Source	(95% CI)
Sequencing = TPM	
Miao.1, Kidney, n = 28	−1.71 [−4.30; 0.88]
Braun, Kidney, n = 139	−0.20 [−1.69; 1.29]
Riaz, Melanoma, n = 33	−0.12 [−2.14; 1.90]
Mariathasan, Kidney, n = 46	0.75 [−2.60; 4.10]
Fumet.2, Lung, n = 41	1.02 [−2.39; 4.43]
Mariathasan, Bladder, n = 133	1.54 [0.09; 2.99]
Van_Allen, Melanoma, n = 39	2.11 [−0.58; 4.80]
Jung, Lung, n = 26	3.78 [0.04; 7.52]
Total	0.66 [−0.32; 1.63]

Heterogeneity: $\chi^2_7 = 10.32$ ($P = .17$), $I^2 = 32\%$ [0%; 70%]

Source	(95% CI)
Sequencing = FPKM	
Nathanson, Melanoma, n = 24	0.49 [−1.53; 2.51]
Liu, Melanoma, n = 112	0.99 [−0.17; 2.15]
Hugo, Melanoma, n = 27	2.07 [−0.34; 4.48]
Total	1.04 [0.12; 1.97]

Heterogeneity: $\chi^2_2 = 0.99$ ($P = .61$), $I^2 = 0\%$ [0%; 90%]

Total: 0.80 [0.18; 1.41]

Heterogeneity: $\chi^2_{10} = 11.77$ ($P = .30$), $I^2 = 15\%$ [0%; 56%]

Test for overall effect: $z = 2.55$ ($P = .01$)

Test for subgroup differences: $\chi^2_1 = 0.32$ ($P = .57$)

Estimated logOR: −2 0 1 2 3 4