

Supplementary Table 2. Gene Ontology (GO) enrichment analysis based on 3 functional groups (biological processes (BP), cellular components (CC), and molecular functions (MF) in EVs isolated from RJ-treated ADSCs.

Molecular functions (MF): Downregulated

Category		Term	Count	%	P-Value	FDR
GOTERM_MF_ALL	GO:0005515~	protein binding	15	75	0.038022687	1