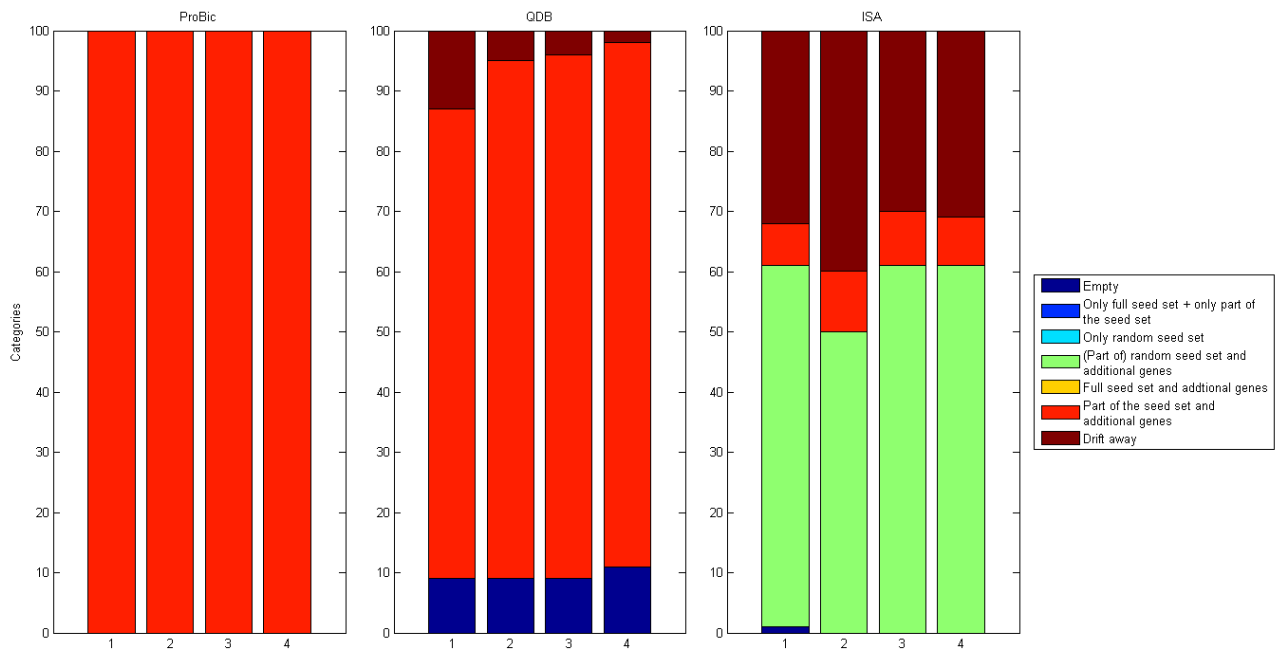
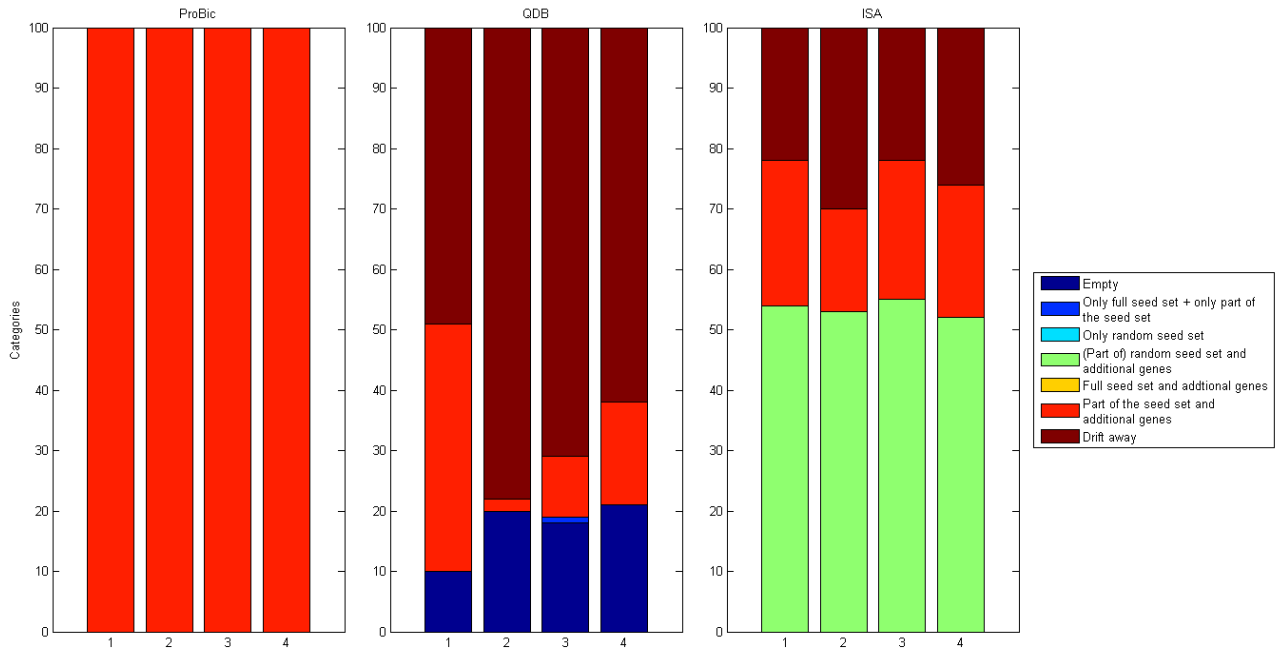


Additional File 7 - Robustness of *ProBic*, QDB and ISA to noisy seed genes of intermediate quality

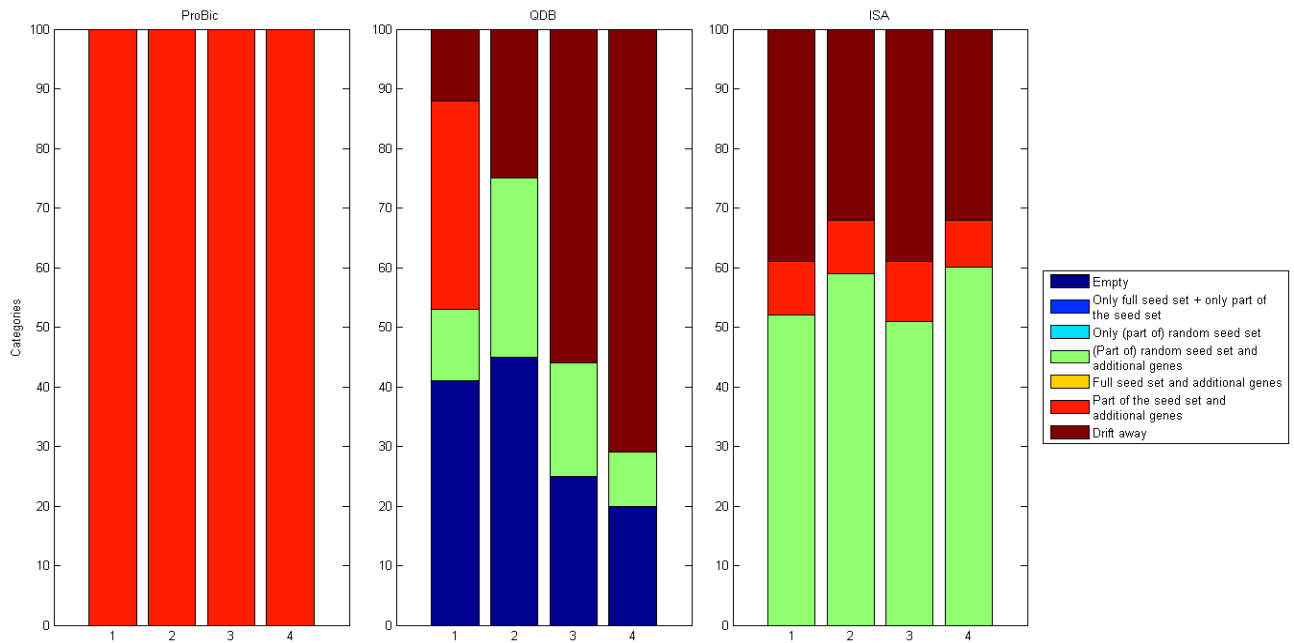
Analysis of biclusters, obtained by the different algorithms after biclustering in the presence of an increasing number of random genes (1 = 20%, 2 = 40%, 3 = 60% and 4 = 80% of random genes), added to the true seed set of three seed sets of intermediate quality (panel a: FadR, panel b: NarL, panel c: OmpR). The procedure of adding random sets was repeated 100 times for each seed set. Legend is as in Figure 3 of the main text.



(a)



(b)



(c)