Electronic Supplementary Information (ESI)

Assessment of the zero-valent iron-based nanotherapeutics for ferroptosis induction and resensitization strategy in cancer cells

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Fig. S1 Characterization of the ultrastructure of different ZVI-based nanoparticles by TEM. TEM images show that all the synthesized NPs used in the experiment had a diameter in the range of 50 to 100 nm. Strong aggregation by the bare ZVI NPs was noted. Gold and CMC coating significantly decrease particle aggregation.



Fig. S2 Characterization of fresh prepared and oxidized ZVI@CMC NPs by XRD and DLS analysis. (A) The freshly prepared ZVI@CMC NPs presented major XRD peak at $2\theta = 44.67$ corresponding to that of the standard Fe⁰ supporting ZVI to be the main component of the NPs. However, the shift of the XRD peak to $2\theta = 35.63$ (iron oxide signal) was discovered in the oxidized NPs. (B) Hydrodynamic size and aggregation were increase in oxidized ZVI NPs.



Fig. S3 Characterization of ZVI@CMC NPs in different storage condition for a period of time by XRD and DLS analysis. (A) These NPs were discovered to be quite stable when preserved in Argon or air up to 28 days even at room temperature as evidenced by well-preserved Fe⁰ signal. (B) There are no significant differences in hydrodynamic size change of ZVI@CMC NPs under Argon or air environment.



Fig. S4 Caspase-dependent apoptosis was not detected in OEC-M1 cells after ZVI NPs treatment. Western blotting revealed that the cleaved caspase 3 and caspase 9 were not detected 24 and 48 h after 5 μ g/mL ZVI NP treatment. Cisplatin (CDDP, 20 μ M) treatment served as a positive control.



Fig. S5 Assessment of NADPH levels paired ZVI-sensitive/resistant cells. The ZVI-refractory cell line OEC-M1 R3 had much higher endogenous NADPH levels than the paired sensitive line OEC-M1.

Table S1. The IC₅₀ of seven OSCC cell lines to four ZVI-based NPs.

		IC ₅₀ (μg/mL)			
		Bare ZVI NPs	ZVI@Au NPs	CMC stabilized ZVI@Au NPs	ZVI@CMC NPs
Sensitive cells Refractory cells	OC3	0.60 <u>+</u> 0.02	3.48 <u>+</u> 0.03	0.58 <u>+</u> 0.01	0.63 <u>+</u> 0.01
	OEC-M1	1.09 <u>+</u> 0.15	1.48 <u>+</u> 0.06	4.58 <u>+</u> 0.38	4.88 <u>+</u> 0.80
	SCC9	0.80 <u>+</u> 0.12	7.05 <u>+</u> 2.73	0.66 <u>+</u> 0.01	0.76 <u>+</u> 0.02
	HSC-3	> 50	> 50	> 50	> 50
	SAS	> 50	> 50	> 50	> 50
	KOSC-3	> 50	> 50	> 50	> 50
	OC2	> 50	> 50	> 50	> 50

	Primer sequence			
Gene	Forward	Reverse		
ACSL4	GCTATCTCCTCAGACACACCGA	AGGTGCTCCAACTCTGCCAGTA		
ZEB1	GCCAATAAGCAAACGATTCTG	TTTGGCTGGATCACTTTCAAG		
GSR	AACATCCCAACTGTGGTCTTCAGC	TTGGTAACTGCGTGATACATCGGG		
AKR1C1	AGTAAAGCTTTAGAGGCCAC	CACCCATGGTTCTTCTCGG		
AKR1C3	GGGATCAACGAGAGACAAACG	AAAGGACTGGGTCCTCCAAGA		
AKR1B1	TGAGTGCCACCCATATCTCA	TGTCACAGACTTGGGGATCA		
AKR1B10	GTGGGGGAAGCCATCCAAGA	CAGCTTCAGGTCCTTGAGGG		
KYNU	TTGCCTGCTGGTGTTCCTAC	TCATGAATGAAGGCACCAGC		
NNMT	GAATCAGGCTTCACCTCCAA	TCACACCGTCTAGGCAGAAT		
GPx-4	GCCTTCCCGTGTAACCAGT	GCGAACTCTTTGATCTCTTCGT		

Table S2. Sequences of the primers used for qPCR analysis