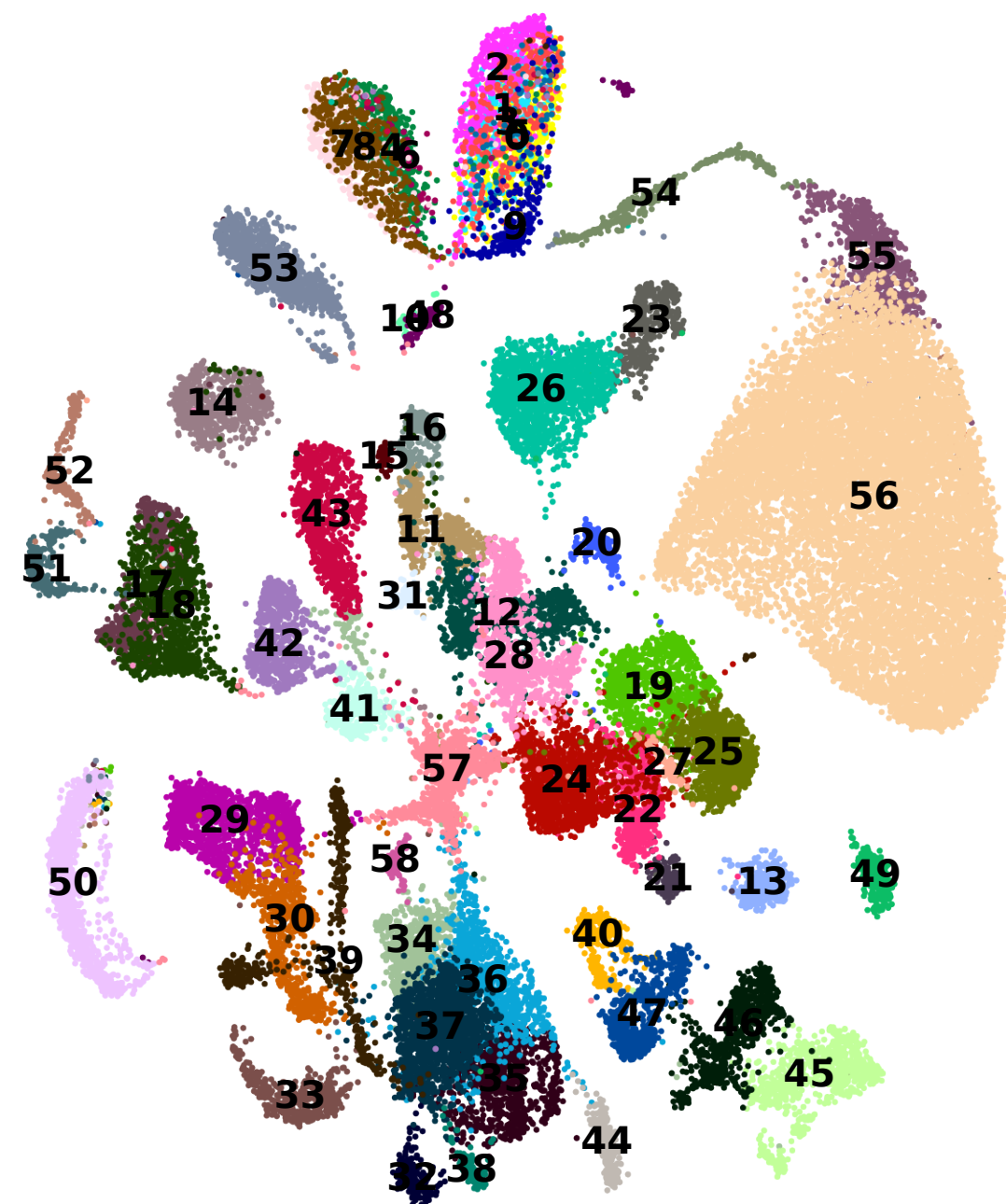
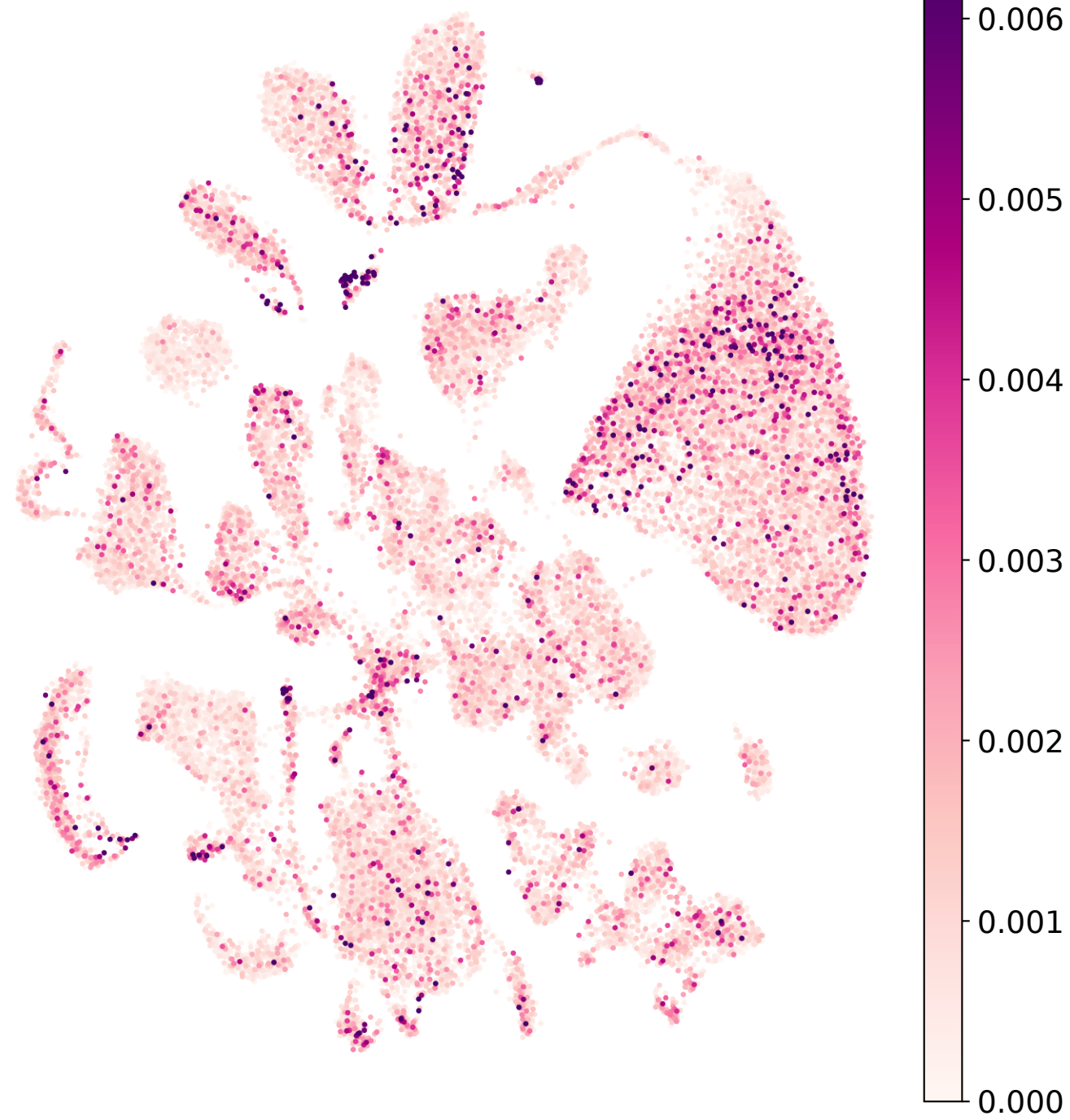


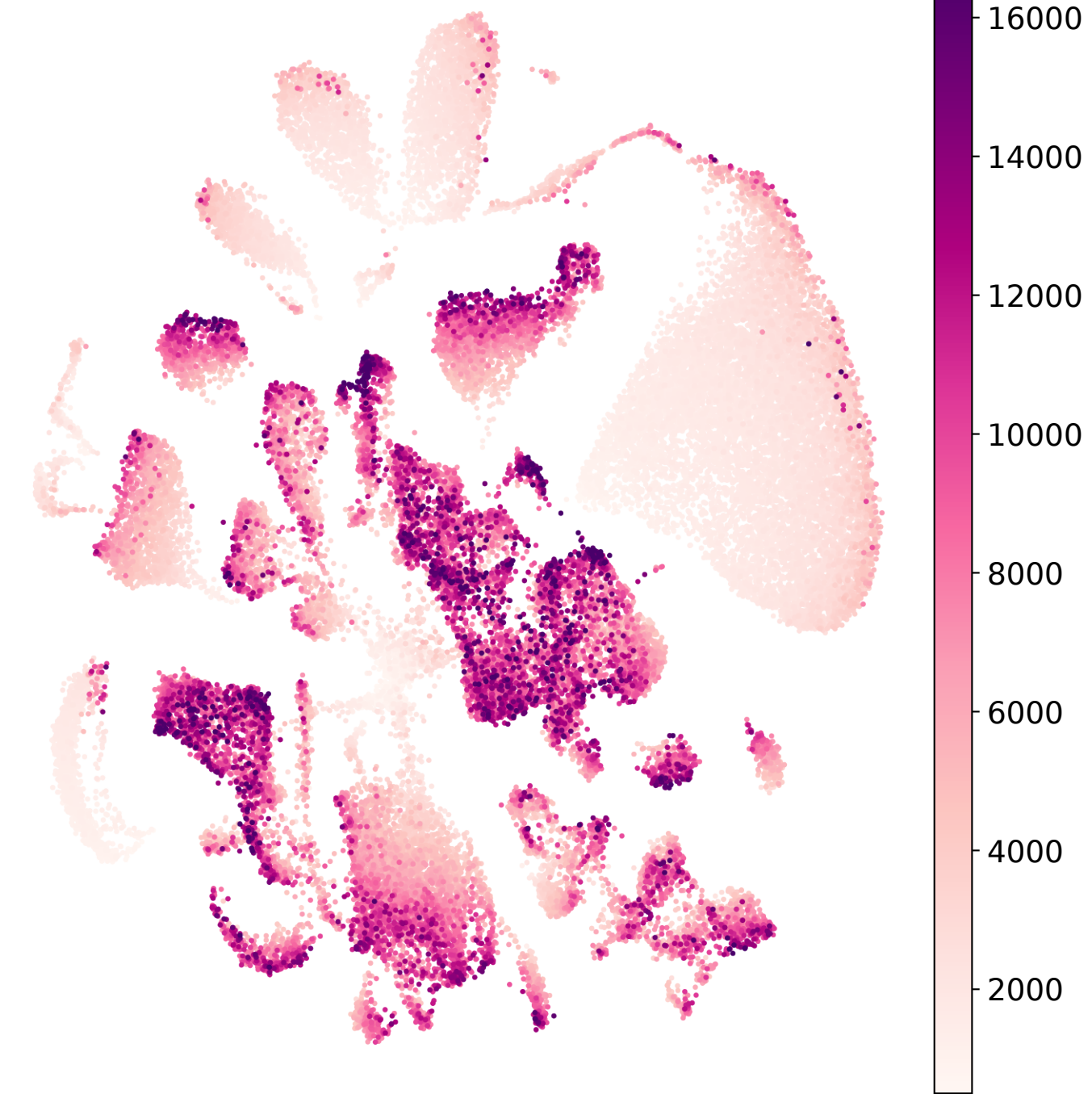
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mt_frac



total_counts



n_genes

