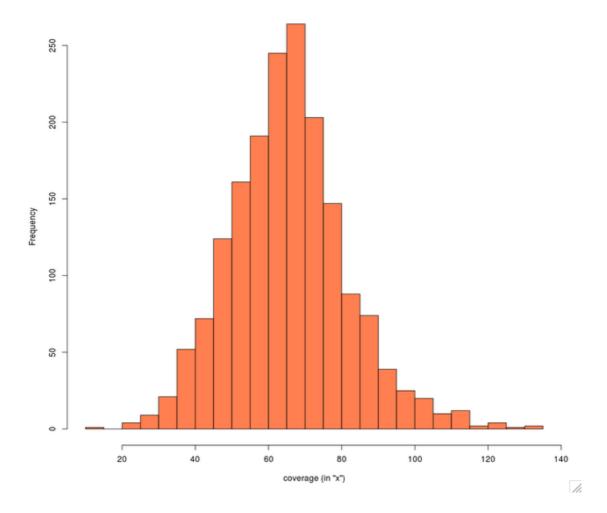
distribution of proband coverages



Supplementary Figure 19: Distribution of proband coverages: We measured coverage in exomes using samtools flagstat to generate the number QC-passed and mapped reads without read-duplicates, multipled this by the sequence read length (75) and divided by the number of bases targeted in the exome, 90 Mb. The mean exome coverage was 69x.