| Study | logHR [95%CI] | P−value |
|-------|---------------|---------|
| DLBC | −1.11 [−2.11; −0.11] | 0.03 |
| CHOL | −0.51 [−1.12; 0.10] | 0.10 |
| PCPG | −0.19 [−0.72; 0.34] | 0.48 |
| THCA | −0.18 [−0.49; 0.13] | 0.27 |
| LIHC | −0.16 [−0.34; 0.02] | 0.09 |
| PRAD | −0.11 [−0.36; 0.14] | 0.43 |
| UCS | −0.08 [−0.55; 0.39] | 0.74 |
| HNSC | −0.03 [−0.21; 0.15] | 0.72 |
| SKCM | −0.01 [−0.17; 0.15] | 0.92 |
| OV | 0.08 [−0.10; 0.26] | 0.34 |
| ESCA | 0.10 [−0.19; 0.39] | 0.52 |
| LUAD | 0.10 [−0.08; 0.28] | 0.26 |
| LUNG | 0.10 [−0.04; 0.24] | 0.13 |
| LUSC | 0.10 [−0.12; 0.32] | 0.38 |
| ACC | 0.11 [−0.32; 0.54] | 0.61 |
| STAD | 0.11 [−0.11; 0.33] | 0.32 |
| SARC | 0.12 [−0.13; 0.37] | 0.33 |
| BLCA | 0.13 [−0.05; 0.31] | 0.17 |
| THYM | 0.14 [−0.43; 0.71] | 0.62 |
| BRCA | 0.15 [−0.07; 0.37] | 0.15 |
| UCEC | 0.15 [−0.28; 0.58] | 0.50 |
| TGCT | 0.18 [−0.21; 0.57] | 0.38 |
| COAD | 0.19 [−0.06; 0.44] | 0.14 |
| KIRC | 0.23 [ 0.05; 0.41] | 0.01 |
| CESC | 0.26 [−0.05; 0.57] | 0.10 |
| MESO | 0.26 [−0.09; 0.61] | 0.15 |
| GBM | 0.34 [ 0.10; 0.58] | 0.01 |
| KIRP | 0.37 [ 0.06; 0.68] | 0.02 |
| READ | 0.39 [−0.18; 0.96] | 0.19 |
| PAAD | 0.40 [ 0.15; 0.65] | 0.00 |
| LGG | 0.46 [ 0.26; 0.66] | 0.00 |
| KICH | 0.56 [−0.15; 1.27] | 0.12 |
| UVM | 0.61 [ 0.08; 1.14] | 0.03 |
| Total | 0.12 [ 0.06; 0.19] | |

−1.5 −0.5 0.5 1 1.5
logHR estimate

Heterogeneity: $\chi^2_{32} = 65.05$ ($P < .001$), $I^2 = 51\%$ [27%; 67%]
Test for overall effect: $z = 3.76$ ($P < .001$)