

Supplementary Table 1. Univariate cox proportional hazard regression analysis results of immunogenic cell death-related lncRNAs in endometrial carcinoma.

id	HR	HR.95L	HR.95H	p-value
AC004585.1	0.137544605	0.028907838	0.654442514	0.012677857
SCARNA9	0.609948644	0.375630052	0.990435524	0.045627828
FAM198B-AS1	0.038476824	0.003259061	0.454261509	0.00969681
FKBP14-AS1	6.85210706	1.262031183	37.20301987	0.025776038
DLEU2	2.951432972	1.008909633	8.634030548	0.048136649
AC010615.2	2.235437172	1.103270917	4.529421806	0.02556599
AL035530.2	6.488149496	1.171351622	35.9380421	0.032270095
AC007611.1	0.109107868	0.014086926	0.845076266	0.033911017
SOCAR	0.007758384	0.000123896	0.485829692	0.021337072
UNQ6494	0.000839063	2.03E-06	0.347128919	0.021214411
AC012645.3	0.042936915	0.002151408	0.856917101	0.039297186
AC007996.1	1.566942263	1.016678532	2.415028919	0.041860121
FBXO30-DT	6.908472159	1.039536648	45.91178932	0.045490537
LINC01943	0.063296412	0.006455071	0.620664841	0.017814751
AL161431.1	1.459764949	1.04847224	2.032398784	0.025071777
AC006369.1	0.016749967	0.000316831	0.885524833	0.043381499

Supplementary Table 2. qRT-PCR primers used in this study

Gene name	Forward primer	Reverse primer
SCARNA9	5'-AAG GGC ATA TGT CTG GTG TGT GTG-3'	5'-CCC CAC CCT CAA TCT CAT TCA TTC C-3'
FAM198B-AS1	5'-GGA ATG AAG CCA CCC TTG ACT GAG-3'	5'-GTC TTC TGT GCC GTG TCT CCT TG-3'
FKBP14-AS1	5'-CGA GAC CAG CCT GAC CAA CAT G-3'	5'-CTA CAG ACA CAT GCC ACC ACA CTC-3'
FBXO30-DT	5'-ACC AGG GTC TGC TTC TGT GAG G-3'	5'-CAG TTT GGC ATG GCT GGA GAG G-3'
LINC01943	5'-TGG CAT CAG TCT GGT TGG ATT ACA C-3'	5'-CAG CAG GTA TCA GGC ACA GCA TC-3'
AL161431.1	5'-GGC TCT CTG GCA AGA ACC ACA AG-3'	5'-GAA CGG TGT TGG GTC TGT CAG TAG-3'
β -actin	5'-ACC GAG CGT GGC TAC AGC TTC ACC-3	5'-AGC ACC CGT GGC CAT CTC TTT CTC G-3'

Supplementary Table 3. ShRNA sequences used for cell transfection.

ShRNA	Sequence (5'-3')
sh-FKBP14-AS1-1	F-CCG GAG AAT CAG ACA GAA AAT CCC TCG AGG GAT TTT CTG TCT GAT TCT TTT TTG R-AAT TCA AAA AAA GAA TCA GAC AGA AAA TCC CTC GAG GGA TTT TCT GTC TGA TTC T

	F-CCG GAG ACC AAA ACA GTA TTT TAC TCG AGT
sh-FKBP14-AS1-2	AAA ATA CTG TTT TGG TCT TTT TTG
	R-AAT TCA AAA AAA GAC CAA AAC AGT ATT TTA
	CTC GAG TAA AAT ACT GTT TTG GTC T
	F-CCG GTC TAC TAA AAA TAC AAA AAC TCG AGT
sh-FKBP14-AS1-3	TTT TGT ATT TTT AGT AGA TTT TTG
	R-AAT TCA AAA AAT CTA CTA AAA ATA CAA AAA
	CTC GAG TTT TTG TAT TTT TAG TAG A
	F-CCG GGG ACC TTT TTT CCT TTC GAC TCG AGT
sh-FBXO30-DT-1	CGA AAG GAA AAA AGG TCC TTT TTG
	R-AAT TCA AAA AAG GAC CTT TTT TCC TTT CGA
	CTC GAG TCG AAA GGA AAA AAG GTC C
	F-CCG GAG AAG CAT AAG CTG CAA TAC TCG AGT
sh-FBXO30-DT-2	ATT GCA GCT TAT GCT TCT TTT TTG
	R-AAT TCA AAA AAA GAA GCA TAA GCT GCA ATA
	CTC GAG TAT TGC AGC TTA TGC TTC T
	F-CCG GGC TGC AAT AAA AAT TTG GTC TCG AGA
sh-FBXO30-DT-3	CCA AAT TTT TAT TGC AGC TTT TTG
	R-AAT TCA AAA AAG CTG CAA TAA AAA TTT GGT
	CTC GAG ACC AAA TTT TTA TTG CAG C
	F-CCG GGG AAA TAT CCC ATT AGA TTC TCG AGA
sh-AL161431.1-1	ATC TAA TGG GAT ATT TCC TTT TTG
	R-AAT TCA AAA AAG GAA ATA TCC CAT TAG ATT
	CTC GAG AAT CTA ATG GGA TAT TTC CTT TTT G
	F-CCG GGC TAG GAT ACA AGA GAA ACC TCG AGG
sh-AL161431.1-2	TTT CTC TTG TAT CCT AGC TTT TTG
	R-AAT TCA AAA AAG CTA GGA TAC AAG AGA AAC
	CTC GAG GTT TCT CTT GTA TCC TAG C
	F-CCG GGC CTG AAA TCA TTT ATT TCC TCG AGG
sh-AL161431.1-3	AAA TAA ATG ATT TCA GGC TTT TTG
	R-AAT TCA AAA AAG CCT GAA ATC ATT TAT TTC
	CTC GAG GAA ATA AAT GAT TTC AGG C

Supplementary Table 4. The risk model for immunogenic cell death-related lncRNAs.

Immunogenic cell death-related lncRNA	Coefficient
SCARNA9	-0.401528645653866
FAM198B-AS1	-1.90526441273924
FKBP14-AS1	2.21856134166291
FBXO30-DT	1.63157340507073
LINC01943	-2.36050864717649
AL161431.1	0.600416817300287

