Study	TE	seTE	HR	95%-CI	Weight (fixed)	Weight (random)
BLCA	-0.00	0.0150	1.00	[0.97; 1.03]	9.5%	9.5%
BRCA	0.01	0.0169	1.01	[0.98; 1.04]	7.5%	7.5%
CESC	0.02	0.0242	1.02	[0.97; 1.07]	3.6%	3.6%
CHOL	0.02	0.0458	1.02	[0.93; 1.11]	1.0%	1.0%
COAD	0.02	0.0226	1.02	[0.97; 1.06]	4.2%	4.2%
ESCA	-0.01	0.0276	0.99	[0.94; 1.05]	2.8%	2.8%
GBM	0.04	0.0272	1.04	[0.99; 1.10]	2.9%	2.9%
HNSC	0.01	0.0139	+ 1.01	[0.98; 1.04]	11.1%	11.1%
KICH	0.14	0.0996	1.15	[0.95; 1.40]	0.2%	0.2%
KIRC	-0.04	0.0251	0.97	[0.92; 1.01]	3.4%	3.4%
KIRP	0.04	0.0303	1.04	[0.98; 1.10]	2.3%	2.3%
LIHC	0.01	0.0166	1.01	[0.98; 1.05]	7.7%	7.7%
LUAD	0.03	0.0149	+ 1.03	[1.00; 1.06]	9.7%	9.7%
LUSC	0.02	0.0142	1.02	[0.99; 1.05]	10.6%	10.6%
PAAD	0.03	0.0217	1.03	[0.99; 1.07]	4.5%	4.5%
PCPG	0.23	0.1545	 + 	[0.93; 1.71]	0.1%	0.1%
PRAD	-0.05	0.0816	0.95	[0.81; 1.11]	0.3%	0.3%
READ	0.01	0.0458	1.01	[0.92; 1.10]	1.0%	1.0%
SARC	0.03	0.0245	1.03	[0.98; 1.08]	3.6%	3.6%
STAD	0.01	0.0170	1.01	[0.97; 1.04]	7.4%	7.4%
THCA	-0.01	0.0507	- 0.99	[0.90; 1.10]	0.8%	0.8%
THYM	0.06	0.0663	1.06	[0.93; 1.21]	0.5%	0.5%
UCEC	0.04	0.0202	1.04	[1.00; 1.08]	5.2%	5.2%
Fixed effect model			1.02	[1.01; 1.03]	100.0%	
Overall effect		_	1.02	[1.01; 1.03]		100.0%
Heterogeneity: I	² = 0% [0%; 31	1%], p = 0.7 0.9	1 2			