Source	(95% CI)
Sequencing = TPM	
Miao.1, Kidney, n = 28	−0.06 [−1.39; 1.27]
Riaz, Melanoma, n = 33	0.29 [−0.94; 1.52]
Fumet.1, Lung, n = 39	0.42 [−0.78; 1.62]
Fumet.2, Lung, n = 41	0.90 [−0.35; 2.15]
Mariathasan, Bladder, n = 133	1.22 [0.57; 1.87]
Snyder, Ureteral, n = 22	1.23 [−0.24; 2.70]
Braun, Kidney, n = 139	1.28 [0.54; 2.02]
Van_Allen, Melanoma, n = 39	1.99 [0.28; 3.70]
Jung, Lung, n = 26	2.11 [0.09; 4.13]
Mariathasan, Kidney, n = 46	5.59 [1.34; 9.84]
Total	1.06 [0.70; 1.42]

Heterogeneity: $\chi^2_9 = 12.53$ ($P = .18$), $I^2 = 28\%$ [0%; 66%]

Sequencing = FPKM	
Nathanson, Melanoma, n = 24	0.49 [−0.84; 1.82]
Hugo, Melanoma, n = 27	1.24 [−0.17; 2.65]
Liu, Melanoma, n = 112	1.59 [0.83; 2.35]
Total	1.28 [0.63; 1.93]

Heterogeneity: $\chi^2_2 = 1.98$ ($P = .37$), $I^2 = 0\%$ [0%; 90%]

Total   1.12 [0.82; 1.43]

Heterogeneity: $\chi^2_{12} = 14.98$ ($P = .24$), $I^2 = 20\%$ [0%; 58%]

Test for overall effect: $z = 7.18$ ($P < .001$)

Test for subgroup differences: $\chi^2_1 = 0.33$ ($P = .56$)

0 1 2 3 4 5 6

Estimated logOR