

Supplementary materials for

# Intracellular Fate of Sub-Toxic Concentration of Functionalized Selenium Nanoparticles in Aggressive Prostate Cancer Cells

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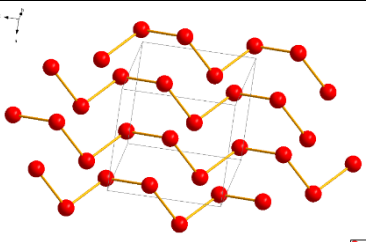
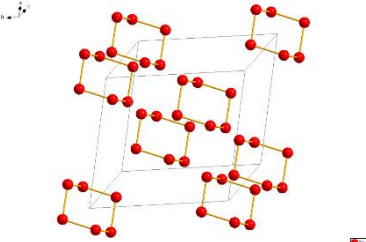
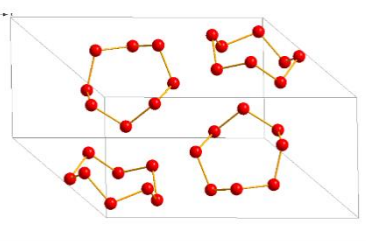
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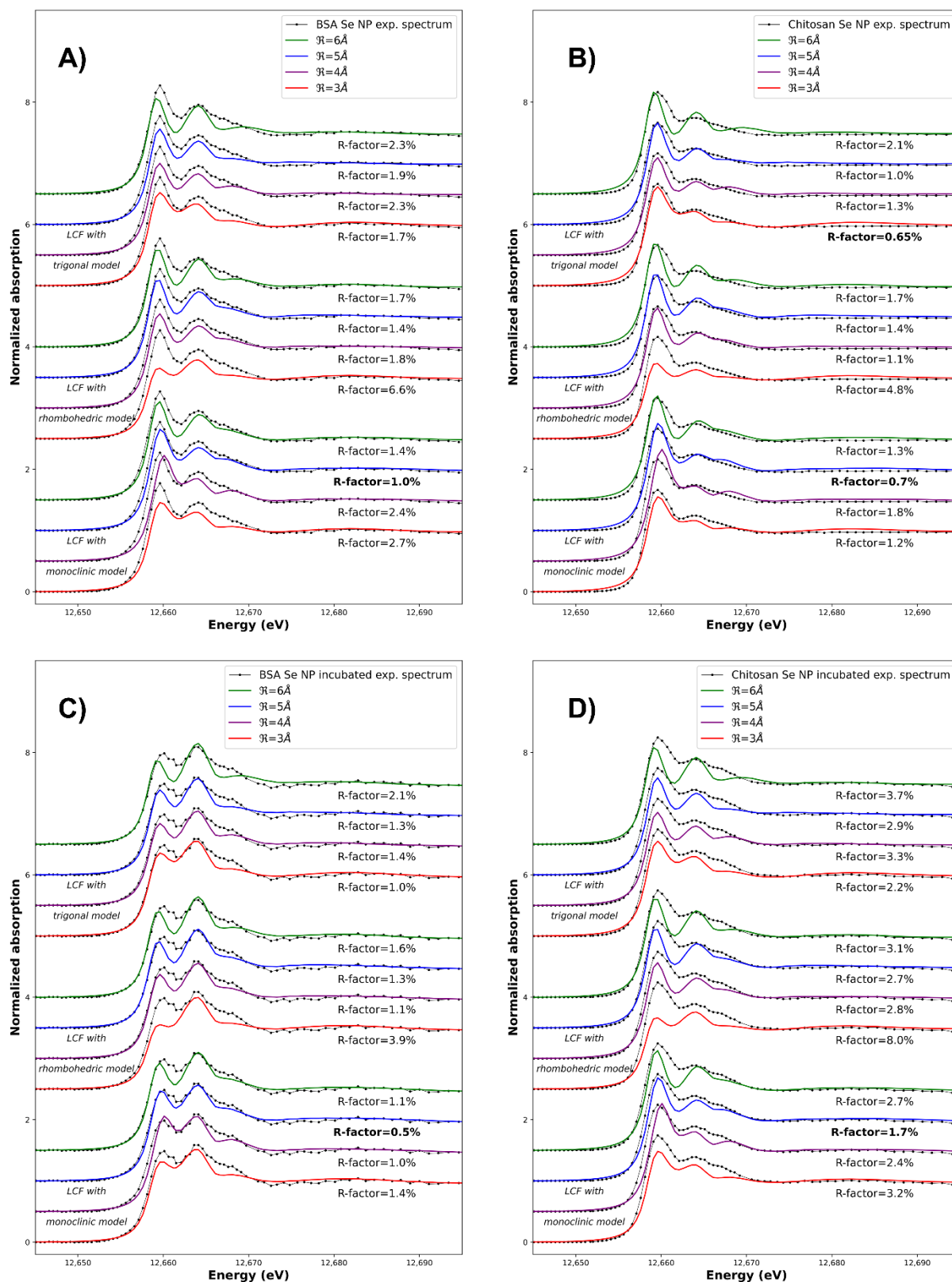
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## 1. XANES calculations

**Table S1:** Crystallographic structures used for the XANES calculations of  $\text{Se}^0$  components. Calculations were performed taking into account cluster radii  $\mathcal{R}$  equals to 3, 4, 5 and 6 Å.

Ref.	Form.	Space Group	Name	Crystallographic scheme
[1]	Se Grey	P3121	Se-6 Trigonal	
[2]	Se-6	R3-H	Se-6 Rhombohedral	
[3]	Se Red	P121/N1	Se-8 Monoclinic	

**Figure S1:** Experimental and calculated spectra for the determination of the Se speciation in the BSA and chitosan nanoparticles, as received and after incubation, by XANES least-square fitting using  $\text{Se}^{\text{IV}}$  experimental and  $\text{Se}^0$  calculated spectra as references: systematic study of the influence of both the radius of calculation ( $\mathcal{R}$ ) and the crystallographic structure (monoclinic, rhombohedral or trigonal) used for the  $\text{Se}^0$  calculation. R-factor of the best fits are in bold for better clarity.



**Table S2:** Values obtained on the determination of the Se speciation in the BSA and chitosan nanoparticles, as received and after incubation, by XANES least-square fitting using Se<sup>IV</sup> experimental and Se<sup>0</sup> calculated spectra as references: systematic study of the influence of both the radius of calculation ( $\mathfrak{R}$ ) and the crystallographic structure (monoclinic, rhombohedric or trigonal) used for the Se<sup>0</sup> calculation. Best fits values are in bold for better clarity.

BSA Se NP				Chitosan Se NP			
	$\mathfrak{R}$ (Å)	Se <sup>0</sup> calc.	Se <sup>IV</sup> exp.		$\mathfrak{R}$ (Å)	Se <sup>0</sup> calc.	Se <sup>IV</sup> exp.
<i>Trigonal model</i>	6	65.4%	34.6%	<i>Trigonal model</i>	6	71.1%	28.9%
	5	69.1%	30.9%		5	75.6%	24.4%
	4	69.8%	30.2%		4	76.4%	23.6%
	3	77.0%	23.0%		<b>3</b>	<b>84.1%</b>	<b>15.9%</b>
<i>Rhombohedral model</i>	6	66.8%	33.2%	<i>Rhombohedral model</i>	6	72.4%	27.6%
	5	68.5%	31.5%		5	74.3%	25.7%
	4	71.6%	28.4%		4	78.1%	21.9%
	3	79.3%	20.7%		3	88.7%	11.3%
<i>Monoclinic model</i>	6	72.1%	27.9%	<i>Monoclinic model</i>	6	78.2%	21.8%
	<b>5</b>	<b>71.3%</b>	<b>28.7%</b>		<b>5</b>	<b>77.4%</b>	<b>22.6%</b>
	4	69.6%	30.4%		4	75.9%	24.1%
	3	77.6%	22.4%		3	85.3%	14.7%
BSA Se NP incubated				Chitosan Se NP incubated			
	$\mathfrak{R}$ (Å)	Se <sup>0</sup> calc.	Se <sup>IV</sup> exp.		$\mathfrak{R}$ (Å)	Se <sup>0</sup> calc.	Se <sup>IV</sup> exp.
<i>Trigonal model</i>	6	54.3%	45.7%	<i>Trigonal model</i>	6	66.6%	33.4%
	5	57.9%	42.1%		5	70.8%	29.2%
	4	58.6%	41.4%		4	71.4%	28.6%
	3	64.5%	35.5%		3	79.1%	20.9%
<i>Rhombohedral model</i>	6	55.5%	44.5%	<i>Rhombohedral model</i>	6	68.1%	31.9%
	5	57.1%	42.9%		5	69.9%	30.1%
	4	60.0%	40.0%		4	73.2%	26.8%
	3	67.2%	32.8%		3	81.0%	19.0%
<i>Monoclinic model</i>	6	60.2%	39.8%	<i>Monoclinic model</i>	6	73.6%	26.4%
	<b>5</b>	<b>59.8%</b>	<b>40.2%</b>		<b>5</b>	<b>73.3%</b>	<b>26.7%</b>
	4	58.8%	41.2%		4	72.0%	28.0%
	3	65.3%	34.7%		3	79.7%	20.3%

Optimum adjustments are always obtained for  $\mathfrak{R}$  =5Å and a monoclinic structure, as for the Se<sup>0</sup> red reference, except for Chitosan Se NP where an adjustment of slightly better quality is obtained for  $\mathfrak{R}$  =3Å and a trigonal structure. Two points factors lead us not to consider this solution as it would induce:

- a structural evolution (from trigonal to monoclinic) with incubation,
- an improvement of the crystallinity of the Se<sup>0</sup> core as the corresponding  $\mathfrak{R}$  radius would increase from 3 to 5Å.

## 2. Cellular Se concentration

**Table S3:** Cellular Se concentrations and speciation measured on a QQQ-ICP-MS (Agilent 8900). n/a stands for non-measured sample. <LD stands for below the limit of detection calculated following IUPAC guidelines as  $LD_i = x_{bi} + k s_{bi}$  where  $k = 3$  (95% confidence level), and  $x_{bi}$  and  $s_{bi}$  are, respectively, the mean and standard deviation of the number of counts measured in blanks (i.e. acids stored and handled with the same material used to test the samples). n include replicate data only (i.e. re-run analyses)

Culture media = supernatant									Cell pellet = solid residue		
Total Se measured on digested samples				Se speciation [ppb, n=1] measured on non-digested samples					Total Se measured on digested samples		
Se (ppb) [wet mass]	2sd	n		SeCys	SeIV	SeMet	Unk.	SeVI	Se [ng/cells]	2sd	n
Ctrl-PC3-1	7.41	0.25	2	0.014	0.036	0.001	1.058	0.001	3.44E-06	n/a	1
Ctrl-PC3-2	n/a	n/a	n/a	0.004	<LD	<LD	1.008	0.003	1.13E-06	n/a	1
Ctrl-PC3-3	8.61	0.71	2	0.036	0.584	0.005	1.018	0.005	n/a	n/a	n/a
Ctrl-PC3-4	n/a	n/a	n/a	0.010	0.011	0.006	1.029	<LD	n/a	n/a	n/a
SeIV-PC3-1	483.54	8.51	2	10.116	86.510	0.644	1.204	0.188	8.97E-05	n/a	1
SeIV-PC3-2	448.48	6.42	2	9.373	72.183	0.552	1.232	0.096	1.00E-04	n/a	1
SeIV-PC3-3	454.86	7.35	2	9.523	71.232	0.685	1.050	0.013	3.64E-05	3.18E-06	2
SeIV-PC3-4	336.01	4.09	2	6.371	78.879	0.034	0.852	0.069	4.63E-05	9.66E-07	2
BSA-PC3-1	540.39	6.55	2	9.695	93.772	0.107	1.122	2.161	4.88E-05	1.39E-06	2
BSA-PC3-2	525.88	9.88	2	9.382	113.452	0.715	1.022	2.103	4.72E-05	1.11E-06	2
BSA-PC3-3	420.09	12.11	2	7.345	109.090	0.455	1.106	1.739	7.89E-05	2.63E-06	2
BSA-PC3-4	518.56	5.87	2	9.775	61.578	<LD	1.345	2.272	4.01E-05	2.32E-06	2
Chito-PC3-1	80.93	0.47	2	1.016	16.542	0.083	1.028	1.268	1.62E-06	3.01E-07	2
Chito-PC3-2	59.25	0.52	2	0.669	16.559	0.025	0.652	0.991	2.30E-06	4.64E-07	2
Chito-PC3-3	53.83	2.03	2	0.579	11.661	0.006	0.536	0.824	8.64E-07	1.07E-07	2
Chito-PC3-4	82.25	0.03	2	1.049	15.601	0.063	1.036	1.274	9.36E-07	1.49E-07	2

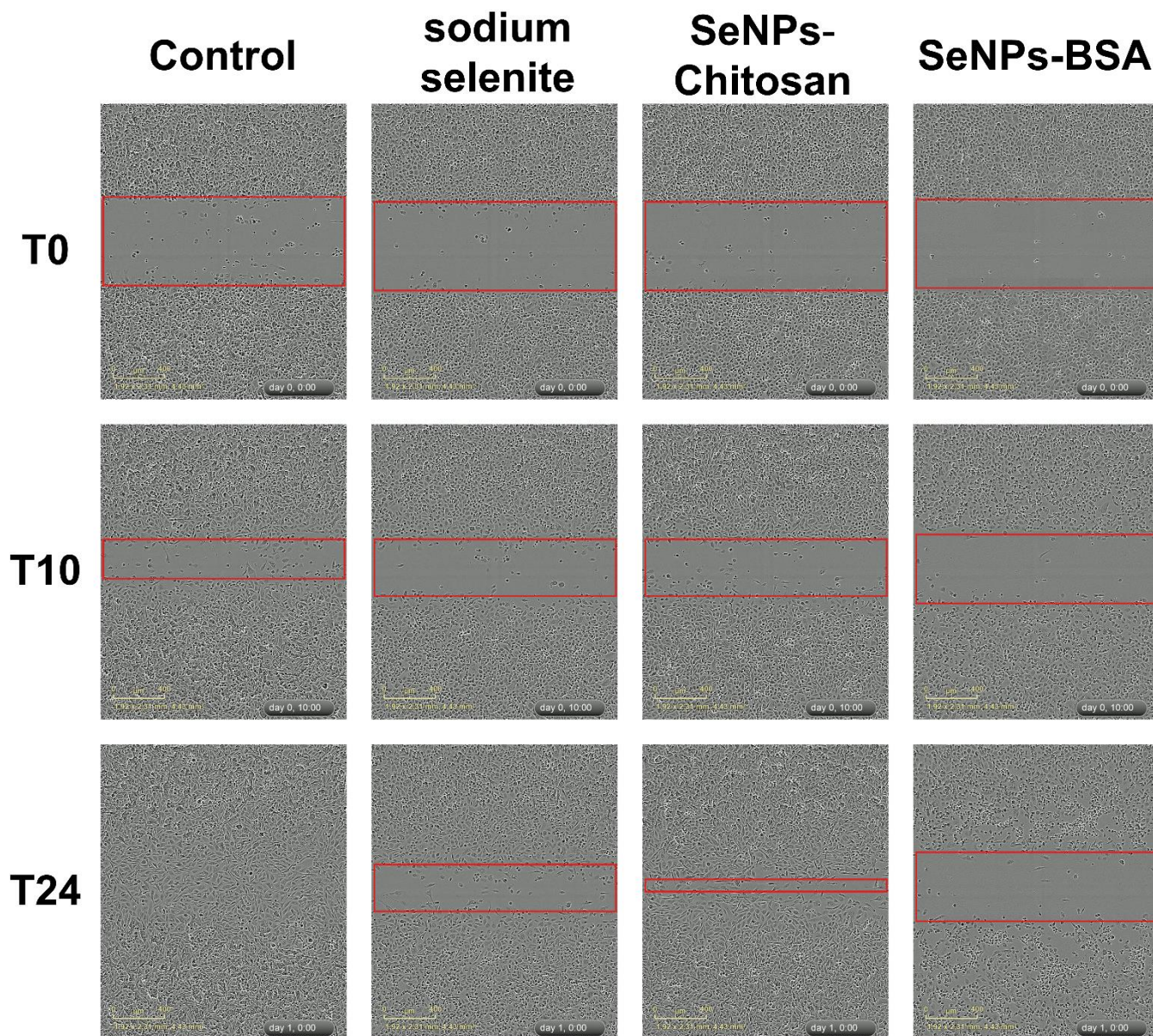
**Table S4:** Selenium concentrations of Se-enriched yeast certified reference material (SELM-1)

Reference material - SELM1			
Sample Name	Se (ppm) [dry mass]	2sd	n
SELM-1	1994	61	3
SELM-2	1981	59	4
SELM-3	1941	55	4
SELM-4	1877	162	3
Certified value (NRC) SELM-1	2031	70	



### 3. Wound healing assay

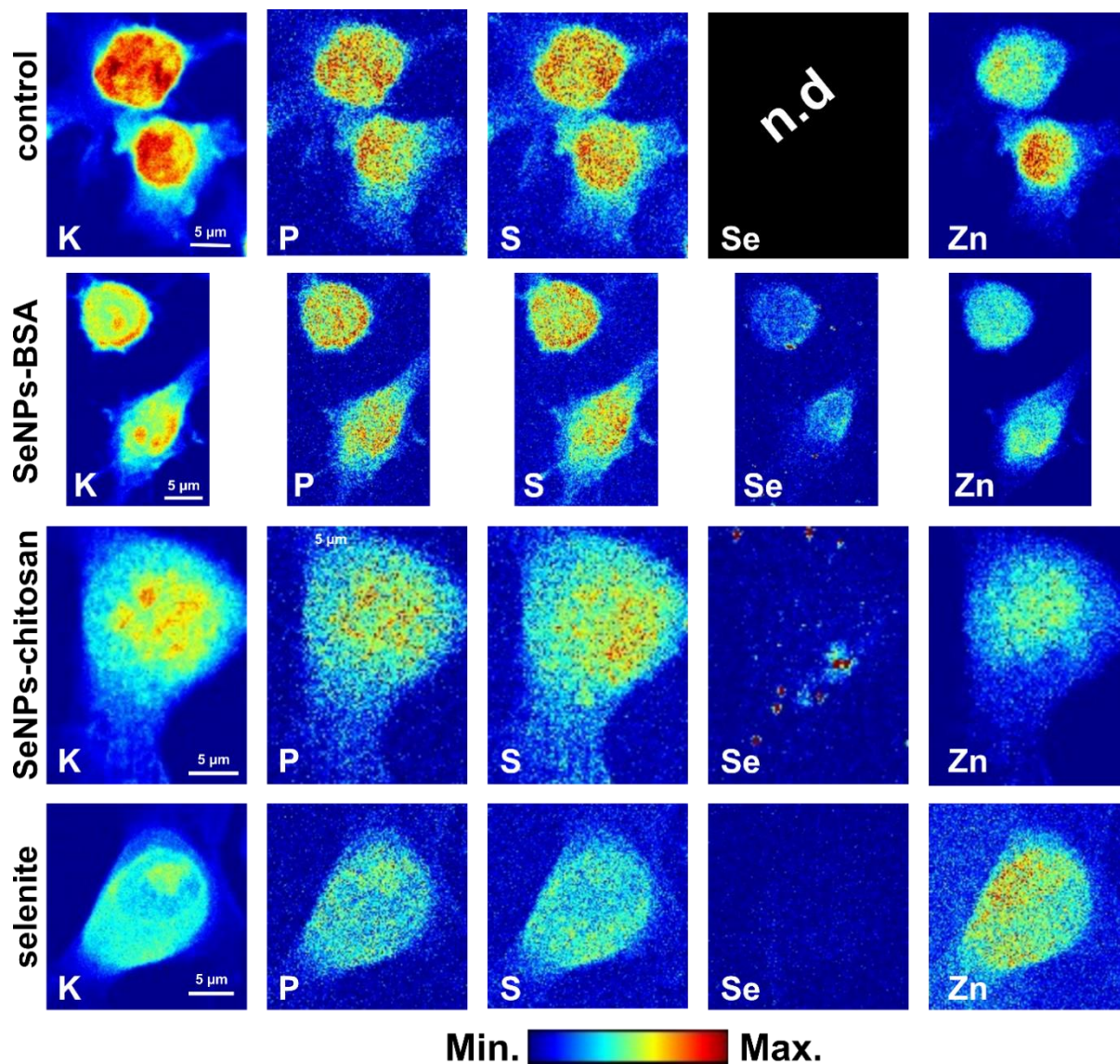
**Figure S2:** High resolution figure wound healing assay of PC3-cells at time 0, 10 and 24h (corresponds to Figure 7.D of the article). Selenium compounds (0 $\mu$ g/ml) was used as the control. Magnification: x10. Red rectangles schematically show the area of the wound (T0), and the regions of proliferation/migration (T10/T24).





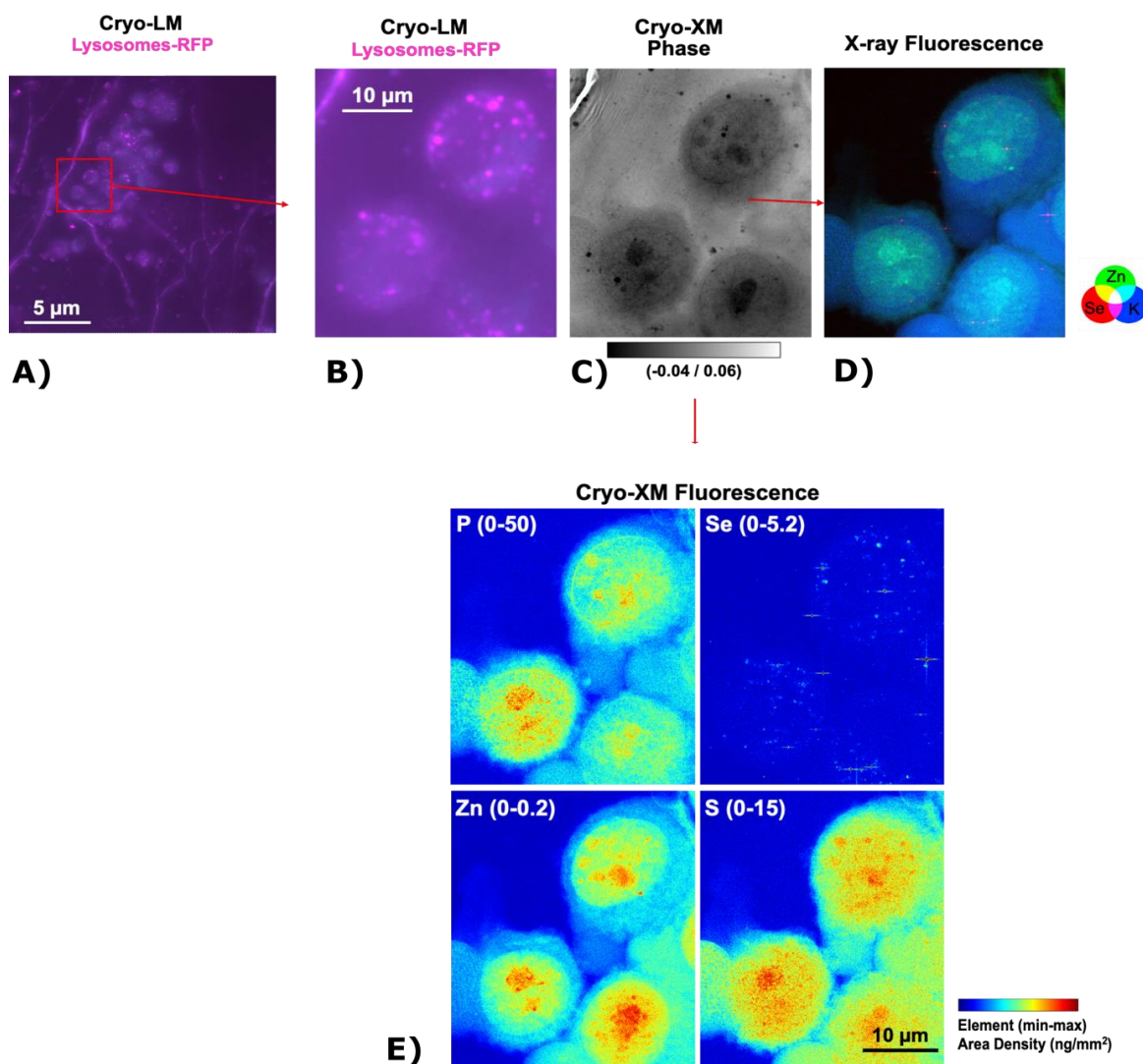
#### 4. X-ray fluorescence elemental mapping

**Figure S3:** Representative X-ray fluorescence imaging at 200nm stepsize at I14 Diamond nanoprobe beamline for plunge-frozen/Freeze-dried PC-3 cells exposed 24h at an IC20 concentration of SeNPs-BSA, SeNPs-Chitosan and sodium selenite aqueous solution. Control PC-3 cells are without any Se treatment. The cellular distribution for potassium (K), phosphorus (P), sulfur (S), selenium (Se) and zinc (Zn) are presented. n.d.: not detected.





**Figure S4.** Cryo-correlative imaging of PC-3 cells exposed to IC20 concentration of SeNPs-Chitosan for 24h. A) Cryo-optical fluorescence imaging (120K) of vitrified PC-3 cells cultured on Si3N4 membranes. Live cells were first incubated overnight with Lysosomes-RFP CellLight™, BacMam 2.0 reagent following manufacturer recommendations, then cells were exposed for 24h to SeNPs-Chitosan at IC20 concentration, followed by plunge-freezing procedure. B) Zoom in the insert of A). C) Hard X-ray phase contrast imaging of the same cells (panel B) at the ID16A cryo-nanoimaging allow to show some structural details mainly the nuclear region and shape of the cell. D) Merge of different element maps (Potassium K - blue, Selenium Se - red, Zinc Zn - green) of the same cells (panel B). K attests of the preservation of cell integrity after freezing. E) Cryo-XRF nanoimaging at 50nm step-size of a panel of different element maps (Phosphorus P, Selenium Se, Zinc Zn and Sulfur S) distribution in the same PC-3 cells exposed to SeNPs-BSA as panel B.



## 4. Genotoxicity

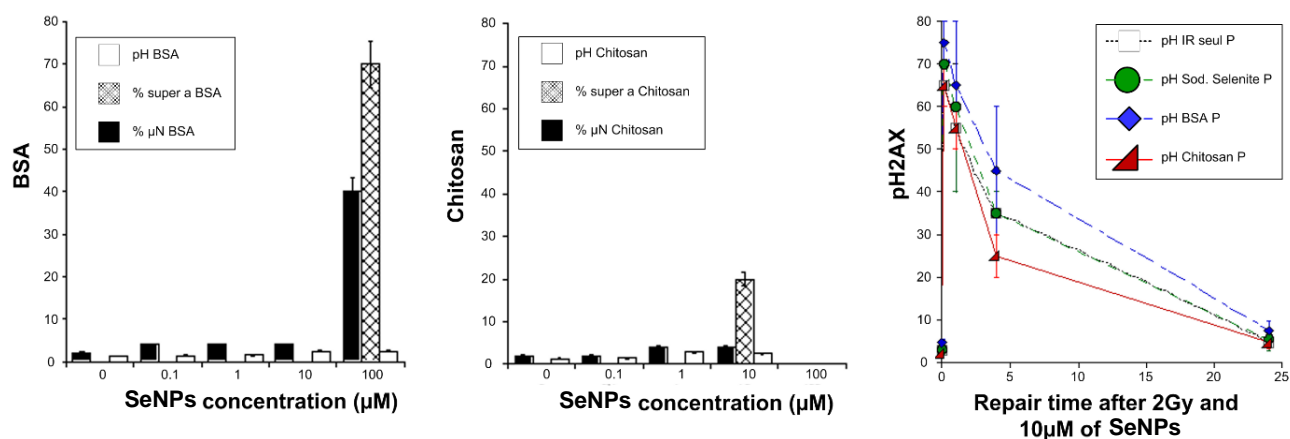
### 4.1. Genotoxicity Assessment of selenium nanoparticles on cancer cells

At day 1, 20  $\mu\text{L}$  of cell solution were plated onto 12-mm diameter coverslip (20000 cells/20 $\mu\text{L}$ ). After 5min at 37°C, 2 mL medium are added to the cells and left overnight in the incubator (37°C, 5% CO<sub>2</sub>). At day 2, the medium and cell debris were removed and replaced by fresh new medium containing SeNPs-BSA or Chitosan or sodium selenite. A concentration range [0.1, 1, 10, 100  $\mu\text{M}$ (Se)] was used. At day 3, after 24h incubation, cells were thoroughly rinsed 3 times with PBS. Further, the 2 Gy  $\gamma$ -irradiation was performed. Dosimetry certifications and the  $\gamma$ -rays irradiations were performed with a 6 MeV  $\gamma$ -rays clinical irradiator (SL 15 Phillips) (dose-rate: 6 Gy.min<sup>-1</sup>) at anti-cancer Centre Léon-Bérard, (Lyon, France) [4, 5]. Control followed the same stages than treated cells except the irradiation. For the kinetic analysis, cells were fixed with 100 $\mu\text{L}$  solution of 4% PFA diluted in PBS for 10 min and rinsed with twice with PBS. Next, Immunofluorescence protocol and foci scoring were followed as described elsewhere [4, 6]. The stabilization was obtained with 200 $\mu\text{L}$  of lysis solution laid on cells for 3 min, and then rinsed with PBS. Coverslips were transferred to a new petri dish. Anti- $\gamma$ -H2AXser139 antibody (clone JBW301; Merck, Millipore, Darmstadt, Germany) was applied at 1:800. Anti-pATMser1981 (clone 10H11.E12; Millipore, Germany) antibodies were used at 1:100. The first antibody, pH2AX, was used in a 1:800 dilution. 30 $\mu\text{L}$  were used per coverslip, incubation at 37°C for 1h, then rinsed with PBS. In a light-preserved room, 30  $\mu\text{L}$  of second Ab, diluted in 1:100 in PBS-3%BSA solution, was applied on coverslips containing cells (20min at 37°C), then rinsed with PBS. Cells on the coverslips are labelled with Vectashield DAPI (H100 abcsys) mounting solution. DAPI allows to see micronuclei and DNA.

### 4.2. Genotoxicity assay

We studied the induced DNA damage (double-strand breaks - DSBs) by  $\gamma$ -H2AX labeling and micronucleus labeling. The kinetics of nuclear pATM foci (repair signaling protein) are used to quantify DSB recognition prior to repair by suturing (the predominant repair mode in mammals). Se<sup>0</sup>NPs or sodium selenite do not generate a significant number of DSBs and, consequently, the rate of persistent DSBs after 24h of exposure remains very low at the concentrations we had chosen, but also for high concentrations such as of 100  $\mu\text{M}$ . At sub-cytotoxic concentrations, the notion of genotoxicity of the soluble sodium selenite species are non-existent and there are no major effects concerning the 2 types of nanoparticles. Figure S4 shows a slowdown in the rate of recognition by DSBs generated by pre-irradiation of PC3 cells following 24h exposure to BSA-coated Se<sup>0</sup>NPs, which (above) caused a slowdown in the proliferation of these highly aggressive prostate cancer cells. This effect on ATM suggests indirectly that internalized and potentially biotransformed Se<sup>0</sup>NPs have an impact intracellularly, leading to a reduction in the migration and hence invasion capacity of these aggressive cells, but also to a slowing down of ATM addressing to the nucleus when DSBs are generated.

**Figure S5:** Plot of DSB number (ph) and micronucleus number ( $\mu\text{N}$ ) for PC-3 cells exposed to increasing concentrations of 0 (control) to 100  $\mu\text{M}$  Se<sup>0</sup>NPs coated with BSA (left) or chitosan (middle). The curves on the right show DSB repair time after irradiation at 2Gy (generating 70-75 double-strand breaks) following exposure to SeNPs or sodium selenite salts (10 $\mu\text{M}$  for 24h).



## 5. References

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