



**Figure S3.** Enrichment of microRNA families modulating DEPs in 3e-treated Skmel-28 cells.

Results from ShyniGO 0.80 microRNA target database are shown for analysis of microRNA putatively regulating the differential expressed proteins (DEPs) obtained from comparison between DMSO-treated and 3e-treated cells for 24h (A), 48h (B) and 72h (C). In bar diagrams, fold enrichment is colour-scale represented (from the lowest blue to the highest red) and microRNA are ranked based on fold discovery rate (FRD) values (longer bar corresponds to lower FDR).