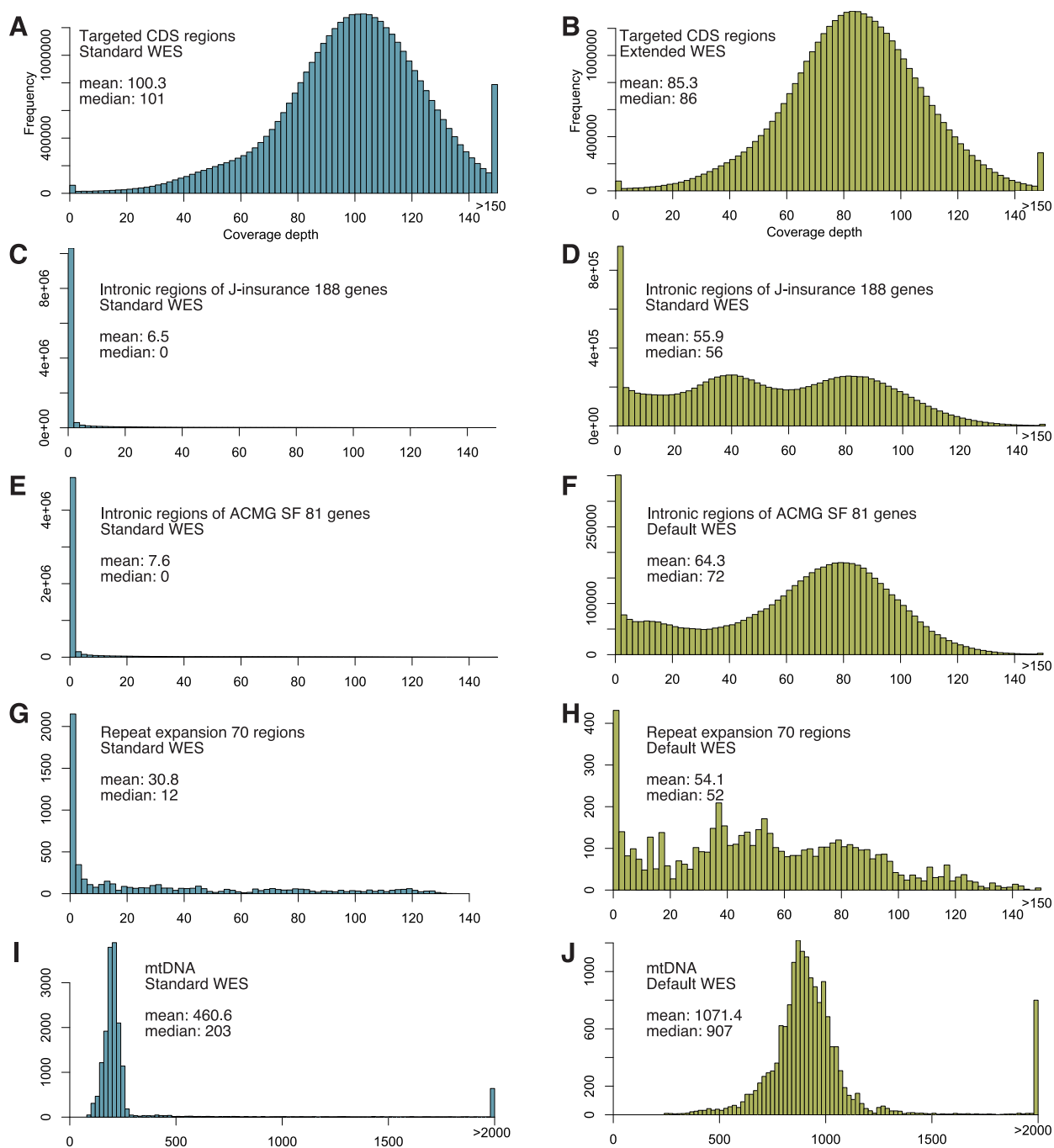




Supplementary Fig. S5 Distribution of coverage depth across different genomic regions in standard and extended WES for HG002 sample. All WES datasets were normalized to 9 Gb of sequencing data prior to comparative analysis. **A, C, E, G, I** Coverage depth distribution for each genomic region in standard WES. **B, D, F, H, J** Coverage depth distribution for each genomic region in extended WES