Figure S25. **Distribution of nonzero subsamples per TF-gene edge.** Distribution of nonzero subsamples per TF-gene interaction, using per-gene and network average instability cutoffs of .05, .1, and .2. Results are shown for the No Prior TRN using 50 subsamples. The dotted-red line marks the instability cutoff in terms of nonzero subsamples, and the average model size at that cutoff appears as text in the upper right hand.