Supplemental Figure 3: Jovanovic et al.

The figure represents a cumulative distribution of mRNA expression data. The horizontal axis shows the log2 ratio of "mir-58" expression compared to wild type (wt), ranging from -1.5 to 1.5. The vertical axis represents the cumulative fraction, ranging from 0 to 1.

The different lines in the graph indicate various datasets and methods:
- RIP-chip (Protein down)
- RIP-chip (Protein up)
- RIP-chip (all)
- TargetScan
- Random