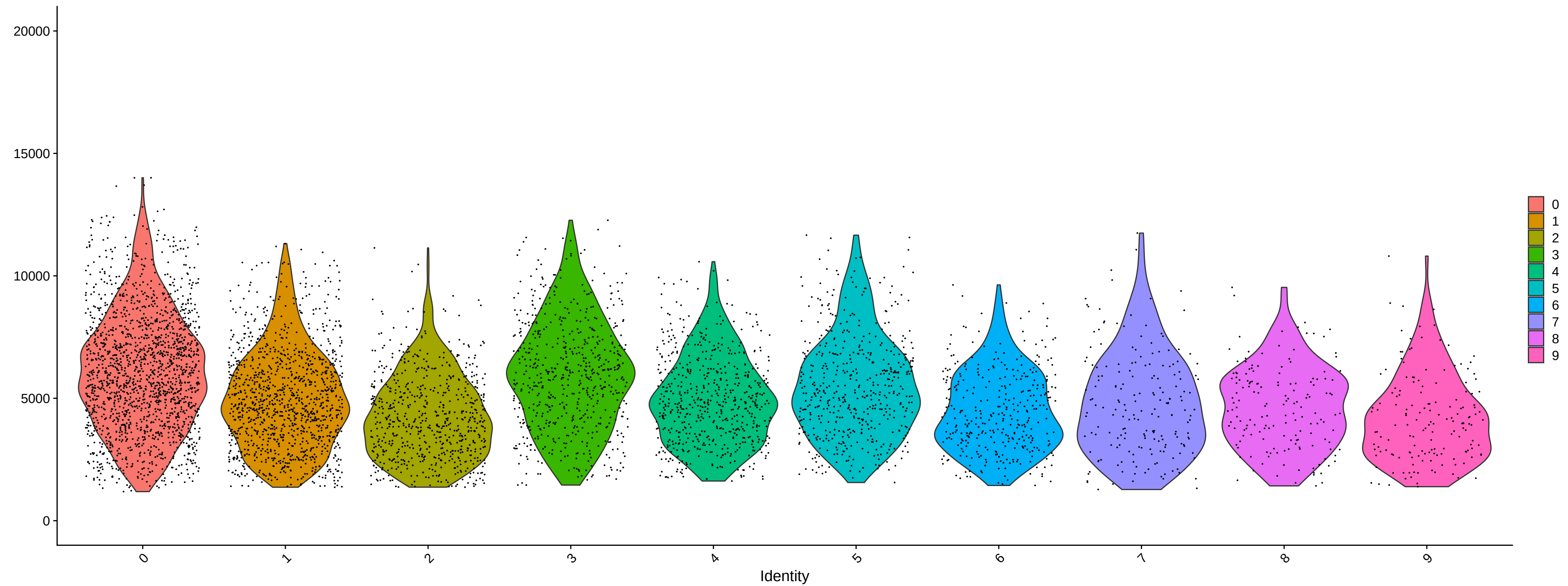
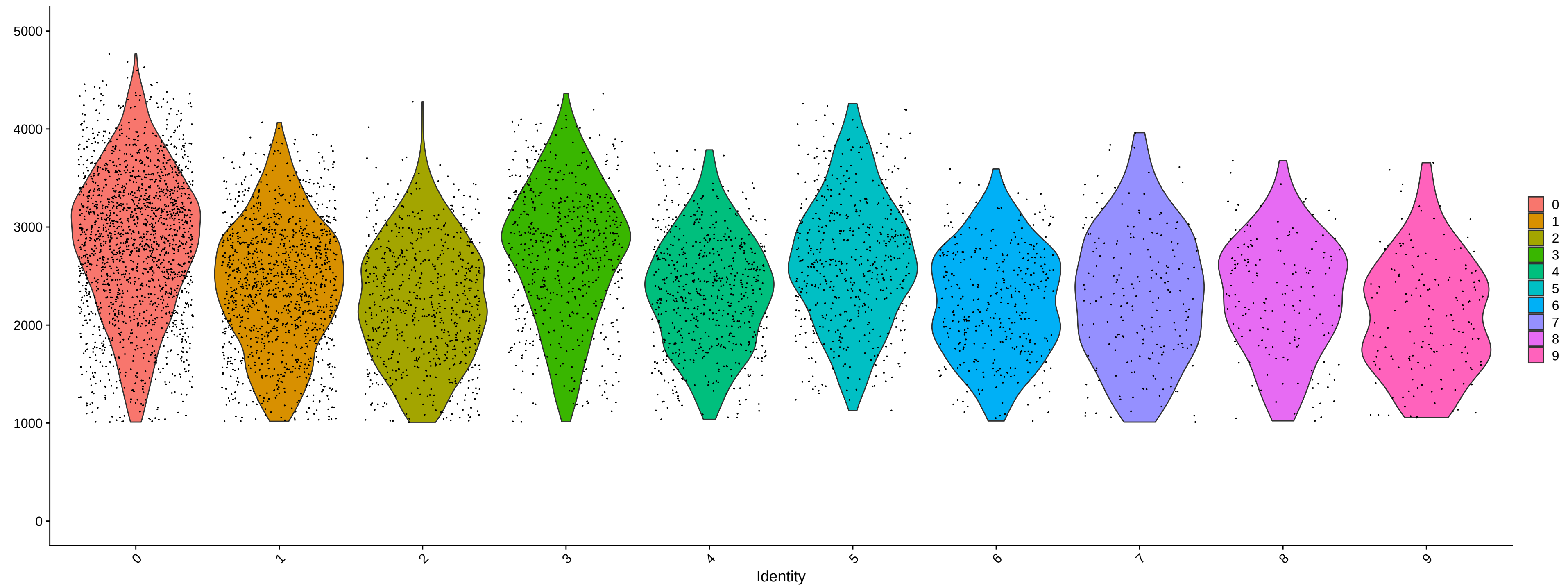


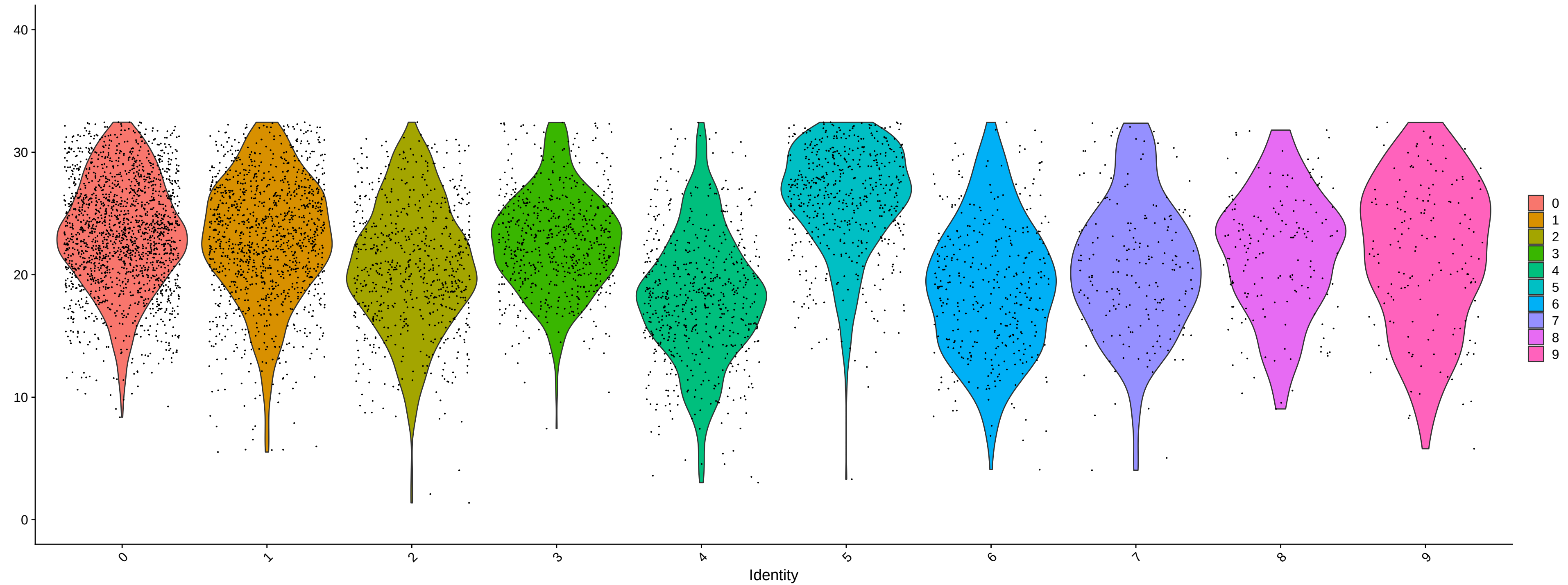
total UMI counts filtered cells



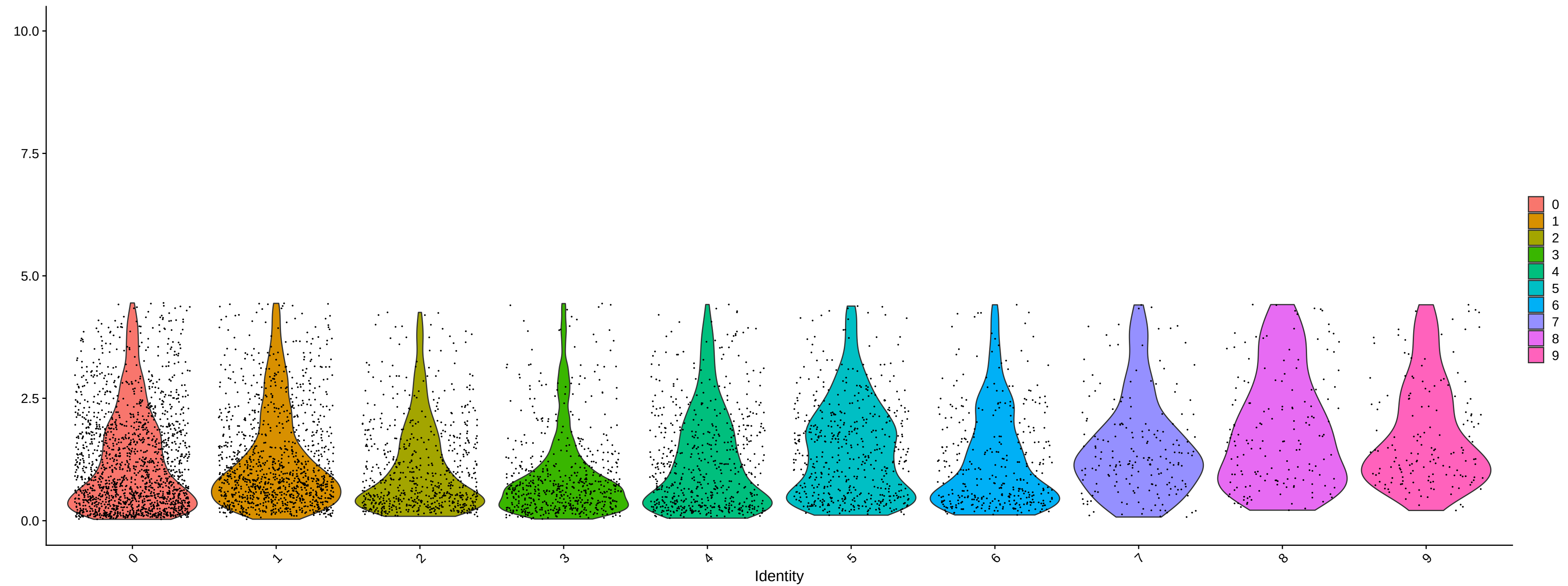
genes measured filtered cells



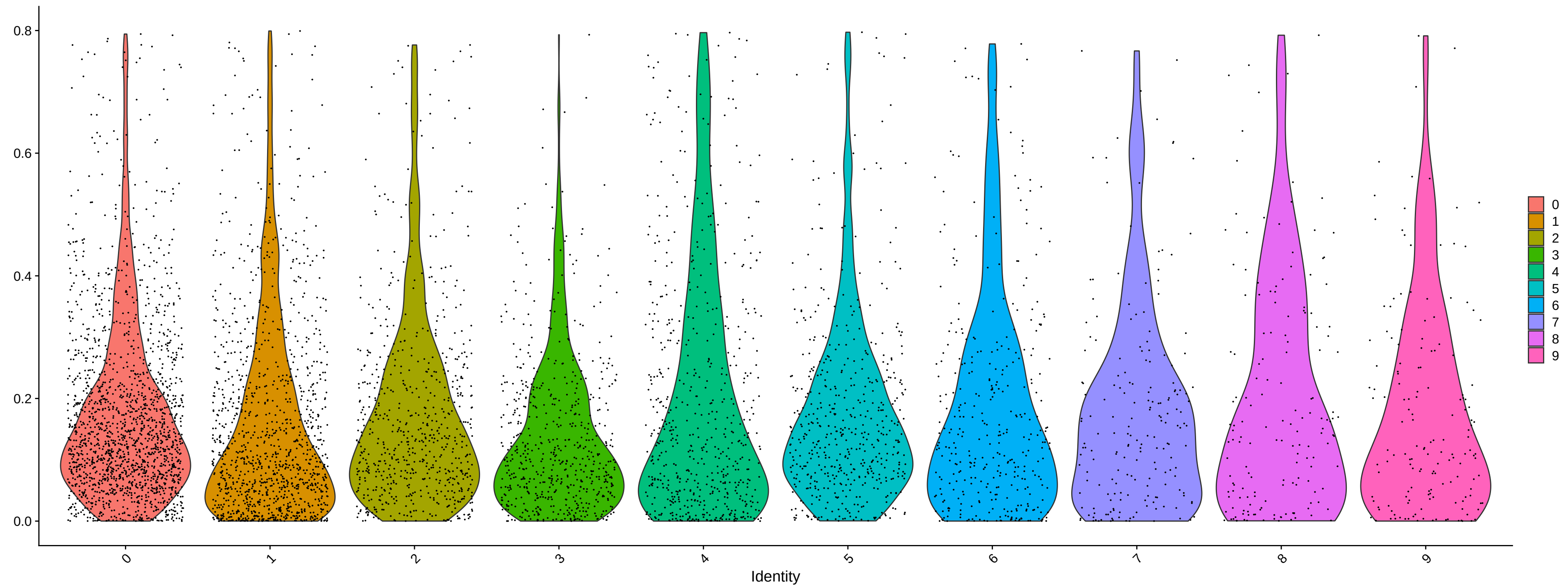
% mitochondrial reads filtered cells



% ERCC reads filtered cells



scDbfFinder_doublet_score



decontX_contamination

