Peptide Sequencing with Top-down Synthesized TiO, Nanowires Using Laser Desorption/Ionization Time-of-Flight Mass Spectrometry

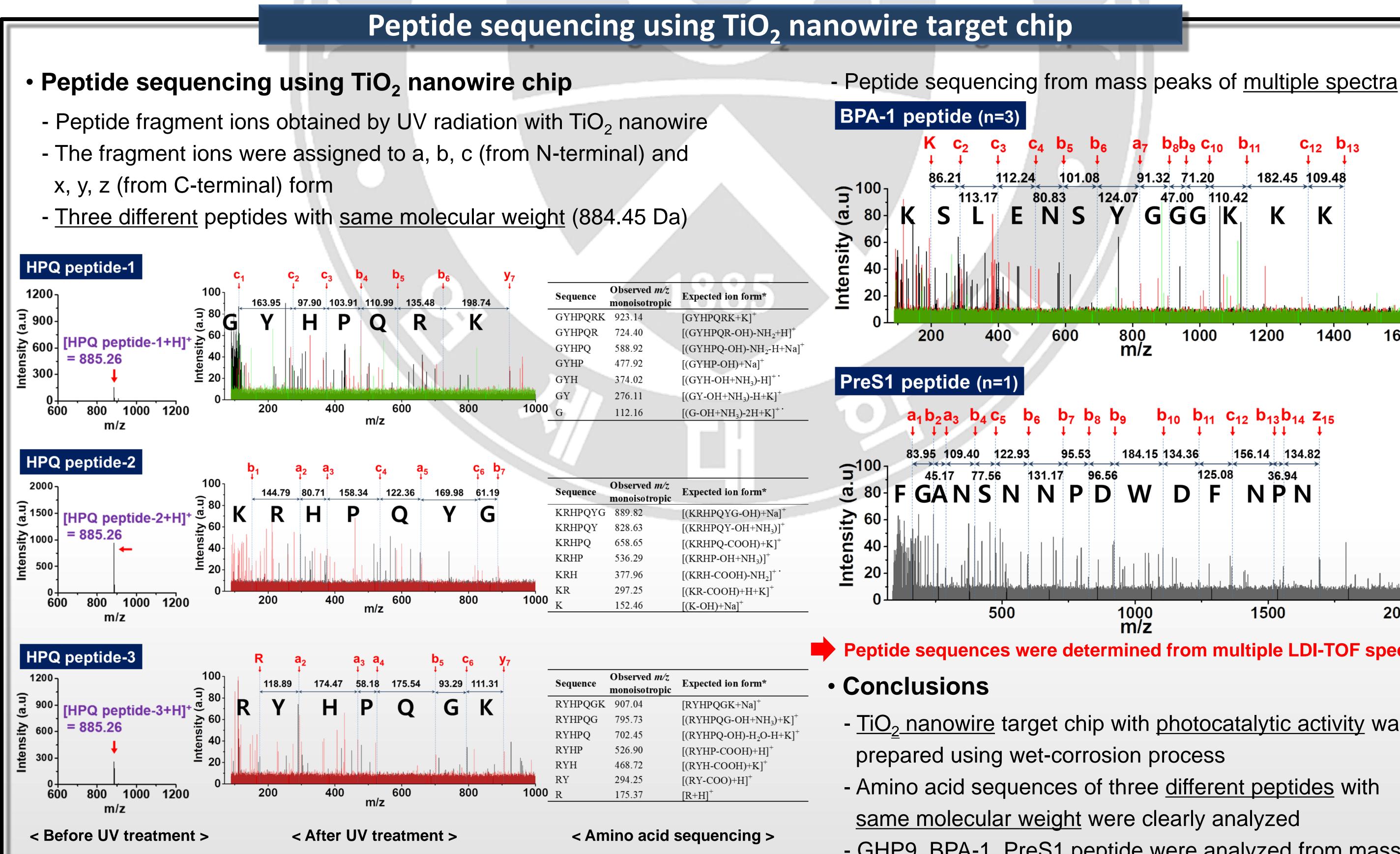
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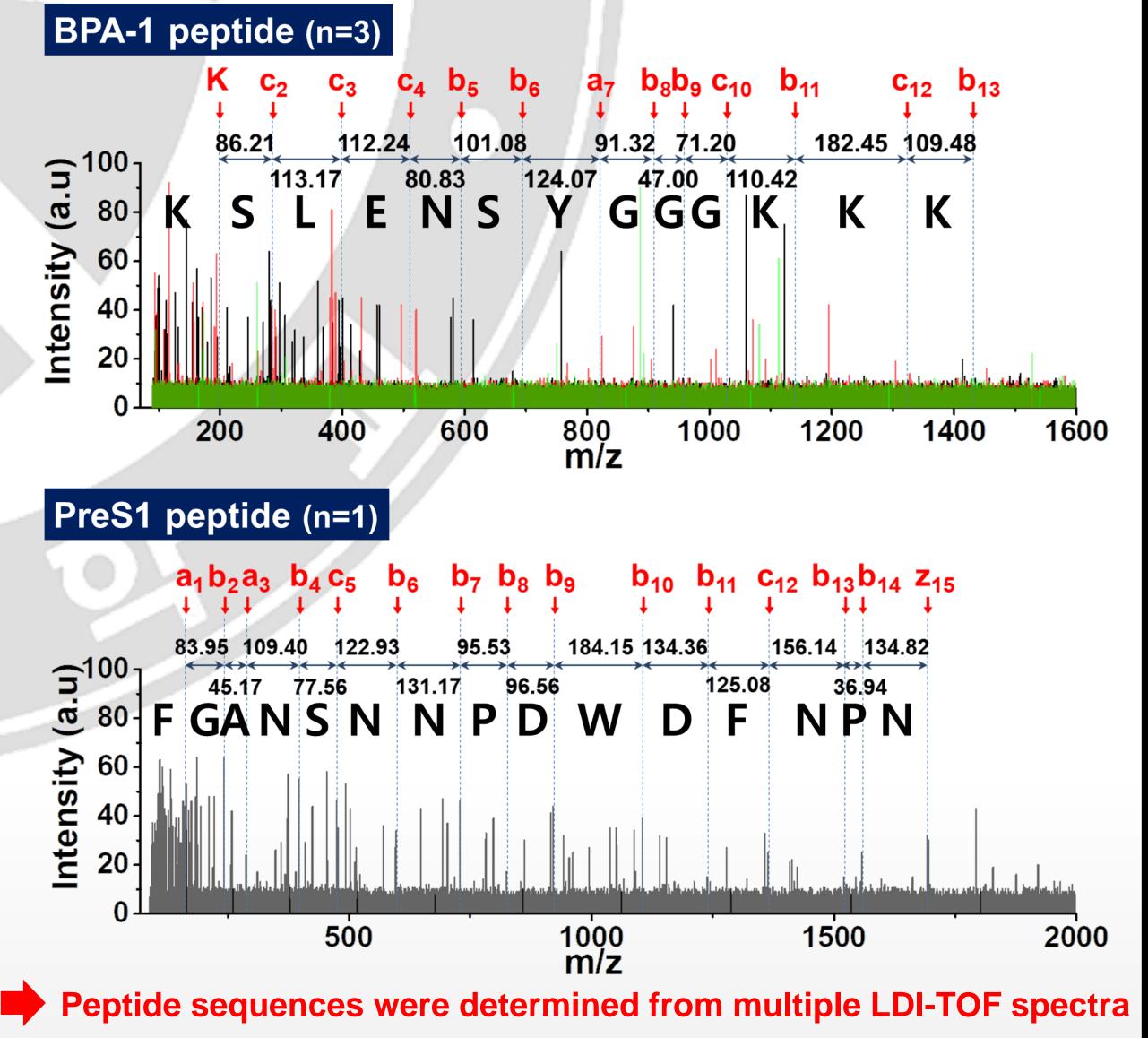
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Introductions Objective Synthesis and apply TiO₂ nanowire target chip to LDI-TOF mass spectrometry for peptide sequencing Laser desorption/ionization time-of-flight (LDI-TOF) - Advantages: Time of flight mass analyzer Short analysis time, high sensitivity, Detector Ion beam low sample consumption Peptide sequencing using TiO₂ nanowire chip - Fragmentation of a peptide by photocatalytic decomposition with UV radiation peptide on TiO₂ nanowires **Before UV treatment** TiO₂ nanowires from **Dropping peptide** Fragmentation of peptide 200 400 600 800 1000 1200 on TiO₂ nanowires by UV radiation (254 nm) wet-corrosion process Fragmentation of peptide by UV radiation (254 nm) TiO₂ NW with photocatalytic activity was used to peptide sequencing

Characterization and Optimization Synthesis of TiO₂ nanowire target chip - Synthesis by mild hydrothermal (wet-corrosion) process - Porous structure was generated after alkali treatment **10 M KOH** H₂O 48 h TiO₂ nanowires from **Polished Ti plate** Water treated TA (TAW) Alkali treated Ti (TA) wet-corrosion process (TAWH) < TiO₂ nanowire synthesis process and SEM images of each process > Optimization of UV radiation time for fragmentation - GHP9 (GHPQGKKKK) peptide was fragmented on TiO2 nanowire plate with UV for different exposure time (30, 60, 120 s) UV treatment (30 s) **Before UV treatment UV treatment (60 s)** UV treatment (120 s) \leftarrow [PQGK+H]⁺ = 430.04 GHPQ-COOH-NH₂+Na]⁺ $[GH-COOH-NH+K]^+ = 1047.23$ Optimized UV exposure time for sequencing was determined to 30 sec



Sequences of three different HPQ peptides were completely determined



Conclusions

- TiO₂ nanowire target chip with photocatalytic activity was prepared using wet-corrosion process
- Amino acid sequences of three different peptides with same molecular weight were clearly analyzed
- GHP9, BPA-1, PreS1 peptide were analyzed from mass peaks from single or three LDI-TOF MS spectra