## Supplementary Information

## Lung-targeted feedback regulation of the mitochondrial ATP synthetic pathway for orthotopic tumors suppression

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Fig. S1 Hydrodynamic diameter of ZIF-90/AIPH/BE.



Fig. S2 Zeta potential diagram of ZIF-90/AIPH/BE.



Fig. S3 UV spectra of ZIF-90, ZIF-90/AIPH, ZIF-90/BE, ZIF-90/AIPH/BE.



Fig. S4 SEM of ZIF-90/AIPH/BE after incubation with 2 mM ATP.



**Fig. S5** Release profiles of RhB when subjected to the mimic extratumoral conditions: (A) the fluorescence spectra; (B) the amount of released RhB.



**Fig. S6** The selectivity of ZIF-90/RhB nanoprobe for ATP. (1: K<sup>+</sup>; 2: Na<sup>+</sup>; 3: Ca<sup>2+</sup>; 4: Mg<sup>2+</sup>; 5: Cu<sup>2+</sup>; 6: Ba<sup>2+</sup>; 7: NH<sub>4</sub><sup>+</sup>; 8: NO<sub>3</sub><sup>-</sup>; 9: PO<sub>3</sub><sup>3-</sup>; 10: SO<sub>4</sub><sup>2-</sup>; 11: SO<sub>3</sub><sup>2-</sup>; 12: CO<sub>3</sub><sup>2-</sup>; 13: HCO<sup>3-</sup>; 14: Br<sup>-</sup>; 15: Cl<sup>-</sup>; 16: Gly; 17: Glu; 18: GMP; 19: AMP; 20: ATP).



**Fig. S7** Fluorescence ratio of DCF through a reaction between DCFH-DA and alkyl radicals from AIPH at different temperature.



Fig. S8 Fluorescence intensity of ZIF-90/RhB in A549 cells upon treatment with DCC and Ca<sup>2+</sup>.



Fig. S9 Standard curve of NADH concentrations detected by Amplite Colorimetric Total NAD and NADH Assay Kit.



Fig. S10 Standard curve of ATP contents in A549 cells detected by ATP assay Kit.



**Fig. S11** ATP content in per cell after A549 cells were treated with different formulations for 24 h. The values of the control group were all set as 100%.



Fig. S12 SEM of mitochondrion-orientated liposome.



Fig. S13 Co-localization experiments involving mitochondrion-orientated liposome@RhB, MitoTracker in A549 cells.



Fig. S14 AM/PI staining confocal microscopy images of A549 or LO2 cells with different treatments.



**Fig. S15** *In vivo and ex vivo* fluorescence imaging after ZIF-90/Cy5.5 NPs was induced in normal Balb/c mice by intravenous injection for 2 h.



**Fig. S16** The top 18 most abundant proteins in the protein corona of different nanoparticles determined. Values were calculated from the molar masses of each protein identified by LC–MS.