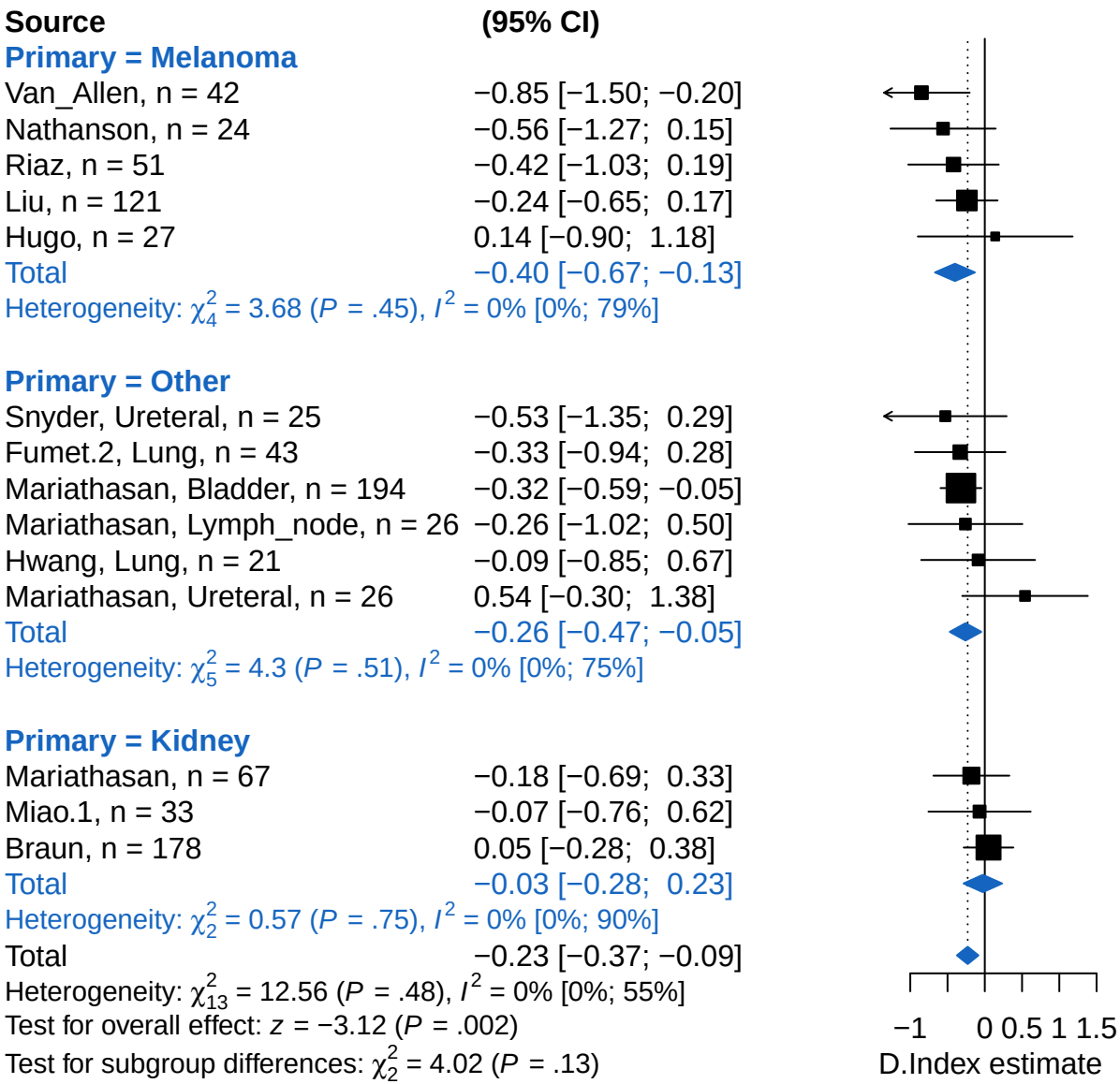
Source
(95% CI)
Primary = Melanoma
Van_Allen, n = 42
−0.85 [−1.50; −0.20]
Nathanson, n = 24
−0.56 [−1.27; 0.15]
Riaz, n = 51
−0.42 [−1.03; 0.19]
Liu, n = 121
−0.24 [−0.65; 0.17]
Hugo, n = 27
0.14 [−0.90; 1.18]
Total
−0.40 [−0.67; −0.13]
Heterogeneity: χ^2_4 = 3.68 (P = .45), I^2 = 0% [0%; 79%]
Primary = Other
Snyder, Ureteral, n = 25
−0.53 [−1.35; 0.29]
Fumet.2, Lung, n = 43
−0.33 [−0.94; 0.28]
Mariathasan, Bladder, n = 194
−0.32 [−0.59; −0.05]
Mariathasan, Lymph_node, n = 26
−0.26 [−1.02; 0.50]
Hwang, Lung, n = 21
−0.09 [−0.85; 0.67]
Mariathasan, Ureteral, n = 26
0.54 [−0.30; 1.38]
Total
−0.26 [−0.47; −0.05]
Heterogeneity: χ^2_5 = 4.3 (P = .51), I^2 = 0% [0%; 75%]
Primary = Kidney
Mariathasan, n = 67
−0.18 [−0.69; 0.33]
Miao.1, n = 33
−0.07 [−0.76; 0.62]
Braun, n = 178
0.05 [−0.28; 0.38]
Total
−0.03 [−0.28; 0.23]
Heterogeneity: χ^2_2 = 0.57 (P = .75), I^2 = 0% [0%; 90%]
Total
−0.23 [−0.37; −0.09]
Heterogeneity: χ^2_{13} = 12.56 (P = .48), I^2 = 0% [0%; 55%]
Test for overall effect: z = −3.12 (P = .002)
Test for subgroup differences: χ^2_2 = 4.02 (P = .13)
−1
0
0.5
1
1.5
D.Index estimate