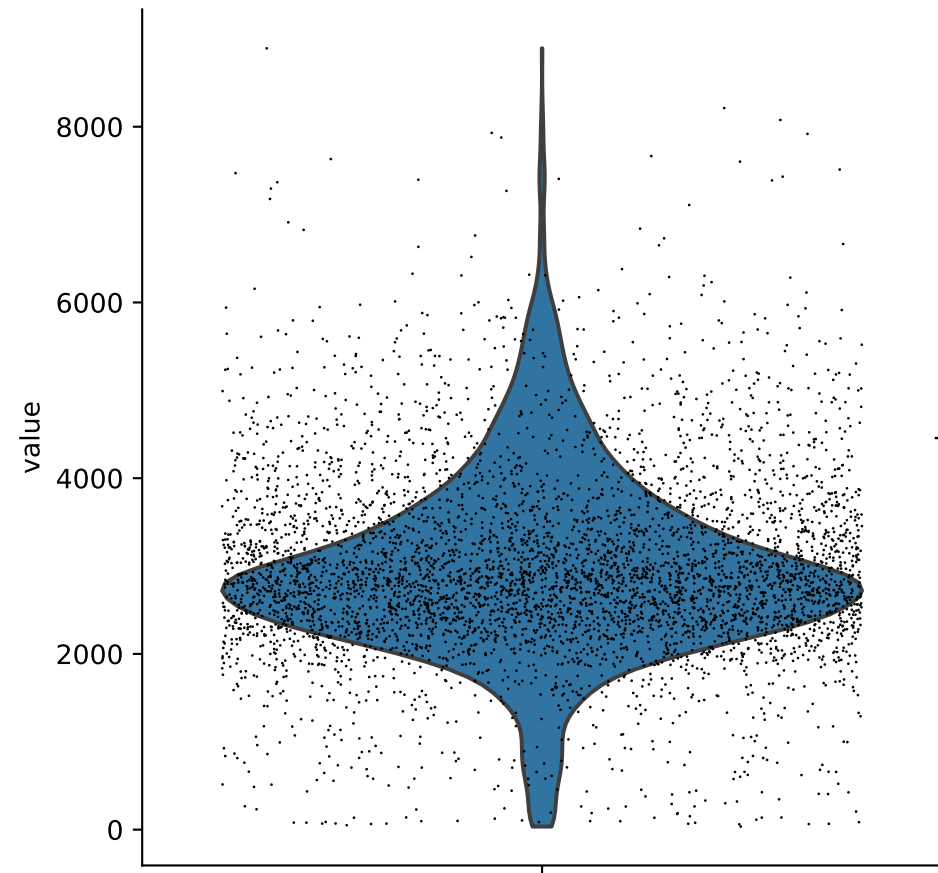
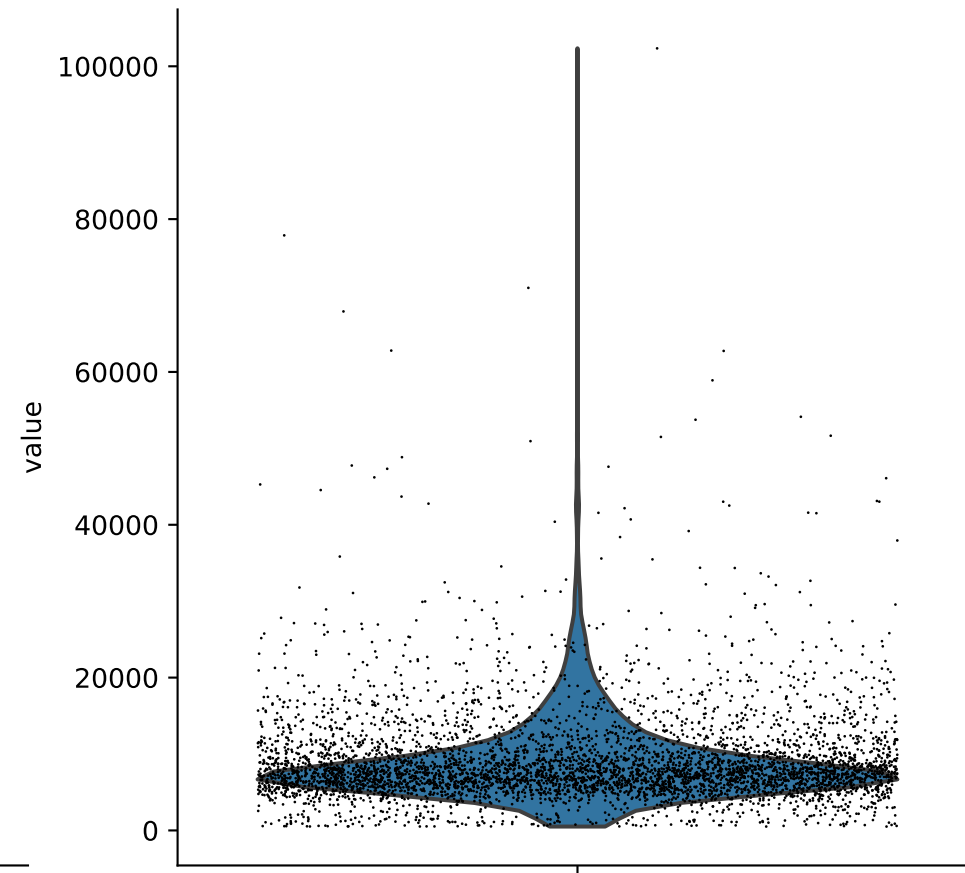


n\_genes\_by\_counts



total\_counts



pct\_counts\_mt

