

Supporting Information

Nanoparticle-Mediated Intracellular Delivery Enables Cryopreservation of Human Adipose- Derived Stem Cells Using Trehalose as the Sole Cryoprotectant

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Table S1. Encapsulation efficiency (EE) and loading content (LC) of propidium iodide (PI) with and without genipin crosslinking using the procedure given in Scheme 1

Feeding ratio PI : NPs	Genipin crosslinking	EE, %	LC, %
1 : 10	No	3.7 ± 0.9	0.4 ± 0.1
1 : 10	Yes	36.4 ± 5.8	4.1 ± 0.2

NPs: Pluronic F127-chitosan nanoparticles

Table S2. Encapsulation efficiency (EE) and loading content (LC) at different feeding ratios of free trehalose (fTre) to Pluronic F127-chitosan NPs in weight and the diameter of the resultant GNCs-encapsulated trehalose (nTre) determined by dynamic light scattering (DLS) at different temperature and pH values. Also shown are the data of size for empty GNPs.

Feeding ratio fTre : GNPs	EE, %	LC, %	Diameter of GNPs/nTre, nm			
			pH 5		pH 7	
			22 °C	37 °C	22 °C	37 °C
0 :GNPs	N/A	N/A	82.2 ± 4.6	37.3 ± 2.2	28.0 ± 3.5	24.8 ± 0.4
1 : 50	63.5 ± 4.8	1.4 ± 0.1	38.1 ± 8.1	24.9 ± 2.1	25.8 ± 8.3	19.9 ± 3.3
1 : 20	59.1 ± 3.1	3.5 ± 0.2	57.0 ± 15.0	30.6 ± 3.6	24.0 ± 7.1	22.1 ± 5.6
1 : 10	40.6 ± 5.8	4.6 ± 0.4	91.5 ± 18.2	33.5 ± 1.5	26.8 ± 6.2	20.1 ± 2.4
1 : 1	4.9 ± 0.6	5.6 ± 0.6	93.4 ± 18.3	35.2 ± 3.1	28.5 ± 5.1	21.8 ± 2.9
10 : 1	1.3 ± 0.1	14.4 ± 1.5	105.4 ± 14.5	47.0 ± 7.2	36.2 ± 2.6	28.7 ± 4.1
20 : 1	1.2 ± 0.1	26.6 ± 0.8	137.5 ± 16.2	49.5 ± 5.6	42.3 ± 6.1	29.5 ± 6.4

All data are presented as mean ± standard deviation and N/A represents “ not applicable”.

Table S3. Genes and the corresponding primers used for real time RT-PCR studies

Gene		Primer
Sox2	Forward	5'- CCC ACC TAC AGC ATG TCC TAC TC -3'
	Reverse	5'- TGG AGT GGG AGG AAG AGG TAA C -3'
Klf4	Forward	5'- GAA ATT CGC CCG CTC CGA TGA -3'
	Reverse	5'- CTG TGT GTT TGC GGT AGT GCC -3'
Nanog	Forward	5'- CCT CCT CCA TGG ATC TGC TTA TTC A -3'
	Reverse	5'- CAG GTC TTC ACC TGT TTG TAG CTG AG -3'
Oct4	Forward	5'- GAA ATT CGC CCG CTC CGA TGA -3'
	Reverse	5'- CAT AGT CGC TGC TTG ATC GCT TG -3'
GAPDH	Forward	5'- GAG TCA ACG GAT TTG GTC GTA -3',
	Reverse	5'- ATG GGT GGA ATC ATA TTG GAA -3'