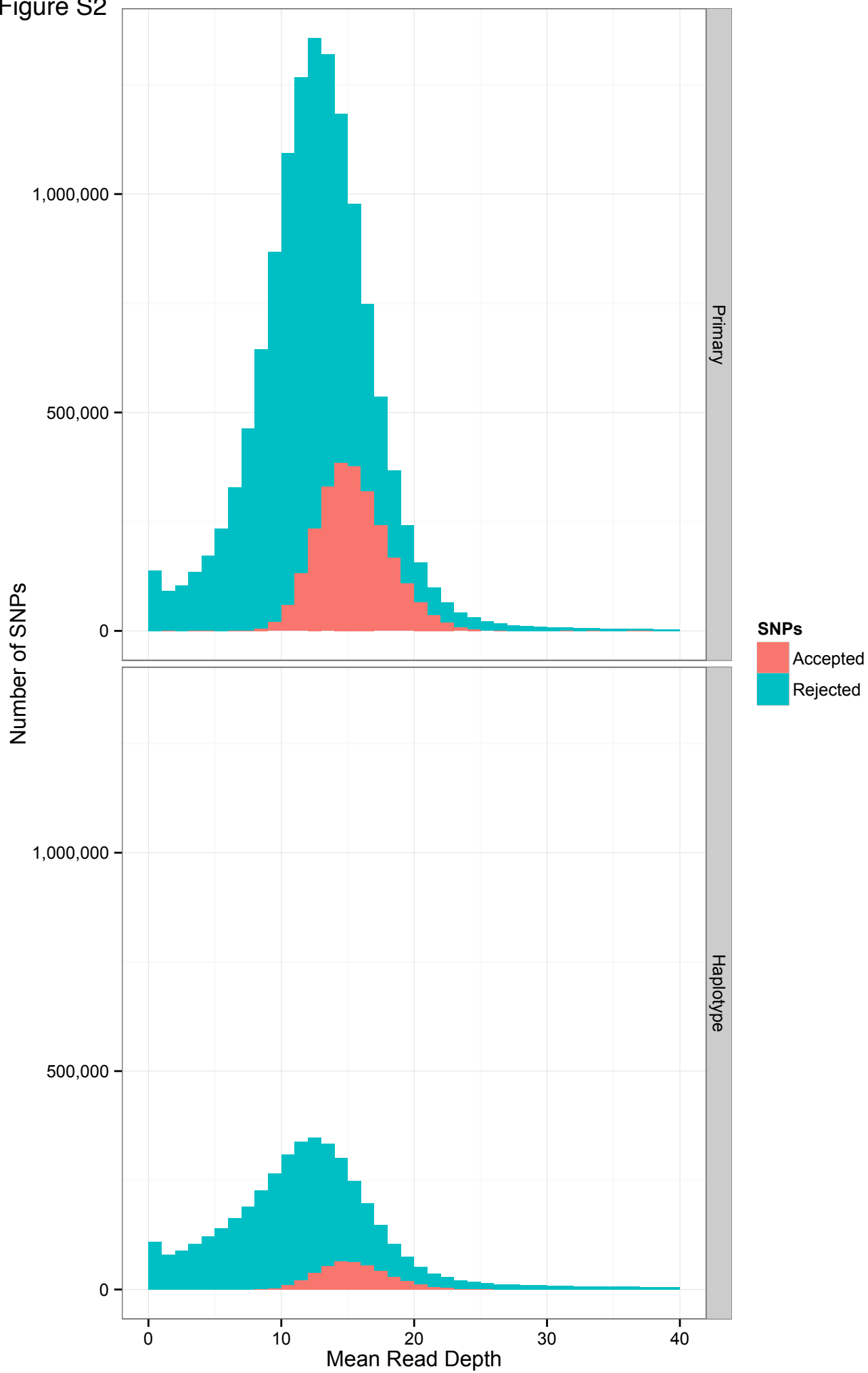


Figure S2



**Figure S2** Stacked bar charts of mean read depths for accepted and rejected SNPs. Read depths calculated per marker, taking a mean for each marker across all individuals. Distributions truncated at maximum 40 reads; maximum read depth was 427.