## Supplementary figure legends

**Figure S1: Chemical structure of the identified enhancers**. Chemical structures of compounds #1 to #24 (**a**), #25 to #48 (**b**) and #49 to #50 (**c**).

**Figure S2:** Determination of cytotoxicity based on cell number. Cell number was determined during the primary screen done in HeLa cells for LNPs (**a**) or Chol-siRNAs (**b**). Cell number was determined during the screen performed in human primary fibroblasts for LNPs (**c**) or Chol-siRNAs (**d**), and in mouse primary hepatocytes for LNPs (**e**) or Chol-siRNAs (**f**). Note that the compounds order in the present graphs are similar to those of the figures 2 and 3.

Figure S3: PD739 and PD740 are transfection reagent. (a) Z-score of GFP intensity versus DMSO treated condition. (b) Number of cells normalized to DMSO treated condition.

Figure S4: Cytotoxicity determination under compounds dose response. The number of cells from the experiments shown in the figure 2d top panels (increasing doses of compounds) was determined for four enhancers of LNPs (a) and of Chol-siRNAs (b).

Figure S5: BADGE and CPW1-J18 are both improving silencing by two different mechanisms. (a) GFP down-regulation 72 h after transfection with LNP-siRNA (40 nM, top panels) and uptake of LNP-siRNA-alexa647 (40 nM, 5 h incubation, bottom panels) after treated with DMSO (right panels), BADGE (middle panels) or CPW1-J18 (right panels) in HeLa and HeLa GFP cells. (b,c) Quantification of the uptake of LNP-siRNA-alexa647 (c) and of GFP down-regulation (b) in HeLa and HeLa GFP cells. Mean  $\pm$  s.e.m. n=3 (\*\*Pvalue<0.01, \*\*\*Pvalue<0.001).

Figure S6: BADGE treatment does not alter the mean number of siRNAgold particles per LNPs. Quantitative analysis of the number of siRNA-gold per LNPs after treatment with DMSO or BADGE. Mean  $\pm$  s.e.m. n=3 (Pvalue=0.88). Figure S7: Profile parameters for QMPIA on chemicals impact on EGF and transferrin endocytosis. Description of the different parameters used to drawn the endocytic profiles and exemplified here by hydroxychloroquine.

Figure S8: Proposed lipid crosslinking mediated by reaction of BADGE with lipid head-groups. Description of the proposed reaction between BADGE and the lipid head-groups.







Figure S1c





PD 739 PD740 0 -2 -2 -4 -6 -6 -10 -12 -14 -16



Fig. S3





b





b



С







Figure S6



Figure S7



Figure S8