

**MiR-199a-5p regulates sirtuin1 and PI3K in the rat
hippocampus with intrauterine growth restriction**

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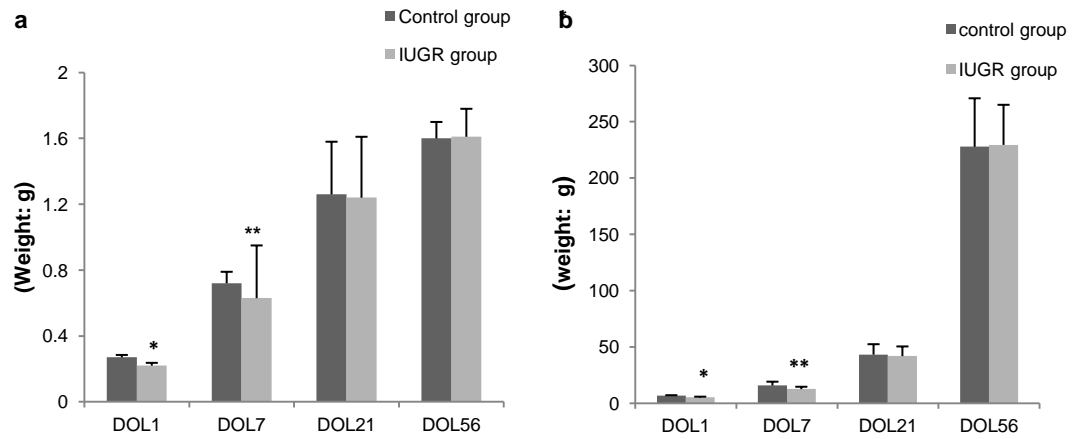
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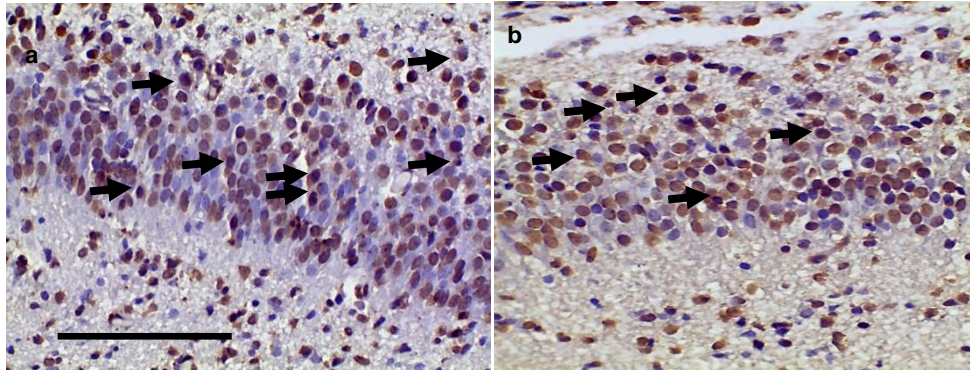
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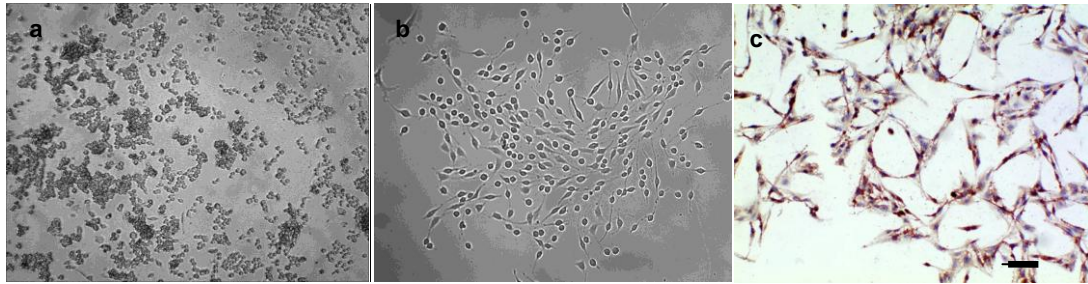
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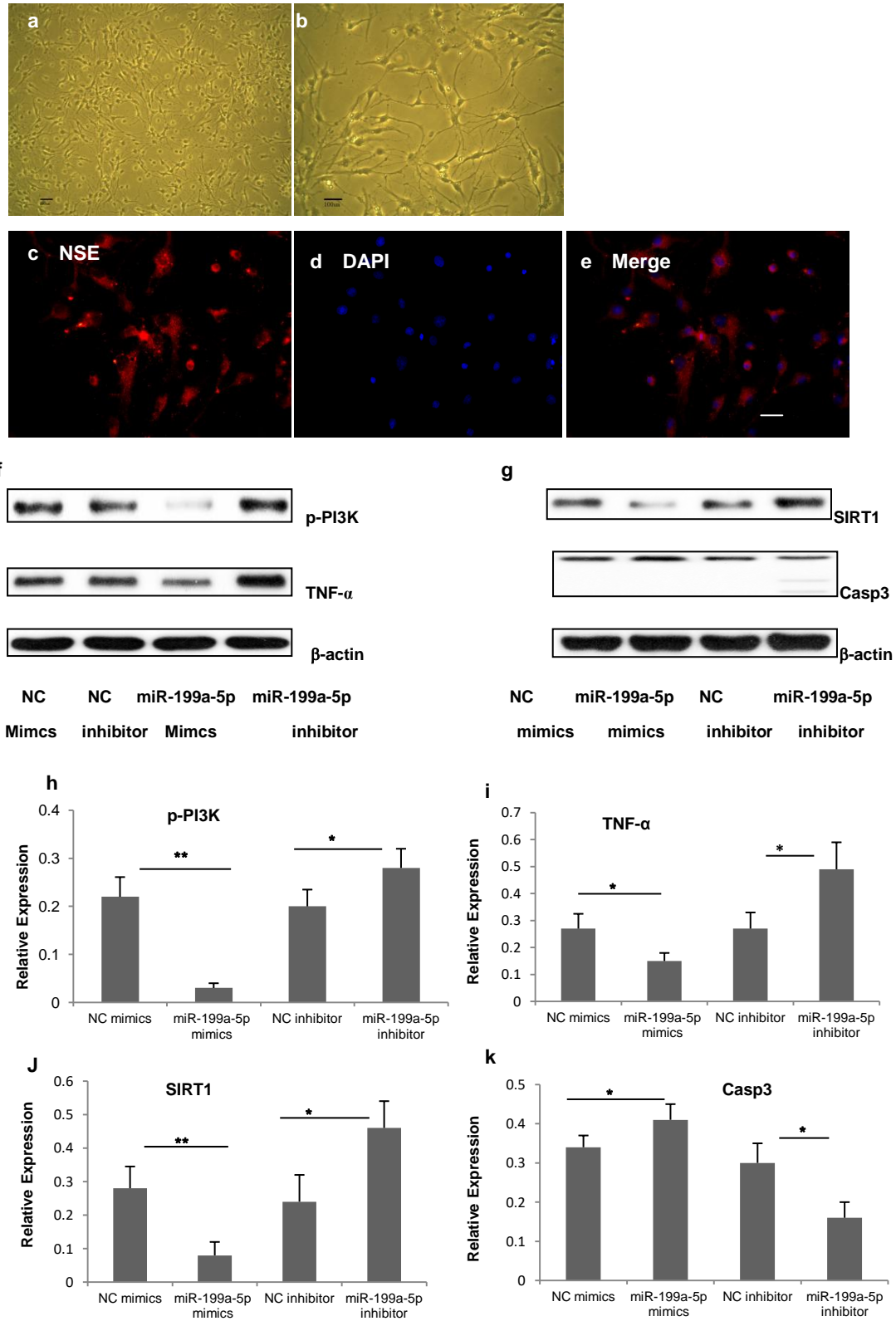
1. Fig. S1. Brain weights and Body weight at different times. (a) The weight of brain between the control and IUGR groups at different times after birth. (b) The body weight between the control and IUGR groups at different times after birth. The values are expressed as the mean \pm SD. * $P < 0.05$, ** $P < 0.01$.



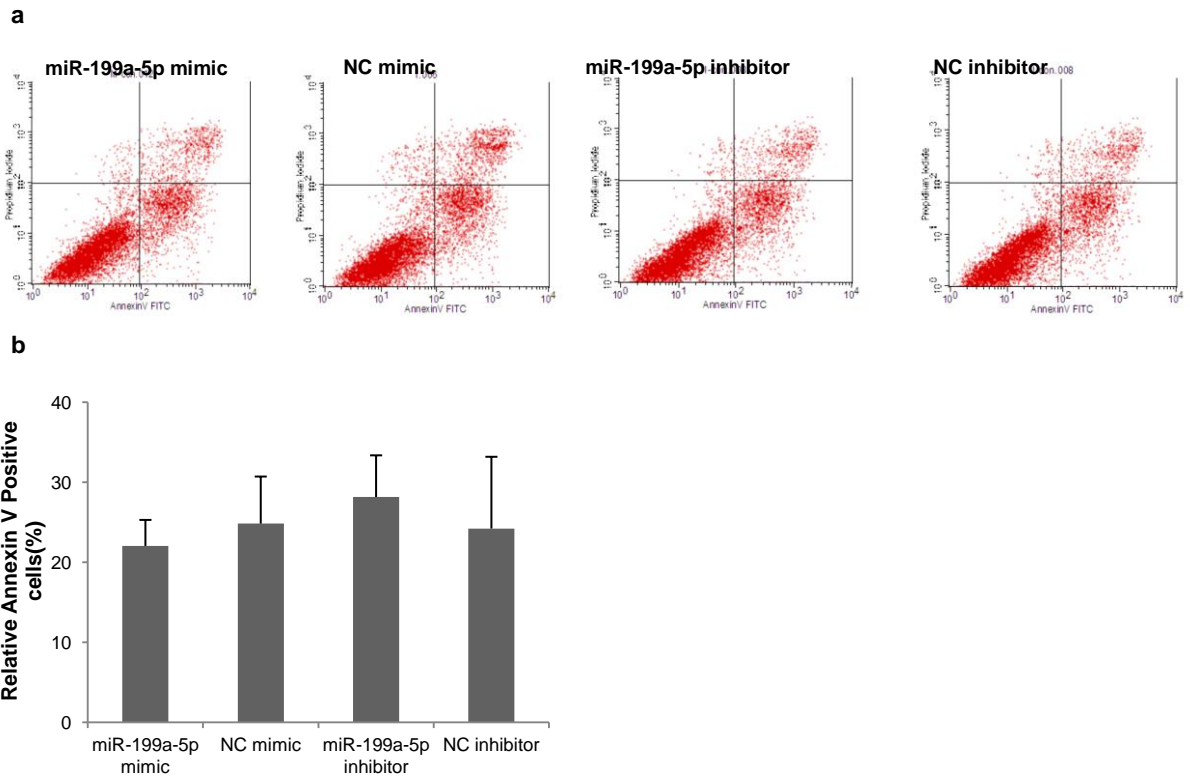
2. Fig. S2. Apoptotic nuclei stained using the TUNEL technique in the hippocampi CA1 region of the control group (a) and the IUGR group (b) of the DOL1 rats. Original magnification $\times 400$. Scale bars = $200\mu\text{m}$.



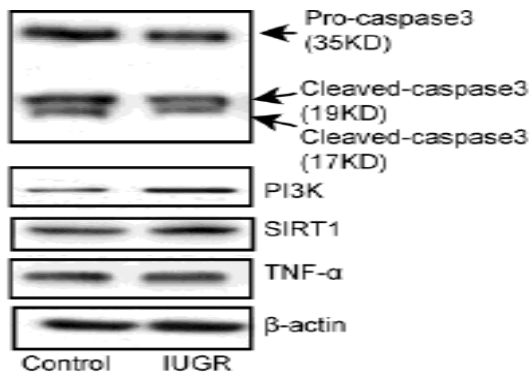
3. Figure S3. PC12 Cell culture and treatment. (a) PC12 cell culture. (b) PC12 cells treated by NGF. (c) PC12 cells staining by NSE. Original magnification $\times 100$.



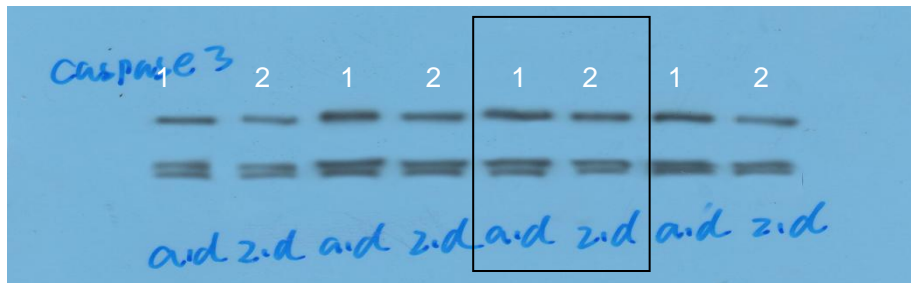
4. Fig. S4. MiR-199a-5p inhibitors PI3k, TNF- α , SIRT1 and Casp3. (a, b). primary neuronal culture. a, $\times 100$; b, $\times 200$. (c, d, e) primary neurons staining by NSE and DAPI. (h, i, j, k) Western blot analysis of p-PI3K, TNF- α , SIRT1 and Casp3 protein expression in primary neurons following transfection. Scale bars = 100 μ m.



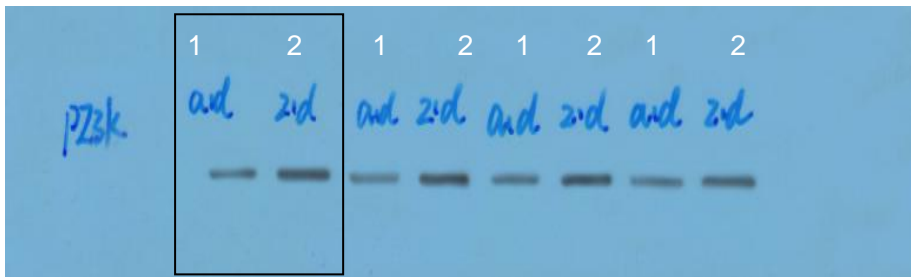
5. Fig. S5. Effect of miR-199a-5p on PC12 cells apoptosis detected by flow cytometer. (a) Flow cytometric analysis of PC12 cells were treated with miR-199a-5p mimics and inhibitors. (b) Data are presented as the percentage of Annexin V positive cells which analyzed by flow cytometer. The values are expressed as the mean \pm SD. * $P < 0.05$, ** $P < 0.01$.



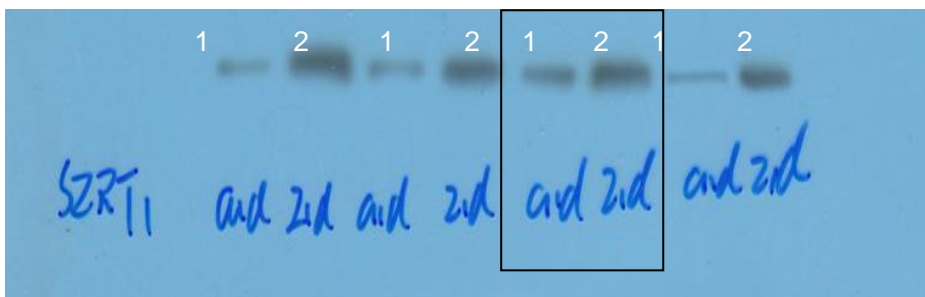
Caspase-3:



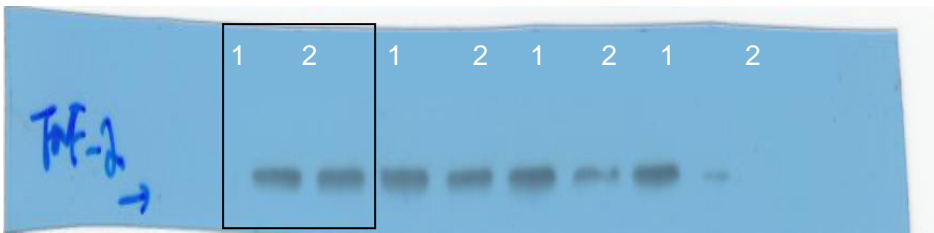
PI3K:



SIRT1:



TNF- α :



β -actin:

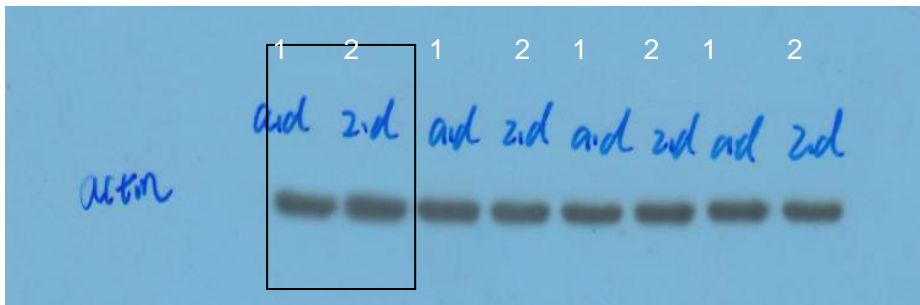
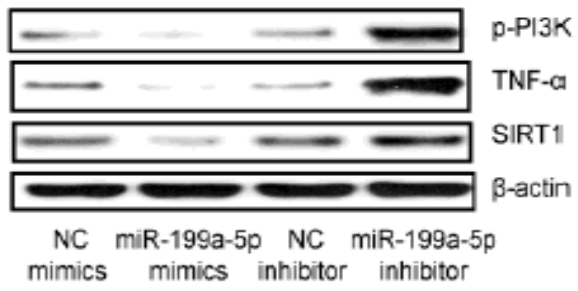
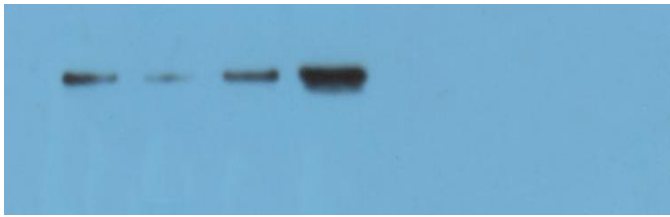


Figure S6. Original Western image for Figure 3a. 1, control group; 2, IUGR group.



p-PI3K



TNF-α



SIRT1



β-actin

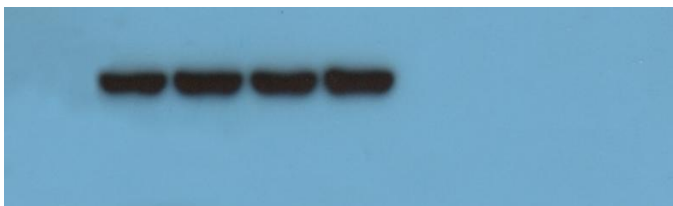


Figure S7. Original Western image for Fig. 4f.

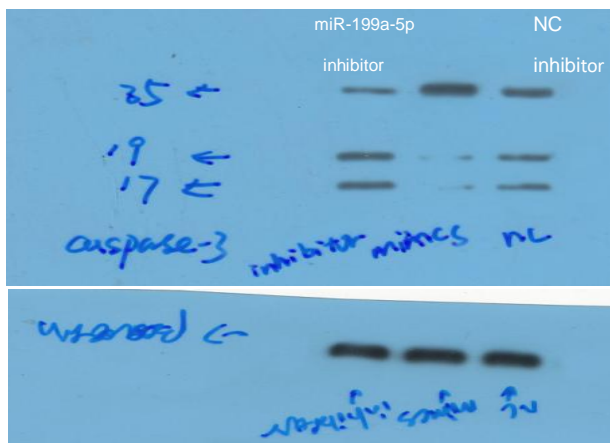
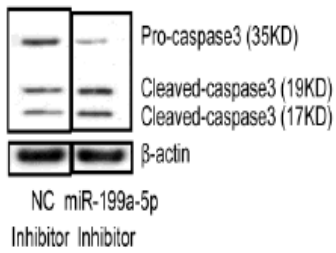
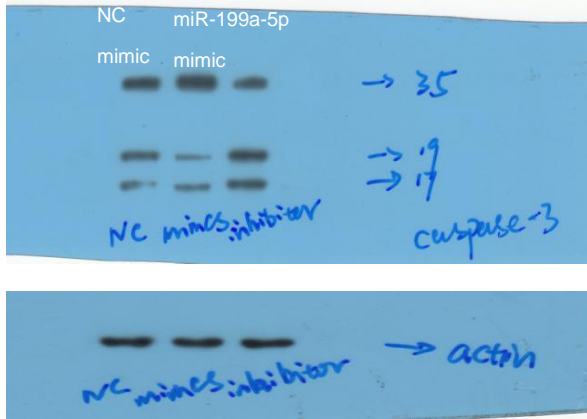
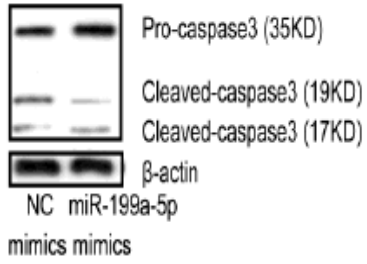
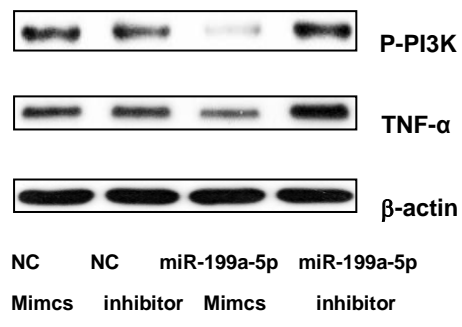


Figure S8. Original Western image for Fig. 5b.



p-PI3K



TNF-α



β-actin

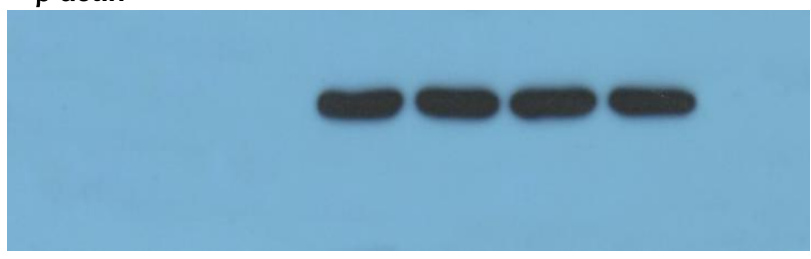
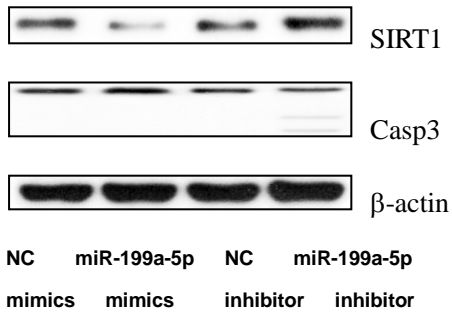
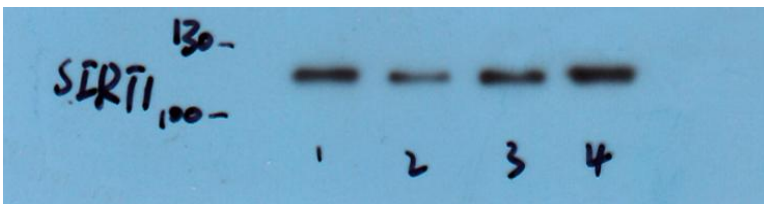


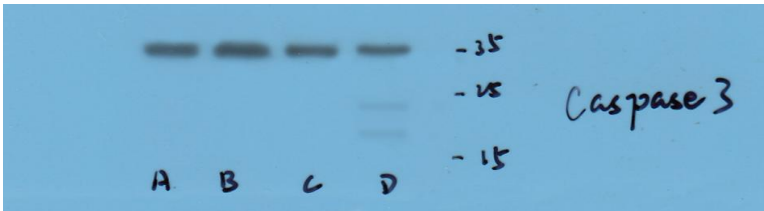
Figure S9. Original Western image for Fig. S4f.



SIRT1



Casp3



β-actin

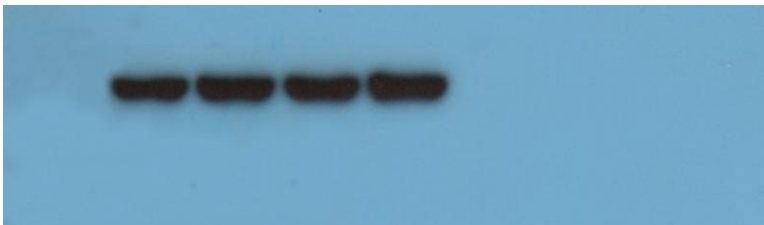


Figure S10. Original Western image for Fig.S4g.

Mature miRNA name	Mature sequences (5'-3')		Predicted target gene
rno-miR-199a	rno-miR-199a-5p	CCCAGUGUUCAGACUACCUGUUC	Sirt1,Atp2,Tnf,casp3, pole3,acer2,Sumo3,pi3k
	rno-miR-199a-3p	ACAGUAGUCUGCACAUUGGUUA	Rab6a,Bre,Aspa,Ankfy1, sgsm2,Depdc5,March10
rno-miR-325	rno-miR-325-5p	CCUAGUAGGUGCUCAGUAAGUGU	Trip11,Pla2g2c,Plk1,Zfp407, Ncbp3,Tnfsf9,Nr2f2
	rno-miR-325-3p	UUUAUUGAGCACCUCUAUCAA	Npc2,vamp7,Fpr2,Prmt1, IkIf1,Nol8,cdn1
rno-mir-219	rno-miR-219a-1-3p	AGAGUUGCGUCUGGACGUCCCG	Bcor,Styl5,Tspan7,Lancl3,cy bb,Dynlt3,Rbm3
	rno-miR-219a-5p	UGAUUGUCCAAACGCAAUUCU	Otc,Hdac6,Apex2,Rrag13, Pdha1,Pola1,Flt3
rno-miR-664	rno-miR-664-1-5p	CUGGCUGGGGAAAAAGAUUGG	Slc25a44, Dicer1, Zbtb34, Ncoa1,Insm2,Dnai2,Slco1b2
	rno-miR-664-3p	UAUUCAUUUACUCCCCAGCCUA	Paip1, Ndr4, Hey2, Nipal1, Aqp1, Slc2a5, Itga9,
rno-miR-551b	rno-miR-551b-5p	GAAAUCAAGCUUGGGUGAGACCU	Nbl1 ,Casp9, Dctn4, Coq4, Klf12, Nhsl1, Usf1, Hoxa2
	rno-miR-551b-3p	GGCGACCAUACUUGGUUCAGU	Srsf5, Btd7, Ipo7, Skap2, Trak1, Cbl11, Ptpbh,Skp2
rno-mir-34C	rno-miR-34c-3p	AAUCACUAACCACACAGCCAGG	Cpsf4, Glrx, Lox, Hdhd2, Polr2c, Rnf2,Lix1, Eepd1
	rno-miR-34c-5p	AGGCAGUGUAGUUAGCUGAUUGC	Bcor,Tspan7,Cybb, Slc38a5, Porcn, Wdr13, Eras, Plp2
rno-miR-195	rno-miR-195-5p	UAGCAGCACAGAAAUAUUGGC	Flot1, Eya3, swt1,Rff1 , Cngn3, Ifitm1, Nodal, Thoc2
	rno-miR-195-3p	CCAAUAUUGGCUGUGCUCCA	Tbp,P2rx4, Szrd1, Lmo1,Cdh8,Sf1,Megf11
rno-miR-204	rno-miR-204-5p	UUCCCUUUGUCAUCCUAUGCCU	Casp3,Sirt1,Zfx,Prima1,Tcea 3,Prcc2b,Pdcl,Rnd3,Sf1
	rno-miR-204-3p	GCUGGGAAGGCAAAGGGACGUU	Pdcl3,Pla2r1,Banf1,Dtd2, Xpo7,Lrnf2,Lmf1,Timp2
rno-miR-483	rno-miR-483-5p	AAGACGGGAGAAGAGAAGGGAG	Bcor,Otc,Xk,Cybb,Lancl3,P orch,Ebp,Wdr13,Rbm3
	rno-miR-483-3p	CACUCCUCCCCUCCGUCUUGU	Sra1,Fech,Galr1,Bap1,Scn1b ,Git1,Hipk2,Wipf3,Tns3
rno-miR-322	rno-miR-322-5p	CAGCAGCAAUUCAUGUUUUGGA	Lrp6,Ovov1,G2e3,Hoxa4, Lamc2,mitf,Cxcl12,Ret
	rno-miR-322-3p	AAACAUGAAGCGCUGCAACA	Wdr26,Klf8,Gnrhr,Polh,Irf5, Nfib,Bdp1,Gpm6a

rno-miR-329	rno-miR-329-5p	AGAGGUUUUCUGGGUCUCUGUUUC	Nadk2,Ccdc2,Strd4,Gfra3, Cd14,Gde1,Scai,Ehd1
	rno-miR-329-3p	AACACACCCAGCUAACCUUUUU	Brat1,Sox6,Wdte1,Pbx3, Rgs8,Tead2,Ankfy1,Etv3
rno-miR-196b	rno-miR-196b-5p	UAGGUAGUUUCCUGUUGUUGGG	Xylt1,Tmx3,Ptcd1,neur11, Nxt2,Cu14b,Wtap,Fv1
	rno-miR-196b-3p	UCGACAGCACGACACUGCCUUCA	Girx5,Cldn2,Zmiz1,Megf11, Gfy,KdeIr1,Smarcc1,Mpp2
rno-miR-449a	rno-miR-449a-5p	UGGCAGUGUAUUGUAGCUGGU	Calr4,Pan3,Mafk,Gpc2,, Traf3,Rftn2,Sumo1,Naif1
	rno-miR-449a-3p	CAGCUAACAUUGCAACUGCUCUC	Nfib,Tnp1,Ikzf2,Sstr1, Smoc2,Mill1,Casp2,Ube2n
rno-miR-324	rno-miR-324-5p	CGCAUCCCCUAGGGCAUUGGUGU	Prr36, Zfp498, Cinp, Laptm5,Nipal3, Poc5, Ctdsp1, Gla
	rno-miR-324-3p	CCACUGCCCCAGGUGCUGCUGG	Pdha1, Baz2a, Nipal3, Enc1, Gypc, Map2, Cytip, Tcf23
rno-miR-448	rno-miR-448-5p	AACAUCCUGCAUAGUGCUGCCA	Atp6ap2, Bcor, Otc, Sytl5, SrpX, Xk, Cybb, Dynlt3
	rno-miR-448-3p	UUGCAUAUGUAGGAUGUCCCA	Bcor, SrpX, Tbc1d25, Rbm3,Ftsj1, Wdr13, , Gata1,Pqbp1
rno-miR-375	rno-miR-375-5p	GCGACGAGCCCCUCGCACAAACC	Mrm2, Sfn,Usp48, Fgf1, Rsad2, C5ar1, Grin2d, Gprin3
	rno-miR-375-3p	UUUGUUCGUUCGGCUCGCGUGA	Mark3, Scamp1, Nme5,Cys1, Asap2, Nr1h2,Gsg2, Relt
rno-miR-702	rno-miR-702-5p	GUGAGUGGGGUGGUUGGCAUG	Akap4, Cinp,Traf3, Stpg1, Lypla2, Kazn, Pla2g2d, Wdr41,
	rno-miR-702-3p	UGCCACCCUUUACCCACUCCA	Xk,Tsr2, Ephb4, Cdk5r2, Arl13a, Zadh2, Rce1, Snx32,
rno-miR-541	rno-miR-541-5p	AAGGGAUUCUGAUGUUGGUCACAC U	Syp, Pan3, Usp48, Shtn1, Smn1, Mat1a, Tnfsf18, Park7
	rno-miR-541-3p	AGUGGCGAACACAGAAUCCAUC	Cpsf2, Cldn2, Atf6, Pep411, Ptpn13, Rhot1, Coa4, Ptprr
rno-mir-218b		CAUGGUUAGAUCAAGCACAA	SMCP,POLR2G, TEX14, H2AFZ, ERI2, TRPV5, PIGW, PPP1R14C, SGK494, MBNL3

Table S1. The sequence and target gene of differentially expressed miRNAs