



Figure S2. The functional network and enriched functional group of GCMN was analyzed using ClueGO, a biological term enrichment analyzer. The following proteins were significantly enriched in GCMN: melanosome (GO_cellular component), neurotrophin signaling pathway (KEGG pathway), downregulated of MTA-3 in ER-negative breast tumors (Biocarta), cell cycle (KEGG pathway), phospholipase inhibitor activity (GO_molecular function), and glycolysis/gluconeogenesis (KEGG pathway).