Figure S1. Treatment with G-Rh2 induced almost no death in End1/e6e7 cells. Flow cytometry was used to detect the apoptotic rate of End1/e6e7 cells after G-Rh2 treatment. The early apoptotic rate was quantified by IDEAS software v6.1. Data are presented as the mean  $\pm$  standard deviation of three independent experiments and were analyzed by one-way ANOVA followed by Dunnet's post-hoc test. ns >0.05 vs. control, indicated no statistically significant difference. G-Rh2, ginsenoside Rh2; ns, not significant.

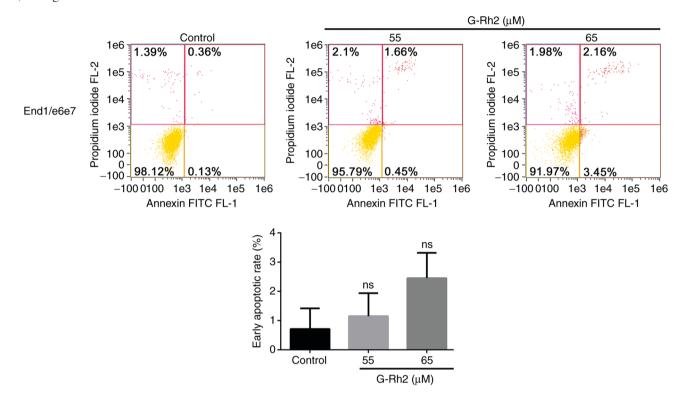


Figure S2. Greater inhibition of mitochondrial OXPHOS than of glycolysis in HeLa cells exposed to G-Rh2. (A) OCR curves after G-Rh2 treatment, as measured by the Seahorse XFp Mitochondrial Stress Test Kit. (B) Related indicators of mitochondrial OXPHOS including basal OCR, ATP-linked, proton leak, maximal OCR and reserve capacity after G-Rh2 treatment were calculated according to the OCR curves. (C) Curves of ECAR rates after G-Rh2 treatment, as detected by the Seahorse XFp Glycolysis Stress Test Kit. (D) Related indicators of glycolysis including glycolysis, glycolytic reserve and glycolytic capacity after G-Rh2 treatment were calculated according to the ECAR curves. The histograms were created separately but then plotted together. Data are presented as the mean ± standard deviation of three independent experiments and were analyzed by one-way ANOVA followed by Dunnet's post-hoc test. \*P<0.05; \*\*P<0.01; \*\*\*P<0.001 vs. control. G-Rh2, ginsenoside Rh2; MMP, mitochondrial membrane potential; mtROS, mitochondrial reactive oxygen species; OCR, oxygen consumption rate; ECAR, extracellular acidification rate; OXPHOS, oxidative phosphorylation; FCCP, carbonyl cyanide-4-(trifluoromethoxy) phenylhydrazone; rot, rotenone (ETC complex I inhibitor); AA, antimycin A (ETC complex III inhibitor); 2-DG, 2-deoxy-d-glucose (glycolysis inhibitor); ETC, electron transport chain.

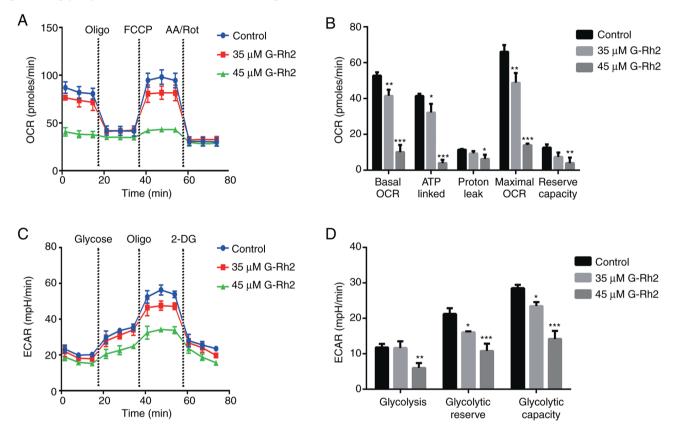


Table SI. Primer sequences used in the study.

Primer purpose/direction	Sequence (5'-3')	Enzyme
pET28a (+)-6xHis-ATP5F1B		
construction		
Sense	CGAGCTCGCGCAAACATCTCCTTCGC	SacI
Antisense	CCAAGCTTTCACGATGAATGCTCTTC	<i>Hin</i> dIII
pCMV-myc-UQCRC1		
construction		
Sense	GGAATTCTAATGGCGGCGTCCGTGG	EcoRI
Antisense	GGTACCCTAGAAGCGCAGCCAGAAC	KpnI
pCMV-10-Flag-NDUFS1		
construction		
Sense	ATAAGAATGCGGCCGCTATGTTAAGGATAC	NotI
Antisense	GGATCCTCAGCATATGGATGGTTCC	<i>Bam</i> HI