Supplemental Data



Suppl. Figure 1

Supplemental Figure 1. N-Exo and CytoD did not affect preadipocyte proliferation, adipogenesis, lipolysis, and adipocyte browning. HPA-v cells were incubated with 50 µg/L of exosomes or 2 µg/mL of CytoD for 24 h (for cell proliferation) or 7 days (for adipocyte differentiation), respectively. Adipocyte differentiation was induced using a standard protocol. (a) Effects of N-Exo and CytoD on cell proliferation. (b) Effects of N-Exo and CytoD on the protein levels of aP2 and ADPN. Mature adipocytes were incubated with 50 µg/L of exosomes or 2 µg/mL of CytoD for 24 h. (c) Effects of N-Exo and CytoD on glycerol concentration in culture medium. (d) Effects of N-Exo and CytoD on UCP1 protein levels. Data are presented as mean \pm SD, n=3. N-Exo: normal (NL20) cell-derived exosomes; aP2: adipocyte protein 2; ADPN: adiponectin; CytoD: cytochalasin D.



Suppl. Figure 2

Supplemental Figure 2. Inhibition of miR-425-3p mitigated the effects of C-Exo on cAMP/PKA signaling and lipophagy. Mature adipocytes were incubated with 50 μ g/L of exosomes for 24 h. (a) Effect of exosomes on the intracellular cAMP concentration. (b) Effect of exosomes on the intracellular PKA activity. (c, d) Effect of exosomes on the protein or phosphorylated protein levels of CREB, HSL, and PLIN 1. (e, f) Effect of exosomes on the protein levels of beclin 1 and LC3. Data are presented as mean \pm SD, n=3; *p< 0.05, **p< 0.01, ***p< 0.001 vs indicated group. N-Exo: normal (NL20) cell-derived exosomes; C-Exo: cancer (A549) cell-derived exosomes; Mir-Inh-C-Exo: cancer (A549) cell-derived exosomes with miR-425-3p inhibition.



Suppl. Figure 3

Supplemental Figure 3. C-Exo and miR-425-3p mimics suppressed insulin signaling. Mature adipocytes were starved serum for 6 h and then incubated with 50 μ g/L of exosomes for 24 h, following by stimulation with 100 nM of insulin for 10 min. (a) Effect of exosomes on Akt1 levels and phosphorylation. (b) Quantitative analysis of Akt1 protein levels in (a). (c) Quantitative analysis of Akt1 phosphorylation in (a). (d) Effect of miR-425-3p mimics on Akt1 levels and phosphorylation. (e) Quantitative analysis of Akt1 protein levels in (d). (f) Quantitative analysis of Akt1 phosphorylation in (d). Data are presented as mean ± SD, n=3; ***p< 0.001 *vs* indicated group. N-Exo: normal (NL20) cell-derived exosomes; C-Exo: cancer (A549) cell-derived exosomes; CNTL: control; INS: insulin.



Supplemental Figure 4. Akt1 overexpression had no effect on the C-Exo-reduced proliferation and differentiation in human preadipocytes. HPA-v cells were overexpressed with Akt1 and then incubated with 50 μ g/L of exosomes for 24 h (for cell proliferation) or 7 days (for adipogenic differentiation), respectively. Adipocyte differentiation was induced using a standard protocol. Oil red O staining and western blot were performed at day 7 of adipogenic differentiation. (a) Effect of exosomes on cell proliferation. (b) Effect of exosomes on oil red O staining. (c, d) Effect of exosomes on the protein levels of aP2 and ADPN. Data are presented as mean ± SD, n=3. N-Exo: normal (NL20) cell-derived exosomes; C-Exo: cancer (A549) cell-derived exosomes; aP2: adipocyte protein 2; ADPN: adiponectin; OE: overexpression.



Supplemental Figure 5. Akt1 overexpression partly mitigated the effects of C-Exo on adipocyte lipolysis and white adipocyte browning. Mature adipocytes were overexpressed with Akt1 and then incubated with 50 µg/L of exosomes for 24 h. (a) Effect of exosomes on oil red O staining. (b) Effect of exosomes on glycerol concentration in culture medium. (c, d) Effect of exosomes on UCP1 protein levels. Data are presented as mean \pm SD, n=3; *p< 0.05, **p< 0.01, ***p< 0.001 vs indicated group. N-Exo: normal (NL20) cell-derived exosomes; C-Exo: cancer (A549) cell-derived exosomes; OE: overexpression.

| 3' UTR WT | 3' UTR Mutant |
|-----------------|--|
| 5'-CUUCCCGAG-3' | 5'-CUCAAAGCG-3' |
| 5'-UUUCCCGAA-3' | 5'-UUCAAAGCA-3' |
| 5'-CUUCCCGAU-3' | 5'-CUCAAAGCU-3' |
| 5'-CUUCCCGAU-3' | 5'-CUCAAAGCU-3' |
| 5'-CUCCCGAAC-3' | 5'-CUAAAAGAC-3' |
| | 3' UTR WT 5'-CUUCCCGAG-3' 5'-UUUCCCGAA-3' 5'-CUUCCCGAU-3' 5'-CUUCCCGAU-3' 5'-CUCCCGAAC-3' |

Supplemental Table 1. Mutations of has-miR-425-3p-binding site