

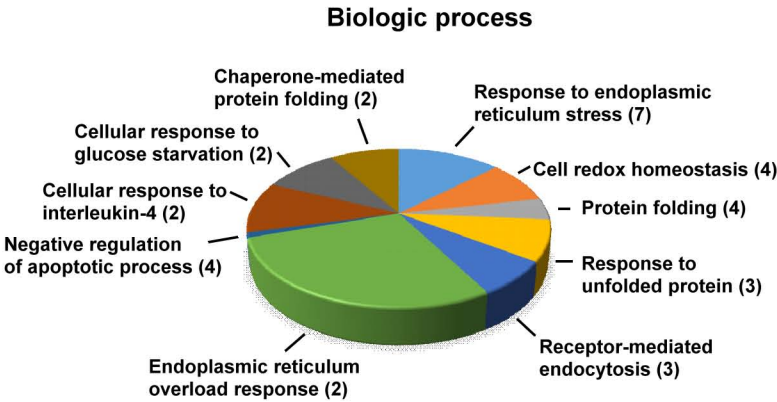
A

mRNA	Gene	Probability exact	Consevation phylop	miRmap score
hsa-miR-92a-1-5p	RCN3	93.46	87.67	90.31

B

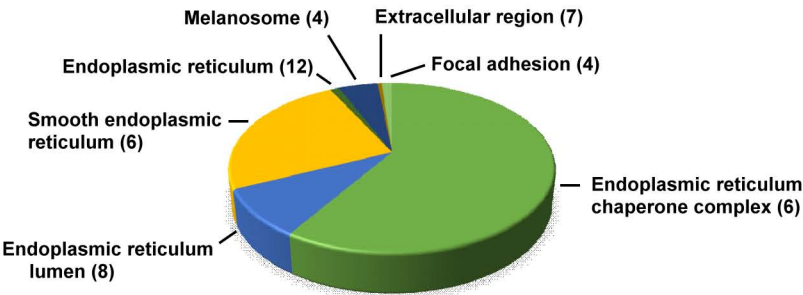
hsa-miR-92a-1-5p	3'.....GUUGGCUA	GGGUUGGA...5'
nt(289-295)		
WT	RCN3	3'UTR
	human	5'.....GAACCGC - CCAACCC...3'
	mouse	5'.....CCAACCCU...3'
	rabbit	5'.....CCUCCCAU...3'

C



D

Cell component



E

KEGG pathway

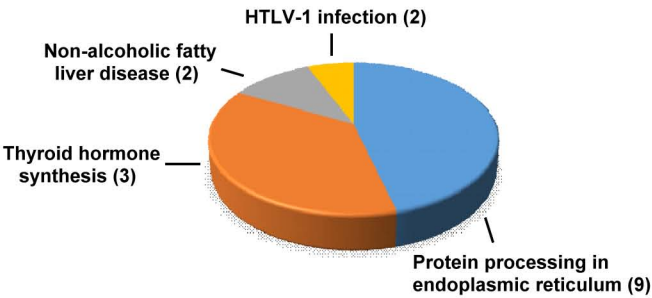


Fig S4