

Figure S1: SMPs-PLGA exhibit extra-lysosomal localization in NRVMs

(A) Transmission electron microscopy (TEM) and (B) Scanning electron microscopy (SEM) of SMPs-PLGA (Scale Bar = 100 nm). (C) Intracellular uptake rate of SMPs-PLGA-FITC in NRVMs (n = 3). (D) Left panel: Representative confocal images of NRVMs transfected with LAMP1-RFP (red) and incubated with SMPs-PLGA-FITC (green) for 8 and 24 h. Hoechst 33258 (blue) was used as counterstaining. Scale Bar = 10 μ m. Right panel: Intensity profile graphs of SMPs-PLGA-FITC and LAMP1-RFP. (E) 3-dimensional reconstruction of stack images of NRVMs transfected with LAMP1-RFP (red) and incubated with SMPs-PLGA-FITC (green) for 24 h. Hoechst 33258 (blue) was used as counterstaining. Scale Bar = 5 μ m for whole images, 2 μ m for magnified areas. (F) Orthogonal projection of stack images of NRVMs transfected with SMPs-PLGA-FITC (green) for 24 h. Hoechst 33258 (blue) was used as counterstaining. Scale Bar = 5 μ m for whole images, 2 μ m for magnified areas. 3258 (blue) for 24 h. Hoechst 33258 (blue) was used as counterstaining. Scale Bar = 5 μ m for whole images, 2 μ m for magnified areas. 2 μ m for magnified areas bar = 5 μ m for whole images, 2 μ m for magnified areas. 2 μ m for magnified areas bar = 5 μ m for whole images, 2 μ m for magnified areas. 2 μ m for magnified areas bar = 5 μ m for whole images, 2 μ m for magnified areas. Data are expressed as means \pm SEM (**p < 0.01, ***p < 0.001 vs t = 0; ##p < 0.01, vs t = 8 h)

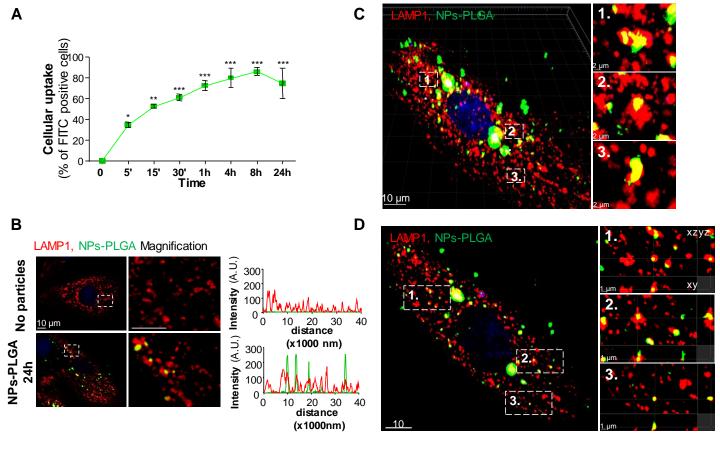


Figure S2: NPs-PLGA address lysosomal compartment in H9C2

(A) Intracellular uptake rate of NPs-PLGA-FITC in H9C2 (n = 3). (B) Left panel: Representative confocal images of H9C2 transfected with LAMP1-RFP (red) and incubated or not with NPs-PLGA-FITC (green) for 24 h. Hoechst 33258 (blue) was used as counterstaining. Scale Bar = 10 μ m. Right panel: Intensity profile graphs of NPs-PLGA-FITC colocalization with LAMP1-RFP. (C) 3-dimensional reconstruction of stack images of H9C2 transfected with LAMP1-RFP (red) and incubated with NPs-PLGA-FITC (green) for 24 h. Hoechst 33258 (blue) was used as counterstaining. Scale Bar = 10 μ m for whole images, 2 μ m for magnified areas. (D) Orthogonal projection of stack images of H9C2 transfected with LAMP1-RFP (red) and incubated with LAMP1-RFP (red) and incubated with LAMP1-RFP (red) and incubated with NPs-PLGA-FITC (green) for 24 h. Hoechst 33258 (blue) was used as counterstaining. Scale Bar = 10 μ m for whole images, 2 μ m for magnified areas. (D) Orthogonal projection of stack images of H9C2 transfected with LAMP1-RFP (red) and incubated with NPs-PLGA-FITC (green) for 24 h. Hoechst 33258 (blue) was used as counterstaining. Scale Bar = 10 μ m for whole images, 2 μ m for magnified areas. (D) Orthogonal projection of stack images of H9C2 transfected with LAMP1-RFP (red) and incubated with NPs-PLGA-FITC (green) for 24 h. Hoechst 33258 (blue) was used as counterstaining. Scale Bar = 10 μ m for whole images, 1 μ m for magnified areas. Data are expressed as means \pm SEM (*p < 0.05**p < 0.01, ***p < 0.001 vs t = 0)

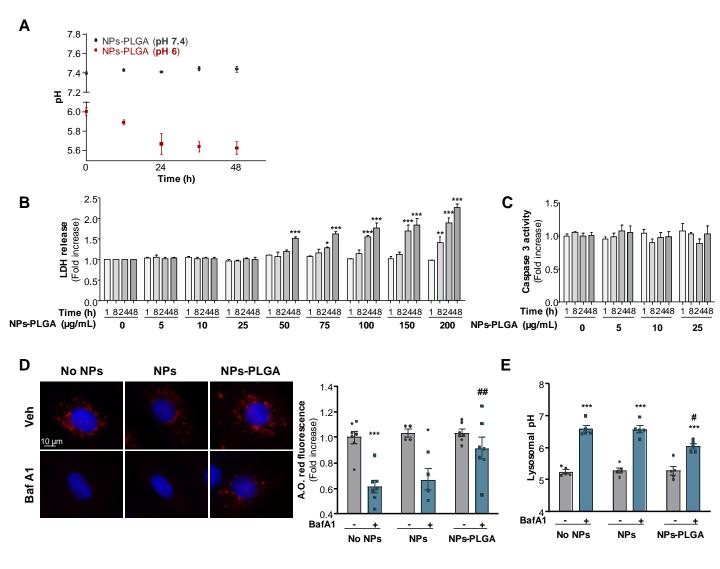


Figure S3: NPs-PLGA prevent BafA1-induced lysosomal dysfunction in H9C2

(A) pH changes of NPs-PLGA in 20 mM pH 6 or 7.4 PBS solution over a 48h period (n = 3) (B) LDH release and (C) Caspase 3 activity in H9C2 incubated with increasing times and concentrations of NPs-PLGA (n = 4). (D-E) H9C2 were pre-incubated O/N with 25 μ g/mL of NPs or NPs-PLGA and treated with Bafilomycin A1 (100 nM, 1 h). (D) Left panel: Representative images of Acridine Orange red fluorescence. Scale Bar = 10 μ m. Right panel: Fluorimeter measurements of Acridine Orange red staining (n = 4-7). (E) Fluorimeter measurements of Lysosensor yellow/blue staining (n = 5). Excitation was measured at 360 nm and the ratio of emission 440/540 nm was calculated. Data are expressed as means \pm SEM (*p < 0.05, **p < 0.01, ***p < 0.001 vs Veh or t = 0; #p < 0.05, ##p < 0.01 vs BafA1)

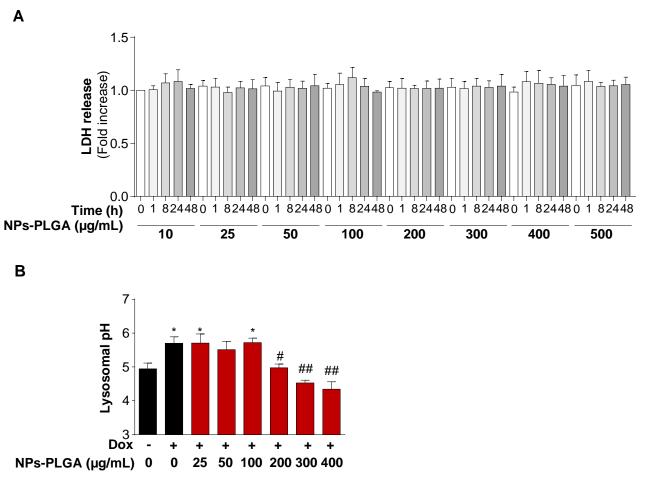


Figure S4: NPs-PLGA are not cytotoxic in NRVMs

(A) LDH release in NRVMs incubated with increasing times and concentrations of NPs-PLGA (n = 4). (B) Fluorimeter measurements of Lysosensor yellow/blue staining in NRVMs incubated O/N with increasing doses of NPs-PLGA in the presence of Dox (1 μ M, 10 h) (n = 8). Excitation was measured at 360 nm and the ratio of emission 440/540 nm was calculated. Data are expressed as means ± SEM (*p < 0.05 vs Veh; #p < 0.05, ##p < 0.01 vs Dox)

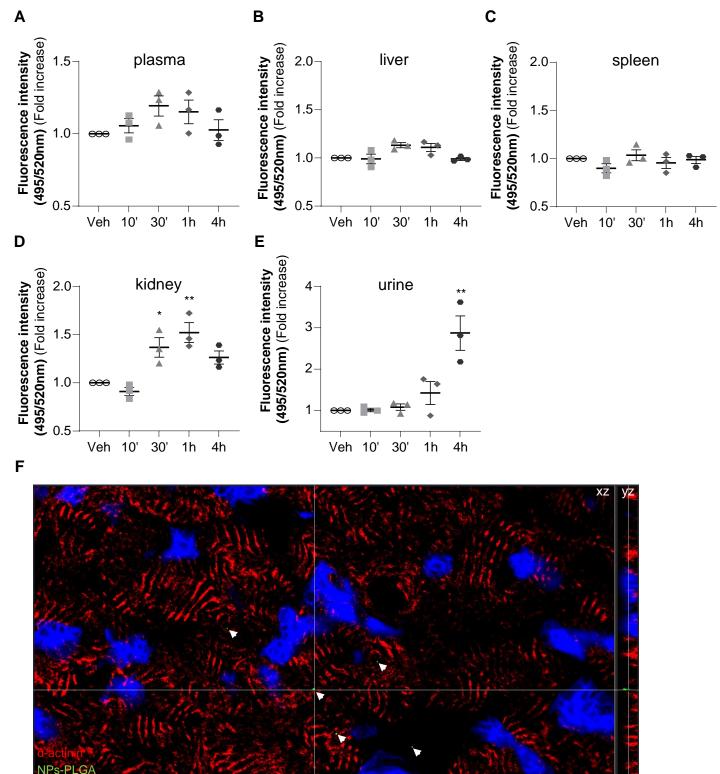


Figure S5: Biodistribution of NPs-PLGA after intratracheal nebulization in mice

<u>8 µm</u>

(A-E) Time-course quantification of green fluorescence in different organs of mice treated with 1 mg/kg of NPs-PLGA-FITC via intratracheal nebulization (n = 3) with (A) plasma, (B) liver, (C) spleen, (D) kidney, and (E) urine. Data are normalized per mg of tissue or per volume. (F) Orthogonal projection of stack images of heart tissues from mice collected 1 h after intratracheal nebulization of Veh or 1 mg/kg of NPs-PLGA-FITC. Green: NPs-PLGA-FITC, red: alpha-actinin, blue: DAPI. Scale bar = 8 μ m. Data are expressed as means \pm SEM (*p < 0.05, **p < 0.01 vs Veh)

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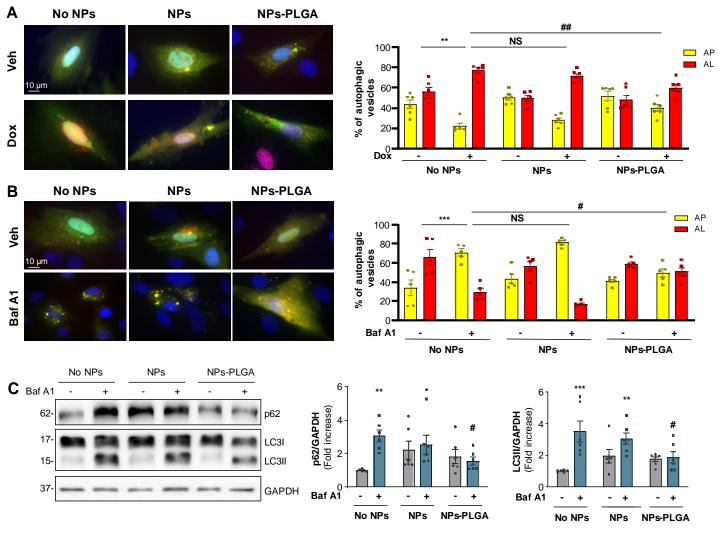


Figure S6: NPs-PLGA prevent Dox and BafA1-induced autophagic flux blockade in H9C2

(A-C) H9C2 were pre-incubated O/N with Veh or 25 μ g/mL of NPs or NPs-PLGA and treated or not with (A) Doxorubicin (1 μ M, 16 h) or (B-C) Bafilomycin A1 (100 nM, 1 h). (A) Left panel: Double immunofluorescence imaging of RFP-GFP-LC3-transfected H9C2. Right panel: Quantification of yellow puncta (AP, autophagosomes) and red puncta (AL, autolysosomes) for each condition (n = 5-6). (B) Left panel: Double immunofluorescence imaging of RFP-GFP-LC3-transfected H9C2. Right panel: Quantification of yellow puncta (AP, autophagosomes) and red puncta (AL, autolysosomes) for each condition (n = 5-6). (B) Left panel: Double immunofluorescence imaging of RFP-GFP-LC3-transfected H9C2. Right panel: Quantification of yellow puncta (AP, autophagosomes) and red puncta (AL, autolysosomes) for each condition (n = 5). (C) Left panel: Representative immunoblots and right panel: quantifications of p62 and LC3-II expression in H9C2 (n = 6). GAPDH was used as a loading control. Data are expressed as means ± SEM (*p < 0.05, **p < 0.01, ***p < 0.001 vs Veh; #p < 0.05, ##p < 0.01 vs Dox or BafA1)

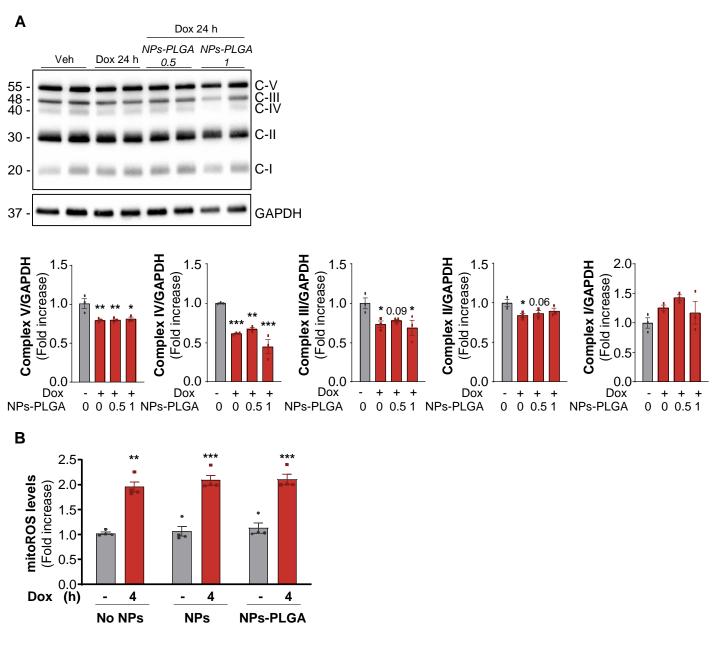


Figure S7: NPs-PLGA do not prevent Dox-induced early mitochondrial dysfunction

(A) Left panel: Representative immunoblots and right panel: quantifications of OXPHOS complexes expression in cardiac homogenates of mice 24 h after Veh or Dox (5 mg/kg) \pm NPs-PLGA (0.5 or 1 mg/kg) administration (n = 3). Dox and NPs-PLGA were administered at the same time. GAPDH was used as a loading control. (B) Quantification of mitoROS assessed by fluorimeter measurements of mitoSOX fluorescence on H9C2 treated with Veh or Dox (1 μ M, 4 h) (n = 4). Data are expressed as means \pm SEM (*p < 0.05, **p < 0.01, ***p < 0.001 vs Veh)

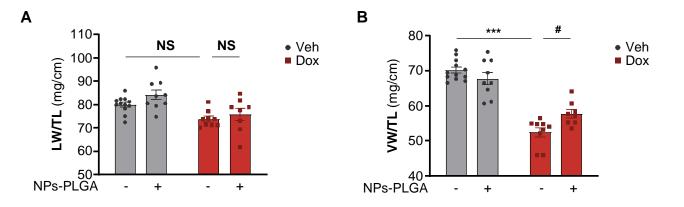


Figure S8: Morphometric parameters of Dox-treated mice after IT nebulization of NPs-PLGA

(A-B) Mice were treated with 4 once-per-week injection of Veh or Dox (5 mg/kg) \pm NPs-PLGA (1 mg/kg). Dox and NPs-PLGA were administered at the same time. Morphometric parameters with (A) Lung weight-to-Tibia length ratio (n = 8-12) and (B) Ventricles weight-to-Tibia length ratio (n = 8-12). Data are expressed as means \pm SEM (***p < 0.001 vs Veh; #p < 0.05 vs Dox)