Source (95% CI)

Primary = Kidney

Miao.1, n = 33-0.79 [-1.71; 0.13] Braun, n = 178-0.07 [-0.46; 0.32] 0.06 [-0.51; 0.63] Mariathasan, n = 67Total -0.11 [-0.42; 0.19]

Heterogeneity: $\chi_2^2 = 2.48$ (P = .29), $I^2 = 19\%$ [0%; 92%]

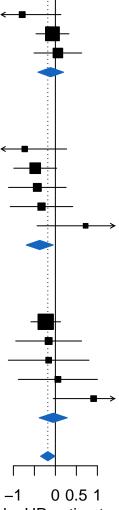
Primary = Melanoma

Nathanson, n = 24	-0.73 [-1.73; 0.27]
Liu, n = 121	-0.48 [-0.99; 0.03]
Riaz, n = 51	-0.43 [-1.12; 0.26]
Van_Allen, n = 42	-0.33 [-1.07; 0.41]
Hugo, $n = 27$	0.72 [-0.44; 1.88]
Total	-0.37 [-0.70; -0.05]
Heterogeneity: $\chi^2 = 4.13$ ($P = .39$), $I^2 = 3\%$ [0%: 80%]	

Primary = Other

Mariathasan, Bladder, n = 194 -0.23 [-0.58; 0.12] -0.16 [-0.94; 0.62] Fumet.2, Lung, n = 43Snyder, Ureteral, n = 25-0.16 [-1.12; 0.80] Mariathasan, Lymph_node, $n = 26 \ 0.06 \ [-0.88; \ 1.00]$ Mariathasan, Ureteral, n = 260.91 [-0.05; 1.87] Total -0.05 [-0.39; 0.28] Heterogeneity: $\chi_4^2 = 4.92$ (P = .30), $I^2 = 19\%$ [0%; 83%] Total -0.18 [-0.35; -0.01] Heterogeneity: $\chi_{12}^2 = 13.44 \ (P = .34), \ I^2 = 11\% \ [0\%; 50\%]$ Test for overall effect: z = -2.03 (P = .04)

Test for subgroup differences: $\chi_2^2 = 2.11$ (P = .35)



logHR estimate