Figure S9 Distribution of mRNA expression levels in interactions predicted by PIP-Lcut_{600} (red) and PIC-Lcut_{600} (blue) for *S. cerevisiae*. PIC-Lcut_{600} shows less bias towards highly expressed ORFs compared to PIP-Lcut_{600}. Expression data are retrieved from yeast reference mRNA expression set introduced by Greenbaum et al. (2002, Bioinformatics 18:585-96).