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ARTICLE TYPE

Supporting information

Ferritin nanocages loaded with gold ions induce oxidative stress and apoptosis in MCF-7 human breast cancer cells

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5 Figure S1. Photo of the solutions of Au₂phen -encapsulated AFt, Auoxo4-encapsulated AFt and of Auoxo3-encapsulated AFt.

Figure S2. Gold binding site region in Au_2phen -encapsulated AFt. Bijvoet difference Fourier map 10 calculated with anomalous data and reported at 4 σ .

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Figure S3. Gold binding site region in **Auoxo4**-encapsulated AFt. 2Fo-Fc electron density maps are reported at 0.8 σ (cyan) and 4 σ (blue). Bijvoet difference Fourier map calculated with anomalous data and reported at 3 σ is in orange in panel B.

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 Table S1. Data collection and refinement statistics

	Au2phen-encapsulated	Auoxo4-encapsulated
	AFt	AFt
Data Collection statistics		
X-ray source	Rotating anode	Rotating anode
Wavelength	λ=1.5418Å	λ=1.5418Å
Space group	F432	F432
Unit-cell parameters a=b=c (Å), α,β,γ (°)	180.97 90	180.88 90
Ft Monomers per a. u.	1	1
Resolution (Å)	104.5-1.82	104.4-2.60
	(1.85-1.82)	(2.64-2.60)
Observed reflections	287733	32931
Unique reflections	23324	8209
Completeness (%)	99.9 (100)	99.0 (99.7)
Rmerge	0.121 (0.680)	0.187 (0.414)
Rpim	0.126 (0.269)	0.145 (0.401)
Ι/σ(Ι)	11.9 (3.2)	5.4 (2.9)
Multiplicity	12.3 (7.2)	4.0 (4.1)
CC1/2	0.857	
Refinement		
Resolution (Å)	104.5-1.82	104.4-2.60
n. of reflections in working set	22173	7881
n. of reflections in test set	1138	391
R factor/Rfree (%)	14.3/17.2	18.4/23.6
n of non-H atoms in the refinement	1822	1605
Occupancy of Au ions	0.50, 0.30	0.50, 0.35
B-factor of Au ions (Å ²)	16.9, 48.1	41.5, 47.1
Overall B-factor	18.6	23.6
Deviations from ideality values R.m.s.d. bonds (Å)	0.022	0.013

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R.m.s.d. angles (Å) 1.86

1.47

Table S2. Comparison of anomalous peak intensity of the most representative Cd ions, i.e. those conserved in the structures of AFts, in **Au2phen-**, **Auoxo4- Auoxo3-**encapsulated AFts and in the structure of CDDP- and CBDCA-encapsulated AFts, used as control. Only data collected using CuK α are compared.

	Auoxo4- encapsulated AFt	Au2phen- encapsulated AFt	Auoxo3- encapsulated AFt	CDDP- encapsulated AFt	CBDCA-encapsulated AFt
Source	CuKa	CuKa	CuKa	CuKa	CuKa
Resolution	λ=1.5418 A 2.6 Å	λ=1.5418 A 1.82 Å	λ=1.5418 A 1.96 Å	λ=1.5418 A 2.06 Å	∧=1.5418 A 1.96 Å
PDB code			5IX6		5MIK
Atom in 5ERK					e/Å ³
CD 1. Close to Glu11	<0.10	0.23	0.22 (CD8)	0.24	0.22
CD 2. Close to Glu53/Glu56	<0.10	0.16	0.21(CD9)	0.16	0.22
CD 3. Close to Glu60	<0.10	0.28	0.20 (CD15)	0.31	0.25
CD 4. At the binary axis. Close to Asp80	<0.10	1.80	1.19 (CD10)	0.92	1.08
CD 5. Close to Glu88	Absent	0.10	<0.10	<0.10	0.12
CD 6. Close to Asp127		0.10	<0.10	0.35	<0.10
CD 7. Close to His132	0.13	0.23	0.25(CD11)	0.28	< 0.10 (0.17 using λ=0.97Å) Interpreted as Pt
CD 8. At the ternary axis. Close to Glu130	0.30	0.51	0.58 (CD12)	0.41	0.39
CD 9. Close to His114 and at 3.7 from Cys126	<0.10	<0.10	0.19	0.25	0.08/0.09
Atoms in 5IX6					
Au1 close to SG Cys126 and His114	Absent	Absent	0.76 Interpreted as Au	<0.10	
Au2 close to Cys48 (and Close to Arg52)	E.d. absent <0.1	E.d. absent <0.1	0.37 Interpreted as Au	<0.10	0.16 Interpreted as Cd
Au3 close to His49	<0.10	< 0.10 Interpreted as water	0.52 Interpreted as Au	<0.10	0.04/0.13 (0.17 using λ=0.97Å) Interpreted as Pt
Au4 close to SG Cys126	0.22 Interpreted as Au	1.06 Interpreted as Au	1.00 Interpreted as Au	<0.10	-
Au5 close to His147	-	< 0.1 Interpreted as water	Interpreted as Au 3.14	<0.10	-
CD 11. Close to Cys48	<0.10	0.16 Interpreted as Cd	0.50 Interpreted as Au6 close to Cys48 (and close to His49	0.25 Interpreted as Cd	0.16 Interpreted as Cd

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Au7 close to His132	E.d. absent <0.10	E.d absent <0.1	Interpreted as Au 0.16	0.21 Interpreted as Pt	
Au close to Cys126	Interpreted as Au 0.13	Interpreted as Au 0.23	Interpreted as water. Close to Au1	<0.10	
CD16	<0.10	Interpreted as water <0.10	0.21	Absent Close to Asp53	

Table S3.Uninterpreted peaks of electron density in the Fo-Fc map:

Auoxo4-encapsulated AFt

	e/Å ³	Reason
Peak 1	0.81	Too close to SG atom of Cys126, CL and Au atoms
Peak 2	0.80	In the middle oft he nanocage. Not bound tot he protein
Peak 3	0.68	Too close to side chain of His149
Peak 4	0.60	Too close to CD20 and wat 266
Peak 5	0.59	Too close to side chain of Arg153
Peak 6	0.56	Too close to side chain of His149
Peak 7	0.51	Too close to CD1

Au_2phen -encapsulated AFt

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e/Å ³	Reason

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Peak 1	0.81	Too close to gold atom and SG atom of Cys126
Peak 2	0.80	Too close to a water molecule
Peak 3	0.68	Too close to side chain of Gln79
Peak 4	0.60	Too close to side chain of Glu163
Peak 5	0.59	
Peak 6	0.56	
Peak 7	0.51	