

sanofi

Protein vs peptide immunocapture: the case study of the quantitation of total sBCMA by IC-LC-MS/MS

> <u>Alexandra Tavernier</u>, Johanna Paris and Sandrine Descloux

Translational Medicine and Early Development, Biomarkers and Clinical Bioanalyses, Montpellier, Sanofi

EBF, Barcelona, November 15th, 2023

Introduction

Case study: Quantification of sBCMA

General backgroundWhy quantifying sBCMA?

- Analytical challenges

Anti-protein vs anti-peptide immunocapture Development of the two approaches Comparison of both approaches: pro and cons

Quantification of a biomarker by IC-LC-MS/MS

Context of Use (CoU): method validation results

Conclusion and perspectives

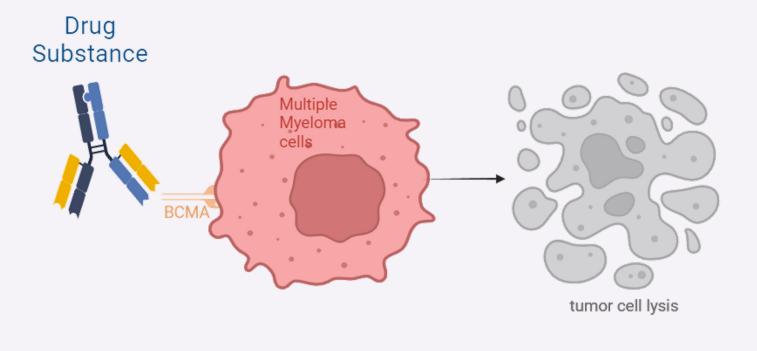
General Background

Overexpression of BCMA in MM patients

Multiple Myeloma (MM) is a disease characterized by a clonal expansion of plasma cells in Bone Marrow



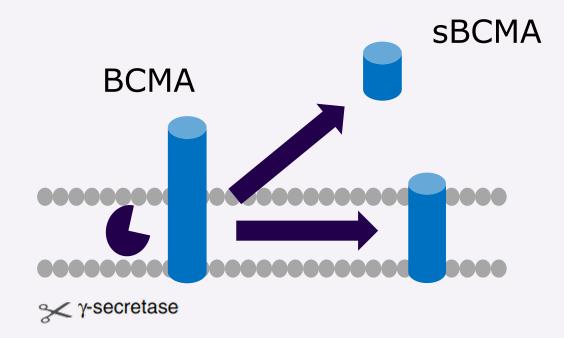
Scientific Animations



- BCMA overexpressed in Multiple Myeloma cells
- BCMA-targeted therapies
- BCMA expression is usually assessed by flow cytometry or immunochemistry

Why quantifying sBCMA?

Soluble B-cell maturation antigen



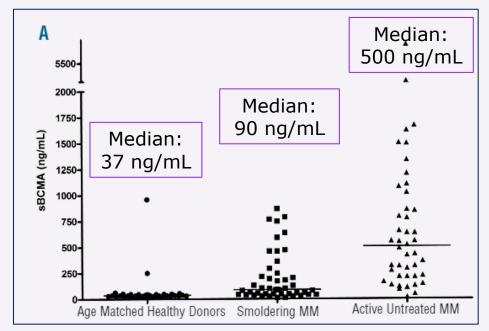
Gamma-secretase mediated shedding

1- sBCMA as a biomarker

Concentrations of sBCMA are high in MM patients, correlated with the proportion of plasma cells in bone marrow biopsies, as well as related to disease progression and treatment efficacy.

Short biomarker half-life (24-36h).

Less invasive than BMA Analysis.



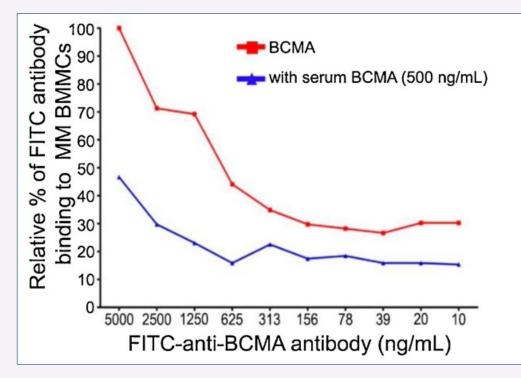
sBCMA levels Ghermezi et al, 2017

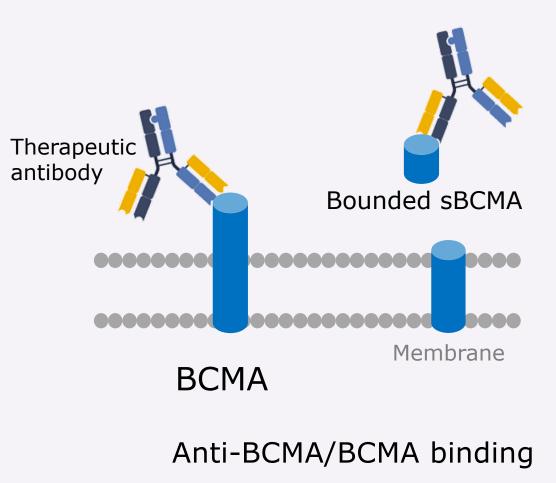
Why quantifying sBCMA?

Off target of the drug substance

2- Interference of sBCMA in treatment

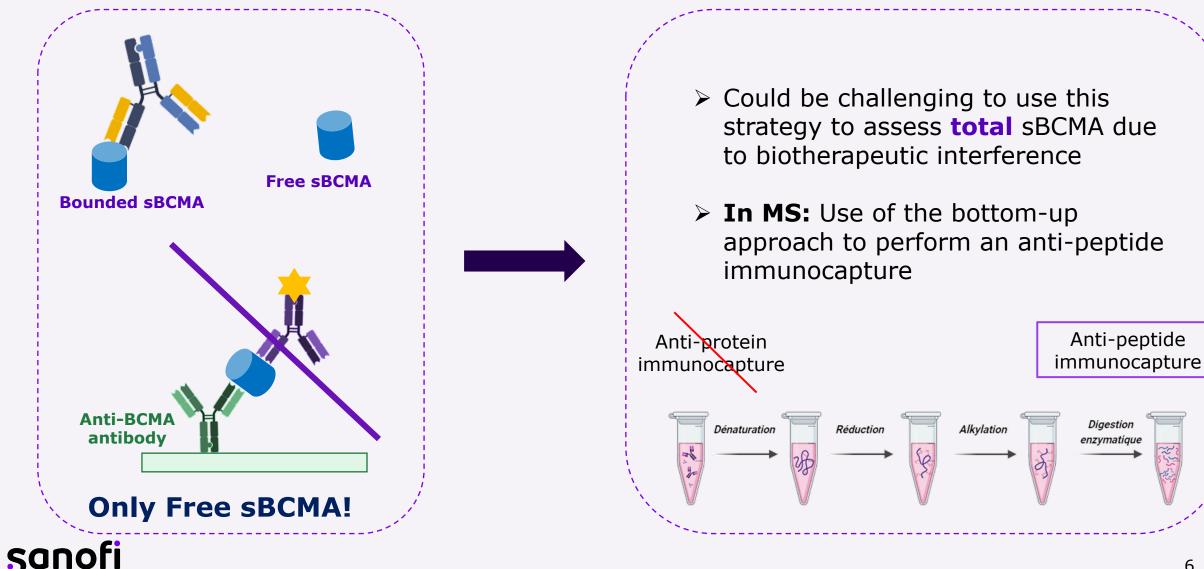
Reduces the efficacy of the treatment





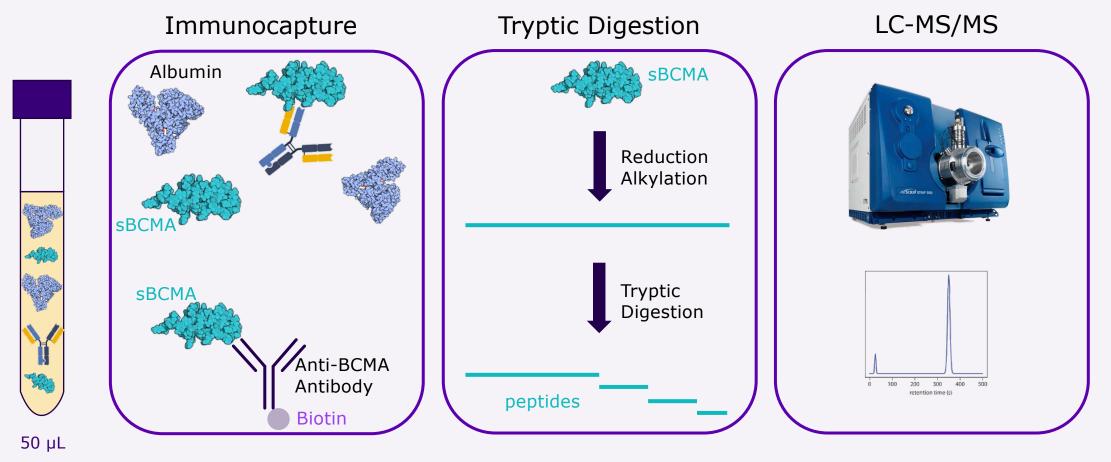
sBCMA interference, Chen et al, 2019

sBCMA Quantification strategies



Hybrid LC-MS/MS Strategy 1 – Protein Capture

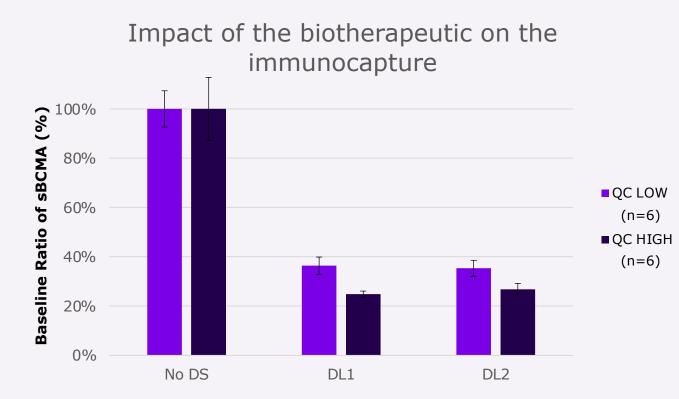
Capture with anti-protein antibody



Competition expected between the capture antibody and the drug

Hybrid LC-MS/MS Strategy 1 – Protein Capture

Drug substance (DS) interference



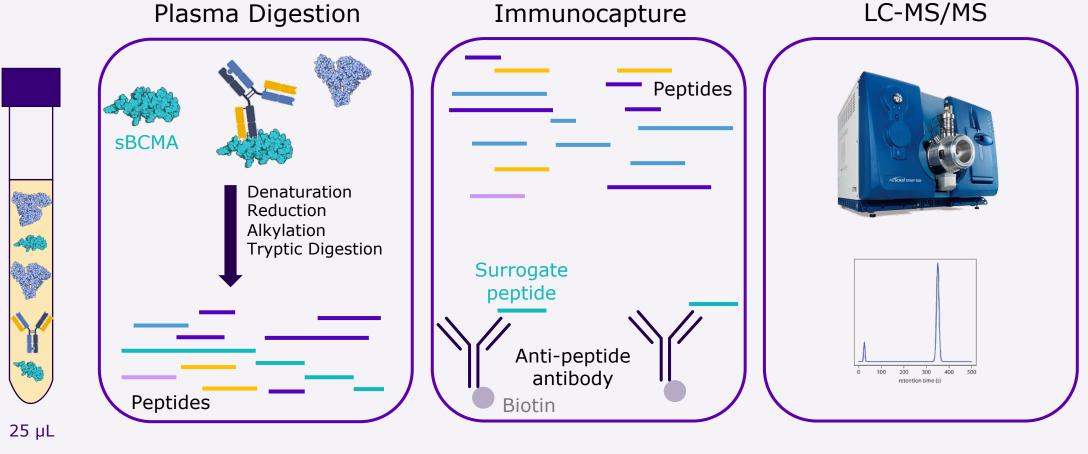
- Immunocapture recovery
- Spiking of two dose level (DL) concentrations of the BCMA-targeting drug substance (DS)
- QC LOW: endogenous level (~25 ng/mL)
- QC HIGH: Spiked at 700 ng/mL sBCMA
- Incubation 4h at 37°C

Strong Interference of the drug substance using this approach



Hybrid LC-MS/MS Strategy 2 – Peptide Capture

Capture with anti-peptide antibody

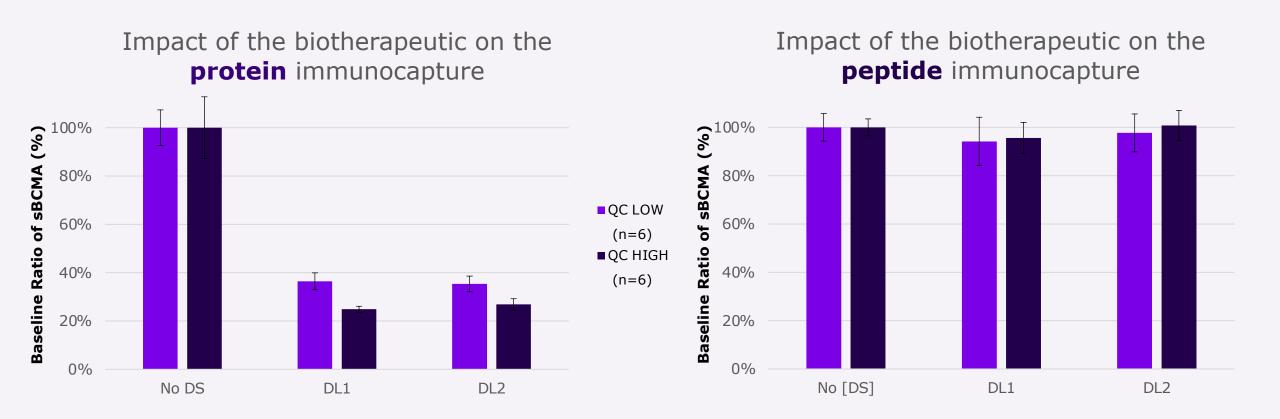


Total sBCMA ?



Comparison of the two approaches

Drug substance (DS) interference



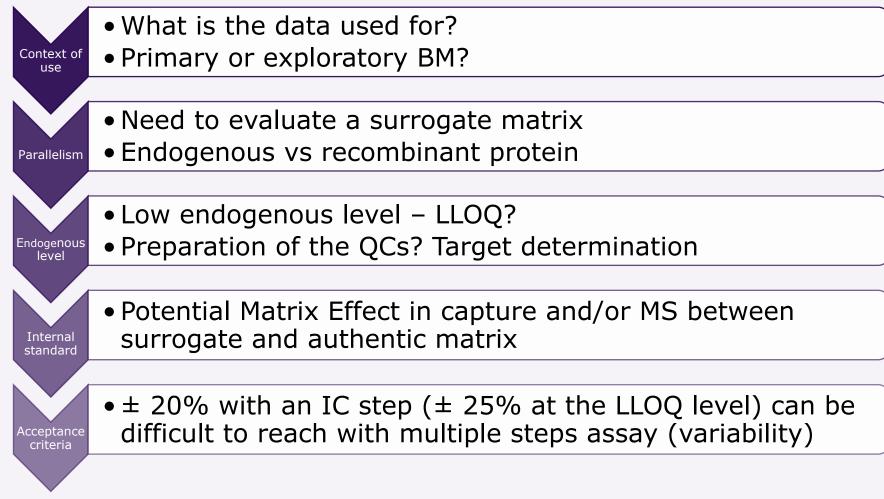
Elimination of the drug substance interference

Comparison of the 2 strategies

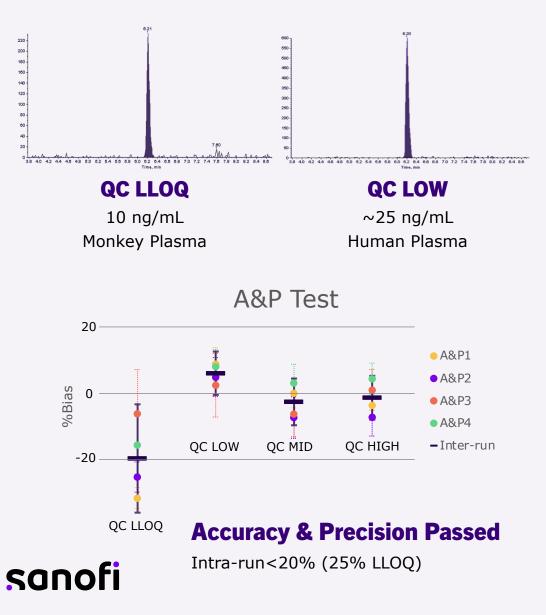
Method development

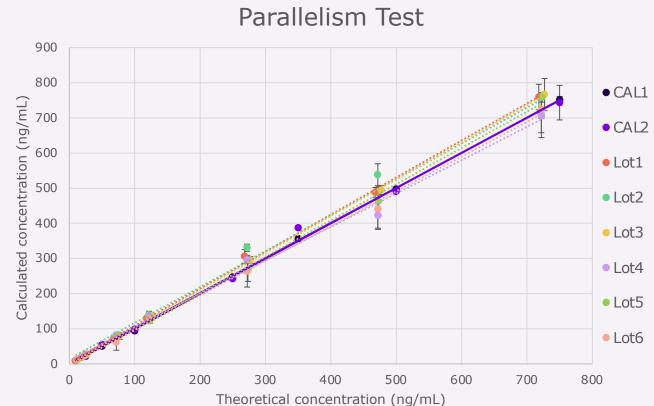
		Anti-protein Approach		Anti-peptide Approach	
Critical reagent Reference standard	Capture reagent		Commercial		Custom made immunization time consuming
	Internal standard	Heavy protein: Unavailable and/or expensive		Flanked heavy peptide: Easy and cheap to produce	
Process	Digestion	Usual protocol		Plasma digestion: Optimization required Denaturation step added	
	Automated Immunocapture		Streptavidin tips		Magnetic Beads Volumes

Challenges of a Biomarker quantification by LC/MS-MS



Bioanalytical method validation





Parallelism Test Passed

2 calibration curves (CAL1 & CAL2 in monkey plasma), 6 human plasma lots tested (spiking of 50, 100, 250, 450, 700 ng/ml of recombinant sBCMA, dilution by two in monkey plasma)

Conclusion

The method was successfully qualified (Context of Use validation)

Interference with drug substance was eliminated with the anti-peptide immunocapture

This bioanalytical strategy can be used:

- In BM quantification: Biotherapeutic Interference
- In PK measurements: ADA Interference
- Needs to be **anticipated** (delays and costs)

Thank you

Johanna Paris • Principal Scientist

Sylvie Bethegnies • Bioanalyst

Sandrine Descloux • Head of BCB Physchem analyses

Stéphane Muccio, Carole Legeay, Catherine Aubert, Mathieu Cyrille and Béatrice Pradeilles

Olivier Fedeli • Head of BCB Montpellier

Thanks to Valerie Boutet, Nathalie Macé, Jacqueline Courta and Hongfang Wang

sanofi

Thank you for your attention



Mass Spectrometry Team in Montpellier, France 15