Supporting Information

Nanoparticle-Based Follistatin Messenger RNA Therapy for Reprogramming Metastatic Ovarian Cancer and Ameliorating Cancer-Associated Cachexia

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Figure S1: Representative histopathological examination of murine liver sections after chronic injection of Luc mRNA LNPs.



Figure S2: Blood analysis for evaluation of systemic toxicity of chronically injected Luc mRNA LNPs. (A) erythrocytes and (B) leukocytes evaluation; (C) blood chemistry; (D) markers of liver and pancreatic inflammation.



Figure S3: Expression of hepatic inflammatory phase reactants in mice receiving 3 daily injections of *Luc* mRNA LNPs.



Figure S4: Proportion and type of cells internalizing GFP mRNA LNPs and expressing GFP. (A) Flow cytometry analysis and gating approach in evaluating in vitro and ex vivo samples, Comp - 561- 1 - A - anti-HLA-ABC antibody; Comp - 628 - 1 - A - anti-GFP antibody. (B)

Analysis of in vitro subculture of ES-2-WT and Raw 264.7 cells (3:1), where controls were treated with PBS and GFP mRNA LNP Tx samples were treated with GFP mRNA LNP formulation. (C) ES-2-WT-bearing mice were treated with GFP mRNA LNPs (2ug mRNA per mouse), control mice were injected with PBS. Collected IP lavage fluid containing mixture of ES-2-WT cells and murine immune cells was subjected to flow cytometry analysis.







Figure S6: In vitro evaluation of gene expression from ES-2-WT spheroids.





Correlation between FST and ActA concentrations in malignant ascites 80 -0.7046 60 FST (ng/mL) 95% CI -0.8683 to -0.4026 R^2 0.4965 40 20 0 2000 3000 1000 4000 ActA (pg/mL)

Figure S8: Correlation between FST and ActA concentrations from pooled samples of healthy and ES-2-WT-bearing mice, as well as mice treated with FST mRNA LNP formulation.

Table S1: Dynamic Light Scattering Properties of Assembled mRNA LNP platforms

LNP	Dynamic Light Scattering			
	d (nm)	PDI	ζ (mV)	
Luc mRNA LNP	84.15	0.148	9.07	
EGFP mRNA LNP	102.00	0.207	-2.38	
FST mRNA LNP	84.57	0.065	-7.78	
EGFP/FST mRNA LNP	86.55	0.217	3.04	
Empty LNP	72.22	0.116	1.55	

Table S2: Estimated mean survival for evaluation of combinatorial treatment with CDDP and FST or *FST* mRNA LNPs

Treatment	Mean	Std.	95%	95%
Group	Survival	Error	LCL	UCL
PBS	12.2	1.16	9.93	14.47
CDDP	21.2	1.66	17.96	24.44
FST _H	17.0	1.34	14.37	19.63
CDDP+FST _H	23.8	1.96	19.96	27.64
FST mRNA LNP	25.8	2.80	20.31	31.29
CDDP+ FST mRNA LNP	33.8	2.35	29.19	38.41

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Table S3: mRNA ORF sequences

FST mRNA ORF (CleanCap®)
ATGGTCCGCGCGAGGCACCAGCCGGGTGGGCTTTGCCTCCTGCTGCTGCTGCTCT
GCCAGTTCATGGAGGACCGCAGTGCCCAGGCTGGGAACTGCTGGCTCCGTCAAGC
GAAGAACGGCCGCTGCCAGGTCCTGTACAAGACCGAACTGAGCAAGGAGGAGTG
CTGCAGCACCGGCCGGCTGAGCACCTCGTGGACCGAGGAGGACGTGAATGACAA
CACACTCTTCAAGTGGATGATTTTCAACGGGGGGCGCCCCCAACTGCATCCCCTGT
AAAGAAACGTGTGAGAACGTGGACTGTGGACCTGGGAAAAAATGCCGAATGAAC
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CCTCTGAGCAATATCTCTGTGGGAATGATGGAGTCACCTACTCCAGTGCCTGCC
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CAGCGAGTGTGCCATGAAGGAAGCTGCCTGCTCCTCAGGTGTGCTACTGGAAGTA
AAGCACTCCGGATCTTGCAACTCCATTTCGGAAGACACCGAGGAAGAGGAGGAA
GATGAAGACCAGGACTACAGCTTTCCTATATCTTCTATTCTAGAGTGGTAA
<i>FLuc</i> mRNA ORF Sequence (CleanCap® FLuc mRNA, L-7202)
AUGGAGGACGCCAAGAACAUCAAGAAGGGCCCCGCCCCUUCUACCCCCUGGA
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UGCCCGGCACCAUCGCCUUCACCGACGCCCACAUCGAGGUGGACAUCACCUACG
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CUGAACACCAACCACCGGAUCGUGGUGUGCAGCGAGAACAGCCUGCAGUUCUU
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EGFP mRNA ORF Sequence (CleanCap® EGFP mRNA, L-7601) AUGGUGAGCAAGGGCGAGGAGCUGUUCACCGGGGUGGUGCCAUCCUGGUCGA GCUGGACGGCGACGUAAACGGCCACAAGUUCAGCGUGUCCGGCGAGGGCGAGG GCGAUGCCACCUACGGCAAGCUGACCCUGAAGUUCAUCUGCACCACCGGCAAG CUGCCCGUGCCCUGGCCCACCUCGUGACCACCCUGACCUACGGCGUGCAGUGC UUCAGCCGCUACCCCGACCACAUGAAGCAGCACGACUUCUUCAAGUCCGCCAU GCCCGAAGGCUACGUCCAGGAGCGCACCAUCUUCUUCAAGGACGACGACU ACAAGACCCGCGCCGAGGUGAAGUUCGAGGGCGACACCCUGGUGAACCGCAUC GAGCUGAAGGGCAUCGACUUCAAGGAGGACGGCAACAUCCUGGGGGCACAAGCU GGAGUACAACUACAACAGCCACAACGUCUAUAUCAUGGCCGACAAGCAGAAGA ACGGCAUCAAGGUGAACUUCAAGGACGCCACAACAUCGAGGACGGCAGCAGGUG CAGCUCGCCGACCACUACCAGCAGAACACCCCCAUCGGCGACGACGGCAGCGUG CUGCCCGACAACCACUACCAGCAGAACACCCCCAUCGGCGACGACGGCAGCGUG CUGCCCGACAACCACUACCUGAGCACCCCAGUCCGCCCGGGACAAGACCCCAAC GAGAAGCGCGAUCACAUGGUCCUGCUGGAGUUCGUGACCGCCGCGGGAUCAC UCUCGGCAUGAACGAGCUGUACAAGUAA