0308		0.91	[0.85; 0.96]	2.9%	4.1%
PCPG	−0.12	0.0905		0.88	[0.74; 1.06]	0.3%	0.7%
PRAD	−0.04	0.0823		0.96	[0.81; 1.12]	0.4%	0.8%
READ	0.02	0.0424		1.02	[0.93; 1.10]	1.6%	2.6%
SARC	−0.01	0.0252		0.99	[0.94; 1.04]	4.4%	5.2%
STAD	0.02	0.0256		1.02	[0.97; 1.08]	4.2%	5.1%
THCA	−0.03	0.1025		0.97	[0.80; 1.19]	0.3%	0.5%
THYM	−0.06	0.0909		0.94	[0.79; 1.13]	0.3%	0.7%
UCEC	0.03	0.0210		1.03	[0.99; 1.08]	6.3%	6.3%
Fixed effect model				**1.00**	**[0.99; 1.01]**	**100.0%**	**−−**
Overall effect				**1.00**	**[0.98; 1.01]**	**−−**	**100.0%**

Heterogeneity: $I^2 = 42\%$ [5%; 65%], $p = 0.02$

0.5 1 2