LDH	(XRD, FTIR, SEM/TEM,	Z-Average	PDI	ζ-	Note
	PSD) Availability	size (nm)		potential	
				(mV)	
1. LDH	(XRD, FTIR, TEM, PSD)	110	0.14	43	
2. sLDH	(XRD, FTIR, TEM, PSD)	50	0.17	39	
3. ILDH	(TEM, PSD)	210	0.18	43	
4. LDH-BSA in	(FTIR, TEM, PSD)	150	0.22	-20	
5. LDH-BSA in PBS		180	0.20	-15	LDH:BSA = 2:25
6. LDH-FITC	(XRD, FTIR, TEM, PSD)	100	0.20	41	10% FITC in total anions exchanged
7. LDH-FITC-BSA in H2O		180	0.29	-23	As above
8. LDH-FITC-BSA in PBS		167	0.24	-13	As above
9. LDH-LMWH	(XRD, FTIR, SEM, PSD)	123	0.23	35	10% LMWH in total anions exchanged
10.LDH-LMWH- BSA in H2O		198	0.26	-19	As above. LDH:BSA = 2:12.5
11.LDH-LMWH- BSA in culture medium		165	0.24	-17	As above. LDH:BSA = 2:12.5
12.LDH-MTX	(XRD, FTIR, SEM, PSD)	103	0.16	40	10% MTX in total anions exchanged
13.LDH-MTX-BSA in H2O		173	0.25	-17	As above.
14.LDH-MTX-BSA in culture medium		179	0.25	-15	As above
15.LDH-BSA-ODN in H2O	(PSD)	170	0.15	-24	LDH:ODN = 40:1
16.LDH-BSA-ODN in PBS	(PSD)	173	0.19	-16	LDH:ODN = 40:1
17.LDH-ODN (10:1)	(XRD, TEM)	200-700		-26 to 40	Different ODNs

Table S1. Supporting data summary for reviewers

a) Following supplementary figures were displayed sequentially;

b) LDH:BSA mass ratio was 2:5, and otherwise indicated.

Supplementary Figures

1. LDH



Refs:

Xu, ZP; Stevenson, G; Lu, CQ; Lu, GQ Max; Bartlett, PF; Gray, PP: *J. Am. Chem. Soc.* **128**, 36-37, 2006; Xu, ZP; Stevenson, G; Lu, CQ; Lu, GQ: *J. Phys. Chem. B* **110**, 16923-16929, 2006; Xu, ZP; Kurniawan, ND; Bartlett, PF; Lu, GQ Max: *Chem. Eur. J.* **13**, 2824-2830, 2007; Xu, ZP; Walker, TL; Liu, K-L; Cooper, HM; Lu, G.Q. Max; Bartlett, PF: *Inter. J. Nanomed.* **2**, 163-174, 2007; Xu, ZP; Jin, YG; Liu, SM; Hao, ZP; Lu, GQM: *J. Colloid Interface Sci.* **326**, 522-529, 2008





A. PSD; B. TEM; C. XRD; D. FTIR

Refs:

Dong HY; Parekh HS; **Xu ZP**: *Journal of Colloid and Interface Science*, **437**, 10-16, 2014; Dong HY; Chen M; Rahman S; Parekh HS; Cooper HM; **Xu ZP**: *Applied Clay Science*, **100**, 66-75, 2014; Chen, M; Cooper, HM; Zhou, JZ; Bartlett, PF; Lu, G.Q. (Max); **Xu, ZP**: *Journal of Colloid and Interface Science* **390**, 275-281, 2013.

3. ILDH



Ref:

Xu, ZP; Stevenson, G; Lu, CQ; Lu, GQ: J. Phys. Chem. B 110, 16923-16929, 2006; TEM not published.

4. LDH-BSA in H2O

TEM (LDH-BSA 5:2 in water)



(Bar length = 200 nm)

(Figure 1E)



PSD: Refer to Fig 1B in the manuscript



5. LDH-BSA in PBS





(Scale bar = 200 nm, not published yet)

PSD: refer to Fig 2A in the manuscript





(A) Infrared spectra of (1) hexagonal LDH-CI; (2) rod-like LDH_{FITC-ROD} (10% Cl⁻ equivalent FITC); (3) hexagonal LDH_{FITC-HEX} (10% Cl- equivalent FITC); and (4) LDH_{FITC} with 50% of Cl⁻ ions exchanged by FITC; (B) XRD patterns of LDH-CI/FITC samples. (1): hexagonal LDH_{FITC-HEX} (10% Cl⁻ equivalent FITC); (2) rod-like LDH_{FITC-ROD} (10% Cl-equivalent FITC); and (3) LDH_{FITC} (50% Cl⁻ equivalent FITC). #: brucite-like phase; *: gibbsite phase.

Ref:

Xu, ZP; Niebert, M; Porazik, K; Walker, TL; Cooper, HM; Middelberg, APJ; Gray, PP; Bartlett, PF; Lu, GQM: *J Control Release.* **130**, 86-94, 2008.

9. LDH-LMWH







XRD, FTIR spectra and TEM of (a) Cl-LDH, (b) LMWH10-LDH, (c) LMWH20-LDH, (d) LMWH50-LDH, (e) LMWH100-LDH, (f) heparin sodium salt (MW = 4^{6} kD).

Ref:

Gu, Z; Thomas, A; Xu, ZP; Campbell, J; Lu, GQ Max: Chem. Mater. 20, 3715-3722, 2008.

12. LDH-MTX

TEM (bar = 100 nm)



PSD



Data were not published yet.

Note that in this LDH, there were 90% Cl⁻ and 10% MTX⁻ in the interlayer to balance the positive charges in the hydroxide layers. In XRD pattern, the major phase was attributed to LDH-Cl (peaks at 14 and 28°) with a small portion of minor phase LDH-MTX (peak at 6°).

As also shown in the XRD patterns for Data 6 and 9, when there was more LMWH/FITC, the XRD peak belonging to LDH-LMWH/FITC became more obvious. However, the aggregation of LDH-LMWH/FITC nanoparticles became severe, increasing the particle size and losing the monodispersivity (PCS data not shown). Therefore, for a particular drug, LDH-drug nanoparticles are normally dispersed well in aqueous suspension when the drug loading in LDH is equivalent to replacing 10-20% of total Cl⁻ in precursor LDH-Cl, which avoids any aggregation while the drug loading is high enough for subsequent biological tests. So in this research, we chose this loading range (Refer to Section 2.2.2 in the main text) in order to obtain well dispersed LDH suspensions.



In the manuscript, Fig.3 was changed to the graph above (BSA-LDH-ODN in PBS was added):

(A)





(A) TEM image of LDHs; (B) XRD patterns of LDH (a) and dsDNA-Htt#1-LDH (b); High resolution TEM images of LDH (C) and dsDNA-Htt#1-LDH (D), Bars; interlayer width without dsDNA intercalation, arrow; expansion due to dsDNA intercalation.

Ref:

Wong, YY; Cooper, HM; Zhang, K; Chen, M; Bartlett; PF; **Xu, ZP**: *J Colloid Interface Sci.* **369**, 453-459, 2012.