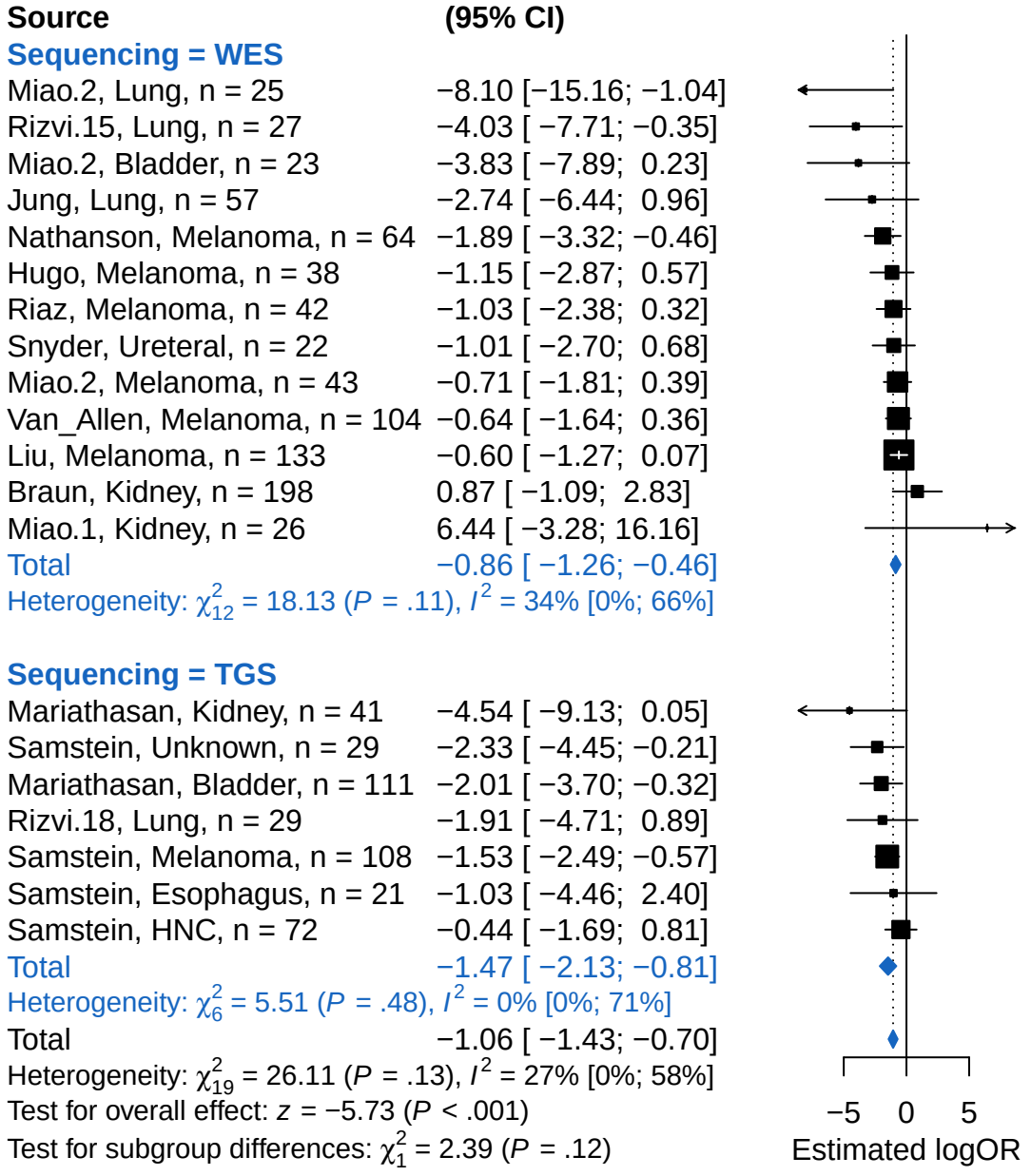
Source
(95% CI)
Sequencing = WES
Miao.2, Lung, n = 25 −8.10 [−15.16; −1.04]
Rizvi.15, Lung, n = 27 −4.03 [−7.71; −0.35]
Miao.2, Bladder, n = 23 −3.83 [−7.89; 0.23]
Jung, Lung, n = 57 −2.74 [−6.44; 0.96]
Nathanson, Melanoma, n = 64 −1.89 [−3.32; −0.46]
Hugo, Melanoma, n = 38 −1.15 [−2.87; 0.57]
Riaz, Melanoma, n = 42 −1.03 [−2.38; 0.32]
Snyder, Ureteral, n = 22 −1.01 [−2.70; 0.68]
Miao.2, Melanoma, n = 43 −0.71 [−1.81; 0.39]
Van_Allen, Melanoma, n = 104 −0.64 [−1.64; 0.36]
Liu, Melanoma, n = 133 −0.60 [−1.27; 0.07]
Braun, Kidney, n = 198 0.87 [−1.09; 2.83]
Miao.1, Kidney, n = 26 6.44 [−3.28; 16.16]
Total −0.86 [−1.26; −0.46]
Heterogeneity: χ^2_{12} = 18.13 (P = .11), I^2 = 34% [0%; 66%]
Sequencing = TGS
Mariathasan, Kidney, n = 41 −4.54 [−9.13; 0.05]
Samstein, Unknown, n = 29 −2.33 [−4.45; −0.21]
Mariathasan, Bladder, n = 111 −2.01 [−3.70; −0.32]
Rizvi.18, Lung, n = 29 −1.91 [−4.71; 0.89]
Samstein, Melanoma, n = 108 −1.53 [−2.49; −0.57]
Samstein, Esophagus, n = 21 −1.03 [−4.46; 2.40]
Samstein, HNC, n = 72 −0.44 [−1.69; 0.81]
Total −1.47 [−2.13; −0.81]
Heterogeneity: χ^2_6 = 5.51 (P = .48), I^2 = 0% [0%; 71%]
Total −1.06 [−1.43; −0.70]
Heterogeneity: χ^2_{19} = 26.11 (P = .13), I^2 = 27% [0%; 58%]
Test for overall effect: z = −5.73 (P < .001)
Test for subgroup differences: χ^2_1 = 2.39 (P = .12)
−5
0
5
Estimated logOR