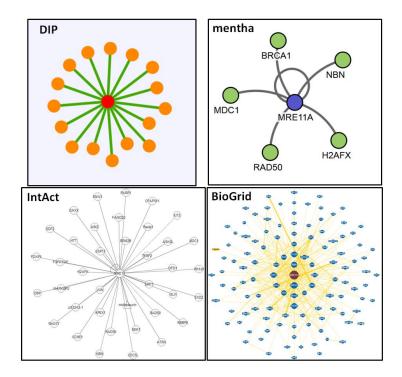
Supplementary figure 1. The query results of different resources are quite different using the human MRE11 gene as example. We queried DIP, mentha, BioGrid and IntAct for interaction data with human gene MRE11 (MRE11 homolog, double strand break repair nuclease).



Supplementary figure 2. MIST supports mapping of cross-species interactions. A. We mapped all the mouse-human PPI from 2591 publications to human genes and compared with independent human-human PPI data. **B.** There is significant overlap between mouse-human and human-human interactions. The overlaps are enriched for highly conserved genes measured by DIOPT score.

