

## Mir-134-3p Driven by Anisomycin Impairs Ovarian Cancer Stem Cell Activity through Inhibiting GPR137 Expression

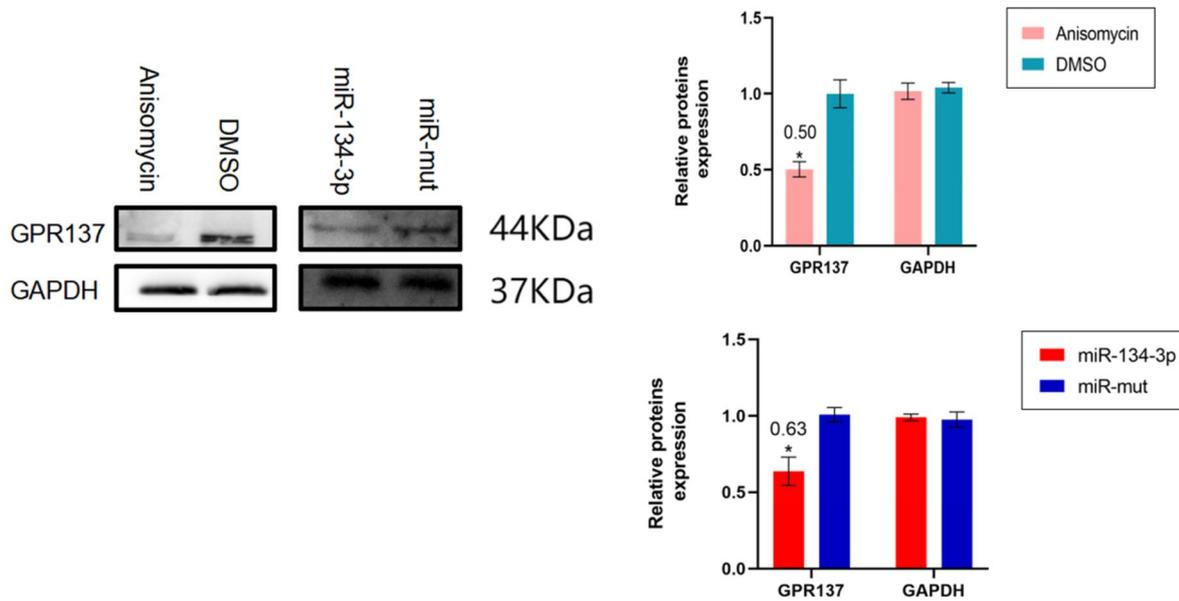


Figure S1 The results of western blot assay.

\*  $P < 0.05$  vs. DMSO, \*  $P < 0.05$  vs. miR-mut.

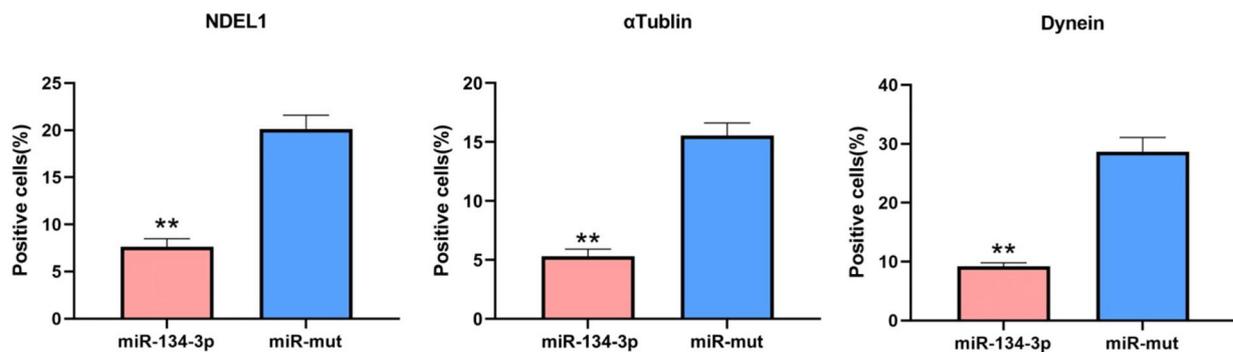


Figure S2 The statistical analysis results of Immunofluorescence staining.

\*\*  $P < 0.01$  vs. miR-mut, n = 3.

**Table S1 The results of qPCR.**

Gene	miR-mut	miR-134-3p	Anisomycin
<b>BASP1</b>	1.00±0.00	0.44±0.02	0.66±0.03
<b>XRCC3</b>	1.01±0.08	0.53±0.05	0.98±0.09
<b>SSR4</b>	1.00±0.07	0.49±0.04	0.54±0.01
<b>OR6V1</b>	1.04±0.18	43.70±0.23	0.47±0.06
<b>FERD3L</b>	1.00±0.02	0.96±0.00	1.99±0.05
<b>NDEL1</b>	1.00±0.00	0.53±0.01	0.39±0.02
<b>HSP12B</b>	1.01±0.10	19.69±0.19	0.83±0.04
<b>18S rRNA</b>	1.01±0.08	1.01±0.09	1.00±0.02

**Table S2 The results of qPCR.**

Gene	Anisomycin	DMSO
<b>TUBA1A</b>	0.71±0.02	1.01±0.09
<b>DYNEIN</b>	0.39±0.02	1.08±0.04
<b>BAX</b>	1.60±0.06	1.01±0.08
<b>BCL2</b>	0.33±0.02	1.01±0.07
<b>CCND3</b>	0.26±0.02	1.00±0.07
<b>CDK2</b>	0.38±0.02	1.00±0.03
<b>18S rRNA</b>	1.00±0.03	1.00±0.03

**Table S3 The results of qPCR.**

Gene	miR-134-3p	miR-mut
<b>GPR137</b>	0.54±0.05	1.00±0.03
<b>TUBA1A</b>	0.88±0.10	1.00±0.05
<b>DYNEIN</b>	0.71±0.05	1.00±0.06
<b>BAX</b>	1.60±0.05	1.00±0.02
<b>BCL2</b>	0.82±0.04	1.01±0.08
<b>CCND3</b>	0.19±0.03	1.00±0.02
<b>CDK2</b>	0.93±0.04	1.00±0.04
<b>18S rRNA</b>	1.01±0.09	1.01±0.07

**Table S4 The results of Western blotting.**

<b>Proteins</b>	<b>Anisomycin</b>	<b>DMSO</b>
<b>α Tubulini</b>	0.60±0.06	1.00±0.08
<b>BAX</b>	4.26±0.48	1.00±0.05
<b>NDEL1</b>	0.45±0.06	1.00±0.21
<b>Dynein</b>	0.46±0.08	1.00±0.04
<b>CDK2</b>	0.40±0.03	1.00±0.02
<b>CCND3</b>	0.50±0.09	1.00±0.11
<b>GAPDH</b>	1.00±0.04	1.00±0.01

**Table S5 The results of Western blotting.**

<b>Proteins</b>	<b>miR-134-3p</b>	<b>miR-mut</b>
<b>α Tubulin3</b>	0.81±0.07	1.00±0.12
<b>BAX</b>	2.51±0.08	1.00±0.08
<b>NDEL1</b>	0.69±0.08	1.00±0.13
<b>Dynein</b>	0.46±0.07	1.00±0.07
<b>CDK2</b>	0.57±0.06	1.00±0.12
<b>CCND3</b>	0.74±0.06	1.00±0.05
<b>GAPDH</b>	1.03±0.09	1.00±0.12