Study	TE	seTE	Н	IR	95%-CI	Weight (fixed)	Weight (random)
BLCA	0.01	0.0386	1.0	01	[0.94; 1.09]	6.6%	6.7%
BRCA	-0.03	0.0576	2.0	97	[0.87; 1.09]	3.0%	4.2%
CESC	0.08	0.0556	 (= − 1.0	80	[0.97; 1.20]	3.2%	4.4%
CHOL	0.03	0.0742	- - 1.0)3	[0.89; 1.19]	1.8%	2.9%
COAD	0.09	0.0649	 	9	[0.96; 1.24]	2.4%	3.6%
ESCA	0.06	0.0644	1.0	06	[0.94; 1.20]	2.4%	3.6%
GBM	0.03	0.0472	- i 1.0)3	[0.94; 1.13]	4.4%	5.4%
HNSC	0.00	0.0309		00	[0.94; 1.07]	10.3%	8.1%
KICH	0.14	0.2115	1.7	14	[0.76; 1.73]	0.2%	0.5%
KIRC	0.12	0.0339	1.1	13	[1.06; 1.21]	8.6%	7.5%
KIRP	0.11	0.0477	1.1	12	[1.02; 1.23]	4.4%	5.4%
LIHC	0.01	0.0220	1.0	01	[0.97; 1.06]	20.5%	9.9%
LUAD	0.02	0.0454	-)2	[0.94; 1.12]	4.8%	5.7%
LUSC	0.01	0.0435	- ‡: 1.0)1	[0.93; 1.10]	5.2%	5.9%
PAAD	-0.13	0.0860	3.0	37	[0.74; 1.04]	1.3%	2.3%
PCPG	0.06	0.1332)7	[0.82; 1.38]	0.6%	1.1%
PRAD	0.16	0.2170	1.1	17	[0.77; 1.80]	0.2%	0.4%
READ	-0.12	0.1547	3.0	38	[0.65; 1.20]	0.4%	0.8%
SARC	0.07	0.0392	1.0)7	[0.99; 1.15]	6.4%	6.6%
STAD	0.04	0.0368	1.0)4	[0.97; 1.12]	7.3%	7.0%
THCA	0.59	0.1654	(→ → 1.8	30	[1.30; 2.48]	0.4%	0.7%
THYM	-0.02	0.1092	2.0	98	[0.80; 1.22]	0.8%	1.6%
UCEC	0.14	0.0455	1.	15	[1.06; 1.26]	4.8%	5.6%
Fixed effect model			∳ 1.0	04	[1.02; 1.06]	100.0%	
Overall effect			<u> </u>	05	[1.02; 1.08]		100.0%
Heterogeneity:	1 ² = 41% [3%; 6	[64%], p = 0	1 2				