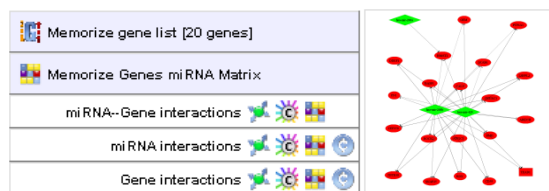


A



gene	Score [average]	Fisher Test [p-value]	miRNA [percent] [number]	Sort by: Fisher Test [p-value]
DUSP1	0.6906	3.9987e-3	6.25% 2/32	3.999e-3
CHURC1	0.8217	7.8990e-3	4.4444% 2/45	
GEM	0.6842	9.3514e-3	4.0816% 2/49	
BASP1	0.8653	9.7331e-3	4% 2/50	
HS6ST2	0.7469	1.1483e-2	2.3622% 3/127	
ATP2C1	0.4607	1.2609e-2	3.5088% 2/57	
CD59	0.9046	1.5353e-2	3.1746% 2/63	
LHFPL2	0.8961	1.5353e-2	3.1746% 2/63	
FN1	0.7163	2.0490e-2	2.7397% 2/73	
RDX	0.8487	2.1600e-2	2.6667% 2/75	
DDEF1	0.9226	2.3316e-2	2.5641% 2/78	
ZBTB38	0.9218	2.5699e-2	2.439% 2/82	
BMI1	0.771	2.8829e-2	2.2989% 2/87	
SNAP25	0.8104	3.0127e-2	2.2472% 2/89	
ROCK2	0.8839	3.2801e-2	2.1505% 2/93	
PTP4A1	0.7675	4.1440e-2	1.9048% 2/105	
TEAD1	0.8925	4.8510e-2	1.7544% 2/114	
CALU	0.8835	7.9536e-2	1.3514% 2/148	
ARID5B	0.7759	7.9536e-2	1.3514% 2/148	
CEP350	0.8324	8.0546e-2	1.3423% 2/149	

B

