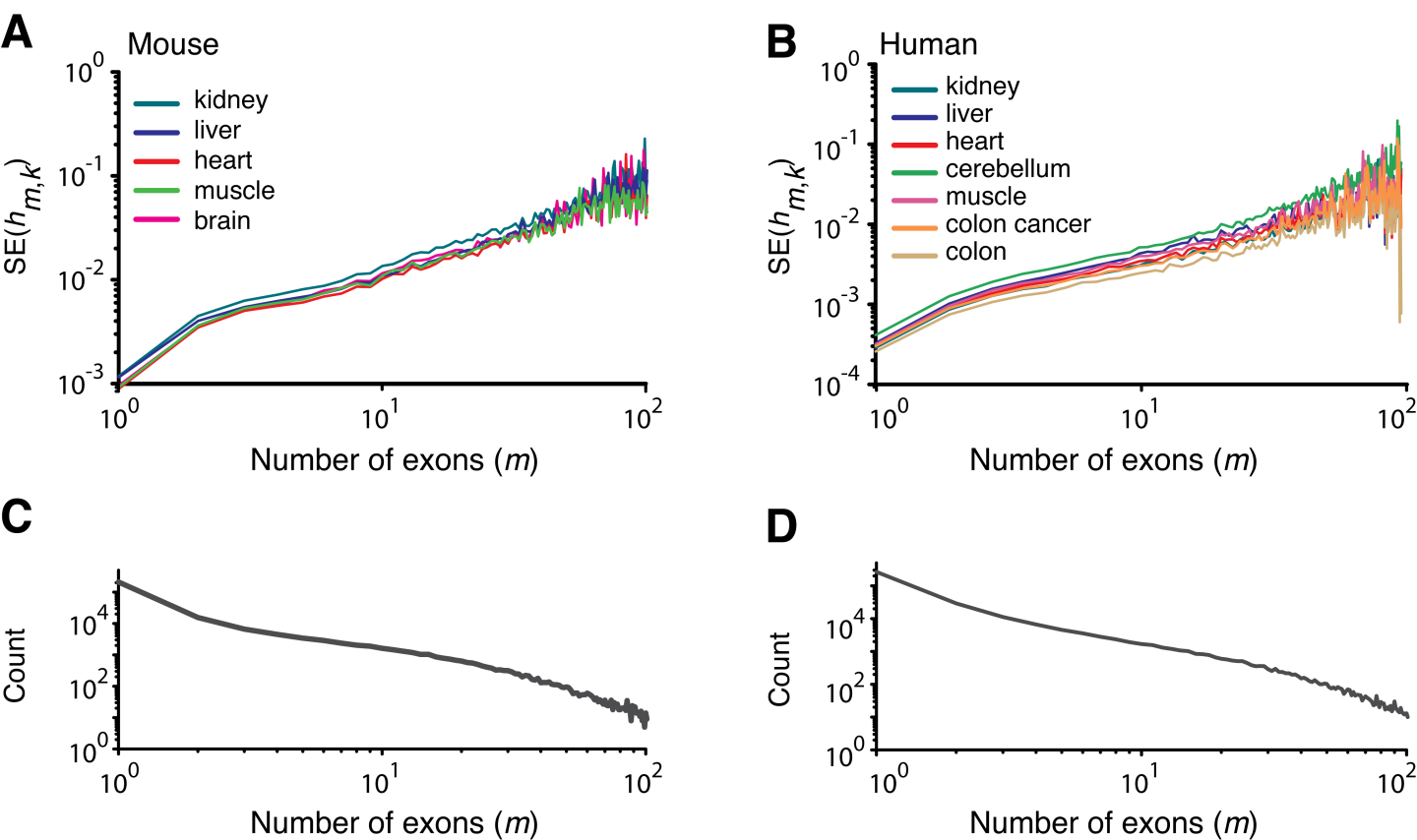# Figure S4

This figure provides complementary data to Figure 6.
A-B. Standard error of $h_{m,k}$ for mouse (A) and human (B).
C-D. Number of transcripts (count) with a given number of exons in mouse (C) and human (D).