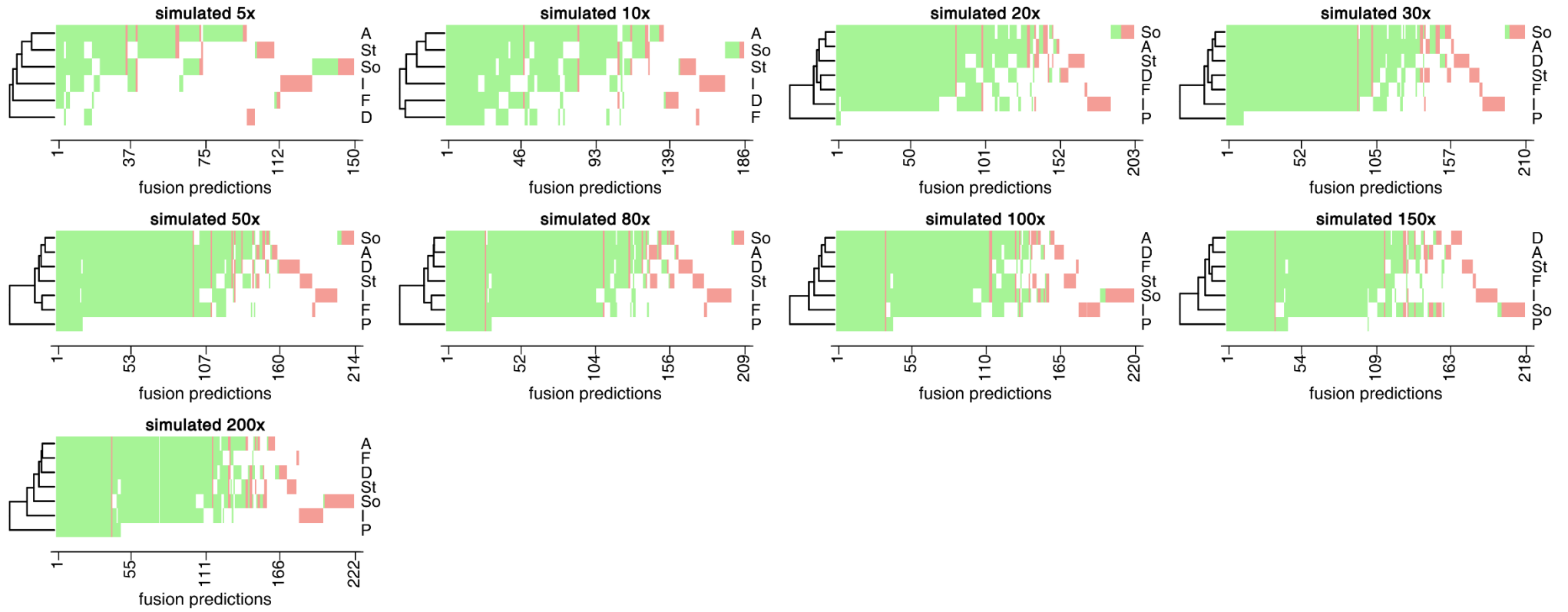
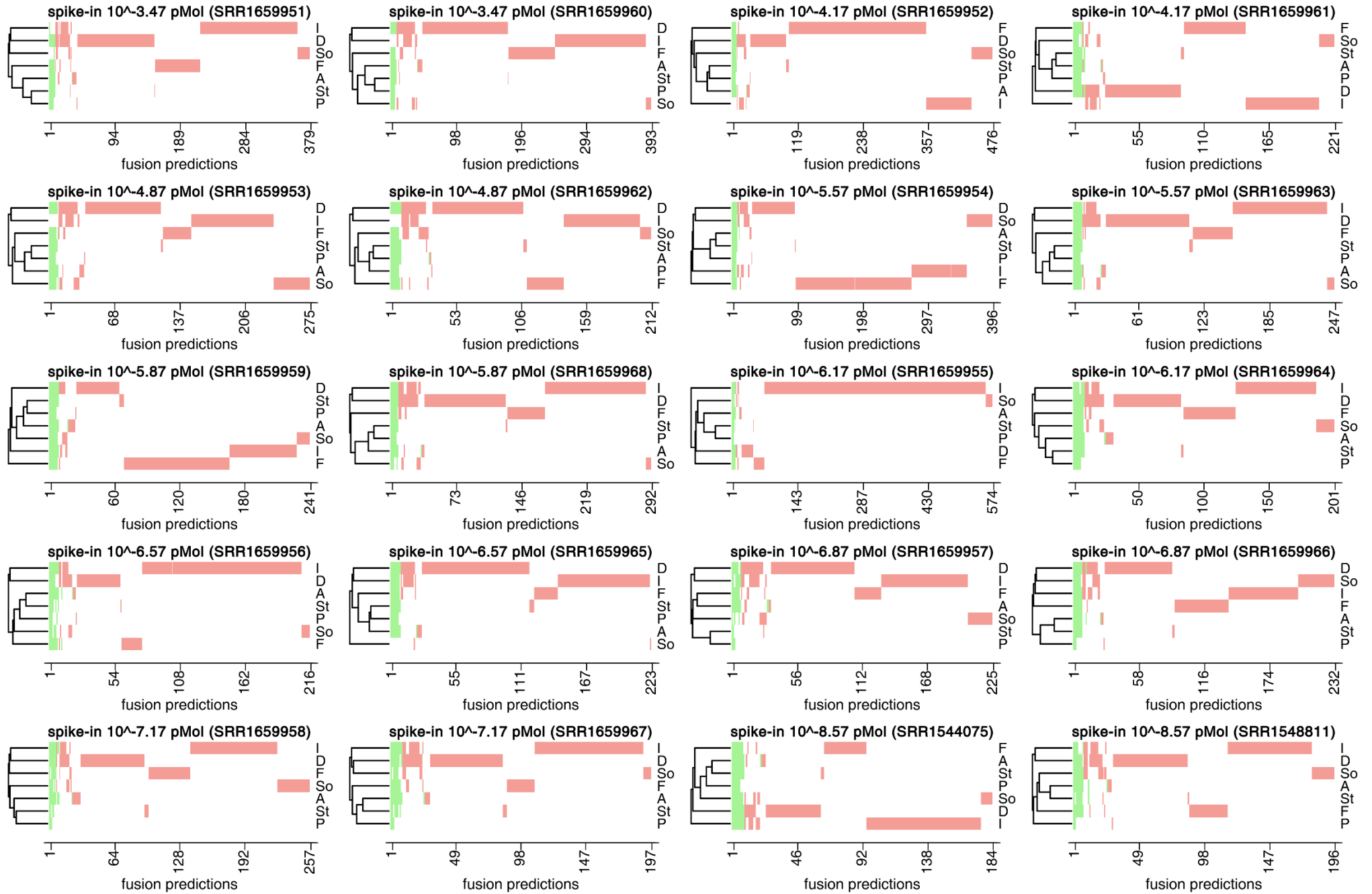
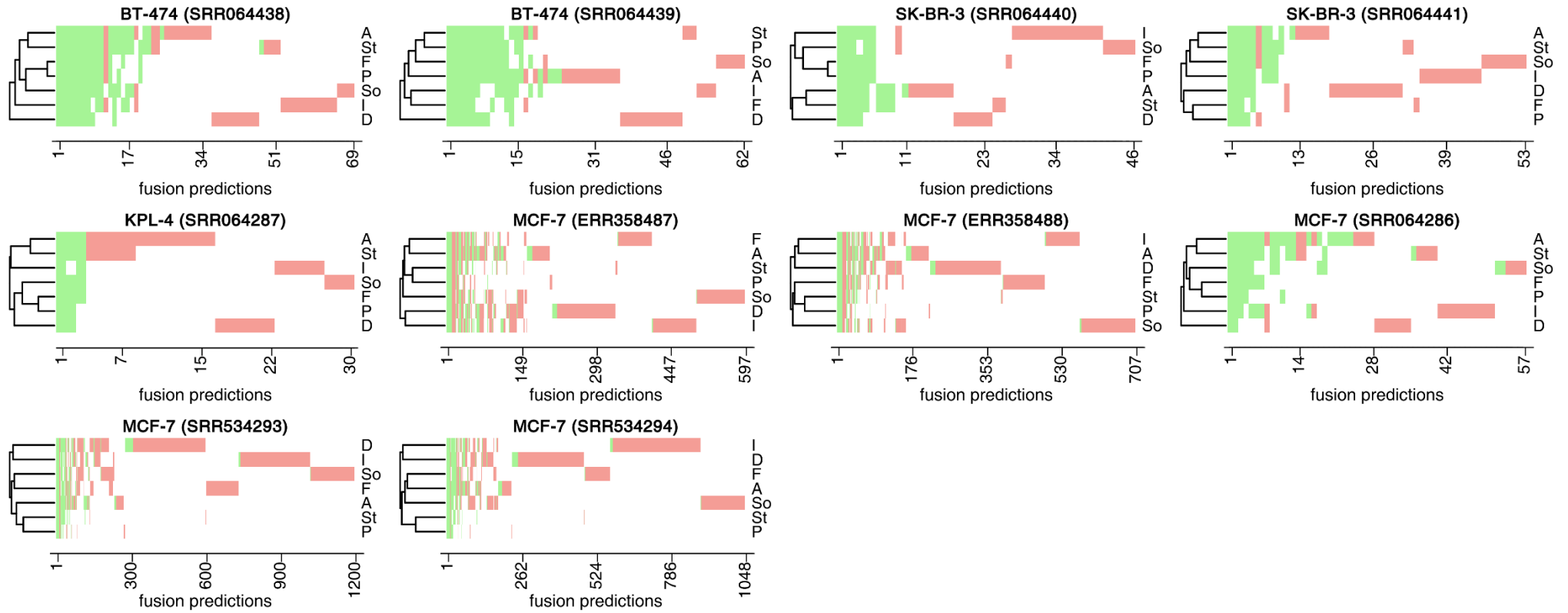


**A**

**B**

**C**

**Supplemental Figure S2: Fusion predictions shared between the methods.**

The fusion predictions are sorted in descending order from left to right by the number of methods that reported them. True positives are shown in green; false positives (simulated dataset) or non-validated predictions (spike-in and cell line datasets) are shown in red. Only gene-to-gene fusions were considered. A, Arriba; D, deFuse; F, FusionCatcher; I, InFusion; P, PRADA; So, SOAPfuse; St, STAR-Fusion. (A) Simulated data. (B) Spike-ins of synthetic fusions. (C) Cell lines.