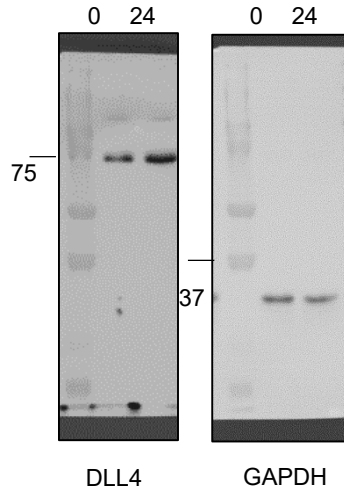


Figure 5

A

0	24
1	8.603955
1	8.083612
1	8.058697

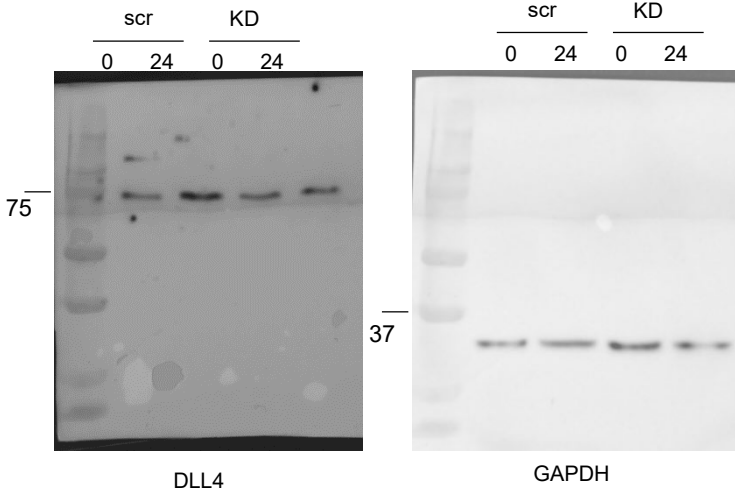
B



C

		scr		
0	1	1	1	
24	8.083612	8.058697	6.475538	
		circHIPK3 KD		
0	0.675974	0.300925	0.320295	
24	3.986161	1.168777	2.049114	

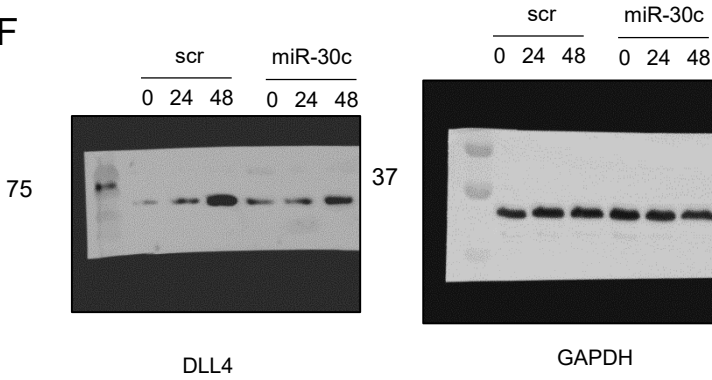
D



E

		scr		
0	1	1	1	
24	7.361501	12.21914	10.28225	
		miR-30c		
0	0.180491	0.181118	0.397768	
24	1.70527	1.068065	0.939523	

F



Scr	0	1	1	1
	24	1.998341	1.266441	1.319536
	48	2.709761	1.681993	2.957234
miR-30c	0	0.242681	0.206852	0.694378
	24	0.469367	0.467275	0.472015
	48	0.781139	0.656711	0.623932

G

GAPDH	DLL4
1	22.7848
1	15.13692
1	12.77277

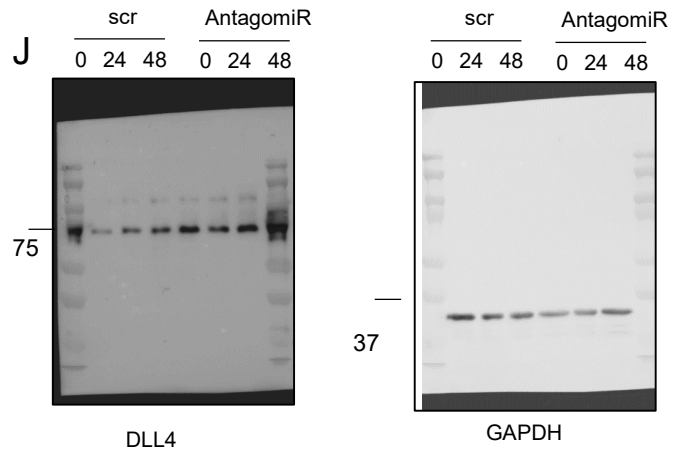
H

		Scr			miR-30c		
10	1	1	1	0.620331	0.630942	0.441696	
20	0.902043	0.810602	1.060757	0.425295	0.496663	0.313365	
50	0.880873	1.135238	1.06284	0.232501	0.253651	0.36737	

I

		Scr		
0	1	1	1	
24	6.110741	5.063026	6.088402	
		miR-30c AntagomiR		
0	1.580083	1.270151	1.515677	
24	7.808254	7.209816	8.723835	

J



K

IgG	Ago2	Ago2 + AntagomiR
0.047102	49.76047	4.948283
0.000912	46.42816	10.03472
0.493776	37.40087	2.842042

L

Scr	20nM	40nM
1	1.233693	1.512903
1	1.138025	1.371298
1	1.118553	1.262132