Figure S2. Expression breadth differences between non-miRNA and miRNA target genes predicted from TargetScanS and RNA22. Distribution of the fraction of miRNA and non-miRNA target genes restrictedly expressed in certain number of tissues for miRNA target genes predicted from TargetScanS (a) and RNA22 (b). p-values: statistical differences of median fractions between miRNA and non-miRNA target genes in each bin by Wilcoxon signed rank tests.