■1.25M NH4OH

—Flowrate

Zirconium(IV) IMAC-based enrichment for mass spectrometry driven phosphoproteomics

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Introduction

Phosphopeptide enrichment is an essential step in large-scale, quantitative phosphoproteomics studies by mass spectrometry.

Several phosphopeptide affinity enrichment techniques exist, such as Immobilized Metal ion Affinity Chromatography (IMAC) and Metal Oxide Affinity Chromatography (MOAC).

We optimized sample loading conditions and compared Zirconium (IV) IMAC (Zr-IMAC) magnetic microparticles to more commonly used Titanium (IV) IMAC (Ti-IMAC) magnetic microparticles and Iron (III) IMAC (Fe-IMAC LC) for phosphopeptide enrichment from complex protein samples prior LC-MS/MS.

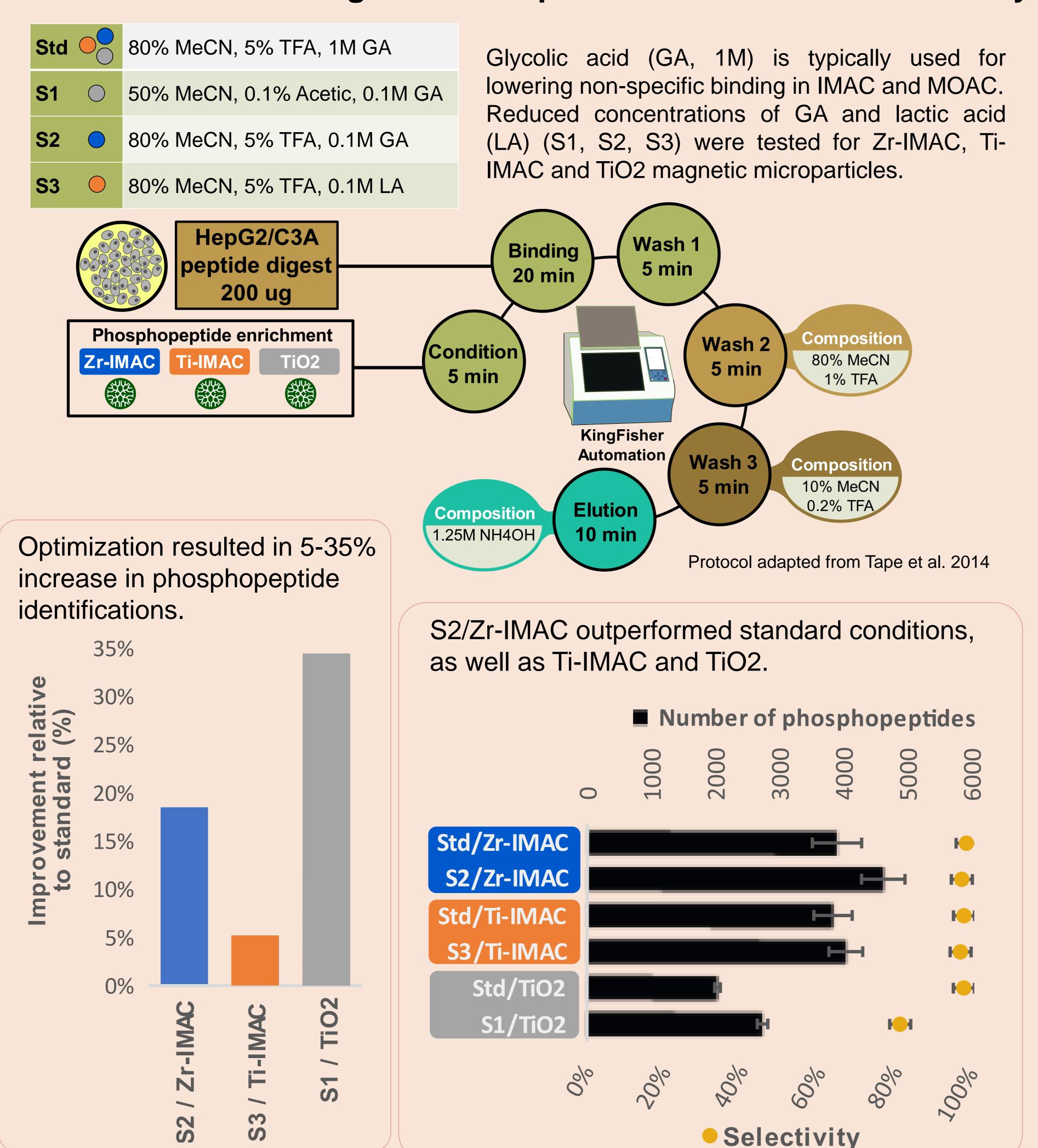
Aims

- To improve phosphopeptide enrichment efficiency of zirconium and titatium based magnetic microparticles.
- Use hydroxy acids to improve phosphopeptide binding to Zr-IMAC, Ti-IMAC and TiO2 magnetic microparticles.
- Assess Zr-IMAC performance in large scale phosphoproteomics experiments.
- Compare Zr-IMAC with wellknown enrichment methods for phosphopeptide enrichment.

Conclusions

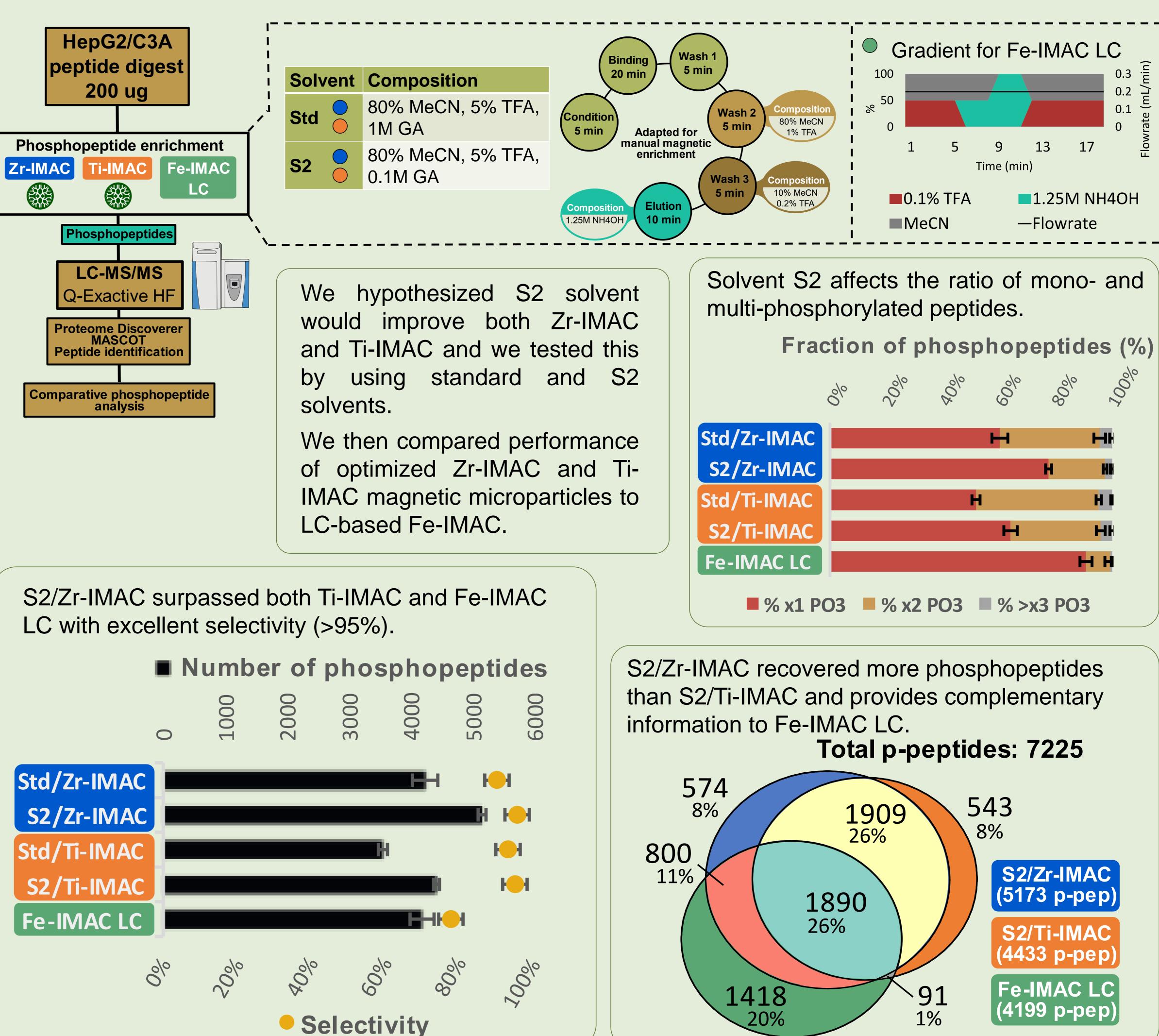
- Zr-IMAC magnetic microparticles selectively and efficiently captures phosphopeptides.
- Optimized Zr-IMAC outperforms more popular methods like Ti-IMAC and Fe-IMAC.
- Optimized Zr-IMAC showed largest complementarity with Fe-IMAC LC.

Hydroxy acid concentration affects phosphopeptide binding to Zr-IMAC/Ti-IMAC/TiO2 magnetic microparticles but not their selectivity



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Optimized Zr-IMAC outperforms Ti-IMAC magnetic microparticles and Fe-IMAC LC



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S2/Zr-IMAC

(5173 p-pep)

