Study	TE seTE	н	R 95%−CI	Weight (fixed)	Weight (random)
BLCA	0.02 0.0376	1.0	2 [0.94; 1.09]	9.2%	7.9%
BRCA	0.08 0.0743	1.0	8 [0.93; 1.25]	2.4%	3.2%
CESC	0.02 0.0477	1.0	2 [0.93; 1.12]	5.7%	6.0%
CHOL	0.09 0.1246	- [9 [0.85; 1.39]	0.8%	1.3%
COAD	0.01 0.0502	- 1.0	1 [0.91; 1.11]	5.2%	5.6%
ESCA	-0.05 0.0757	0.9	5 [0.82; 1.10]	2.3%	3.1%
GBM	0.14 0.0867	1.1	5 [0.97; 1.36]	1.7%	2.4%
HNSC	0.04 0.0385	1.0	5 [0.97; 1.13]	8.8%	7.8%
KICH	0.14 0.1310	1.1	5 [0.89; 1.49]	0.8%	1.2%
KIRC	0.22 0.0532	1.2	4 [1.12; 1.38]	4.6%	5.2%
KIRP	0.11 0.0485	1.1	1 [1.01; 1.22]	5.6%	5.9%
LIHC	0.01 0.0351	1.0	1 [0.94; 1.08]	10.6%	8.5%
LUAD	-0.00 0.0340	1.0	0 [0.93; 1.07]	11.3%	8.8%
LUSC	0.02 0.0373	1.0	2 [0.94; 1.09]	9.4%	8.0%
PAAD	0.14 0.0551	$\frac{1}{t}$ 1.1	5 [1.03; 1.28]	4.3%	5.0%
PCPG	0.04 0.2362	1.0	5 [0.66; 1.66]	0.2%	0.4%
PRAD	0.12 0.1768	1.1	3 [0.80; 1.59]	0.4%	0.7%
READ	0.06 0.0997	1.0	7 [0.88; 1.30]	1.3%	1.9%
SARC	0.07 0.0463	1.0	7 [0.98; 1.18]	6.1%	6.3%
STAD	0.12 0.0559	1.1	2 [1.01; 1.25]	4.2%	4.9%
THCA	0.45 0.1688		7 [1.13; 2.19]	0.5%	0.7%
THYM	0.01 0.2112		1 [0.67; 1.53]	0.3%	0.5%
UCEC	0.07 0.0561	1.0	8 [0.96; 1.20]	4.1%	4.8%
Fixed effect model		1.0	5 [1.03; 1.08]	100.0%	
Overall effect		<u></u>	6 [1.03; 1.09]		100.0%
Heterogeneity: I ² =	: 29% [0%; 58%], <i>p</i> = 0				