

Extended Data Figure 4 | **Endogenous aldehydes mutate the HSC genome.** Circos plots showing the mutations observed in all sequenced HSC clones (wild type, n = 3; $Aldh2^{-/-}$, n = 3; $Fancd2^{-/-}$, n = 4; and $Aldh2^{-/-}$ $Fancd2^{-/-}$, n = 5 HSC genomes). Substitutions, indels and rearrangements are plotted.