

## Electronic supplementary information

of

### A translational approach to assess the metabolomic impact of stabilized gold nanoparticles by NMR spectroscopy

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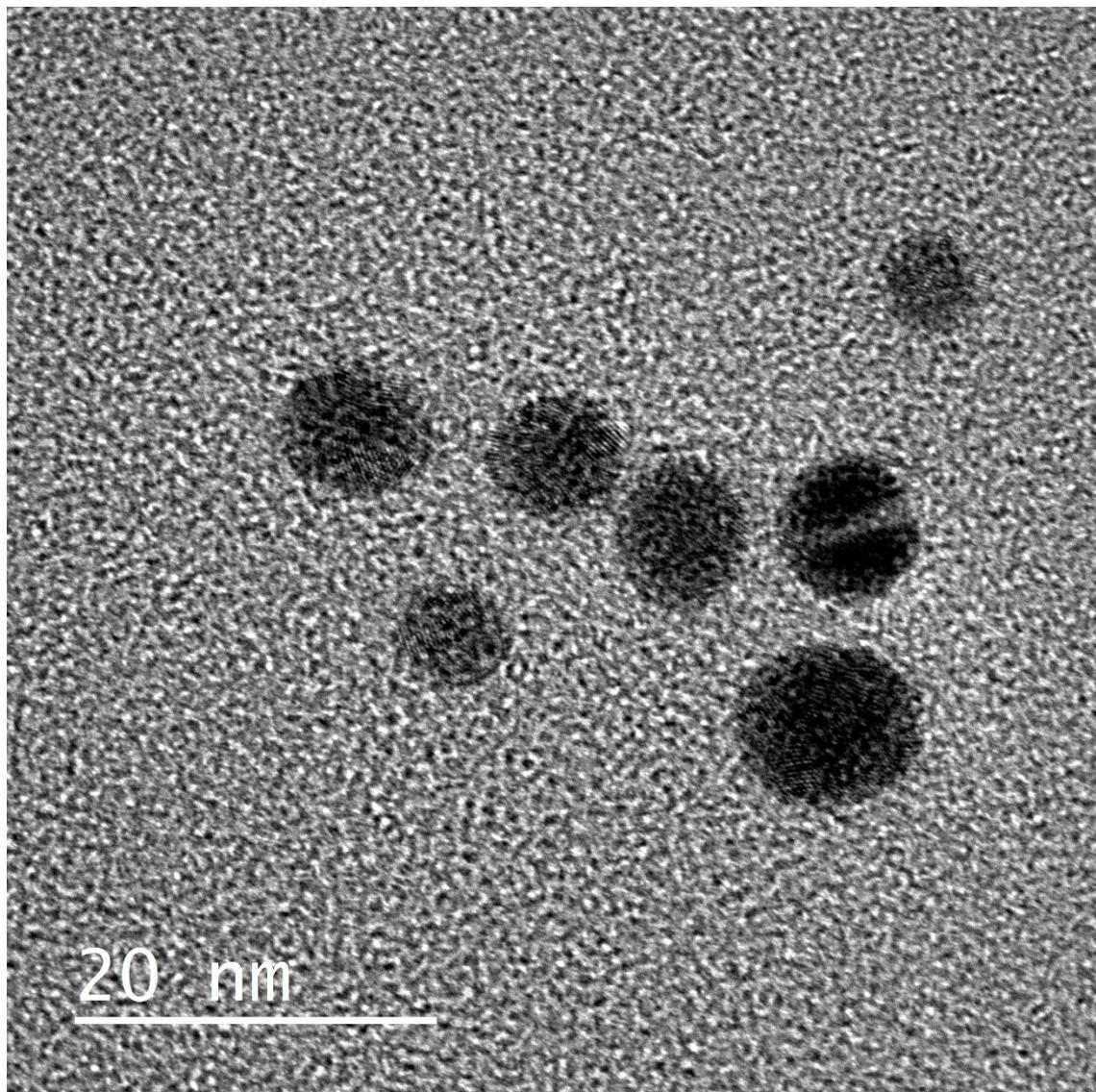
d Instituto de Investigación Sanitaria La Fe, Hospital Universitario i Politécnico La Fe, Valencia, Spain.

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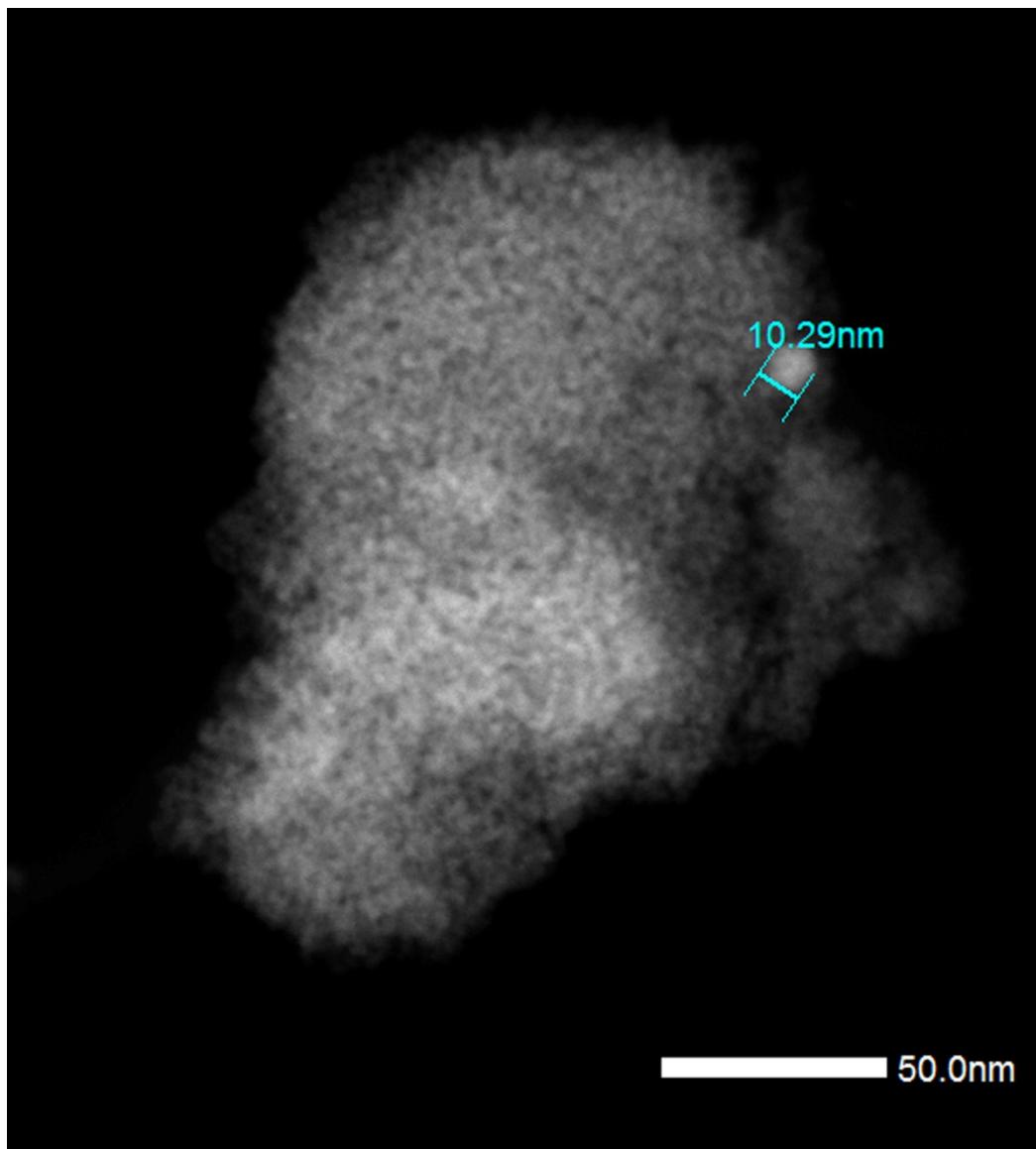
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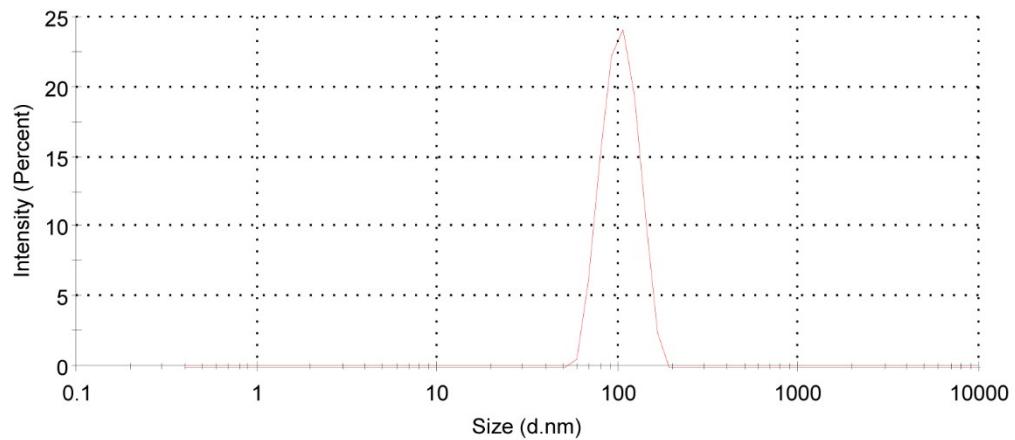
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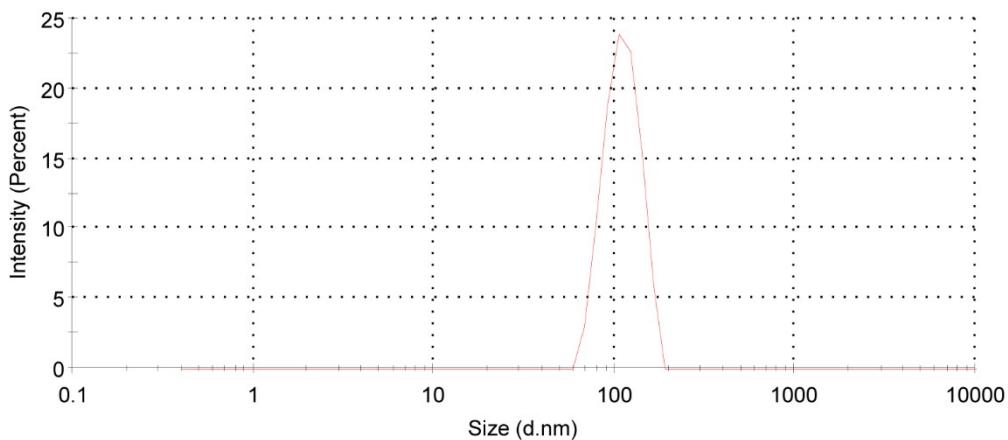
**Supplementary Figure 1.** TEM recorded for AuChi. The average size of Au NPs coated with chitosan was determined with 100 measures of NPs.



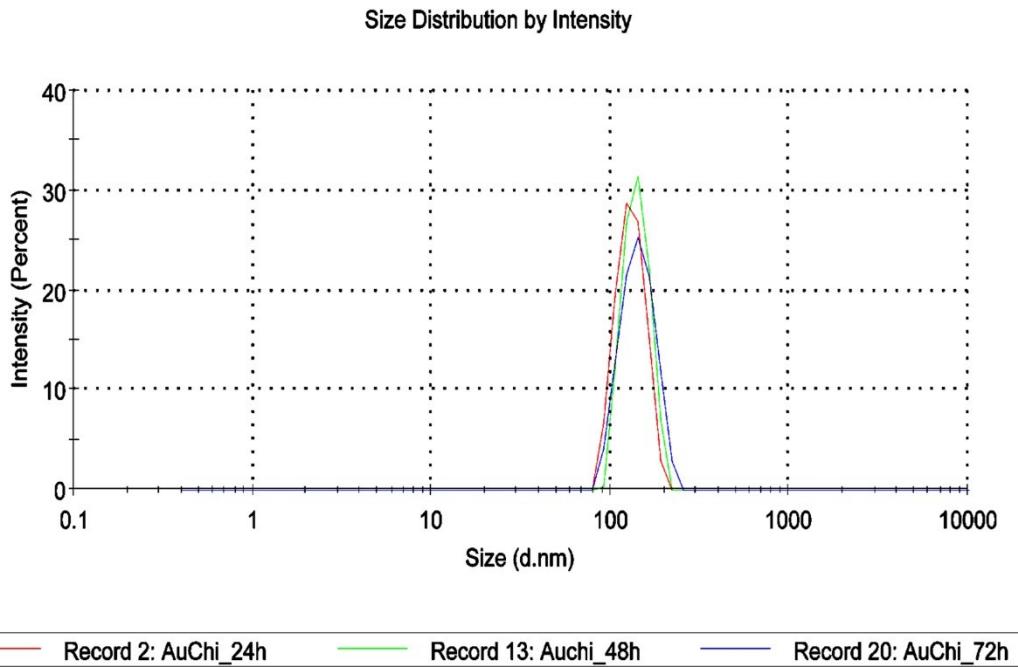
**Supplementary Figure 2.** TEM recorded for  $\text{AuCeO}_2$ . The average size of Au NPs coated with chitosan was determined with 100 measures of NPs.



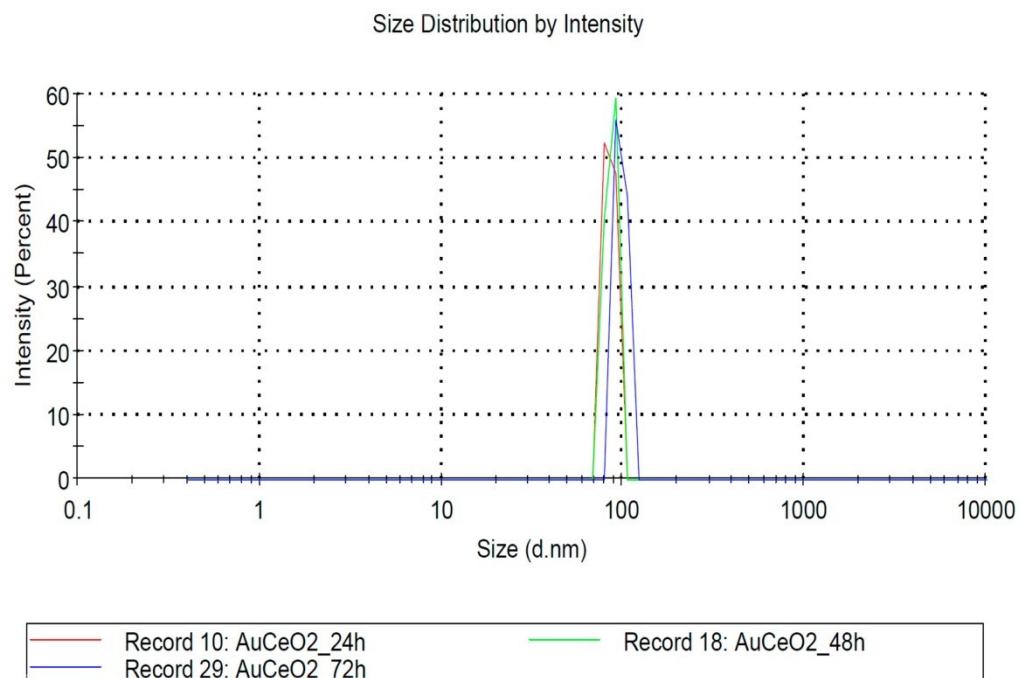
**Supplementary Figure 3.** DLS recorded for AuChi. The analysis performed to determine the hydrodynamic size was done using a 20  $\mu\text{g}/\text{mL}$  solution of AuChi in PBS.



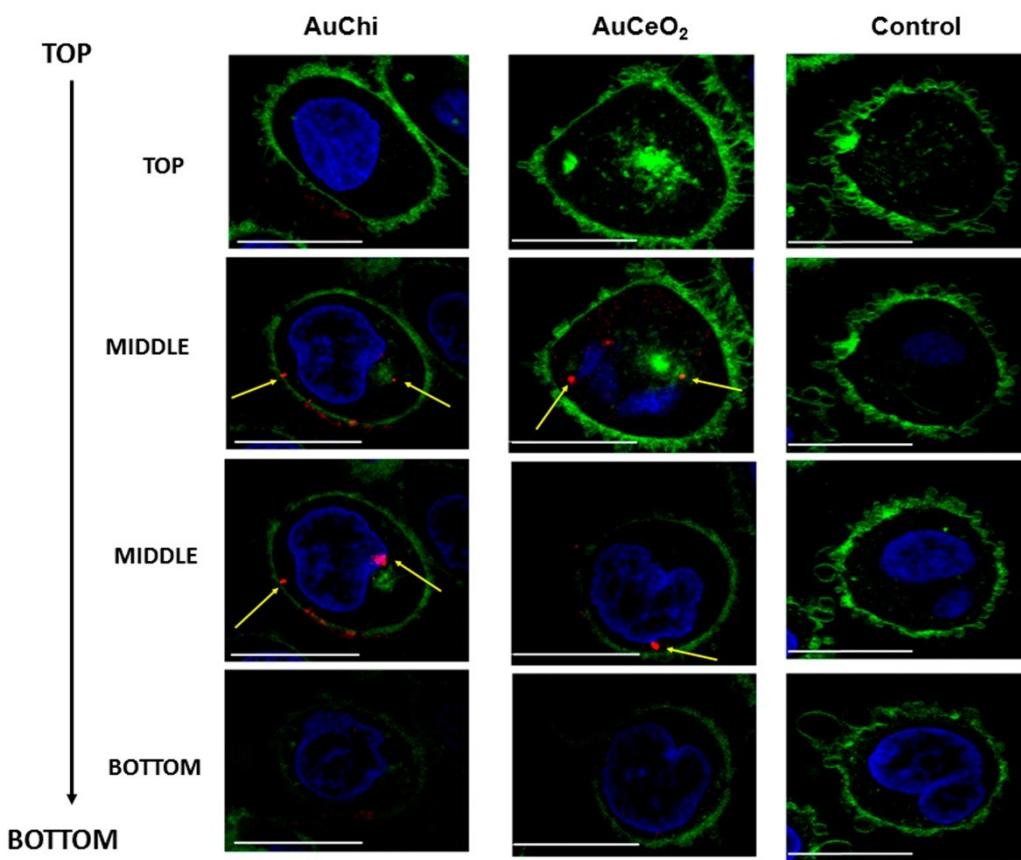
**Supplementary Figure 4.** DLS recorded for AuCeO<sub>2</sub>. The analysis performed to determine the hydrodynamic size was done using a 20  $\mu\text{g}/\text{mL}$  solution of AuCeO<sub>2</sub> in PBS.



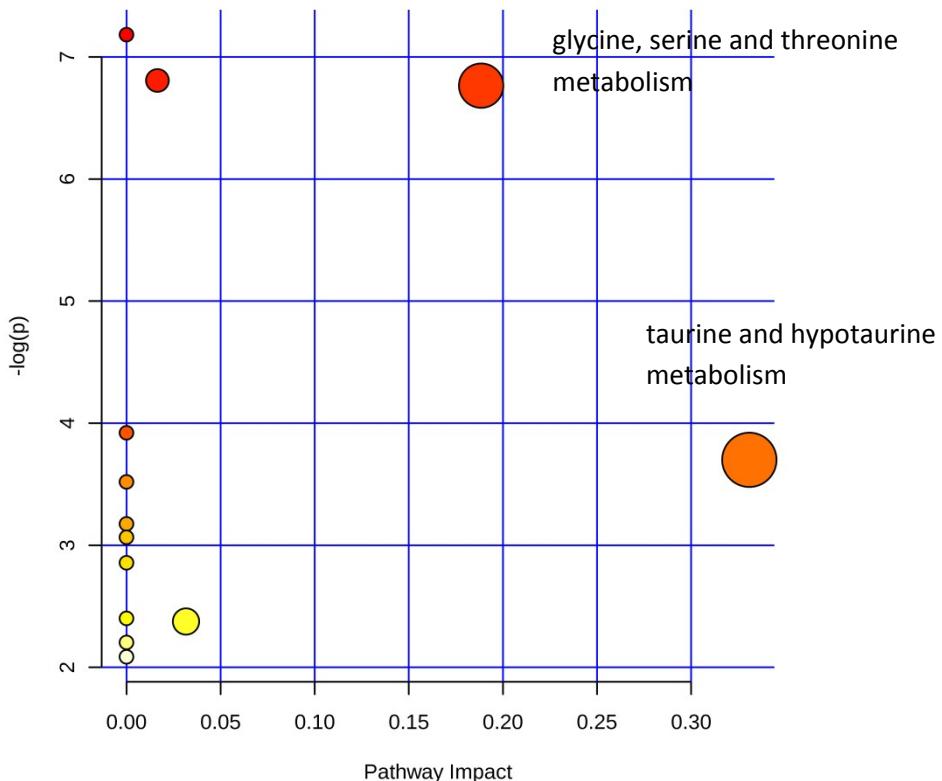
**Supplementary Figure 5.** Dynamic light scattering spectra to determine the hydrodynamic size and the stability of AuChi nanoparticles at 24h, 48h and 72 hours diluted in PBS, final concentration 20 µg/ml.



**Supplementary Figure 6.** Dynamic light scattering spectra to determine the hydrodynamic size and the stability of AuCeO<sub>2</sub> nanoparticles at 24h, 48h and 72 hours diluted in PBS, final concentration 20 µg/ml. A) AuChi, B) AuCeO<sub>2</sub>.

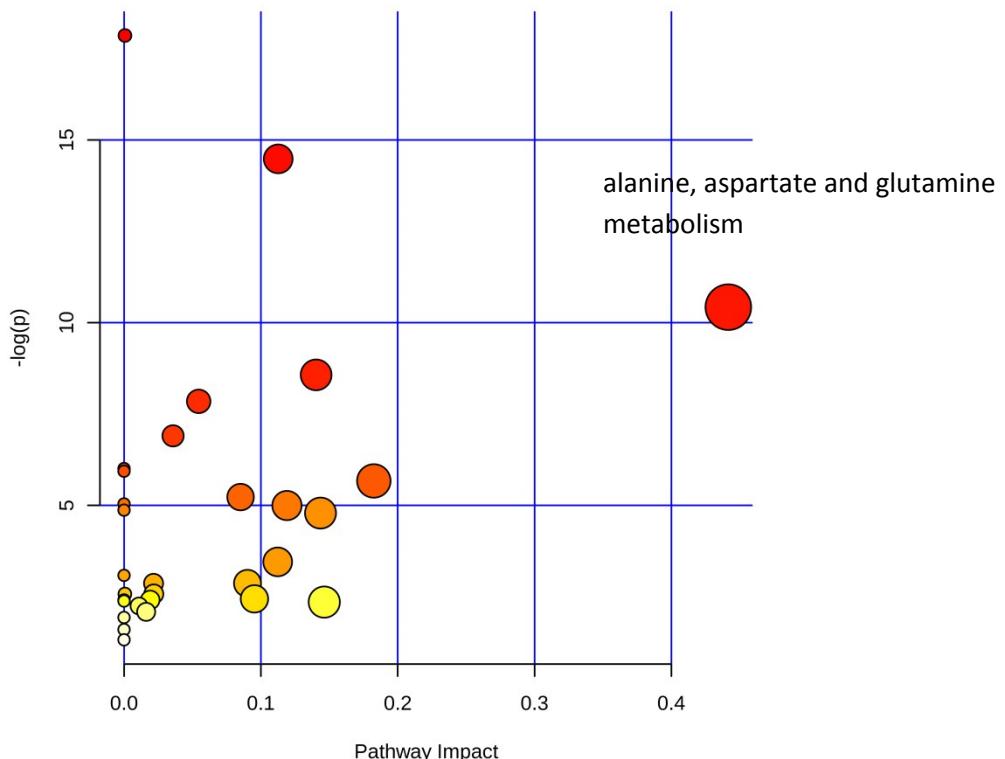


**Supplementary Figure 7.** Z-axis scanning images of AuChi, AuCeO<sub>2</sub> and without NPs (control) in HeLa cells by laser confocal microscope obtained merging the blue (Hoechst) to label the nucleous, and green (CellMask) to label the cellular membrane. Fluorescence were obtained exciting at 405 nm (exciting Hoechst, blue), and 488 nm (exciting CellMask, green) and the emission was collected from 415 to 550 nm in separate channels. The red spots indicating the location of NPs after irradiating at 633 nm (yellow arrows). Scale bars are 20  $\mu$ m in all cases.



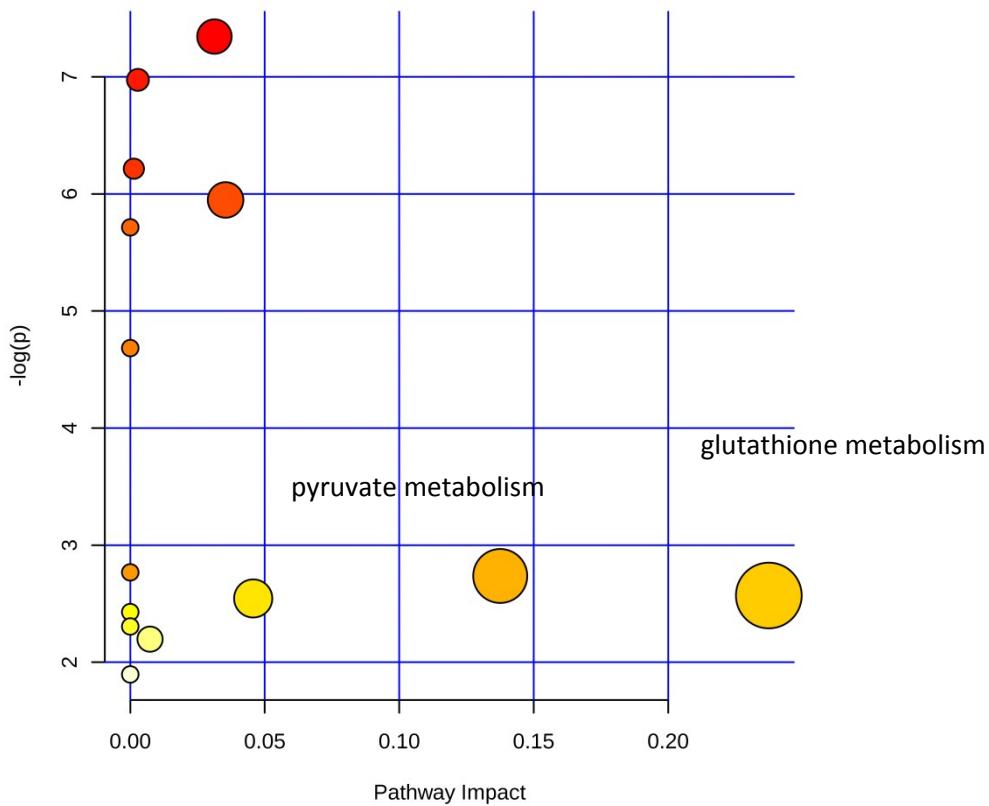
Pathway	Expected	Total	Hits	Raw p	-log(p)	Holm adjust	FDR	Impact
Nitrogen metabolism	39	0,048608	2	0,00075984	7,1824	0,060787	0,030767	0
Primary bile acid biosynthesis	47	0,058579	2	0,001106	6,807	0,087374	0,030767	0,01644
Glycine, serine and threonine metabolism	48	0,059826	2	0,0011538	6,7647	0,089993	0,030767	0,18845
Cyanoamino acid metabolism	16	0,019942	1	0,019818	3,9212	1	0,39504	0
Taurine and hypotaurine metabolism	20	0,024927	1	0,024731	3,6997	1	0,39504	0,33094
Thiamine metabolism	24	0,029913	1	0,029628	3,519	1	0,39504	0
Methane metabolism	34	0,042376	1	0,041798	3,1749	1	0,46637	0
Glutathione metabolism	38	0,047362	1	0,046637	3,0654	1	0,46637	0
Lysine degradation	47	0,058579	1	0,057466	2,8566	1	0,51081	0
Aminoacyl-tRNA biosynthesis	75	0,093477	1	0,090631	2,401	1	0,67615	0
Arginine and proline metabolism	77	0,09597	1	0,09297	2,3755	1	0,67615	0,03163
Purine metabolism	92	0,11467	1	0,11038	2,2038	1	0,73589	0
Porphyrin and chlorophyll metabolism	104	0,12962	1	0,12415	2,0863	1	0,76401	0

**Supplementary Figure 8.** Metabolic routes altered by AuChi and AuCeO<sub>2</sub> particles, at similar time points.



Pathway	Expected	Total	Hits	Raw p	$-\log(p)$	Holm adjust	FDR	Impact
Nitrogen metabolism	39	0,11342	5	1,76E-08	17,855	1,41E-06	1,41E-06	0,00067
Aminoacyl-tRNA biosynthesis	75	0,21811	5	5,15E-07	14,48	4,07E-05	2,06E-05	0,11268
Alanine, aspartate and glutamate metabolism	24	0,069796	3	2,97E-05	10,424	0,0023186	0,00079267	0,4416
Histidine metabolism	44	0,12796	3	0,00018967	8,5702	0,014605	0,0037935	0,14039
Cysteine and methionine metabolism	56	0,16286	3	0,00039102	7,8468	0,029717	0,0062563	0,05455
Arginine and proline metabolism	77	0,22393	3	0,0010047	6,903	0,075355	0,013396	0,03582
Pantothenate and CoA biosynthesis	27	0,078521	2	0,0024587	6,0081	0,18194	0,026441	0
beta-Alanine metabolism	28	0,081429	2	0,0026441	5,9354	0,19302	0,026441	0
Pyruvate metabolism	32	0,093062	2	0,0034503	5,6693	0,24842	0,030669	0,18254
Butanoate metabolism	40	0,11633	2	0,0053657	5,2277	0,38096	0,042925	0,08516
Nicotinate and nicotinamide metabolism	44	0,12796	2	0,0064714	5,0404	0,453	0,045086	0
Phenylalanine metabolism	45	0,13087	2	0,006763	4,9963	0,46664	0,045086	0,11906
Glycine, serine and threonine metabolism	48	0,13959	2	0,0076735	4,87	0,5218	0,047222	0
Glyoxylate and dicarboxylate metabolism	50	0,14541	2	0,0083101	4,7903	0,55678	0,047486	0,14359
D-Glutamine and D-glutamate metabolism	11	0,03199	1	0,031594	3,4548	1	0,1685	0,1123
Cyanoamino acid metabolism	16	0,046531	1	0,045669	3,0863	1	0,22835	0
Taurine and hypotaurine metabolism	20	0,058164	1	0,056803	2,8682	1	0,25246	0,02158
Citrate cycle (TCA cycle)	20	0,058164	1	0,056803	2,8682	1	0,25246	0,09024
Phenylalanine, tyrosine and tryptophan biosynthesis	27	0,078521	1	0,076019	2,5768	1	0,292	0,00062
Valine, leucine and isoleucine biosynthesis	27	0,078521	1	0,076019	2,5768	1	0,292	0,02173
Glycolysis or Gluconeogenesis	31	0,090154	1	0,086848	2,4436	1	0,292	0,0953
Pentose phosphate pathway	32	0,093062	1	0,089539	2,4131	1	0,292	0
Lysine biosynthesis	32	0,093062	1	0,089539	2,4131	1	0,292	0
Vitamin B6 metabolism	32	0,093062	1	0,089539	2,4131	1	0,292	0,01914
Terpenoid backbone biosynthesis	33	0,09597	1	0,092222	2,3836	1	0,292	0
Methane metabolism	34	0,098878	1	0,094899	2,3549	1	0,292	0,14633
Glutathione metabolism	38	0,11051	1	0,10554	2,2487	1	0,31271	0,01095
Ascorbate and aldarate metabolism	45	0,13087	1	0,1239	2,0883	1	0,354	0,01617
Pentose and glucuronate interconversions	53	0,15413	1	0,14449	1,9346	1	0,39858	0
Tyrosine metabolism	76	0,22102	1	0,20138	1,6026	1	0,53702	0
Porphyrin and chlorophyll metabolism	104	0,30245	1	0,26624	1,3234	1	0,68707	0

**Supplementary Figure 9.** Metabolic routes altered by AuChi and AuCeO<sub>2</sub> particles, at different time points (48 h for AuChi and 72 h for AuCeO<sub>2</sub>).



Pathway	Expected	Total	Hits	Raw p	-log(p)	Holm adjust	FDR	Impact
Citrate cycle (TCA cycle)	20	0,041545	2	0,00064639	7,3441	0,051711	0,037434	0,0313
Alanine, aspartate and glutamate metabolism	24	0,049855	2	0,00093584	6,9741	0,073931	0,037434	0,00285
Propanoate metabolism	35	0,072705	2	0,001999	6,2151	0,15592	0,052192	0,00134
Butanoate metabolism	40	0,083091	2	0,0026096	5,9486	0,20094	0,052192	0,03545
Phenylalanine metabolism	45	0,093477	2	0,0032983	5,7144	0,25067	0,052773	0
Tyrosine metabolism	76	0,15787	2	0,0092504	4,6831	0,69378	0,12334	0
Glycolysis or Gluconeogenesis	31	0,064396	1	0,062809	2,7677	1	0,62795	0
Pyruvate metabolism	32	0,066473	1	0,064781	2,7367	1	0,62795	0,13756
Glutathione metabolism	38	0,078936	1	0,076545	2,5699	1	0,62795	0,23743
Glycerophospholipid metabolism	39	0,081014	1	0,078494	2,5447	1	0,62795	0,04571
Nicotinate and nicotinamide metabolism	44	0,0914	1	0,08819	2,4283	1	0,64138	0
Glyoxylate and dicarboxylate metabolism	50	0,10386	1	0,099717	2,3054	1	0,66478	0
Cysteine and methionine metabolism	56	0,11633	1	0,11113	2,1971	1	0,68386	0,00735
Arginine and proline metabolism	77	0,15995	1	0,15016	1,8961	1	0,85803	0

**Supplementary Figure 10.** Metabolic routes altered in an opposite way by AuChi and AuCeO2 particles.

**Supplementary Table 1.** Normalized concentration values of selected metabolites after 24 hours of treatment. P-values were obtained by the Student's t-test.

metabolite	Time (h)		24		24		24		24		24	
	Nanomaterial		Control	Control	AuChi	AuChi	AuChi	AuCeO2	AuCeO2	AuCeO2	P	
	region (ppm)	mean	SEM	mean	SEM	P	mean	SEM	P			
3-hydroxyisovalerate	1,247	1,2719	43,792	5,738	40,774	5,785	0,716	59,060	8,042	0,142		
acetate	1,91	1,93	8,473	0,692	8,267	0,665	0,833	11,912	0,862	0,007		
ADP/ATP	8,53	8,545	1,882	0,162	1,622	0,125	0,223	1,934	0,210	0,845		
alanine	1,47	1,51	14,499	0,865	13,518	0,714	0,395	15,777	1,319	0,430		
aspartate	2,79	2,84	3,717	0,346	3,554	0,326	0,737	4,018	0,368	0,559		
ATP	4,39	4,43	5,617	0,491	5,789	0,539	0,817	0,838	4,158	0,270		
creatine/phocrea	3,0347	3,053	6,376	0,909	6,442	0,482	0,949	3,907	0,256	0,019		
formate	8,45	8,47	2,397	0,223	2,222	0,225	0,589	3,367	0,361	0,037		
fumarate	6,51	6,53	0,760	0,152	0,645	0,126	0,570	1,023	0,241	0,369		
glutamate	2,33	2,38	16,956	1,088	18,011	1,112	0,508	15,833	1,038	0,466		
glutamine	2,44	2,48	6,942	0,750	7,272	1,046	0,801	6,907	0,431	0,968		
glycerol	3,566	3,5711	0,798	0,068	0,846	0,107	0,709	1,571	0,533	0,169		
glycerol derivative	3,46	3,49	5,440	0,763	5,189	1,043	0,848	6,074	0,986	0,618		
glycine	3,5508	3,566	5,511	0,341	5,667	0,308	0,738	5,642	0,522	0,836		
GSSG	3,295	3,3084	1,140	0,080	1,313	0,171	0,374	1,432	0,205	0,203		
histidine	7,08	7,09	0,480	0,058	0,429	0,054	0,533	0,677	0,137	0,205		
isoleucine	1,01	1,03	2,953	0,268	3,417	0,400	0,349	4,292	0,435	0,018		
lactate	1,31	1,37	79,900	8,651	77,429	9,282	0,848	109,693	14,925	0,103		
methionine	2,017	2,027	2,768	0,126	2,554	0,107	0,216	2,684	0,233	0,757		
NAD	8,82	8,86	2,606	0,316	2,357	0,243	0,541	3,335	0,559	0,273		
phenylalanine	7,3	7,45	11,584	1,274	10,969	1,046	0,714	15,630	1,619	0,067		
phosphocholine	4,15	4,19	20,037	2,848	24,396	3,138	0,319	11,863	1,520	0,022		
pyruvate	2,38	2,4	5,125	0,375	5,280	0,443	0,793	6,798	0,497	0,016		
succinate	2,4	2,414	4,272	0,291	4,286	0,501	0,981	4,828	0,337	0,230		
taurine	3,407	3,445	6,485	0,449	6,632	0,890	0,885	6,071	0,839	0,669		
UDP der	4,36	4,39	6,253	0,577	6,370	0,336	0,863	3,635	2,161	0,259		
UDP-NAG	5,5	5,54	1,466	0,180	0,982	0,236	0,122	1,977	0,406	0,267		
valine	1,04	1,06	4,282	0,394	4,889	0,521	0,367	6,024	0,589	0,026		

**Supplementary Table 2.** Normalized concentration values of selected metabolites after 48 hours of treatment. P-values were obtained by the Student's t-test.

metabolite	Time (h)		48		48		48		48		48	
	Nanomaterial		Control	Control	AuChi	AuChi	AuChi	AuCeO2	AuCeO2	AuCeO2	P	
	region (ppm)	mean	SEM	mean	SEM	P	mean	SEM	P			
3-hydroxyisovalerate	1,247	1,2719	20,496	2,299	37,770	5,155	0,007	24,948	2,426	0,201		
acetate	1,91	1,93	5,991	0,349	8,780	0,913	0,012	6,632	0,472	0,292		
ADP/ATP	8,53	8,545	2,435	0,305	1,826	0,311	0,181	2,361	0,218	0,846		
alanine	1,47	1,51	15,965	0,515	13,806	0,542	0,011	13,899	0,530	0,013		
aspartate	2,79	2,84	2,379	0,228	3,117	0,362	0,104	2,464	0,222	0,793		
ATP	4,39	4,43	8,125	0,736	5,957	1,094	0,120	7,601	0,664	0,604		
creatine/phocrea	3,0347	3,053	12,934	1,452	6,658	1,889	0,018	8,582	0,717	0,016		
formate	8,45	8,47	1,653	0,166	2,764	0,396	0,020	1,944	0,201	0,280		
fumarate	6,51	6,53	0,573	0,100	0,685	0,115	0,475	0,550	0,102	0,872		
glutamate	2,33	2,38	16,111	0,890	15,664	0,973	0,739	12,545	0,337	0,002		
glutamine	2,44	2,48	3,275	0,250	4,445	0,427	0,031	3,465	0,319	0,644		
glycerol	3,566	3,5711	1,315	0,096	1,482	0,273	0,571	1,718	0,232	0,128		
glycerol derivative	3,46	3,49	4,593	0,641	4,820	0,843	0,833	6,885	1,357	0,146		
glycine	3,5508	3,566	8,146	0,395	6,625	0,279	0,006	7,333	0,303	0,122		
GSSG	3,295	3,3084	1,154	0,054	1,693	0,140	0,002	1,121	0,069	0,710		
histidine	7,08	7,09	0,368	0,043	0,471	0,067	0,214	0,392	0,042	0,688		
isoleucine	1,01	1,03	2,330	0,236	2,888	0,311	0,172	2,171	0,155	0,580		
lactate	1,31	1,37	161,599	6,528	116,774	14,096	0,011	203,415	10,241	0,003		
methionine	2,017	2,027	3,779	0,254	2,762	0,360	0,035	3,138	0,167	0,051		
NAD	8,82	8,86	2,118	0,144	2,624	0,310	0,159	2,351	0,213	0,378		
phenylalanine	7,3	7,45	8,690	0,794	12,327	1,195	0,022	9,384	0,835	0,555		
phosphocholine	4,15	4,19	18,547	1,692	21,426	2,668	0,376	14,758	1,347	0,099		
pyruvate	2,38	2,4	4,233	0,298	6,223	0,650	0,013	4,689	0,390	0,366		
succinate	2,4	2,414	6,281	0,244	5,170	0,332	0,016	6,395	0,253	0,749		
taurine	3,407	3,445	6,369	0,515	4,911	0,358	0,034	4,587	0,185	0,005		
UDP der	4,36	4,39	13,729	0,935	10,651	1,322	0,076	13,023	0,982	0,610		
UDP-NAG	5,5	5,54	2,768	0,376	2,357	0,376	0,451	2,813	0,409	0,937		
valine	1,04	1,06	3,690	0,330	4,625	0,360	0,074	3,343	0,192	0,376		

**Supplementary Table 3.** Normalized concentration values of selected metabolites after 72 hours of treatment. P-values were obtained by the Student's t-test.

	c	Nanomaterial		72	72	72	72	72	72	72	72
metabolite	region (ppm)	Control	Control	mean	SEM	mean	SEM	P	mean	SEM	P
3-hydroxyisovalerate	1,247	1,2719	24,012	1,291	24,463	3,677	0,909	0,00	40,862	3,113	0,00
acetate	1,91	1,93	6,498	0,222	5,835	0,789	0,431	0,002	9,653	0,801	0,002
ADP/ATP	8,53	8,545	2,309	0,227	1,911	0,322	0,327	0,055	1,715	0,176	0,327
alanine	1,47	1,51	16,812	0,305	14,378	1,865	0,216	0,00	15,784	0,969	0,327
aspartate	2,79	2,84	2,548	0,096	2,132	0,296	0,199	0,018	3,328	0,279	0,018
ATP	4,39	4,43	9,273	0,481	7,621	1,128	0,197	0,00	5,690	0,352	0,00
creatinine/phocrea	3,0347	3,053	13,233	0,481	11,700	1,508	0,347	0,00	6,307	0,305	0,00
formate	8,45	8,47	1,650	0,074	1,437	0,209	0,352	0,003	2,647	0,279	0,003
fumarate	6,51	6,53	0,687	0,048	0,519	0,087	0,112	0,035	0,874	0,065	0,035
glutamate	2,33	2,38	9,995	0,963	7,635	1,092	0,125	0,861	10,216	0,781	0,861
glutamine	2,44	2,48	3,594	0,124	2,973	0,439	0,192	0,025	5,089	0,593	0,025
glycerol	3,566	3,5711	1,397	0,245	0,987	0,146	0,171	0,818	1,464	0,149	0,818
glycerol derivative	3,46	3,49	5,044	0,422	4,609	0,629	0,574	0,009	6,714	0,365	0,009
glycine	3,5508	3,566	8,370	0,201	7,214	0,928	0,241	0,001	6,914	0,289	0,001
GSSG	3,295	3,3084	1,125	0,083	0,900	0,127	0,156	0,623	1,180	0,071	0,623
histidine	7,08	7,09	0,449	0,018	0,422	0,060	0,666	0,00	0,605	0,031	0,00
isoleucine	1,01	1,03	3,068	0,177	2,675	0,345	0,326	0,045	3,059	0,113	0,968
lactate	1,31	1,37	207,204	11,667	182,754	24,312	0,378	0,581	219,994	19,509	0,581
methionine	2,017	2,027	3,396	0,146	3,233	0,382	0,695	0,002	2,722	0,114	0,002
NAD	8,82	8,86	2,360	0,059	2,129	0,294	0,454	0,038	2,916	0,239	0,038
phenylalanine	7,3	7,45	10,175	0,347	9,119	1,257	0,430	0,029	12,849	1,061	0,029
phosphocholine	4,15	4,19	7,842	0,310	6,620	0,899	0,217	0,267	7,043	0,622	0,267
pyruvate	2,38	2,4	3,821	0,264	3,242	0,508	0,327	0,037	5,734	0,796	0,037
succinate	2,4	2,414	5,568	0,122	4,308	0,557	0,042	0,045	5,062	0,197	0,045
taurine	3,407	3,445	6,936	0,150	6,536	0,848	0,649	0,00	4,942	0,121	0,00
UDP der	4,36	4,39	18,774	0,673	15,691	2,055	0,173	0,00	11,341	0,699	0,00
UDP-NAG	5,5	5,54	5,231	0,139	3,594	1,114	0,164	0,00	3,468	0,164	0,00
valine	1,04	1,06	4,501	0,277	4,005	0,524	0,415	0,854	4,569	0,232	0,854