# New methodology to monitor the oxidation of MET due to LC separation

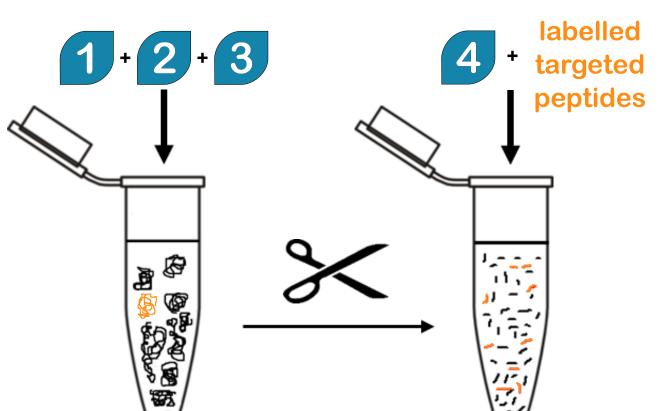
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# Framework

In the context of biomarker discovery and their absolute quantification in complex samples, a standardization strategy aiming to control the entire sample preparation before LC-MS analysis would be extremely valuable. Our approach involves the design of a kit containing a chimeric protein and different levels of its heavy peptides spiked at opportune moment during sample processing.



Chimeric protein with two domains (<sup>15</sup>N)

# **Oxidation sensor**

SPASGIP

Among the peptides included in the protein, one containing a methionine is inserted.

> Why? To monitor artifactual methionine oxidation induced by the whole sample preparation process

**EMSGSPASGIPVK** used as a standard  $\longrightarrow$  conditions to be a good standard:

 $\bigcirc$  stable

O low percentage of oxidation at the moment of the spike

université

**Tryptic peptides (heavy \*\*\*)** 

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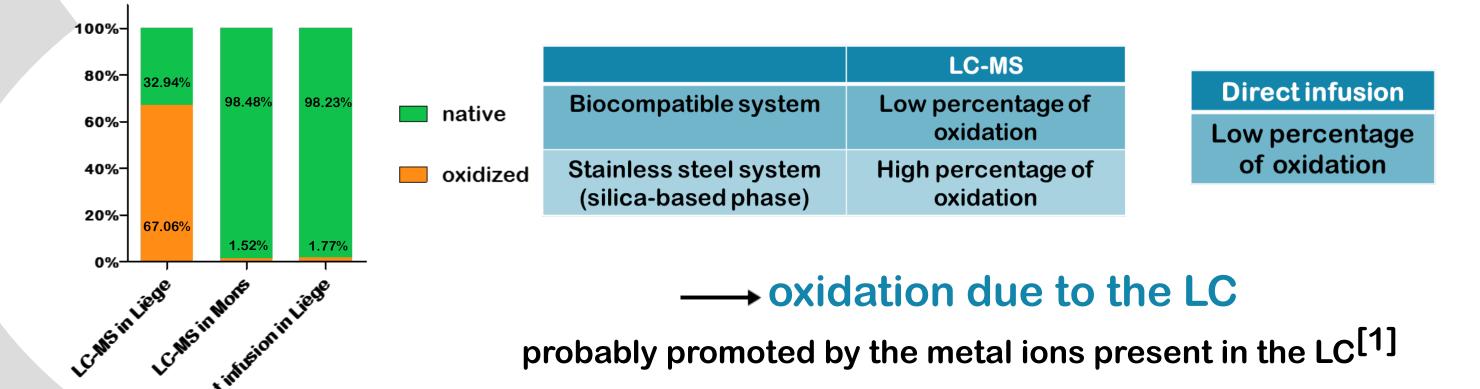
flanking sequences (heavy \*)

**Tryptic peptides with** 

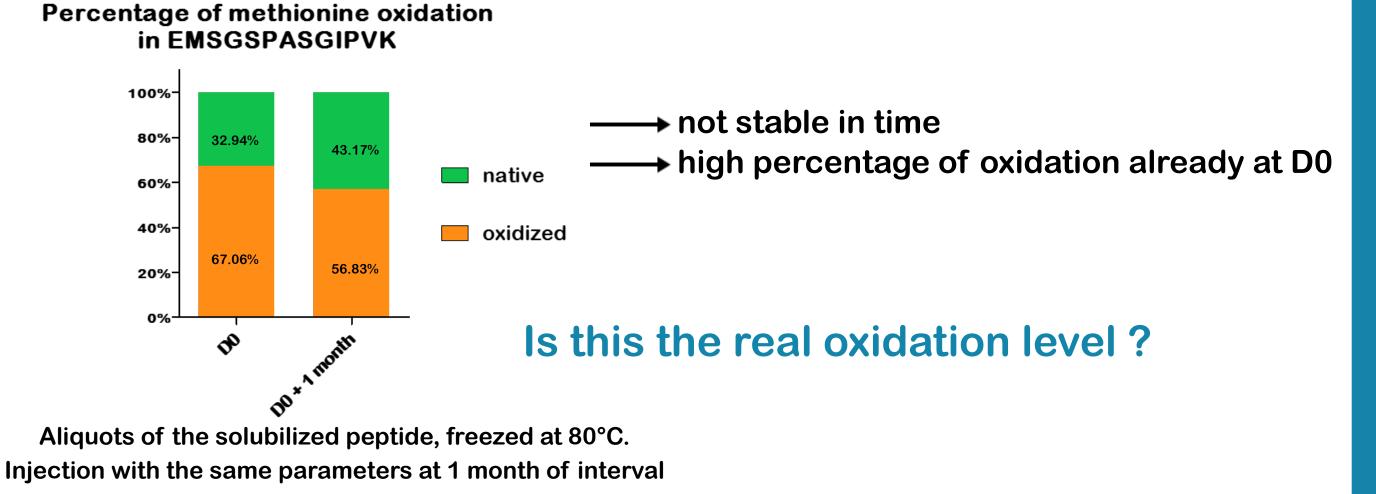
# Is this a good standard?

**First results** 

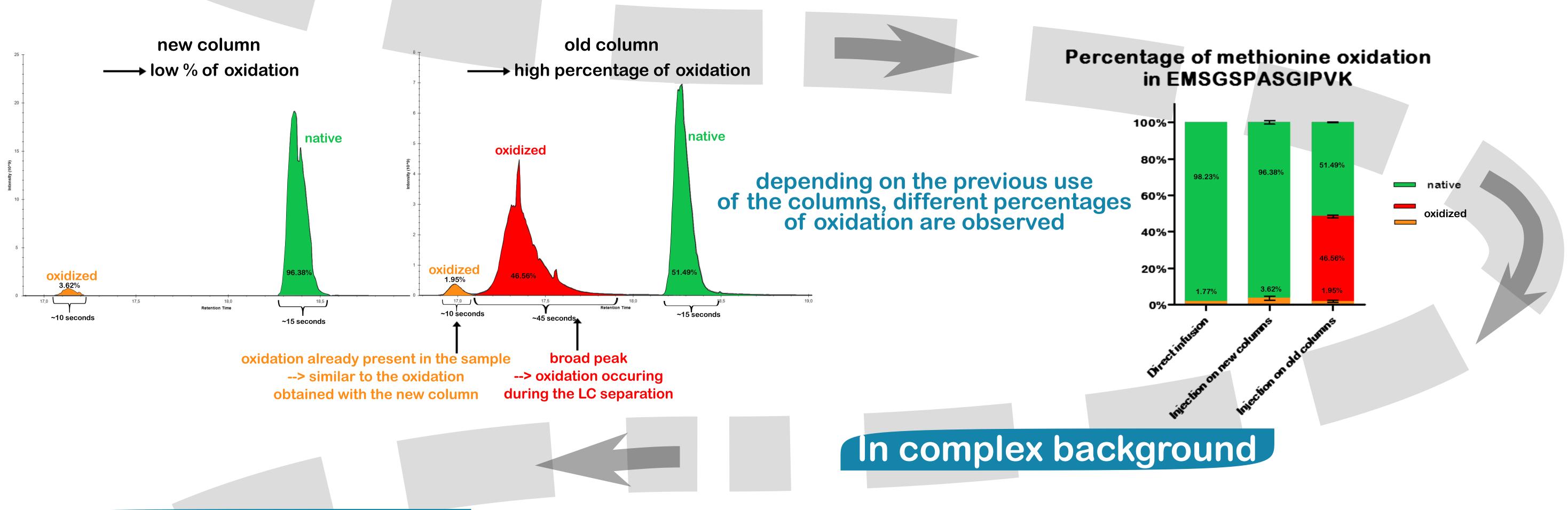
Comparison of percentage of oxidation of EMSGSPASGIPVK between LC-MS in Liège (stainless steel system), LC-MS in Mons (biocompatible system) and direct infusion in Liège



Injection on M-Class system coupled with QExactive<sup>TM</sup> Hybrid Quadrupole-Orbitrap<sup>TM</sup>



### **Unexpected results**



Percentage of methionine oxidation in plasma peptides: set of oxidized and

# **Conclusions/perspectives**

Difference in percentage of oxidation between old and new columns (lifetime ? nature of the previous injection ?)

Peptide with methionine introduced in the kit: • will allow to monitor the degree of oxidation of peptides present in the sample studied • will allow to evaluate the LC system

Strategy to reduce the oxidation within the columns: • chelating agents in the recommended pH range of the LC columns ○ washing

O competitivity between our peptides and another compound more easily oxidized than the peptides

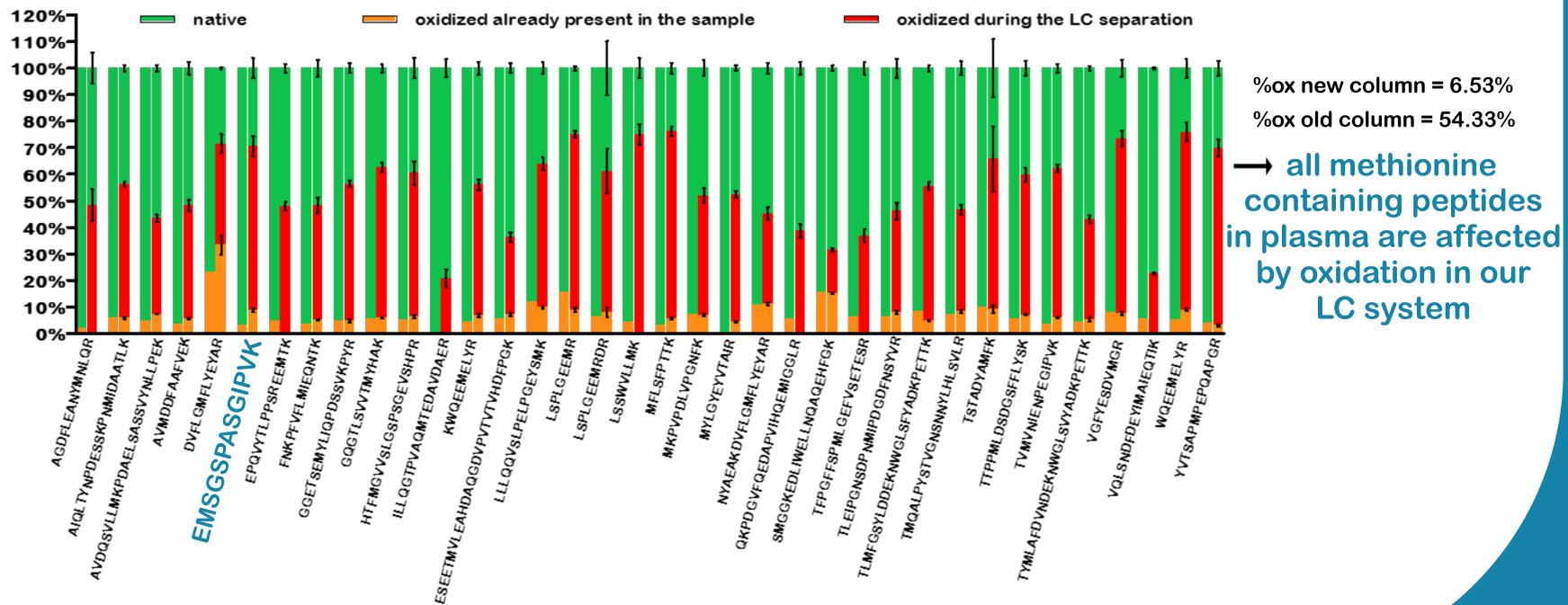
Research on how metal ions can catalyze the oxidation of methionine





#### non-oxidized peptides including our target peptide

left: new column; right: old column



[1] L. Zang et al., "Residual metals cause variability in methionine oxidation measurements in protein pharmaceuticals using LC-UV/MS peptide mapping," Journal of Chromatography B, vol. 895–896, pp. 71–76, May 2012.