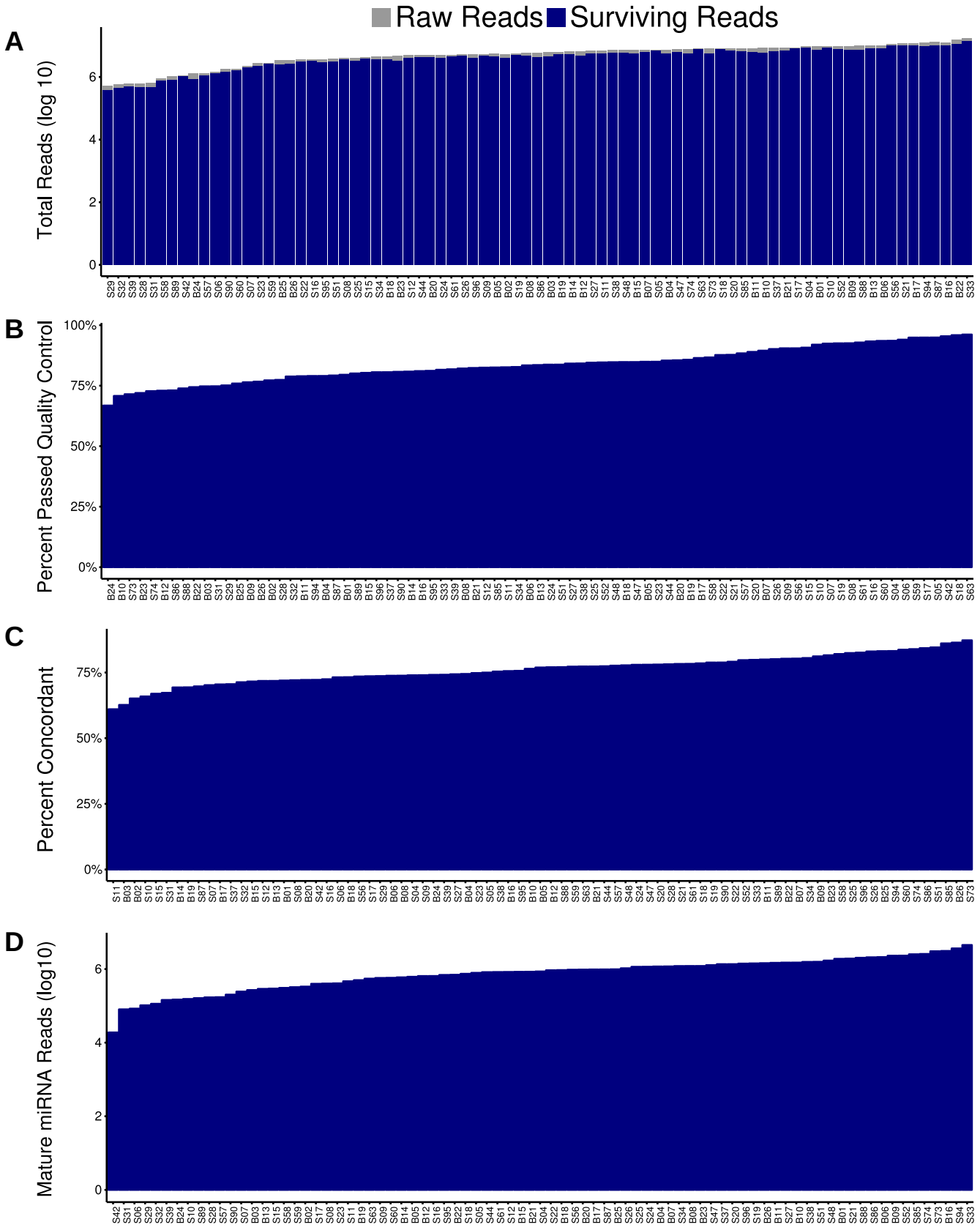

Supplementary Fig. 2. Bar plot of quality control of raw reads. **a.** Total number of raw reads on log10 scale per sample (grey) and percent of reads surviving quality control (blue). **b.** Percent of reads surviving quality control per sample. Bar plot of **c.** percent of concordant paired-end reads per sample and **d.** total number of mapped mature miRNAs per sample on log10 scale.