

Supplementary Table S3: Top 20 circRNA candidates from screening results.

Gene	E-Value	Coordinates
SETD2	0	chr3:47113875-47124548 SETD2
FAT1	0	chr4:186706562-186709845 FAT1
FOXO3	0	chr6:108663454-108664889 FOXO3
AFF1	5E-177	chr4:87046165-87047594 AFF1
ELK4	5E-174	chr1:205616477-205623891 ELK4
CTCF	3E-148	chr16:67610823-67612121 CTCF
AFF4	2E-140	chr5:132892163-132897240 AFF4
ATF1	3E-66	chr12:50814009-50814439 ATF1
MYH9	3E-66	chr22:36341369-36349255 MYH9
PIK3CA	3E-49	chr3:179203543-179204588 PIK3CA
MSH2	2E-44	chr2:47475024-47478519 MSH2
CREBBP	2E-43	chr16:3749626-3770986 CREBBP
CIC	3E-41	chr19:42291344-42292698 CIC
EZR	5E-38	chr6:158766661-158770801 EZR
FBXW7	4E-36	chr4:152346929-152412529 FBXW7
CRTC3	8E-36	chr15:90619740-90629532 CRTC3
NF1	7E-35	chr17:31155982-31182665 NF1
KMT2C	3E-34	chr7:152309965-152315338 KMT2C
ABL2	7E-33	chr1:179131310-179131481 ABL2
XPO1	9E-33	chr2:61485767-61490776 XPO1

Coordinates were mapped to the human reference genome (GTCh38/hg38).