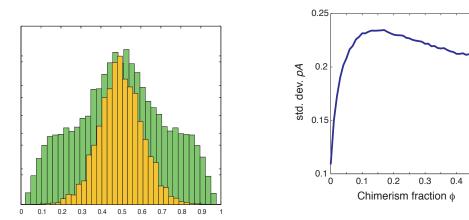
a b



Supplementary Figure 2 | Summary of Chimerism Analysis Data. a. Simulated distributions of the proportion of allele A reads out of the total number of reads at a locus for a marmoset with chimeric fraction $\Phi = 0.33$ (green) and $\Phi = 0$ (i.e., not chimeric; yellow). b. Standard deviation in the proportion of allele A reads across loci vs. chimerism fraction Φ , by simulation. Excess variance in pA is expected for reads from chimeric samples.

0.5