Figure 3: For each regulator in *E. coli* with known interactions inferred: (a) the number of interactions in the reference network (green) and the number of true positives in LeMoNe (red) and CLR (blue); (b) the number of interactions inferred (green) and the number of true positives (red) in LeMoNe, and the number of interactions inferred (yellow) and the number of true positives (blue) in CLR. LeMoNe and CLR networks are both at 30% precision threshold. Regulators are sorted by the difference $TP_{\text{LeMoNe}} - TP_{\text{CLR}}$. The total number of true positives is 171 for LeMoNe and 180 for CLR. For clarity, the $x$-axis in (a) is truncated, the true number of targets for Fis and Fnr is respectively 111 and 173. The number of interactions inferred only counts targets that belong to the reference network.