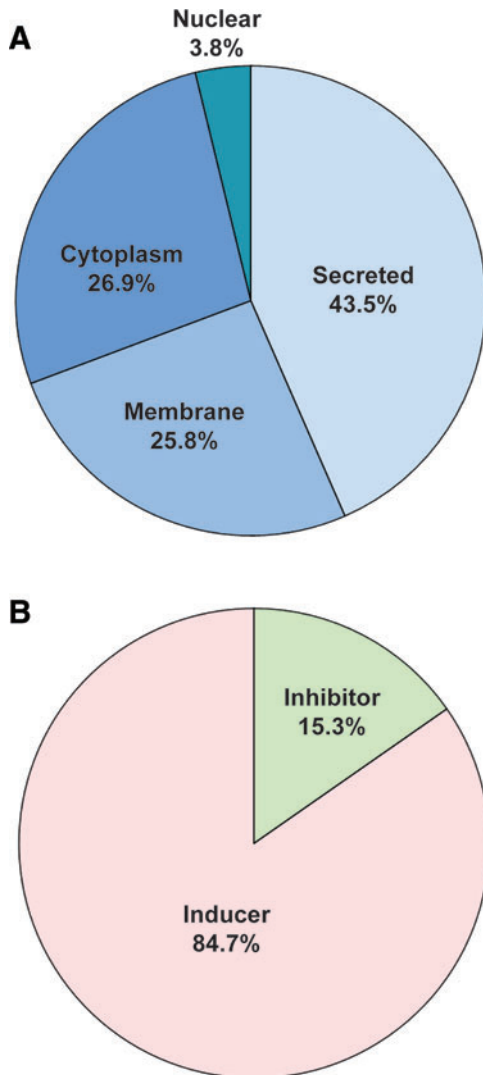# Supplementary Data

SUPPLEMENTARY FIG. S1. pMEX proliferation-associated proteins enriched for secretory proteins. **(A)** HiRIEF LC-MS/MS proteomic analysis followed by IPA analysis determined that the most relatively abundant subcellular classification of pMEX proteins was secretory in nature. **(B)** IPA analysis showing relative abundance distribution of pMEX proteins known to be inducers vs inhibitors of proliferation. $n = 3$, FDR1%. IPA, Ingenuity Pathway Analysis; pMEX, primed mesenchymal stem cells secrete exosomes.