

Figure 1

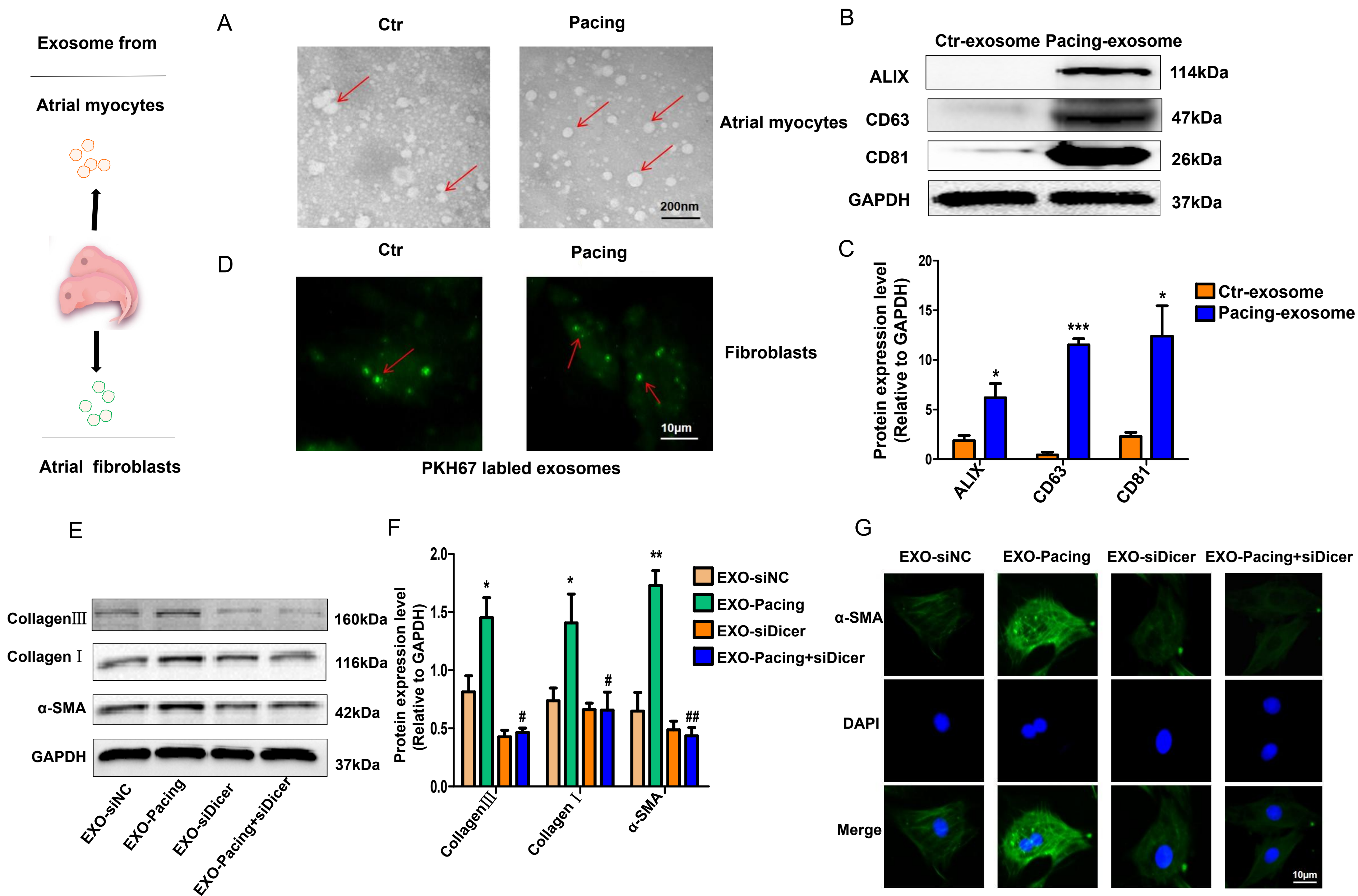




Figure 2

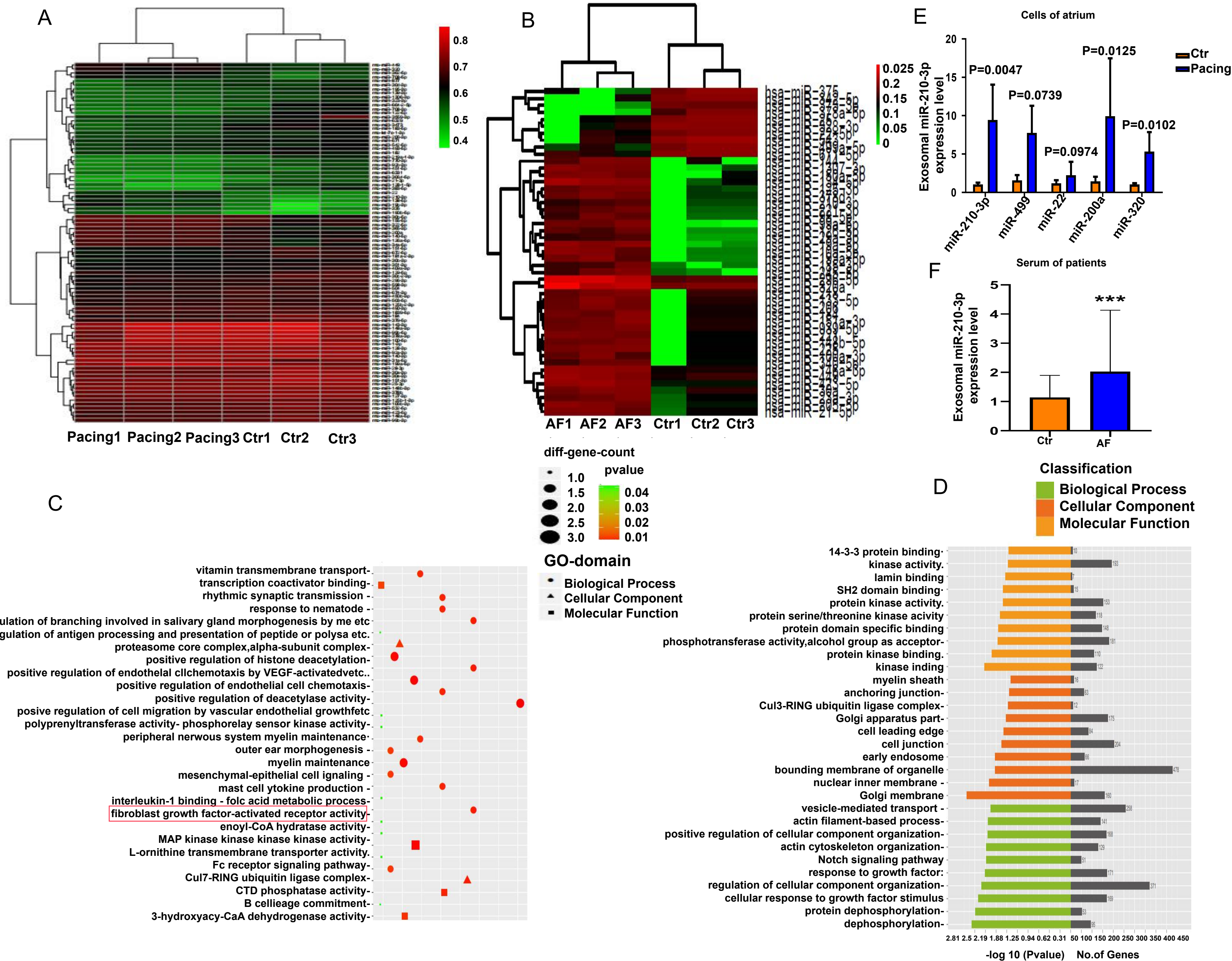




Figure 3

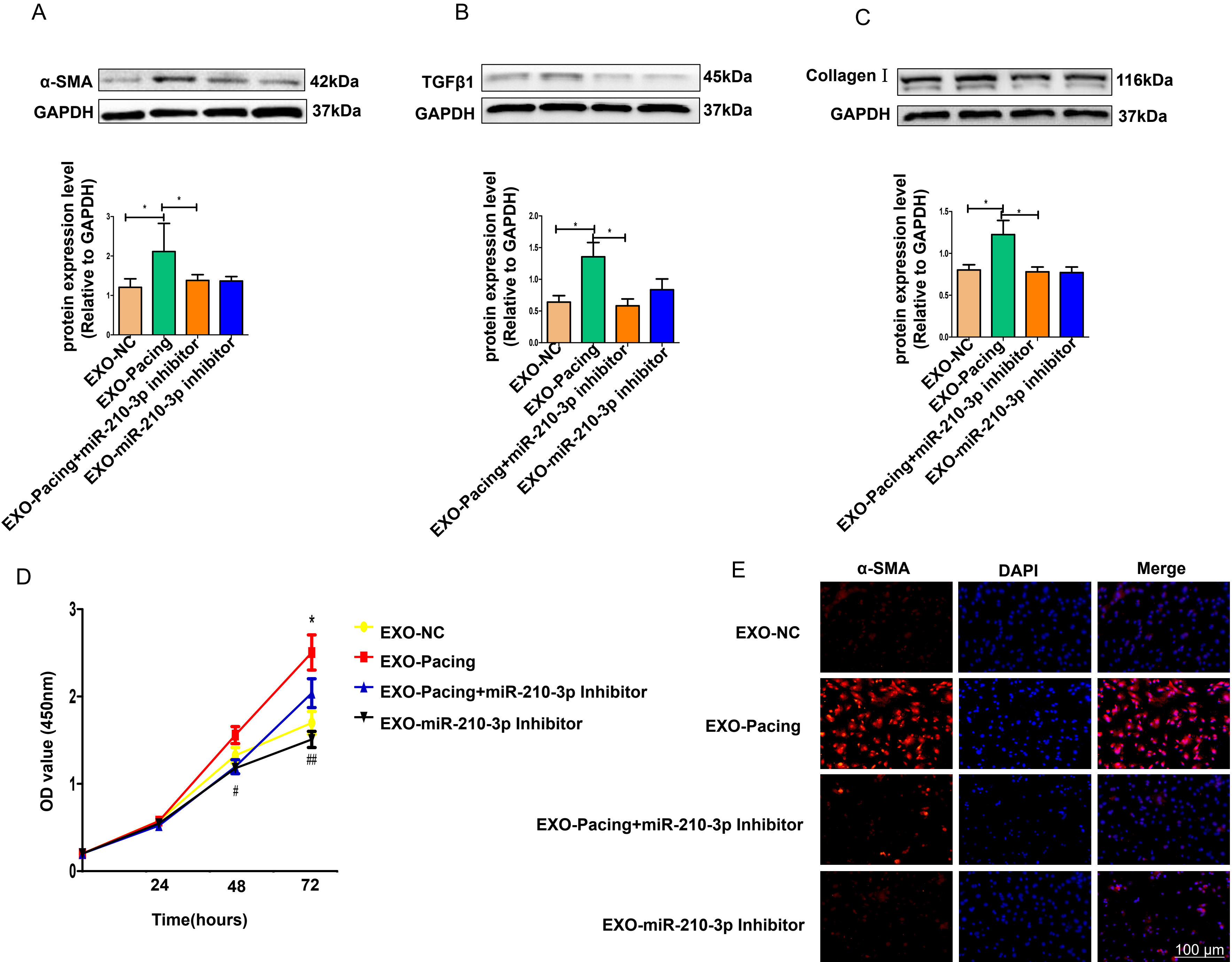




Figure 4

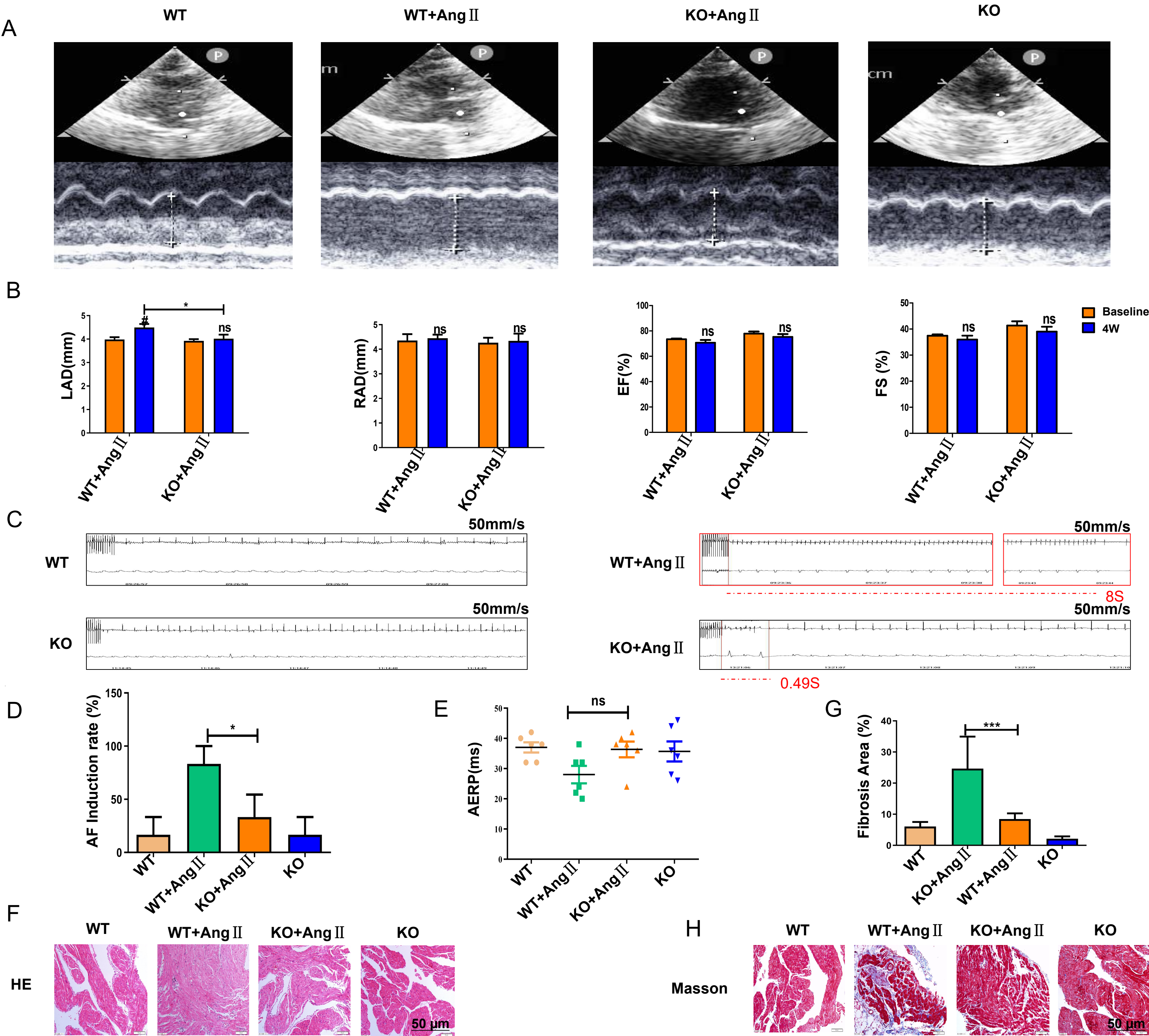


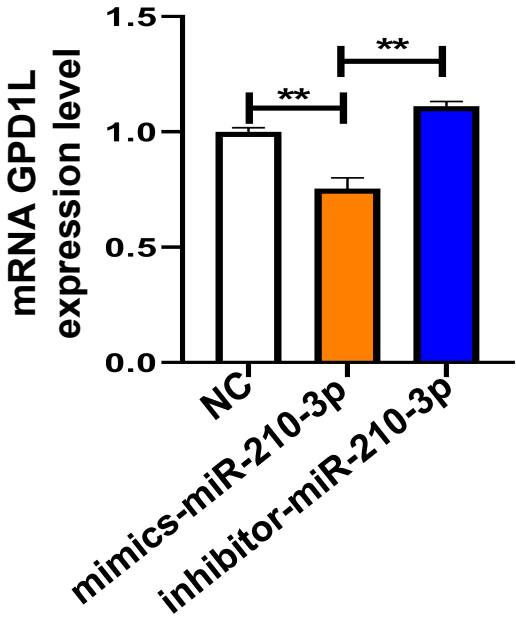


Figure 5

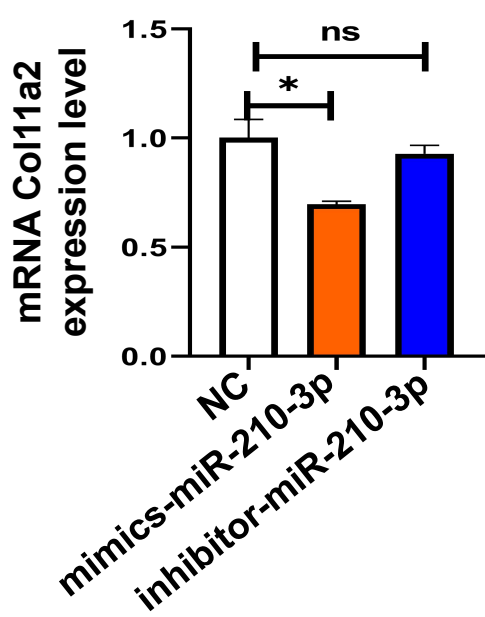
A

	Predicted consequential pairing of target region (top) and miRNA (bottom)	Site type	Context++ score
Position 1632-1638 of GPD1L 3' UTR	5' ... GUCAGCUUUGCUGAA--ACGCACAU... 3' ... AGUCGGCGACAGUGUGCGUGUC 	7mer-m8	-0.35
Position 1605-1611 of Gpd1l 3' UTR	5' ... GUGAACUCACUGAA--ACGCACAC... 3' ... AGUCGGCGACAGUGUGCGUGUC 	7mer-m8	-0.31

C



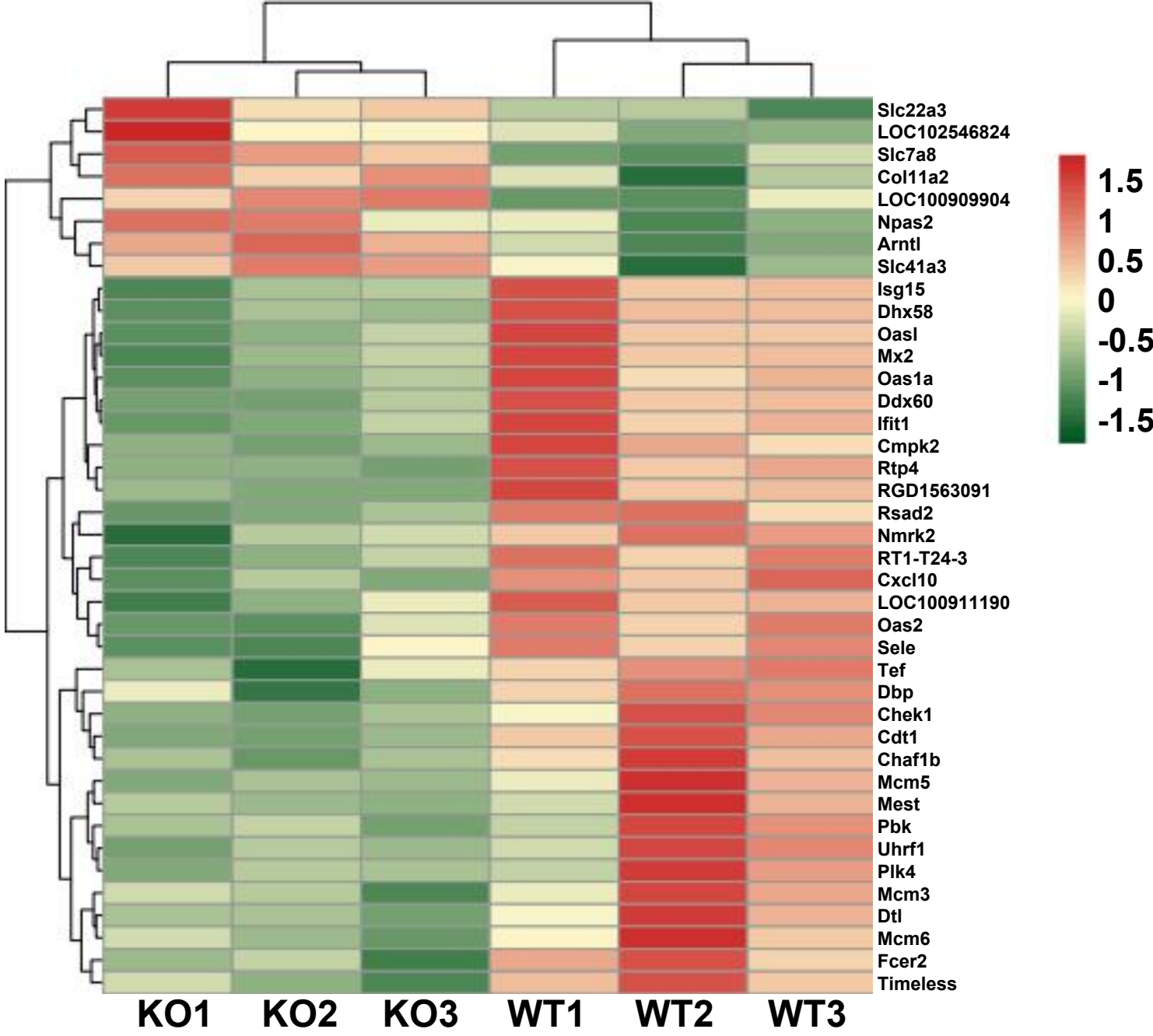
D



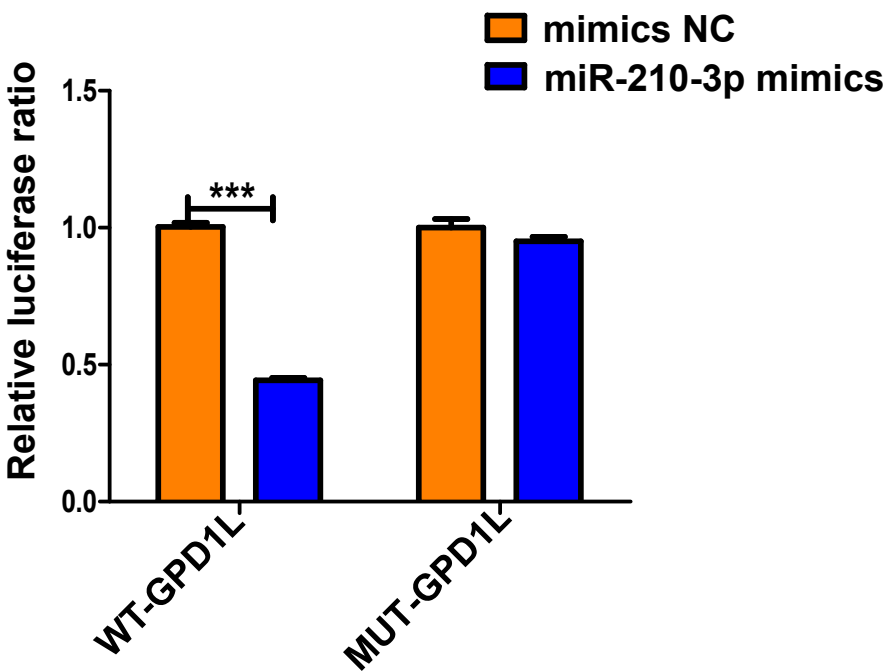
E

WT-GPD1L 5'...1607...TGAACCTCACTGAAACGCACAC...1613...3'  
rno-miR-210-3p 3'...AGUCGGCGACAGUGUGCGUGUC...5'  
MUT-GPD1L 5'...1607...TGAACCTCACTGAATGCGGTGTC...1613...3'

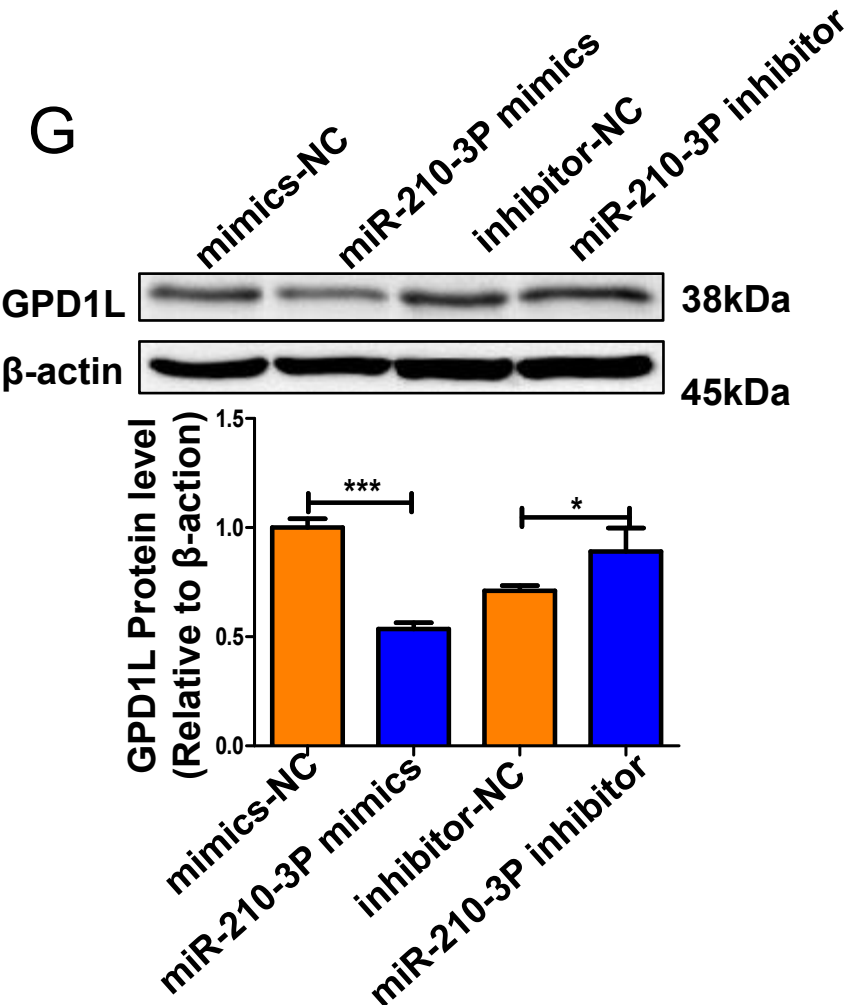
B



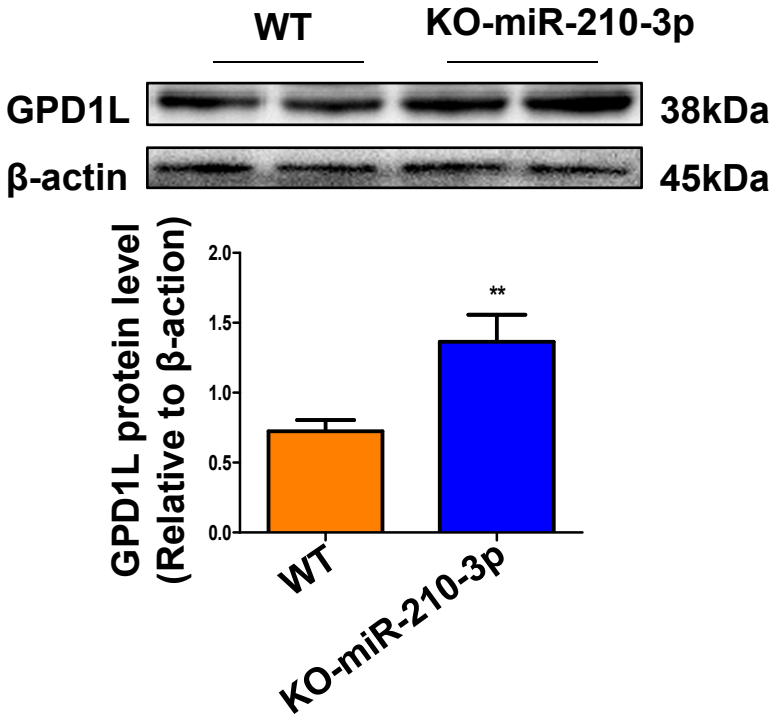
F



G



H



I

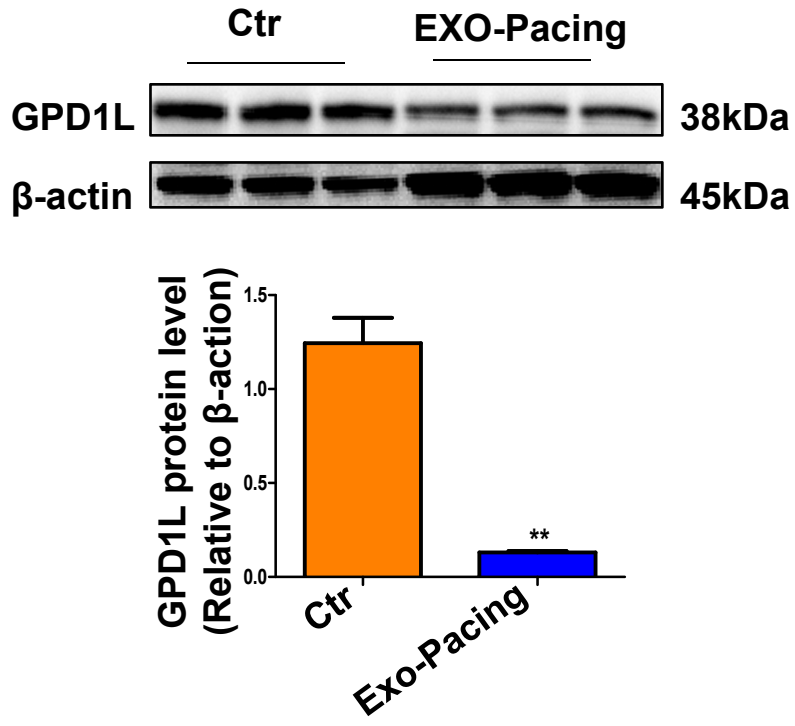


Figure 6

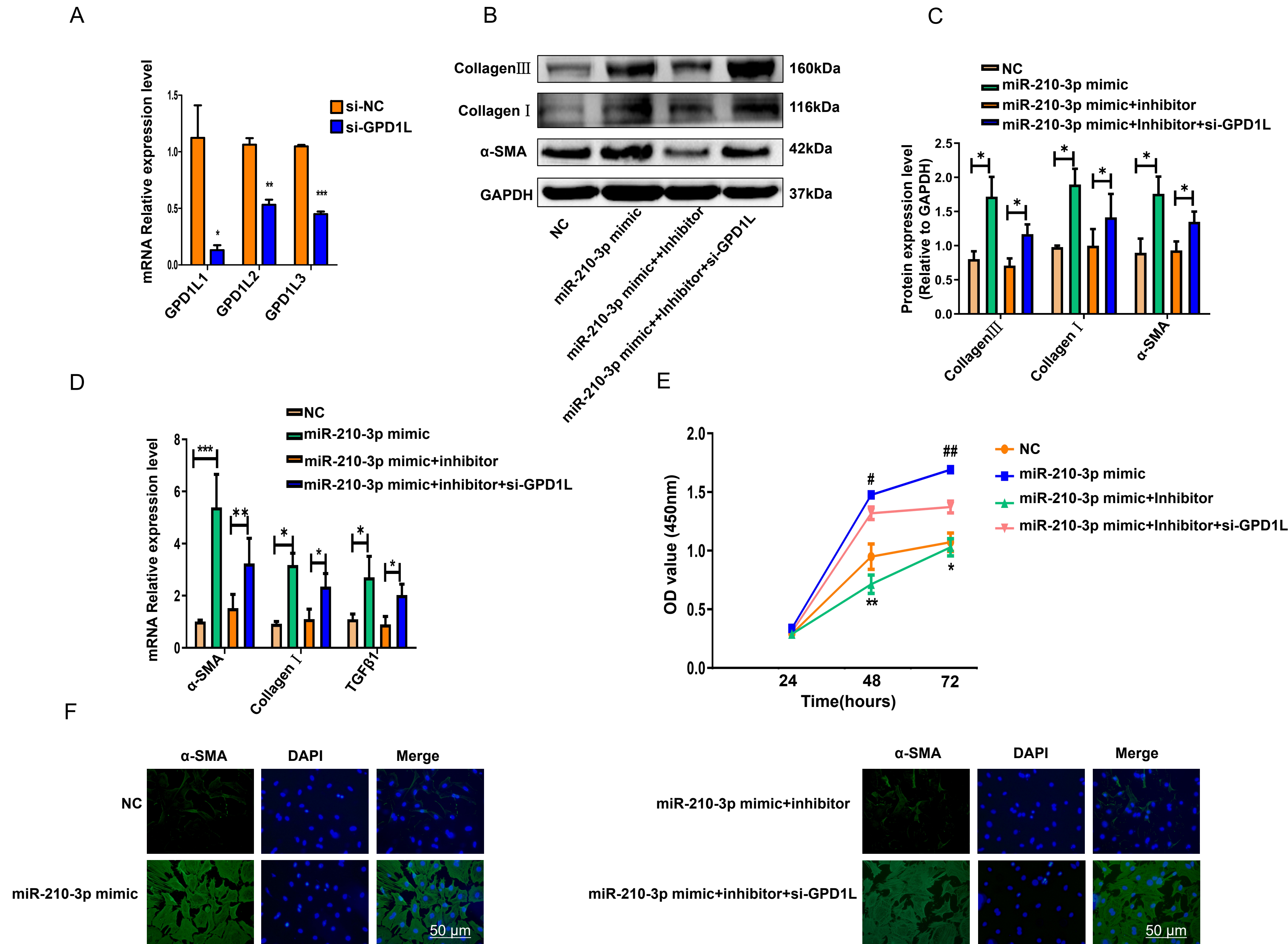




Figure 7

