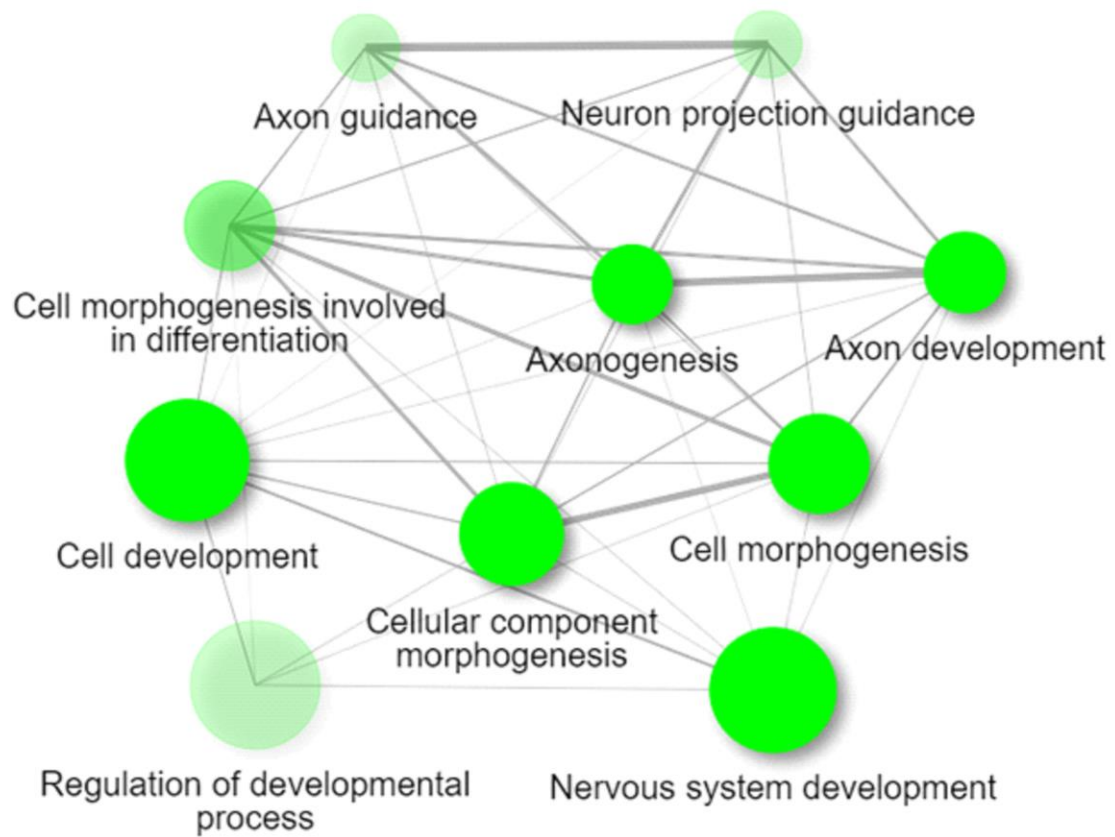




**SUPPLEMENTAL FIG.2 GO enrichment analysis for miR-512-3p target genes.** Gene Ontology terms (biological processes) enriched among the miR-512-3p target genes in MB as revealed by the ShinyGo tool. The color intensity level of the circles agrees with the p value; and the size of the circles indicates the fold enrichment. Only terms with p values < 0.05 were considered