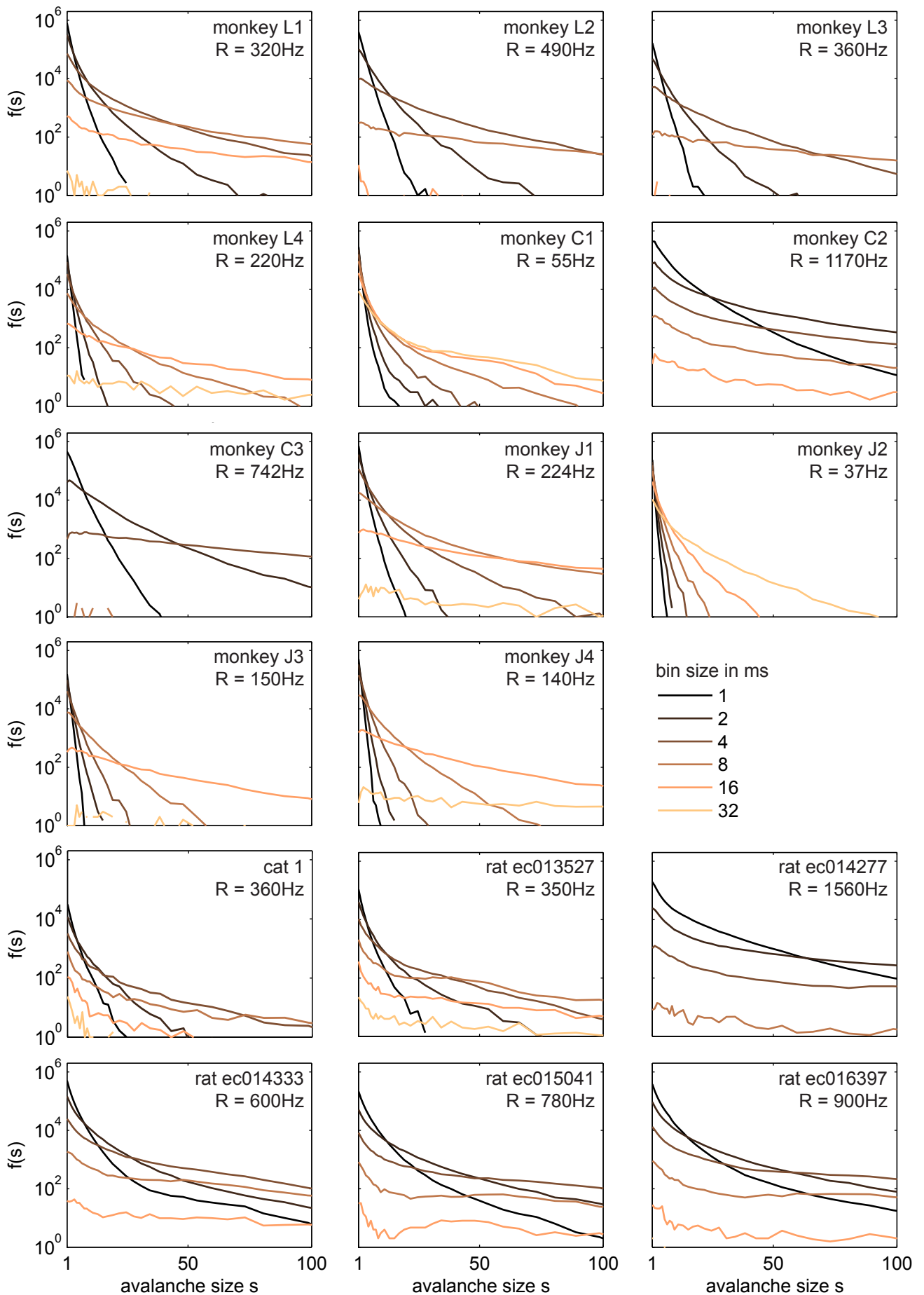


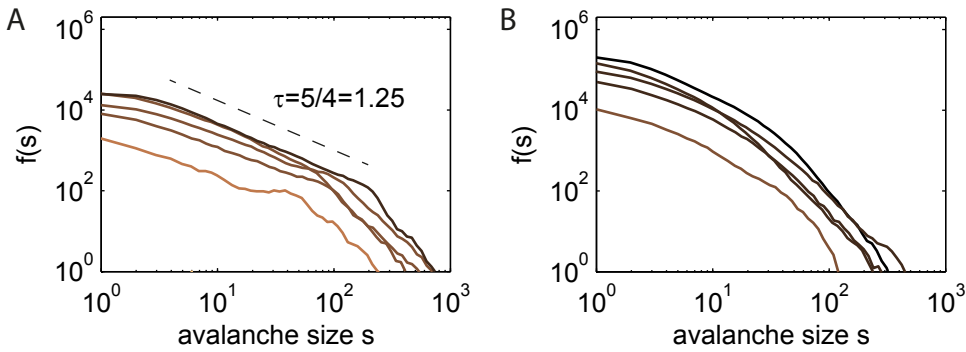
**Figure S1**

These panels show the spike  $f(s)$  for each of the *in vivo* recordings. The experiment ID and the population rate  $R$  is indicated in each plot. For the rat and cat recordings, the name is the same one used on the CRCNS data sharing platform (Blanche, 2009; Mizuseki et al., 2009), from which we obtained the data.



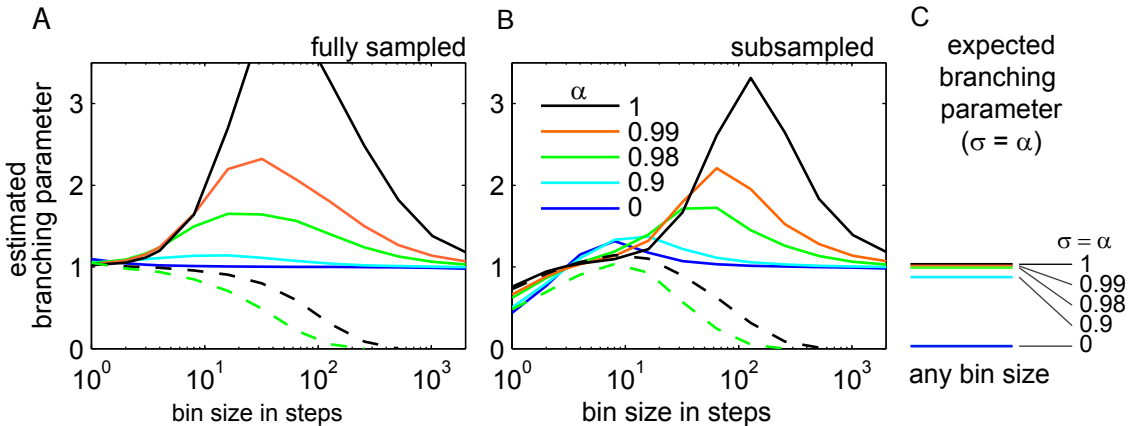
**Figure S2**

This figure is the same as Figure S1, but with log-linear axes instead of double logarithmic axes. It illustrates that none of the  $f(s)$  follows an exponential distribution (i.e. a straight line here).



**Figure S3**

**A.** For each of the *in vivo* rat recordings, the one avalanche size distributions  $f(s)$  was selected, which resembled a power law best. The bin sizes of these  $f(s)$  ranged between 2 ms and 8 ms, corresponding approximately to 3 average inter event intervals ( $\langle IEI \rangle$ ). Assuming a power law, the slope would be 1.25 or less (dashed line). **B.** For each of the  $f(s)$  in A, the corresponding  $f(s)$  with half the bin size was plotted (approximately 1.5  $\langle IEI \rangle$ ). These  $f(s)$  clearly deviated from power laws.



**Figure S4**

This figure shows the same as Figure 7 in the main text, but for the stochastic branching model. In this model, the branching parameter equals the synaptic strength, i.e.  $\sigma = \alpha$ . However, the *estimated* branching parameter  $\sigma^*$  differed from  $\alpha$ . **A, B.** For the driven models (full lines), the spike rate was fixed to  $r = 5$  Hz, while for the model with separation of time scales the drive was infinitesimal small ( $h \rightarrow 0$ ; dashed lines). **A.** Results for the fully sampled model. **B.** Results for the subsampled model ( $N = 100$  neurons). **C.** In the stochastic branching model, the branching parameter  $\sigma$  is a model parameter, and is therefore independent of the bin size. Its value is depicted using the same axes as in A, B.