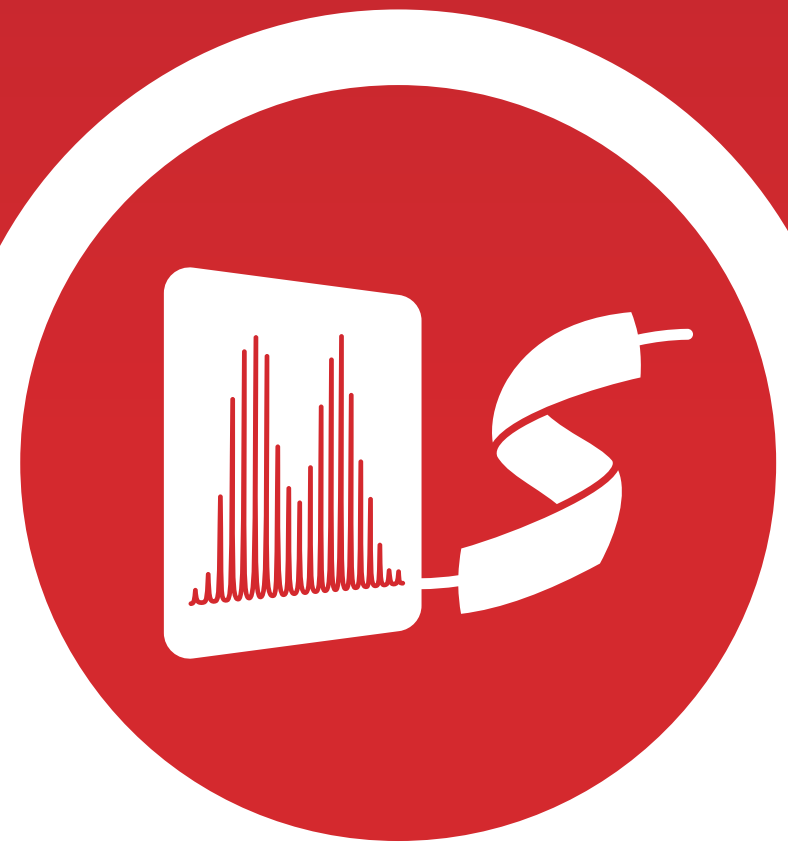


# FAST PHOTOCHEMICAL OXIDATION OF PROTEINS BY SINGLET OXYGEN

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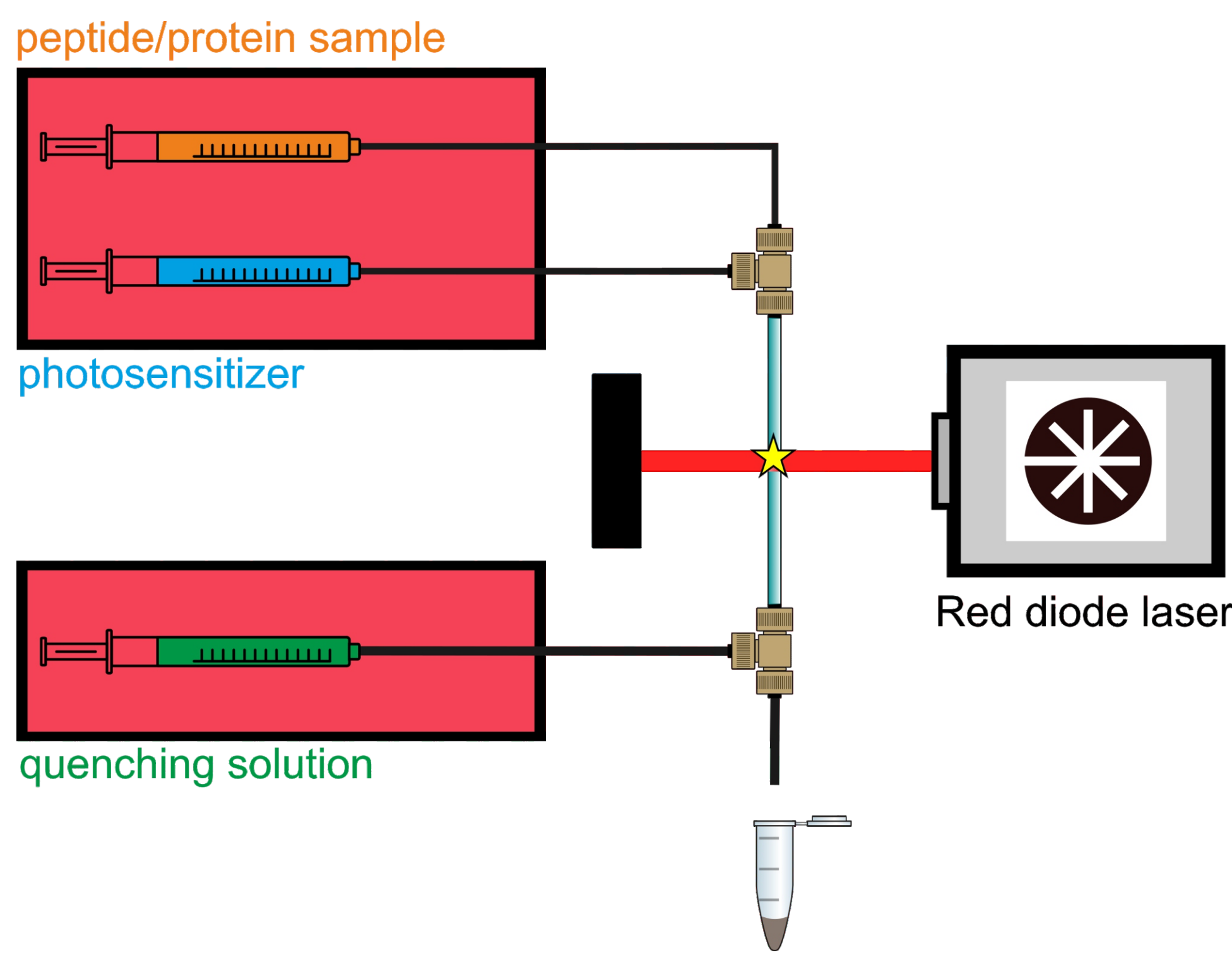
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## INTRODUCTION

- Protein footprinting is a powerful tool for structural characterization of proteins and their complexes that examine conformational changes and ligand binding by determining the solvent accessibility of the backbone or side chain structures of protein molecules [1]. By comparing the footprint of a protein in two different structural states, changes in the protein topography can be detected and interpreted in the context of other structural information [2].
- There are many footprinting methods used with the mass spectrometric detection. One popular approach is based on monitoring extent of oxidation of protein side chains due to exposure of oxidative reagents from solution.
- We examined the possibility to achieve footprinting by exposing model peptides or proteins to molecular singlet oxygen, a known reactive oxygen species.
- Singlet oxygen, O<sub>2</sub>(<sup>1</sup>Δg) or <sup>1</sup>O<sub>2</sub>, is the lowest excited electronic state of molecular oxygen. It is generated by the transfer of energy to ground (triplet) molecular oxygen and it exhibits significantly greater reactivity towards organic compounds than triplet oxygen [3].
- Singlet oxygen reacts with a wide range of biological targets, including DNA/RNA, proteins, and all classes of lipids. Kinetic data are consistent with proteins being a major target for the singlet oxygen, with modifications occurring preferentially at Trp, His, Tyr, Met, and Cys side-chains. It has been also described that reaction between proteins and singlet oxygen generates peroxides in high yield, mostly on Tyr, Trp, and His residue sites. [4,5]

## METHODS

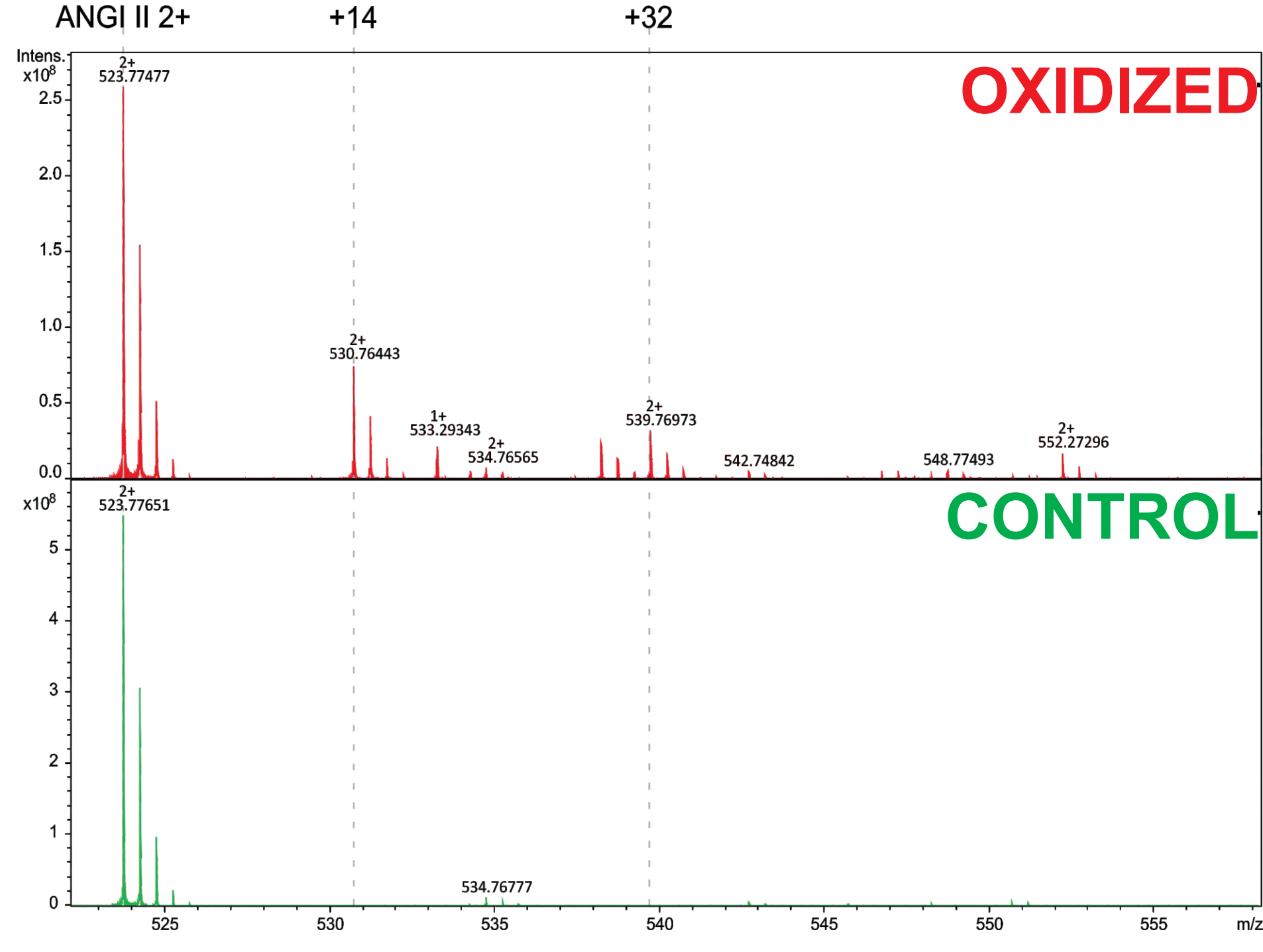


**Schema 1: Apparatus and experimental steps**  
1/ Mixing sample and photosensitizing compound  
2/ Irradiation with red laser light, generation of singlet oxygen and reaction with the analyte  
3/ Quenching the reaction and sample collection  
4/ Detection of collected sample by HPLC-MS

## METHODS

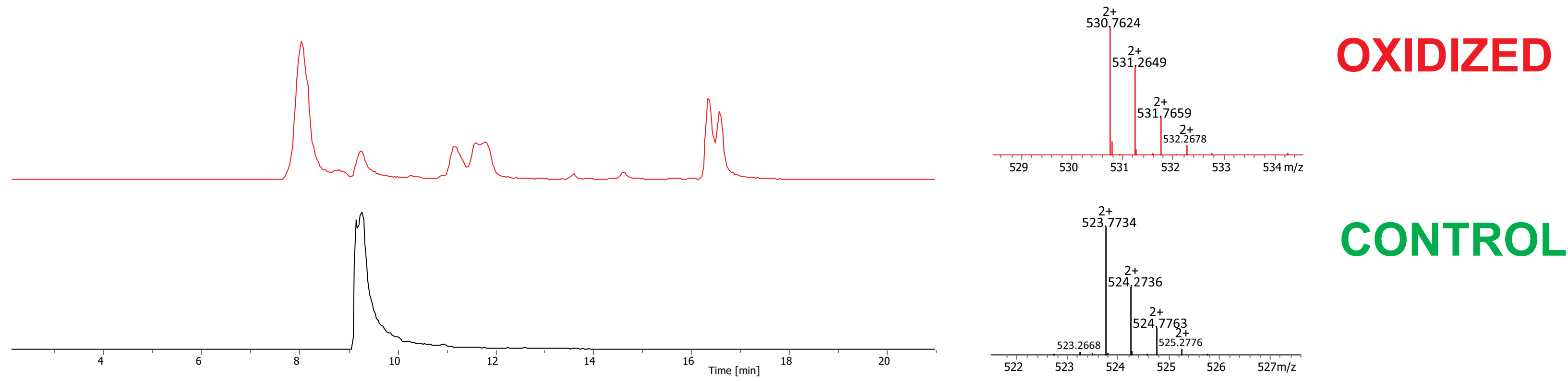
- Quench-flow setup consisted of 3 syringe pumps, separately delivering solution of the protein/peptide to be examined, solution of the photosensitizer compound and quenching solution (Schema 1)
- Highly reactive singlet oxygen was generated after laser irradiation by a photodynamic effect of phthalocyanine - a process in which excited photosensitizer interacts with a ground state oxygen dissolved in water, and creates singlet oxygen [6].
- The collected samples were analyzed by direct infusion into solarix XR 15T, Bruker Daltonics
- The LC analysis was performed using Evosep One EV-1000 (Evosep) and timsTOF SCP (Bruker Daltonics)

## RESULTS



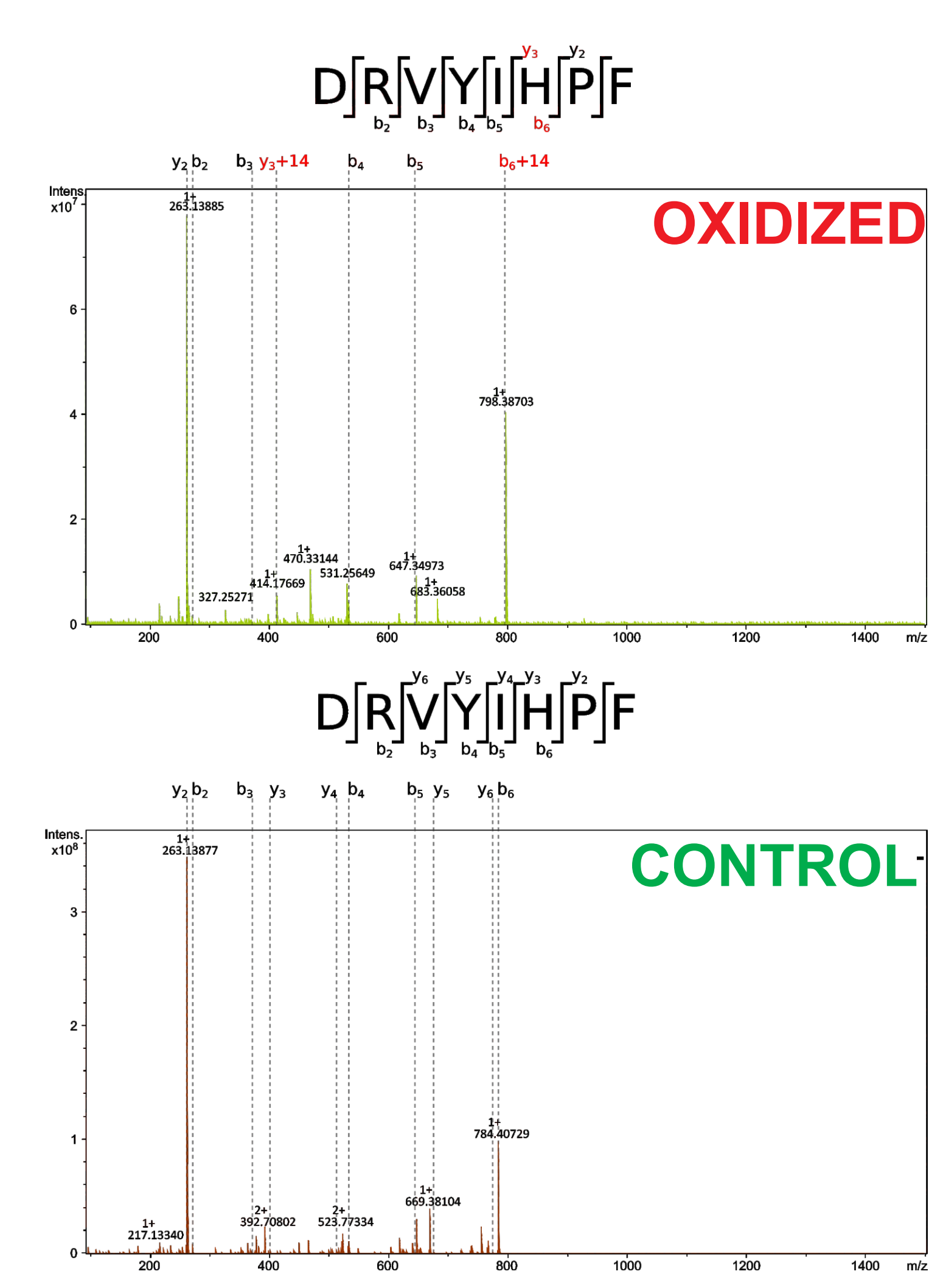
**Figure 1: Mass spectrum of Angiotensin II; sequence=DRVYIHPF**  
Top: After exposure to singlet oxygen  
Bottom: Control (intact Angiotensin II, doubly charged ion at m/z=523)

Mass spectrum of Angiotensin II showed mainly the doubly charged ion at m/z=523. Sample exposed to singlet oxygen showed new doubly charged peaks at m/z=530 (addition of 14Da), corresponding to addition of O atom and loss of two H atoms (or addition of 2O and loss of water) and m/z=539 (addition of 32Da), corresponding to addition of 2O atoms.



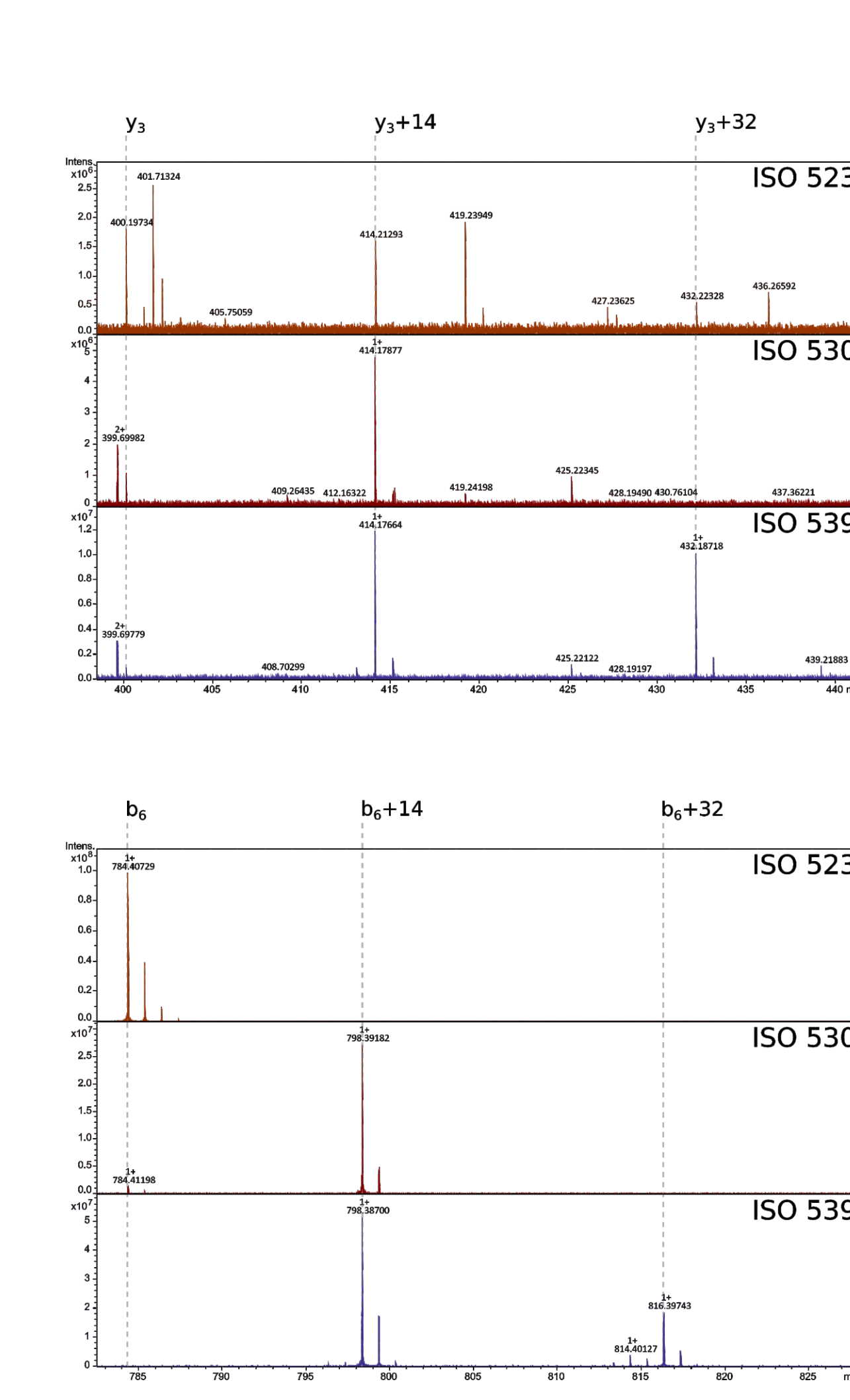
**Figure 2: LCMS analysis of Angiotensin II oxidized by singlet oxygen**  
Top: XIC (m/z=530.7) of oxidized sample  
Bottom: XIC (m/z=523.7) of control sample

## RESULTS

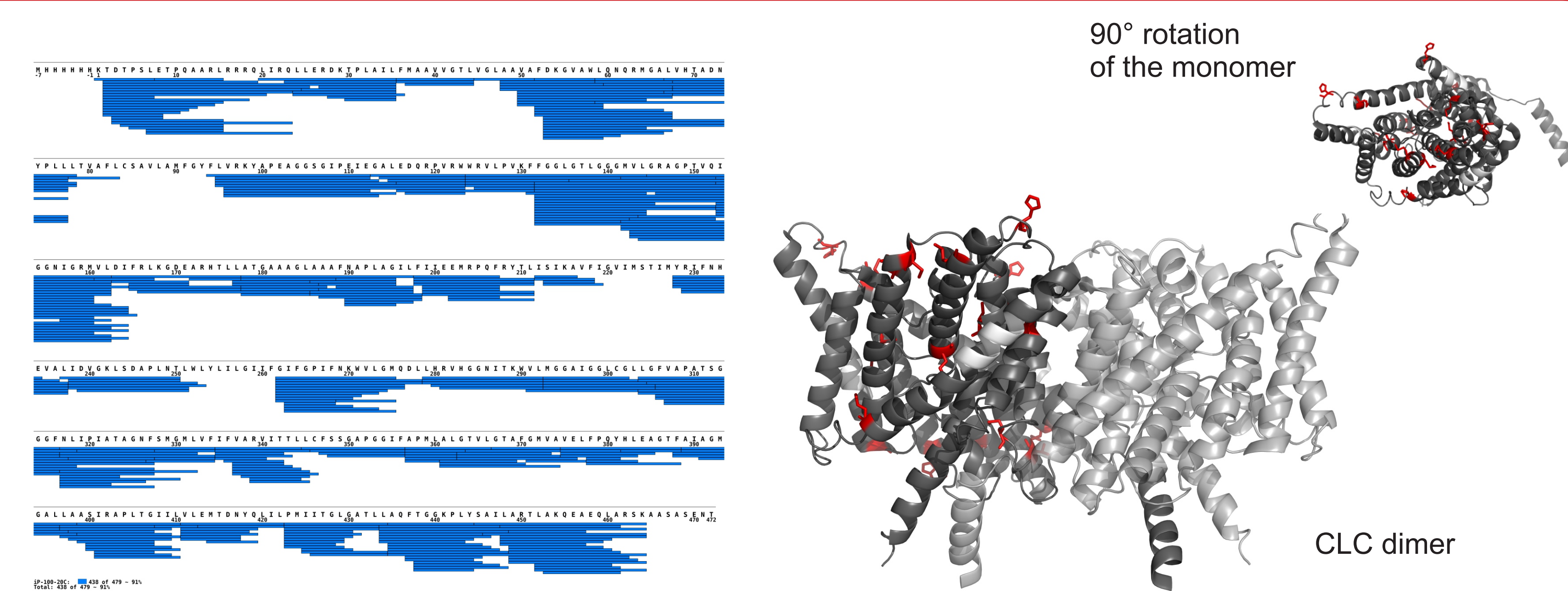


**Figure 3: CID MSMS spectrum of doubly charged Angiotensin II precursors**  
Top: Oxidized precursor; MSMS of 530  
Bottom: Control (intact Angiotensin II); MSMS of 523

The MSMS spectrum shows that the oxidative modification in the sequence begins with fragments y3 and b6, which localizes it to histidine



**Figure 4: CID MSMS spectrum of doubly charged Angiotensin II precursors**  
Top: Fragment y3 and its modifications  
Bottom: Fragment b6 and its modifications

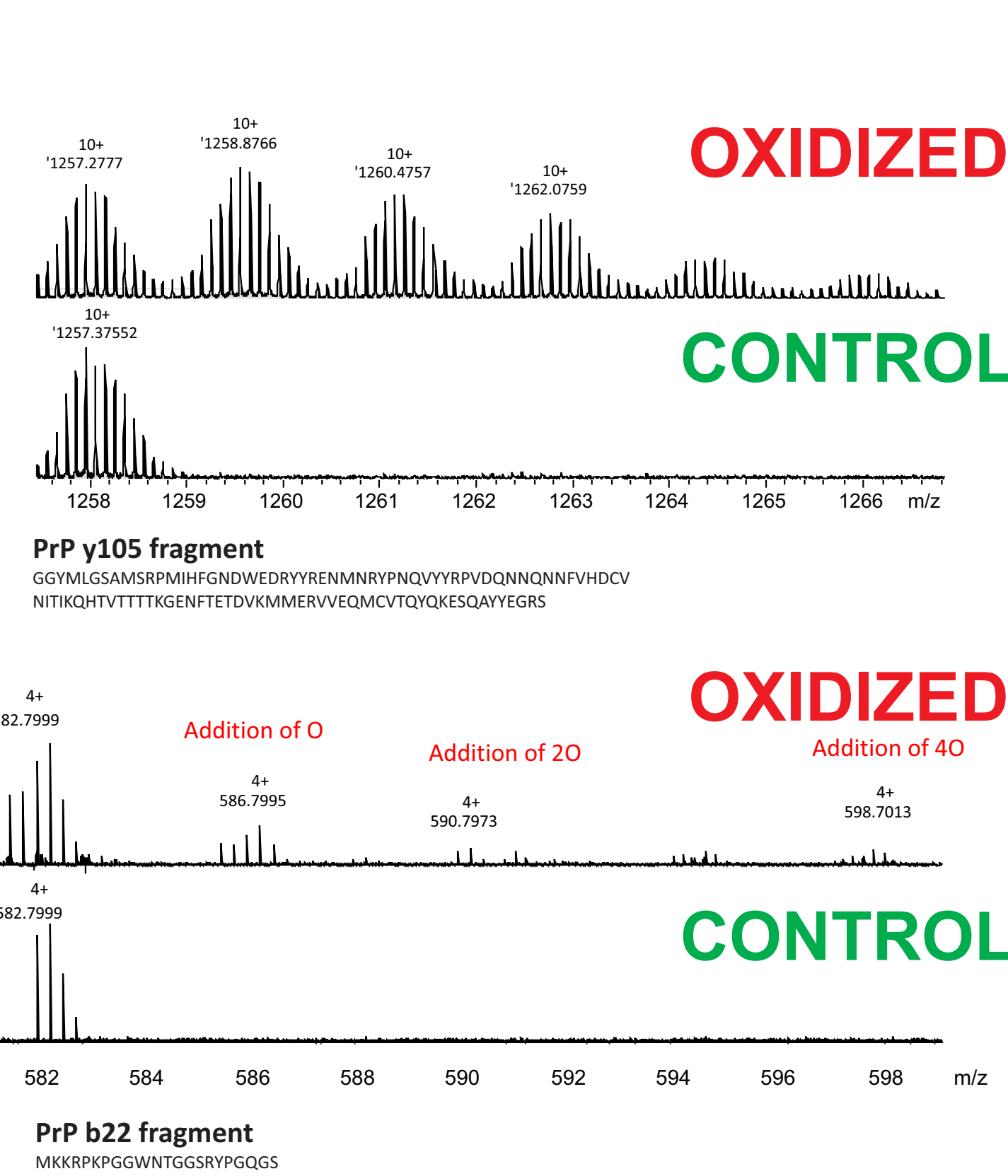


**Figure 7: Singlet oxygen labeling of CLC-ec1 using in-house built quench-flow apparatus.** Summary of sequence coverage (left) and visualization of modified sites on the CLC-ec1 structure (right)

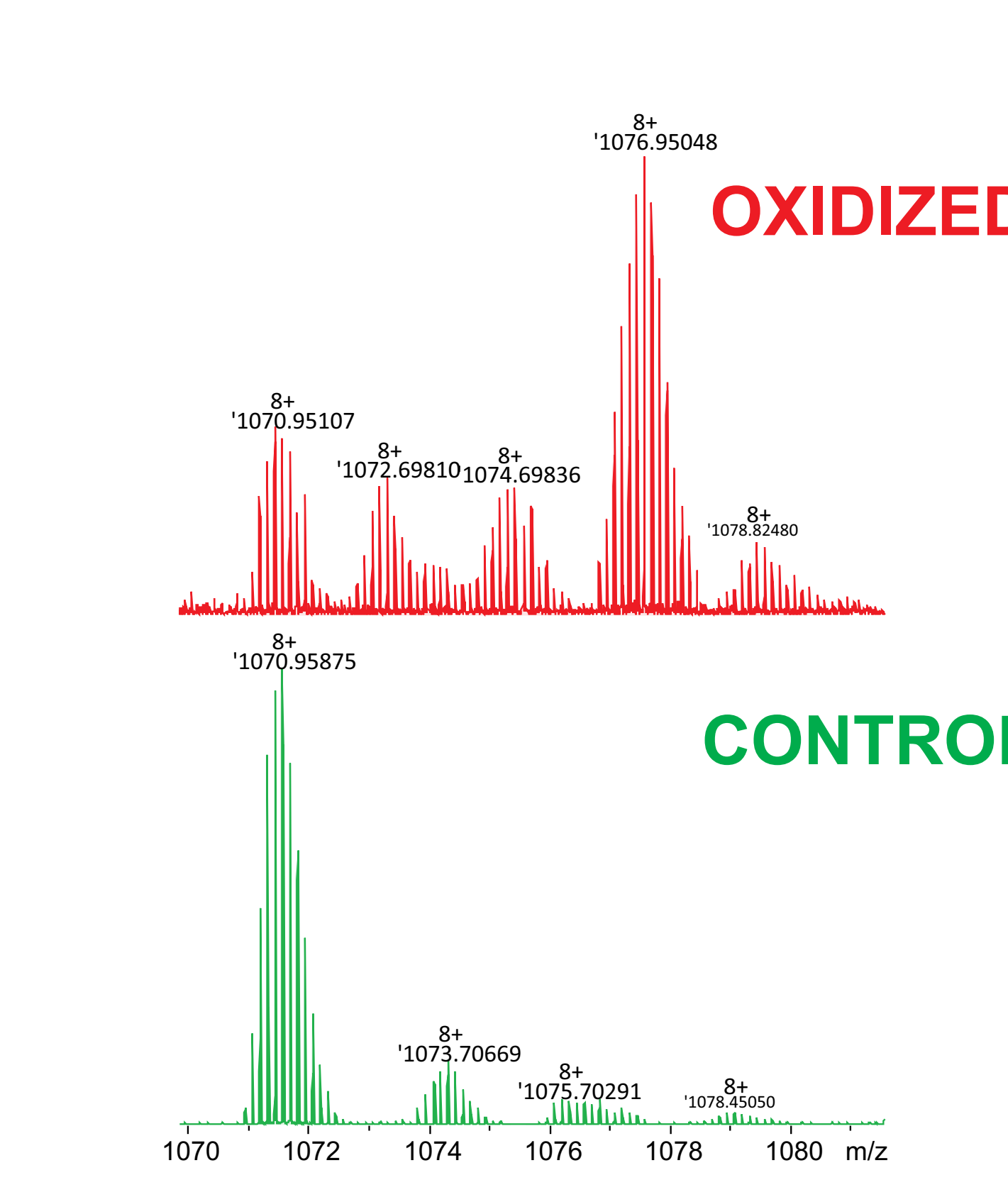
## REFERENCES

- [1] Xu G and Chance M.R.: Hydroxyl Radical-Mediated Modification of Proteins as Probes for Structural Proteomics, Chem. Rev. 2007, 107, 3514-3543.
- [2] Wang L and Chance M.R.: Protein Footprinting Comes of Age: Mass Spectrometry for Biophysical Structure Assessment, Mol Cell Proteomics. 2017 May; 16(5): 706–716.
- [3] Ogilby, P.R.: Singlet oxygen: there is indeed something new under the sun, Chem. Soc. Rev., 2010, 39, 3181-3209.
- [4] Di Mascio et al.: Singlet Molecular Oxygen Reactions with Nucleic Acids, Lipids, and Proteins, Chem. Rev. 2019, 119, 3, 2043–2086.
- [5] Michaeli A., Feitelson J.: Reactivity of Singlet Oxygen Towards Amino Acids and Peptides, Photochemistry and Photobiology, 1994, 59, 284-289.
- [6] Rosenthal I. et al.: The role of molecular oxygen in the photodynamic effect of phthalocyanines. Radiat Res. 1986, 107, 136-42.

## RESULTS



**Figure 5: Top-down analysis of recombinant prion protein (PrP) oxidized by singlet oxygen**  
Top: Fragment y105; oxidized and control  
Bottom: Fragment b22; oxidized and control



**Figure 6: Oxidation of Ubiquitin by singlet oxygen**  
Top: Peaks of oxidized Ubiquitin 8+ species  
Bottom: Peak of control Ubiquitin 8+

## CONCLUSIONS

- Singlet oxygen generated by photodynamic effect was tested as a reagent for protein footprinting
- Experimental conditions were optimized for quench-flow conditions
- Model peptides and proteins were oxidized by singlet oxygen and analyzed by MS
- Oxidative modifications were detected by direct infusion and LC-MS and localization was performed by top-down and bottom-up

## ACKNOWLEDGMENTS

The work was supported by the Czech Science Foundation, project 22-27695S, by the EU H2020 (EPIC-XS), project number 82383 and by the EU H2020 (EU\_FT-ICR\_MS), project number 731077. The authors would also like to thank Petr Man and Zdenek Kukacka for help with the data processing and useful discussion of the results.

