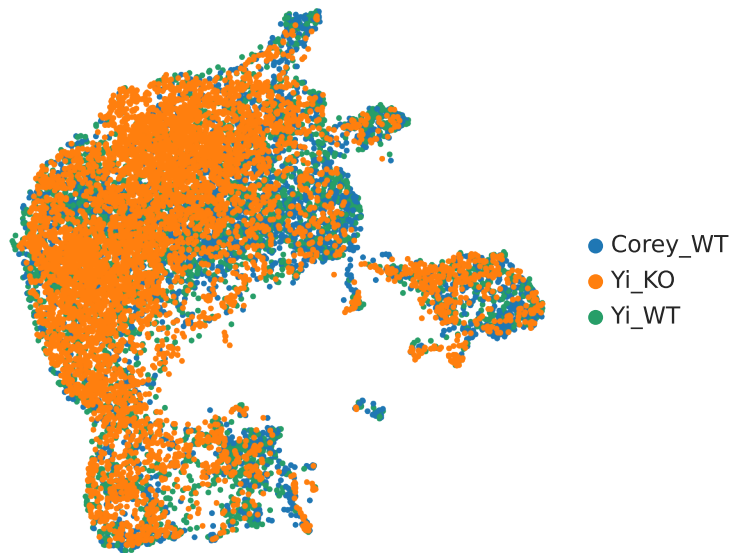
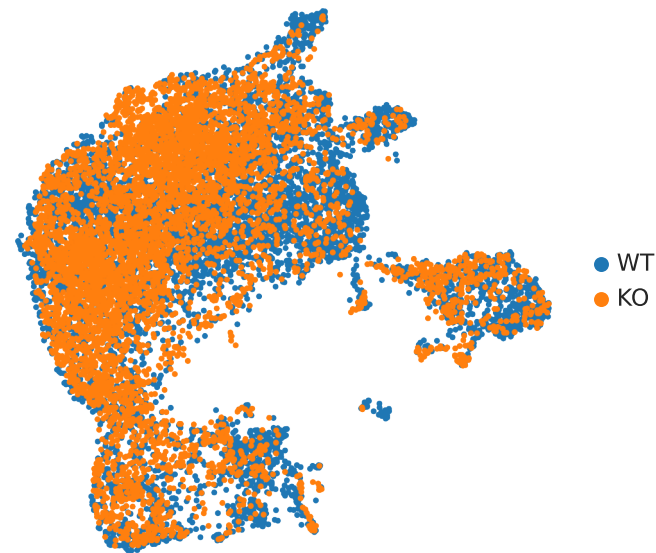


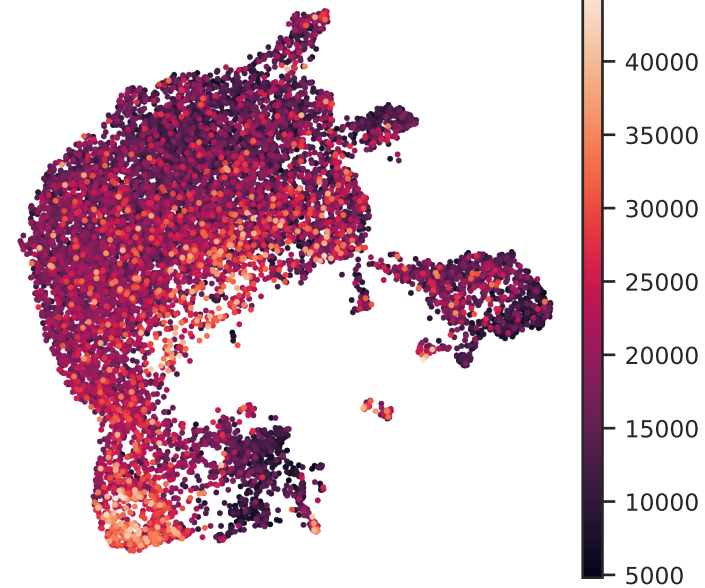
batch



genotype



total\_counts



n\_genes\_by\_counts

