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Structural bases for rational design of new biomaterials based $\triangleleft \gg$ on metallodrug/ β -lactoglobulin adducts

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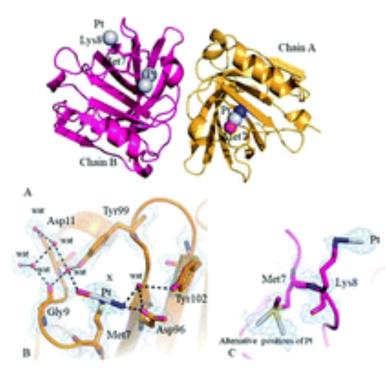
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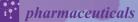
Structural bases for rational design of new biomaterials based on metallodrug/β-lactoglobulin adducts

Graphical Abstract









Abstract:

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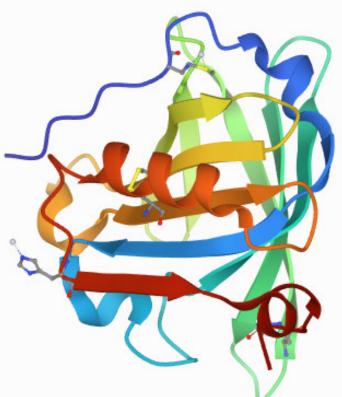
 β -lactoglobulin is a whey carrier protein of 18.4 kDa. Due to its high solubility, safe status, biodegradable nature, gel forming ability, abundance, stability at acidic pH and stability against gastric pepsin, β -lactoglobulin can be considered as a good system for the preparation of micro- or nanoparticles for pharmaceutical industry. It has been demonstrated that β -lactoglobulin-pectin nanoparticles are able to transfer cytotoxic Pt compounds to cancer cells. With the aim to unveil the molecular basis of the metallodrug recognition by β -lactoglobulin, we are analyzing the interactions between this protein and several metallodrugs. The interaction between cisplatin (CDDP), the most used Pt-based anticancer agent, and β lactoglobulin have been investigated both in solution and at solid state. UV-vis absorption spectroscopy and circular dichroism data indicate that the protein retains its conformation upon CDDP binding. X-ray crystallography analysis reveals that β-lactoglobulin interacts with CDDP through coordination of Pt fragments to the side chains of Met7, His146 and Lys8, with the number of binding sites increasing over time. ESI-MS data indicate that $[Pt(NH_3)_2Cl^+]$, $[Pt(NH_3)_2OH^{2+}]$ and $[Pt(NH_3)_2^{2+}]$ fragments interact with the protein. These results open the way for a rational design of new biomaterials based on metallodrug/β-lactoglobulin adduct nanoparticles.



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Introduction



- β-lactoglobulin is the major whey protein in cow's milk of 18.4 kDa (162 amino acids)
- It is a globular protein composed by 8stranded antiparallel β-barrel with a 3turn α-helix on the outer surface.

 $\sqrt{NH_3}$

Cisplatin: the most used Pt-based anticancer agent

The interactions between β -lactoglobulin and cisplatin have been investigated both in solution and at solid state

E. Dufour et al. Biochimica et Biophysica Acta, 1994, **1205**(1), 105. Z. Teng et al. Food Chemistry, 2016, **204**, 391. Z. Teng et al. Food Chemistry, 2014, **159**, 333. Z. Izadi et al. Chem Biol and Drug Design 2016, **88**(2), 209. Z.H. Siddik, Oncogene 2003, **22**, 7265. L. Kelland, Nature Reviews Cancer 2007, **7**, 573. M.H. Green, J. Nat Cancer Inst. 1992, **84**, 306.



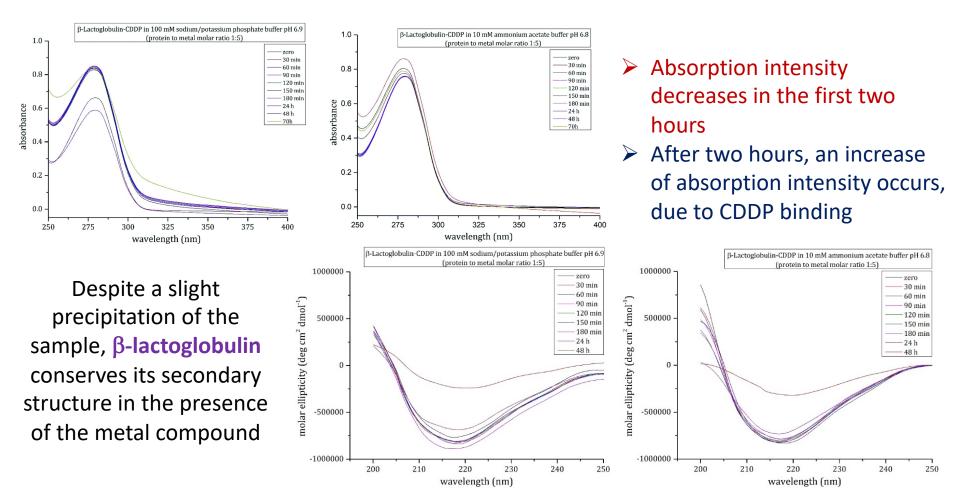
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Results and discussion

In solution characterization of CDDP binding to β -lactoglobulin



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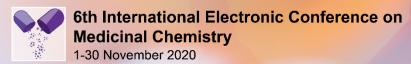
Results and discussion

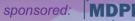
CDDP soaking for 18h

Crystallographic studies

The structure of the CDDP/ β -lactoglobulin adducts contain two molecules in the asymmetric unit

Pt Lys8 Chain A Tyr99 Asp11 Met7 Alternative positions of Pt wat Lys8 Met7 Tyr10: wat Asp96 Gly9 wat wat Glu51* Chain B Tyr99 wat Asp11 Met7 Lys8 wat х Pt His146 Tyr102 His146 Gly9 Asp96 Met7 Alternative positions of Pt





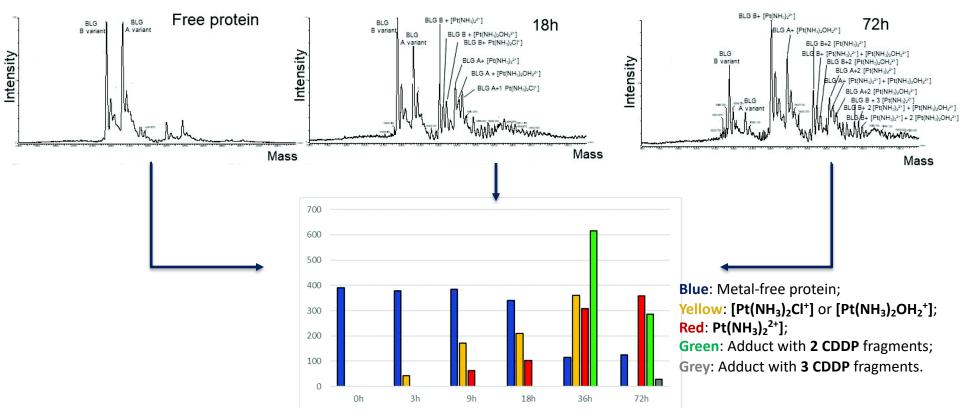
CDDP soaking for 72h

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Results and discussion

Mass spectrometry analysis

Electrospray ionization mass spectra of the CDDP/ β -lactoglobulin are collected as a function of time

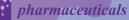


Bar graph summarizes the results of mass spectrometry course experiments. Two forms of native protein are observed for A and B variants of the protein. After three hours the first Pt/β -lactoglobulin is formed. Longer reaction times (9h) give rise to another protein-metal fragment. Only after 36h the presence of two binding Pt sites is observed, while 72h are needed for observing adducts of the protein with three CDDP molecules



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Conclusions



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- The first structural study on the interaction of β-lactoglobulin with an anticancer metallodrug is reported;
- In solution data demonstrate that the protein binds CDDP without affect its overall conformation;
- Crystallographic studies underline the presence of binding sites of CDDP close to the side chain of Met7, Lys8 and His146. β-lactoglobulin can form adducts with one Pt-containing fragment coordinating protein residues side chains;
- ESI-MS analysis reveals the presence of [Pt(NH₃)Cl⁺], [Pt(NH₃)₂OH₂⁺] and [Pt(NH₃)₂²⁺] fragments bond to the protein;
- The number of CDDP fragments embedded by protein increases with time;
- Monodentate and bidentate mode of Pt binding to the protein are possible, the former preceding the latter in coordination of the protein;
- Up to 3 CDDP molecules can bind the protein.

The interactions β -lactoglobulin with other metallodrugs is under investigation. Further efforts will be spent toward a rational design of new biomaterials based on metallodrug/ β -lactoglobulin adduct nanoparticles.

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Acknowledgments

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