



Supplementary Figure 6: Association between whole genome doubling (WGD) and mirrored subclonal allelic imbalance (MSAI). Median proportions by Patient WGD Status (cWGD = 0.02, n=54 tumours; sWGD = 0.07, n=13 tumours; nWGD = 0, n=32 tumours). Kruskal-Wallis p-value is shown ($p=2e-04$). Data is shown for n=99 tumours from the pan-cancer cohort showcased in Figure 5 for which MEDICC2 was used to infer WGD status. Proportion of genome data is assessed by Refphase. cWGD = clonal WGD; sWGD = subclonal WGD; nWGD = non-WGD.