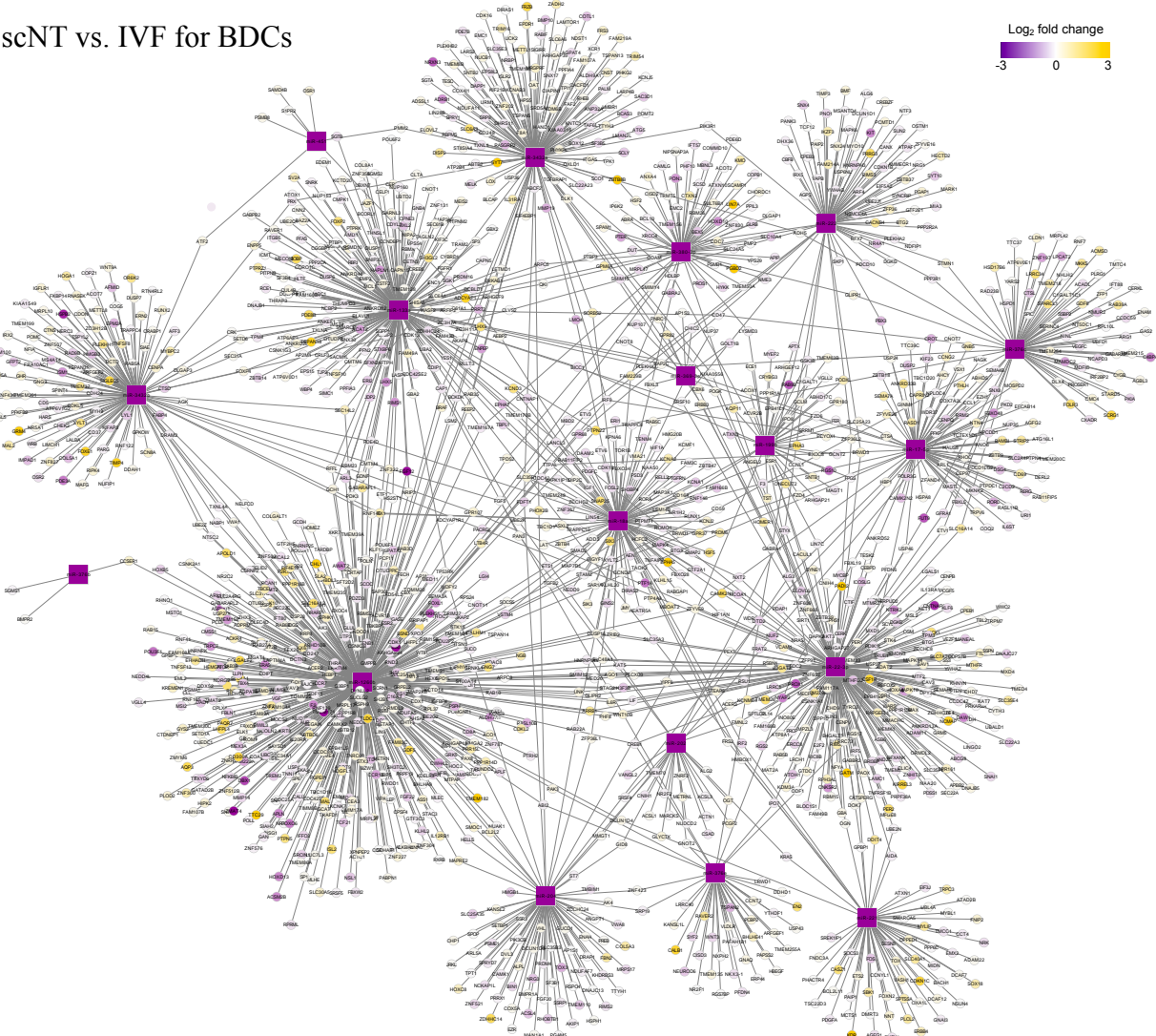


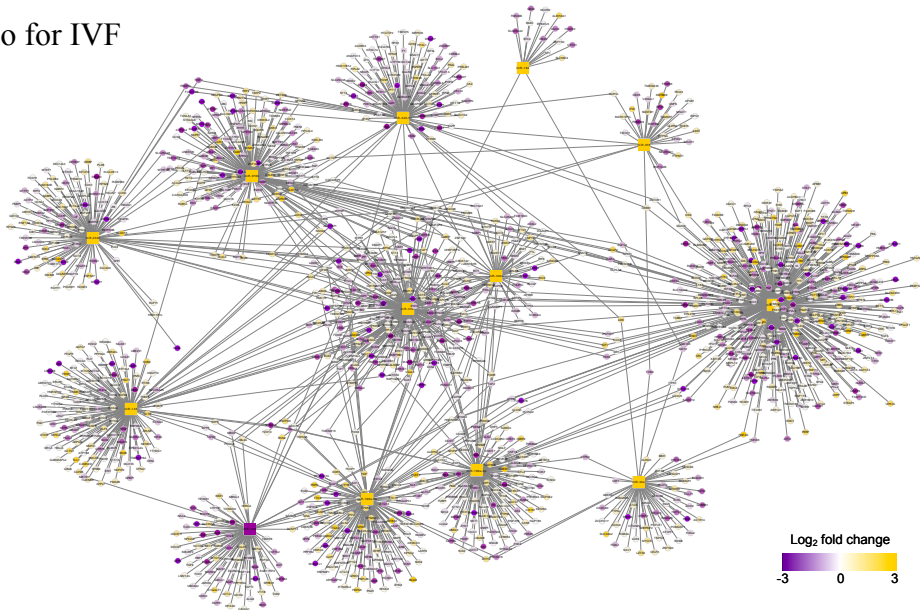
Supplementary File 4. Cytoscape networks for differentially expressed miRNAs and their predicted mRNA targets

Networks depicted on the following pages were created using data for differentially miRNAs and their predicted mRNA targets (TargetScan total context ++ score < -0.35) using Cytoscape with a force-directed layout. Only miRNAs within families annotated by TargetScan as conserved or broadly conserved were included in the network analysis. MiRNAs are represented by squares and target mRNAs as circles and both are shaded according to their \log_2 fold change for the comparison indicated. Edges between miR and mRNA nodes indicate that the connected mRNA is a predicted target of the miRNA in *Bos taurus*.

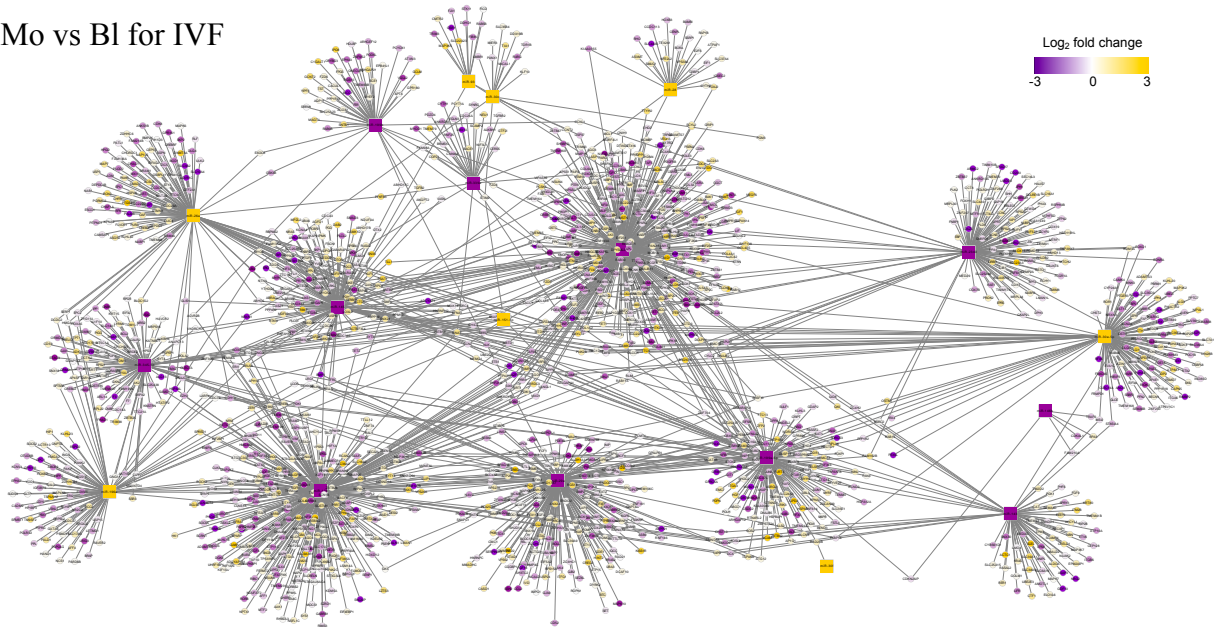
scNT vs. IVF for BDCs



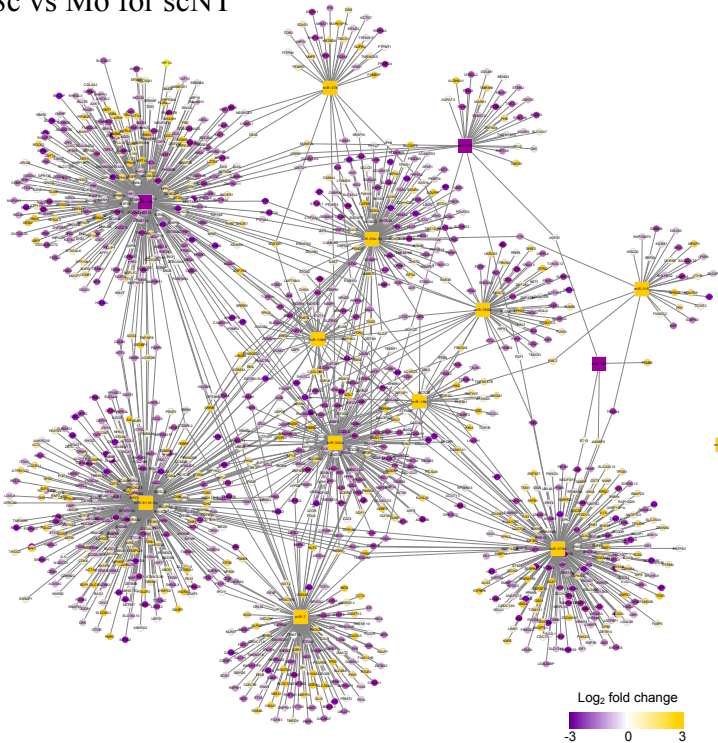
8c vs Mo for IVF



Mo vs Bl for IVF



8c vs Mo for scNT



Mo vs BI for IVF

