

## Considerations for the Application of Multi-Attribute-Method (MAM) by Mass Spectrometry for QC Release and Stability Testing of Biopharmaceuticals

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#### OUTLINE

Considerations for the Application of Multi-Attribute-Method (MAM) by Mass Spectrometry for QC Release and Stability Testing of Biopharmaceuticals

- EFPIA topic team "MAM as QC tool"
- MAM by LC-MS peptide mapping as QC tool
- **enefits and challenges**
- Regulatory pathways
- Supporting elements



### EFPIA TOPIC TEAM "MAM AS QC TOOL"

## Why this initiative?

- Multi-attribute-method (MAM) by mass spectrometry is well established across the industry in non-GMP environments for product and process characterization purposes
- The majority of pharmaceutical companies and many instrument providers are currently working on the extension of MAM to QC labs
- The use of MAM for lot release and stability testing according to GMP is not well established across the industry due to:
  - ongoing evolution and alignment of best practices
  - complexity of method (sample preparation, instrumentation, data analysis)
  - limited experience with filing of MAM as a QC tool
  - regulatory unfamiliarity with MAM as QC tool



## **Mission and vision**

- Team of 25 representatives from 17 pharmaceutical companies
- Founded in March 2021 under the umbrella of EFPIA MQEG<sup>1</sup> to promote

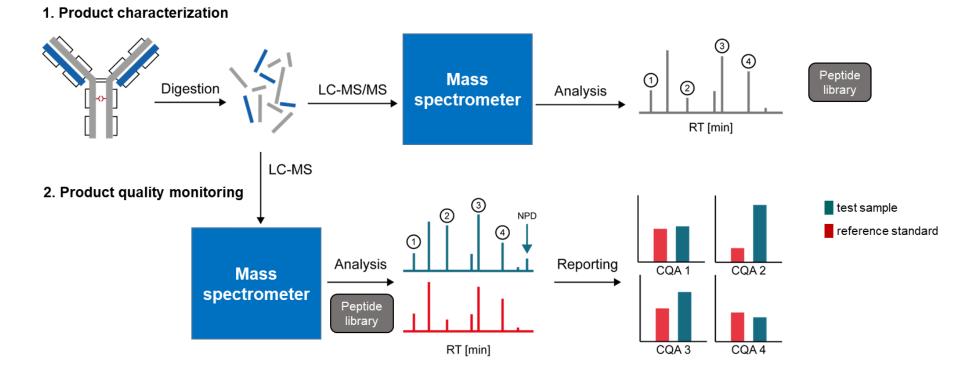
Global acceptance of MAM addressing multiple product quality attributes in a single method for QC release and stability, replacing conventional QC methods (e.g. purity / identity)

- $\rightarrow$  Share and align on best practices across the industry
- → Promote & encourage regulatory filing of MAM for lot release and stability testing under GMP
- $\rightarrow$  Reduce regulatory unfamiliarity and obtain acceptance by health authorities



#### **TECHNICAL CONTEXT**

## **Prototypical MAM by LC-MS peptide mapping workflow**



Targeted monitoring of critical quality attributes + 'New Peak Detection' (NPD) are required to establish MAM by LC-MS peptide mapping as purity assay in a QC environment.



#### **TECHNICAL CONTEXT**

# MAM by LC-MS peptide mapping has the proven <sup>1</sup> capability to replace multiple conventional HPLC / CE-based QC methods

<b>Quality Attribute</b>	<b>Conventional method</b>	<b>Replacement method</b>	
Charge variants	IEX, cIEF, CZE		
Fragments	rCE-SDS	MAM by LC-MS peptide	
Glycans	2-AB HILIC, HPAEC PAD	mapping	
Oxidation	RPC, HIC, peptide mapping LC-UV		
Identity	ELISA, peptide mapping LC-UV		

- The technology is well-advanced with instruments and software solutions being developed from several vendors allowing routine use in a GMP environment.
- Implementation of MAM is supported by established and draft guidelines (e.g. ICH Q2, ICH Q6B, ICH Q14) and will facilitate advanced control strategies in line with ICH Q8.



## Improved quality control testing and shortened development timelines through enhanced product and process understanding

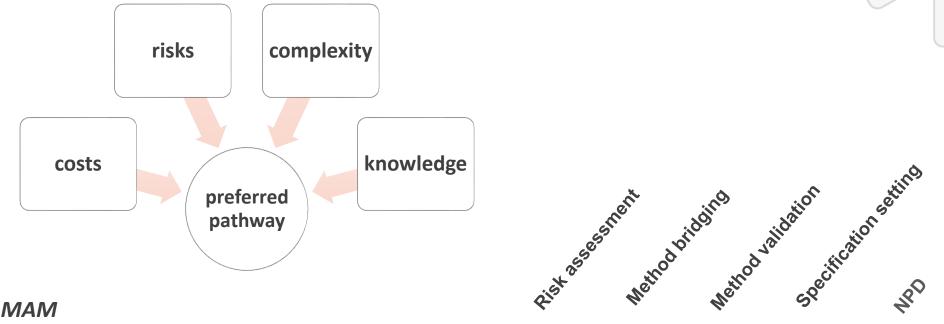
- Provides quantitative information on individual site-specific CQAs therefore enabling more specific control of the safety and efficacy of the drug
- Increase speed by leveraging MAM as platform method with a potential for automation
- De-risks accelerated development by retrospective assessment of newly identified (p)CQAs using previous data sets

## MAM is so far not widely accepted for lot release and stability testing under GMP due to regulatory unfamiliarity & potential business risks

- Limited experience with filing MAM as QC method, replacing conventional methods
- Diverse and unclear regulatory landscape as potential business risks
- Increased effort and risk by parallel testing using MAM & conventional methods
- *Limited experience to validate New Peak Detection (NPD) and set appropriate specifications*



#### **REGULATORY PATHWAYS & SUPPORTING ELEMENTS**



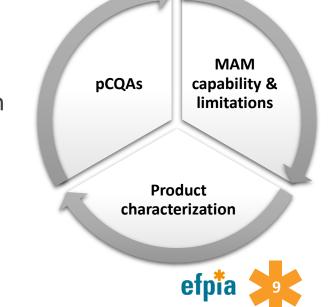
Introduction of MAM

Prior to FIH studies – instead of conventional methods	yes	no	yes	yes	yes
After FIH / prior to registration – in addition to conventional methods	no	no	yes	yes	no
During development / as LCM activity - replace conventional methods		yes	yes	yes	yes



## **Risk assessment**

- "Evaluate the capacity and performance of MAM in the context of the CQAs of the candidate product and its overall control strategy"<sup>1</sup>, which requires thorough understanding of the
  - 1. capabilities & limitations of MAM by LC-MS peptide mapping
  - 2. pCQAs of the product obtained by structure elucidation and forced degradation studies
- typical limitations of MAMs by LC-MS peptide mapping include
  - 1. clipping site (degradation) = clipping site (sample preparation)
  - 2. peptide fragments are too small to be retained on the LC column
  - 3. bottom-up approach: modification on peptide vs intact level
  - 4. potential risk of sample preparation-induced artefacts





- bridging data package as for any other conventional method, i.e. analysis of
  - clinical batches  $\rightarrow$  link to clinical experience / safety and efficacy
  - stability samples incl. forcibly-degraded samples  $\rightarrow$  coverage of all relevant pCQAs
  - IPC samples  $\rightarrow$  absence of matrix interferences (if relevant)
- reportable results for MAM and conventional methods will be different
  - not meaningful to compare absolute values, only trends and rate of evolution of the CQA during stability (long term, accelerated conditions) should be compared
- leverage the performance characteristics as defined in the Analytical Target Profile (ATP)

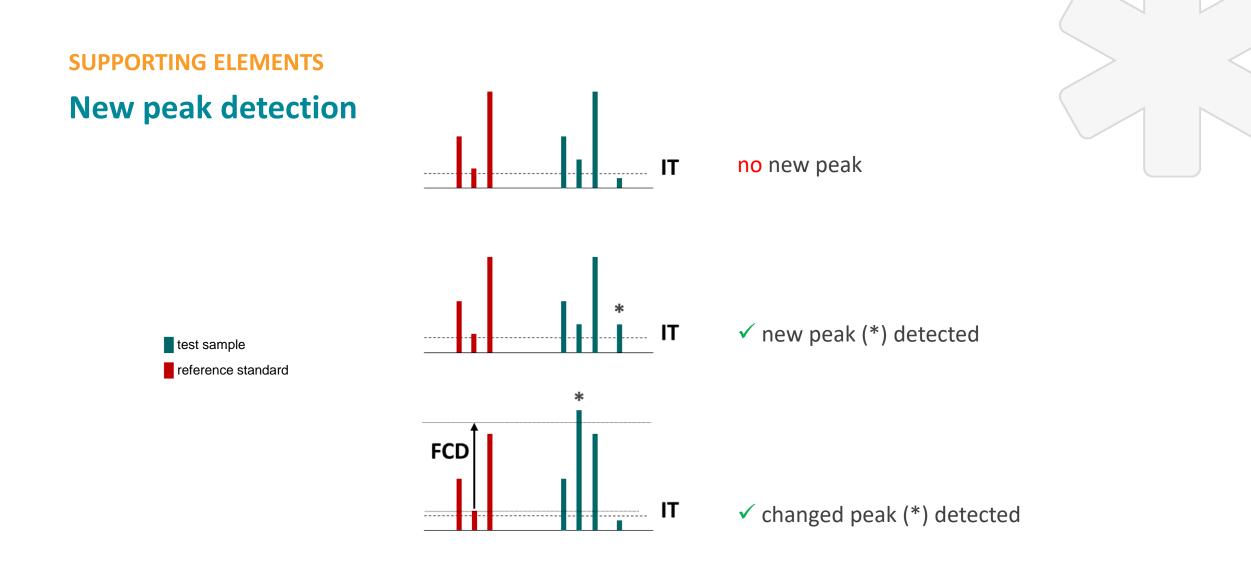


#### **SUPPORTING ELEMENTS**

## **Method validation**

- targeted monitoring according to ICH Q2(R1) quantitative testing for impurities
  - accuracy, precision (repeatability, intermediate precision), specificity, quantitation limit, linearity and range
- NPD parameters (if applicable)
  - fold-change & intensity threshold, mass & retention time windows
- *leverage MAM platform in line with ICH Q14, i.e. consider to apply* 
  - platform robustness data to streamline product specific validation
  - prior knowledge e.g. same peptide / same modification
  - risk-based approaches e.g. similar peptides / same modification
  - performance requirements as defined in the ATP



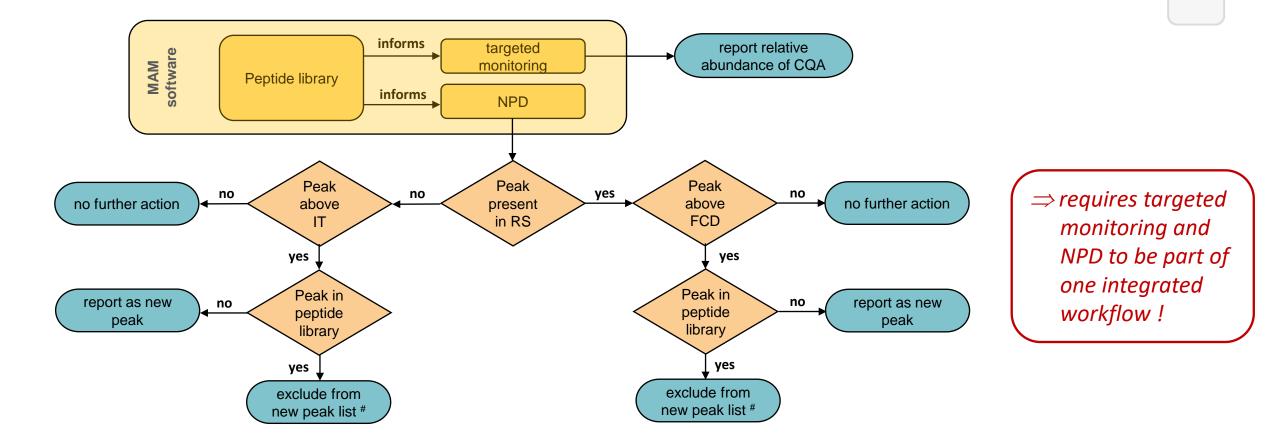


IT: intensity threshold, minimum signal threshold FCD: fold-change detection threshold



#### **SUPPORTING ELEMENTS**

## **New peak detection**



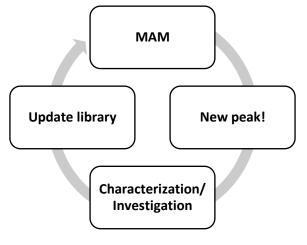
- RS: product specific reference standard
- IT: intensity threshold, minimum signal threshold
- FCD: fold-change detection threshold
- # selected peaks / product variants may be trended to support internal process consistency monitoring



#### **SUPPORTING ELEMENTS**

## **Specification setting**

- *for the targeted monitoring according to ICH Q6B as for conventional methods* 
  - specification limits only for CQAs
  - limits informed by clinical experience, criticality assessment of quality attributes and performance characteristics of the MAM
- for NPD in a stage-appropriate manner as library and NPD parameters are expected to evolve
  clinical development
  - lower warning limit  $\rightarrow$  characterize / update library
  - higher action limit  $\rightarrow$  OOS investigation <u>commercial</u>
  - detection of unknown peak above validated NPD thresholds  $\rightarrow$  OOS investigation





- Replacement of conventional methods with MAM by LC-MS enables improved CQA-centric quality control testing in line with QbD principles
- Depending on the regulatory pathway for the introduction of MAM in a QC lab, different elements such as risk assessment, method bridging and NPD will be required
- The development and validation of a robust NPD workflow is considered a pre-requisite for the use of MAM by LC-MS as a purity assay and the successful replacement of conventional methods
- Method validation and specification setting for the target monitoring should follow established regulatory guidelines, such as ICH Q2 / ICH Q6B and will benefit from upcoming ICH Q14
- Introduction of MAM by LC-MS as a QC method prior to FIH studies is expected to reduce complexity but requires frontloading in terms of early product characterization studies



#### ACKNOWLEDGEMENT

### **Current members of the EFPIA topic team «MAM as a QC tool»**

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