Study	TE	seTE	HR	95%-CI	Weight (fixed)	Weight (random)
BLCA	-0.07	0.0454	0.93	[0.85; 1.02]	12.9%	9.3%
BRCA	-0.06	0.0703	0.94	[0.82; 1.08]	5.4%	6.0%
CESC	0.08	0.0861	1.08	[0.91; 1.28]	3.6%	4.5%
CHOL	0.27	0.1626	<u> t</u>	[0.95; 1.80]	1.0%	1.6%
COAD	0.13	0.0781	1.14	[0.98; 1.33]	4.3%	5.2%
ESCA	-0.17	0.0937	0.85	[0.71; 1.02]	3.0%	4.0%
GBM	0.25	0.1009	1.29	[1.06; 1.57]	2.6%	3.6%
HNSC	0.02	0.0523	1.02	[0.92; 1.13]	9.7%	8.2%
KICH	-0.08	0.3113	0.92	[0.50; 1.70]	0.3%	0.5%
KIRC	-0.03	0.0455	0.97	[0.89; 1.06]	12.8%	9.3%
KIRP	-0.10	0.1436		[0.68; 1.20]	1.3%	2.0%
LIHC	0.05	0.0672	1.05	[0.92; 1.20]	5.9%	6.3%
LUAD	0.10	0.0597	<u>i </u>	[0.98; 1.24]	7.4%	7.2%
LUSC	0.02	0.0534	1.02	[0.92; 1.14]	9.3%	8.1%
PAAD	-0.09	0.1000	0.91	[0.75; 1.11]	2.6%	3.6%
PCPG	-0.27	0.2329	+ <u>r</u> 0.76	[0.48; 1.20]	0.5%	0.8%
PRAD	0.05	0.4338	`` + → 1.05	[0.45; 2.45]	0.1%	0.2%
READ	0.03	0.1669	1.03	[0.74; 1.43]	0.9%	1.5%
SARC	0.03	0.0727	1.03	[0.90; 1.19]	5.0%	5.7%
STAD	-0.02	0.0578	0.98	[0.87; 1.10]	7.9%	7.5%
THCA	1.22	0.5619	6	1.12; 10.15]	0.1%	0.1%
THYM	0.25	0.3330		[0.67; 2.45]	0.2%	0.4%
UCEC	0.11	0.0912	1.12	[0.94; 1.34]	3.2%	4.2%
Fixed effect model			1.01 [[0.98; 1.04]	100.0%	
Overall effect			1.02 [[0.98; 1.06]		100.0%
Heterogeneity: I ²	= 32% [0%; 5	9%], p =	1 1 2			