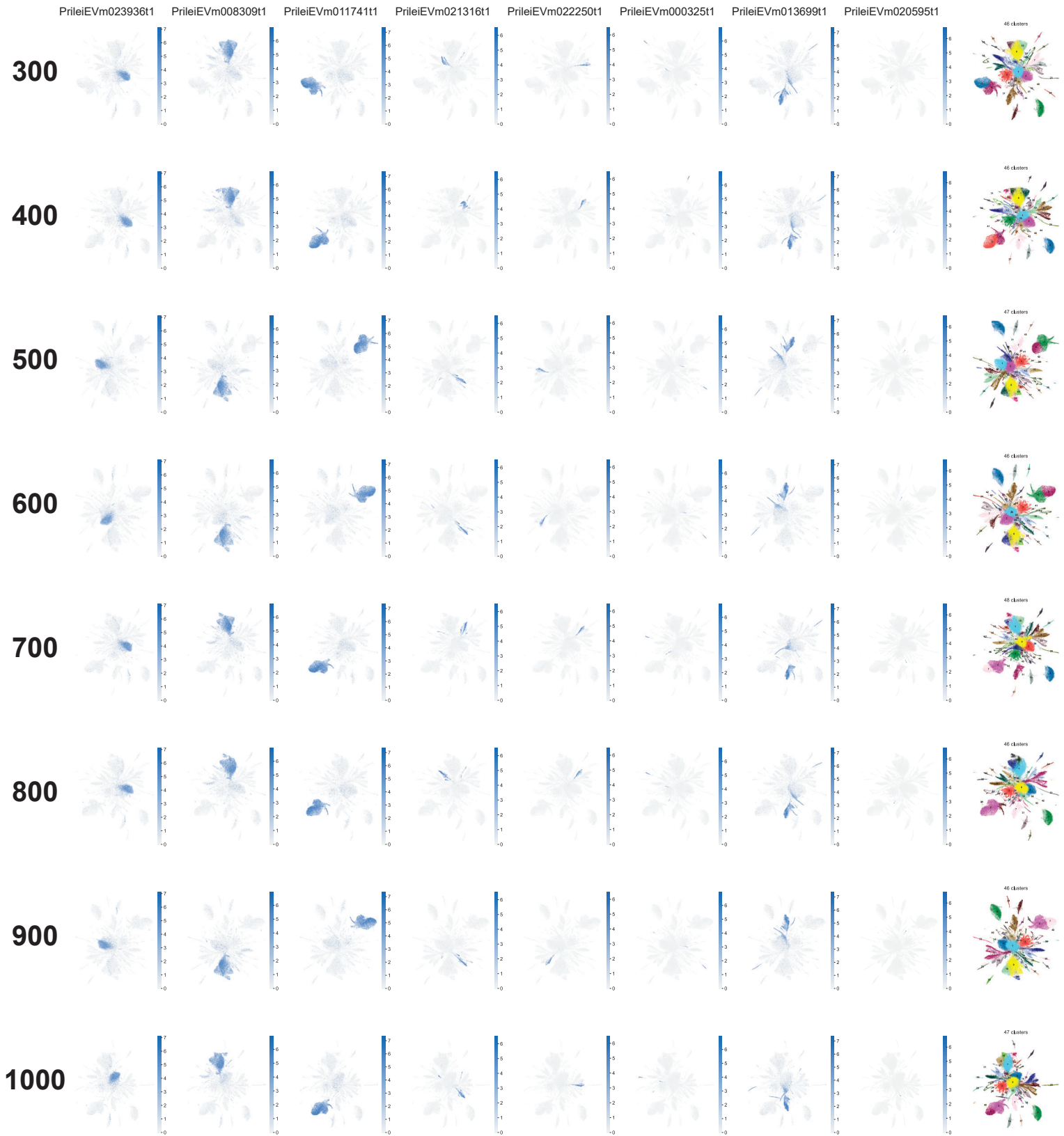


minimum gene counts



maximum number of genes



maximum number of counts

