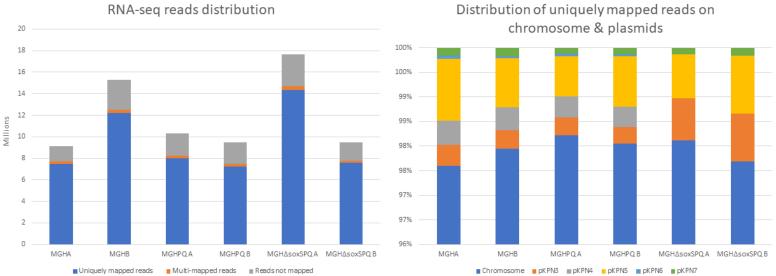
Supplementary figures



Distribution of uniquely mapped reads on

Figure S1. Distribution of RNA-seq reads in each individual dataset.