

S4 Table. Reduction of *Csk*, *Ulk3*, and *Cyp1a2* mRNA levels in cells after treatment with siRNAs.

siRNA Name	Target sequence (5'-3')	Reduction rate (%)	Cell line
Scrambled control siRNA	ACGUGACACGUUCGGAGAA		
<i>Csk</i> #1	GAAUGUAUUGCCAAGUACA	11.7	
<i>Csk</i> #2	ACAAGAAGUACGAAUCUUA	29.3	B16F10
<i>Csk</i> #3	CUGGCCAUCCGGUACAGAA	40.1	
<i>Ulk3</i> #1	UAUCUACCUCAUCAUGGAG	No reduction	
<i>Ulk3</i> #2	ACGAAACAUUCUCACUUGGA	73.4	B16F10
<i>Ulk3</i> #3	GGUUAUUUCUAAAGUUAGA	74.5	
<i>Cypla2</i> #1	CUGAACAUCGUGAAUAACA	22.1	
<i>Cypla2</i> #2	GAUUGUCAACAUUGUCAAU	44.9	NIH3T3
<i>Cypla2</i> #3	GGUUUAAGACCUUCAAUGA	44.9	