

Table SI. Sequences of siRNAs used in the present study.

siRNA ID	Target gene symbol	Sense sequence (5'-3')	Antisense sequence (5'-3')
s9491	NFE2L2	GAAUGGUCCUAAAACACCAtt	UGGUGUUUUAGGACCAUUCtg
s9492	NFE2L2	CGUUUGUAGAUGACAAUGAtt	UCAUUGUCAUCUACAAACGgg
s9493	NFE2L2	CAGUCUUCAUUGCUACUAAtt	UUAGUAGCAAUGMGACUGgg
Negative control siRNA	None	GCGCGCUUUGUAGGAUUCGtt	CGAAUCCUACAAAGCGCGCtt
NFE2L2, NFE2 like BZIP transcription factor 2 (Nrf2).			

Table SII. Characteristics of the 104 patients evaluated for Nrf2, ABCB1 and p-Fyn expression using immunohistochemistry.

Characteristics		No. of patients (n=104)
Age, years		
	Median	65
	Range	36-76
Sex, n (%)		
	Male	59 (57)
	Female	45 (43)
Tumor histology, n (%)		
	Adenocarcinoma	69 (66)
	Squamous cell carcinoma	26 (25)
	Large cell carcinoma	3 (3)
	Adenosquamous cell carcinoma	3 (3)
	Pleomorphic carcinoma	3 (3)
Pathological stage (UICC, 7th edition), n (%)		
	Stage IIA	18 (17)
	Stage IIB	32 (31)
	Stage IIIA	54 (52)
Immunohistochemistry, n (%)		
	Nrf2-positive	47 (45)
	ABCB1-positive	53 (50)
	p-Fyn-positive	48 (46)