

cobas®

Roche

Fueling a new era of innovation
in mass spectrometry
cobas® Mass Spec solution

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Please contact your local representative to confirm the current availability of the cobas[®] i 601 analytical unit, specific Ionify[®] reagents, and their regulatory status in your region.

Chapter 1

The challenge: mass spectrometry today

The medical and clinical need for mass spectrometry has been present for a long time.



From 2008 to 2018, the number of publications related to mass spectrometry tripled¹ and a consistent increase of samples tested using mass spec has been reported in literature for several years.²⁻⁴

The main reason behind this shift is that there are limitations in standard testing methods like immunochemistry and clinical chemistry. For specific patient cohorts, liquid chromatography combined with tandem mass spectrometry (LC-MS/MS) provides better analytical sensitivity, better specificity and/ or accuracy, while being less impacted by matrix effects, interferences or cross-reactivity.⁵

LC-MS/MS has consequently become a widely used technology within clinical reference and referral laboratories worldwide and has also started to be used in some hospitals and regional clinical laboratories.⁴

Mass spec is well established in clinical and academic research and is also used predominantly in the following fields: therapeutic drug monitoring (TDM), urine drug testing (UDT), newborn screening, endocrinology, vitamins, and peptide and protein markers. Mass spec technology is well-known for complex and sophisticated reference methods. Today, mass spec is found in separate, specialized testing sections of the laboratory and there are hurdles that prevent more labs from adopting mass spectrometry in routine operation. Two of the main challenges are that the process can be time-consuming and requires high levels of expertise to be carried out successfully.

Typical weaknesses of existing mass spec solutions were previously described by Vogeser & Zhang in 2018.⁵

- High complexity of instrumentation
- Hardware & software are designed for highly specialized research labs, but not for clinical routine diagnostics
- Lack of automated solutions from sample preparation to sample analysis to result reporting
- Poor integration of mass spec with laboratory information systems
- Operation and troubleshooting of instrumentation requiring highly skilled staff
- Poor standardization across instruments and laboratories
- Long turnaround times which are not suitable for urgent care
- Complex data handling



There are limitations in standard testing methods like immunochemistry and clinical chemistry

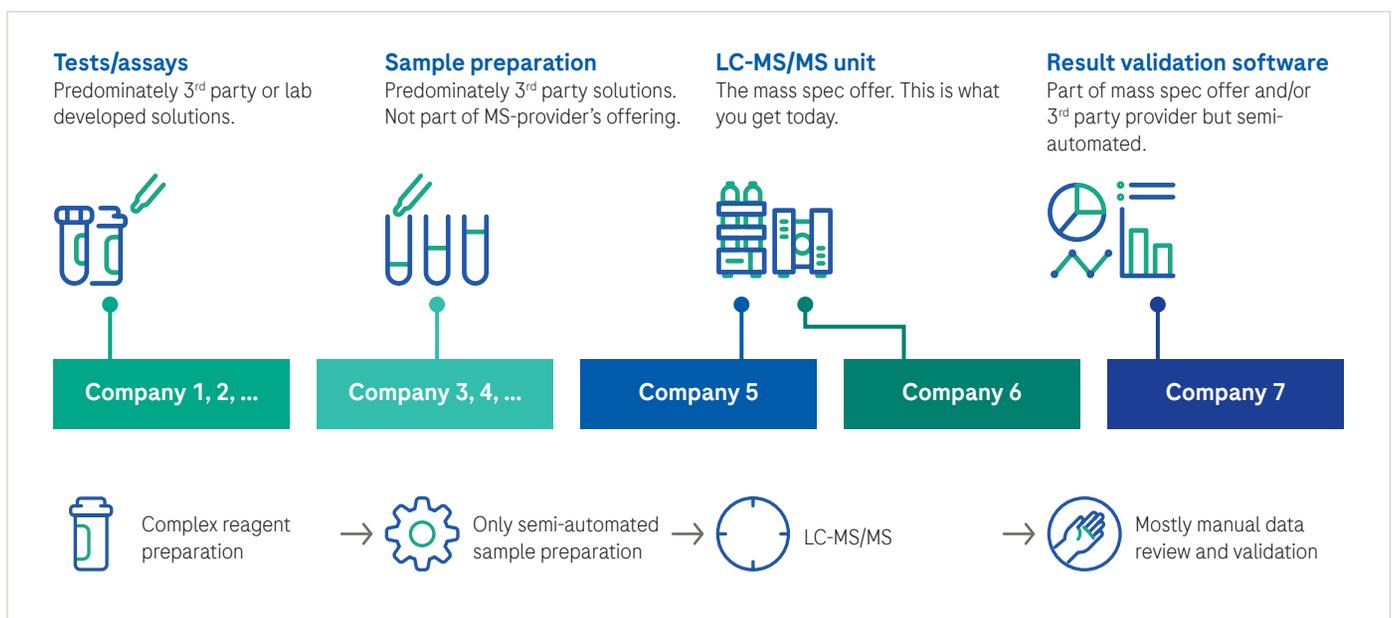
Today's mass spectrometry-based testing often suffers from low throughput and long turn-around times (TAT). Labs typically have to batch samples, setting aside specimens until they have enough to run an analysis. This practice can lead to significant delays in results. Furthermore, running more challenging assays requires lengthy liquid chromatography separation, which severely limits the number of samples that can be processed. Finally, most existing systems lack routine, service-friendly support concepts to quickly address technical issues and minimize downtime.

Due to the lack of a total solution provider, laboratories typically need to interact with numerous different companies to obtain everything required for mass spec. In the most extreme cases, labs may need to deal with up to 7 vendors to provide all the necessary materials and services need to offer a mass spec based testing service. See Figure 1.

Different vendors may need to be engaged to deal with the following:

- Commercially available reagents for calibrators & quality control (QC) or reference material for in-house preparation of calibrator and QC material
- Reagents and consumables for sample preparation and mobile phases for the LC system
- Semi-automated sample preparation solutions
- Instrumentation for liquid chromatography (LC) & mass spectrometry (MS), typically sourced from the same provider
- Software solutions for semi-automated result interpretation and validation

Figure 1: A key challenge for mass spec testing today is the lack of a total solution provider



Chapter 2

The vision: the future of mass spectrometry

The vision for the future of mass spec solutions is clear and has been outlined in publications for many years. In 2015, Zhang & Rockwood had the vision of developing mass spectrometry into “a total automation, high throughput, continuous random-access platform.”

The authors envisioned that an “automated mass spectrometry platform can in the future either be a standalone floor model or integrated into the core laboratory function as one of the automated platform along with chemistry analysers, haematology analysers, coagulation analysers and others.”⁶

Stone & Fitzgerald further specified this vision of future clinical mass spectrometry in 2018: “Such a system would have ease of use similar to automated clinical chemistry analysers – random-access workflow, minimal down-time, 24/7 service and support, and validated and ready-to-use reagents and calibrators supplied by the vendor. These systems would not require specialised end-user skills for operation and would have sampling and software that permit integration to track systems along with ASTM/HL7 interfaces to laboratory information systems.”⁷



As a logical consequence of the envisioned automated solutions which would be integrated into routine laboratory track systems, Greaves et al. predicted in 2019 that there would be an increase in the number of clinical laboratories adopting mass spectrometry based testing, especially in the fields of endocrinology and drug testing.¹



Zhang & Rockwood had the vision of developing mass spectrometry into “a total automation, high throughput, continuous random-access platform.”



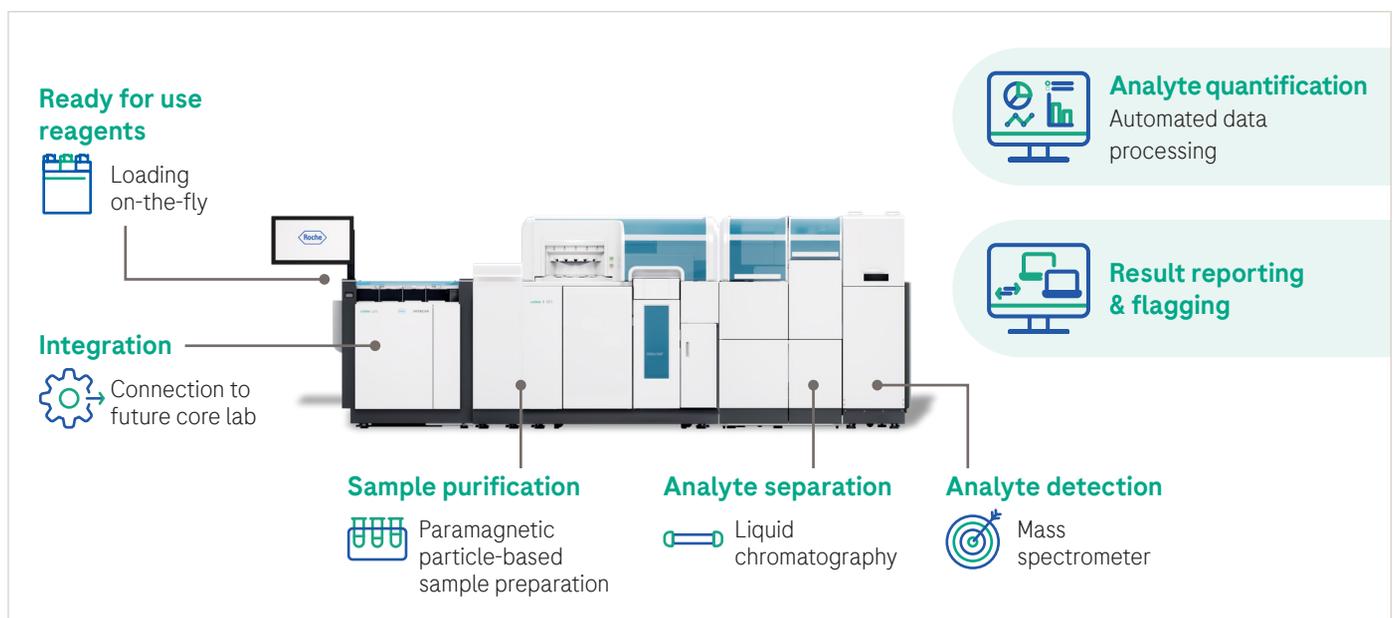
Chapter 3

The answer: Roche vision of cobas® Mass Spec

The aforementioned visions will be made a reality by the **cobas®** Mass Spec solution. Roche Diagnostics is investing in developing a solution for clinical mass spec testing, which is designed to be compliant with IVD regulations, addressing unmet needs and providing answers to the current challenges faced by clinical mass spec operations:

- **Full automation** from sample preparation to result interpretation with random access operation mode
 - **Seamless integration** into clinical chemistry and immunochemistry testing via future **cobas® pro** integrated solutions
 - **A high throughput** of up to 100 tests/ hour to fit the needs of the core laboratory for fast and predictable turnaround times
 - **A broad assay menu** of over 60 analytes planned in two staggered launch waves consolidated on a single platform
- The **cobas®** Mass Spec analyser, which will be called the **cobas® i 601** analytical unit, is designed to transform a complex technology into an integral part of routine clinical testing. It is designed to be a total solution from sample to result for clinical mass spec testing and will include the following, as seen in Figure 2:
- Ready-to-use reagent cassettes with the same design as the well-received **cobas® c** and **cobas® e** reagent packs
 - An automated paramagnetic particle-based sample preparation, ensuring efficient sample purification
 - Fully automated liquid chromatography and mass spectrometry with a **cobas®** look and feel
 - Fully automated data processing, automated result reporting and result flagging to replace manual analyte quantification
 - Integration into **cobas® pro** integrated solutions to provide connection to the central laboratory*

Figure 2: cobas® Mass Spec solution, a sample-to-result total solution for clinical mass spec testing



*At launch, cobas Mass Spec will be in a stand-alone configuration only. All other configurations, including clinical chemistry and immunochemistry analytical units, will be available at a later date.

Chapter 4

The future: cobas® Mass Spec

Here are a few examples of how **cobas®** Mass Spec intends to provide clinical laboratories with a number of benefits:

Fully automated mass spec testing can help to increase efficiency in the laboratory workflow and reduce human errors

cobas® Mass Spec will provide a fully-automated solution from sample preparation to result reporting. After preparing the system, the only action for a lab operator is to load the samples into the analyser. The system will automatically conduct sample preparation, separation and detection, validating the results and finally transferring to the laboratory information system (LIS).

As demonstrated in Figure 3, this will reduce the potential for human error, as well as hands-on time, which in turn will allow the overall efficiency of the testing workflow to be improved.⁶

Convenient and simplified reagent handling can free up staff time

In current mass spec offerings, reagent preparation can be a complex process. Reagents are typically not ready to use (e.g. internal standards and mobile phases need to be prepared). Additionally, appropriate chemistry knowledge and a suitable laboratory is required. With **cobas®** Mass Spec, a streamlined reagent concept will be offered, which will allow fast and easy handling. This will free up time for laboratory staff and will minimise the risk of human error. Product-specific reagents, as well as multianalyte reagents will be provided in easy-to-handle reagent packs or containers, as illustrated in Figure 4.

Figure 4: Reagent handling process will be simplified and more convenient

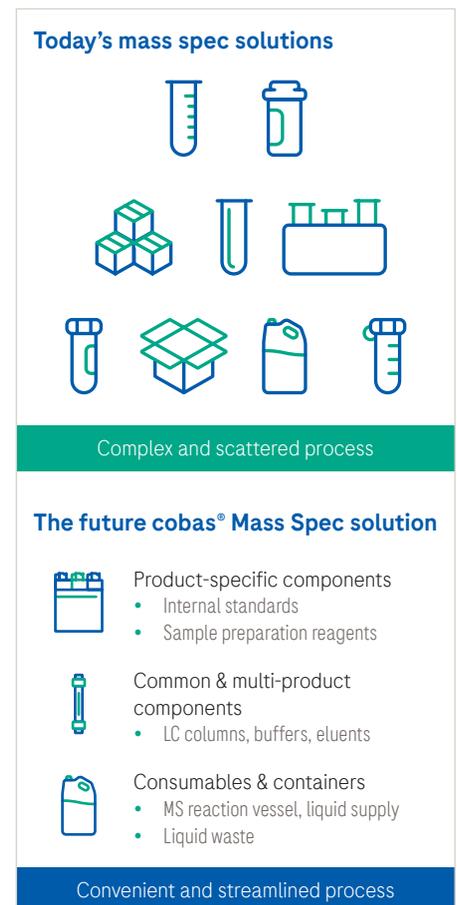
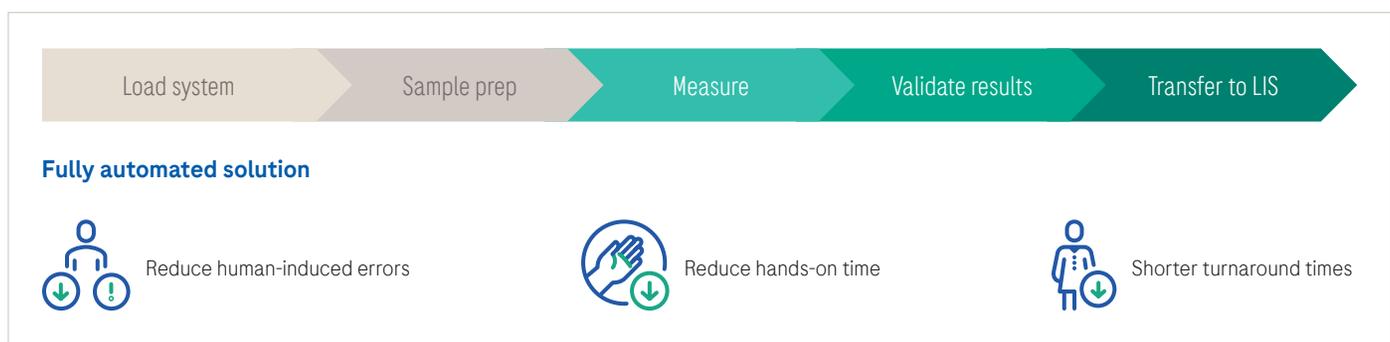


Figure 3: Fully automated mass spec testing saves time and resources



A fully automated IVD solution reduces validation efforts

Currently, to develop and validate LC-MS/MS based methods, a number of different steps need to be considered, and guidelines have to be followed to validate them. These are time-consuming and require high expertise, which can add strain to lab staff.^{8,9}



cobas[®] Mass Spec is designed to be a solution compliant with IVD regulations, covering the workflow as a whole

cobas[®] Mass Spec is designed to be a solution compliant with IVD regulations, covering the workflow as a whole, including the instrumentation, software and reagents. Performance data will be provided in assay method sheets, based on internal testing, as well as both internal and external validation and verification data. Mass spec experts will be relieved from the cumbersome test development process and can focus on more esoteric methods or methods used in fast-changing environments like urine drug testing.

With a random access analyzer patient results can be provided earlier

With **cobas**[®] Mass Spec, random access will be introduced to the world of mass spec testing. Traditionally, users need to change the instrument condition, exchange reagents and columns to run a specific application. When switching to another application, the system has to be re-engineered. This means that the system is currently limited to batch-mode testing which is time-consuming and inefficient.¹⁰

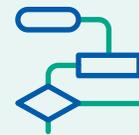


With **cobas**[®] Mass Spec, random access will be introduced to the world of mass spec testing

cobas[®] Mass Spec will allow loading and processing samples as they arrive in the laboratory. No batch mode testing will be required, which will allow both the efficiency and flexibility of the process to increase and this in turn will allow earlier release of patient results.

Automated result validation and reporting simplifies processes

In the current offering in the market, reviewing and releasing results does not only require high expertise but it is also labor-intensive.⁶



cobas[®] Mass Spec will introduce a sophisticated algorithm to automatically process the data

cobas[®] Mass Spec will introduce a sophisticated algorithm to automatically process the data generated from the mass spectrometer. Data will be processed, validated and calculated, and results will be reported directly without visually controlling every single peak. In the case of a flagged result, there will be an alert which signals that the result needs to be reviewed before sending it to the LIS.

A broad variety of analytes in the launch menu will increase the accessibility to mass spec in routine clinical testing

As illustrated in Figure 5, assays for more than 60 analytes are in development and will be launched in a successive waves. The first wave comprises steroids, vitamin D metabolites and TDM tests, including the whole-blood based immunosuppressant drugs. The assays will be provided in 14 ready for use multi-reagent packs.

Standardized and traceable methods are prerequisites for consistent patient results

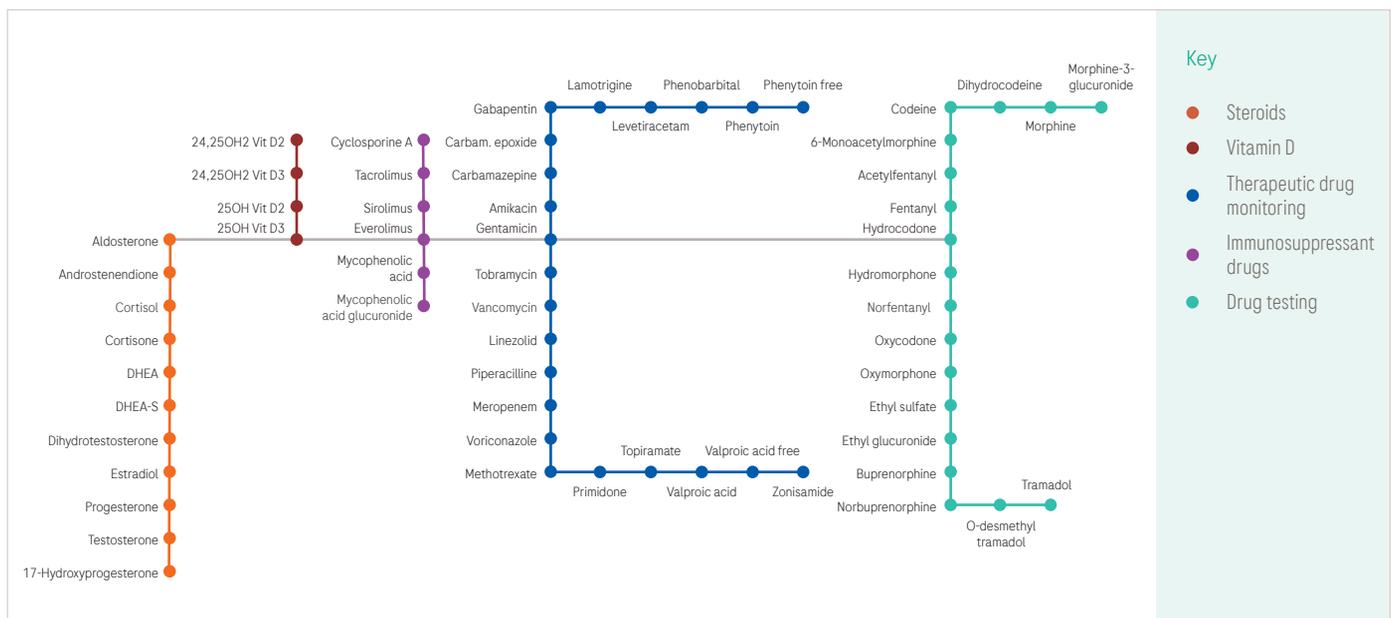
A significant challenge in today's mass spectrometry based clinical testing is the lack of assay standardization. Current MS based tests are almost exclusively Laboratory Developed Tests (LDTs). Because there are no total solution providers, inter-laboratory variability in test results may be high. This is a result of the numerous instrument configurations, pre-analytic and analytic methods, data acquisition conditions, and diverse calibration and quality control concepts.

This makes it challenging to compare the results between labs and sometimes even between instruments within one lab. Furthermore, there is a general lack of

standardization due to the lack of suitable reference measurement procedures or reference materials.^{9,11}

Roche Diagnostics is developing standardized reference methods by collaborating with external mass spec experts. All assays included in the **cobas**[®] Mass Spec portfolio will be traceable to reference methods, which allows standardization of mass spec testing. This, in combination with **cobas**[®] Mass Spec, which includes all process steps required from sample to result, ensures that results can be compared with other labs using the same solution.

Figure 5: Proposed Ionify assay menu.*



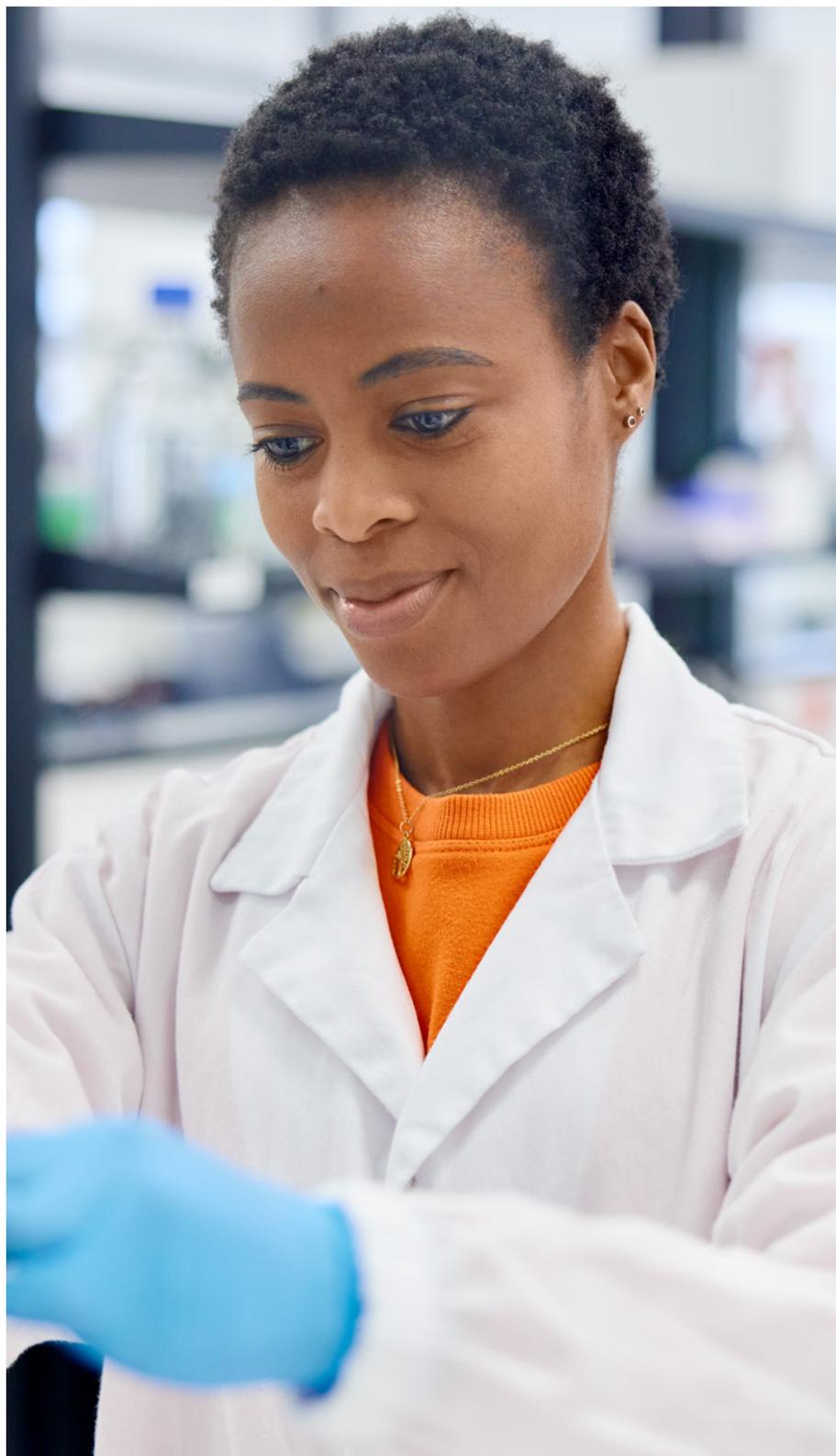
Simplified calibration concept

The current standard of clinical mass spec LDT is characterized by batch mode analysis and typically 6 to 8 calibrator levels included at least once, if not several times per batch. This results in a significant number of calibrators which need to be prepared or thawed and tested with every batch run.

The proposed traceability concept allows a simple calibration of **cobas**[®] Mass Spec applications. The **cobas**[®] Mass Spec calibration follows the concept established in today's immunoassay solutions. Instead of calibration with 6 to 8 calibrator levels, a 2-level re-calibration of a prefabricated calibration (or factory calibration) curve will be applied. Intensive testing done at Roche will significantly increase calibration robustness and lead to a clearly longer calibration interval.



More than **60 analytes**
are in development

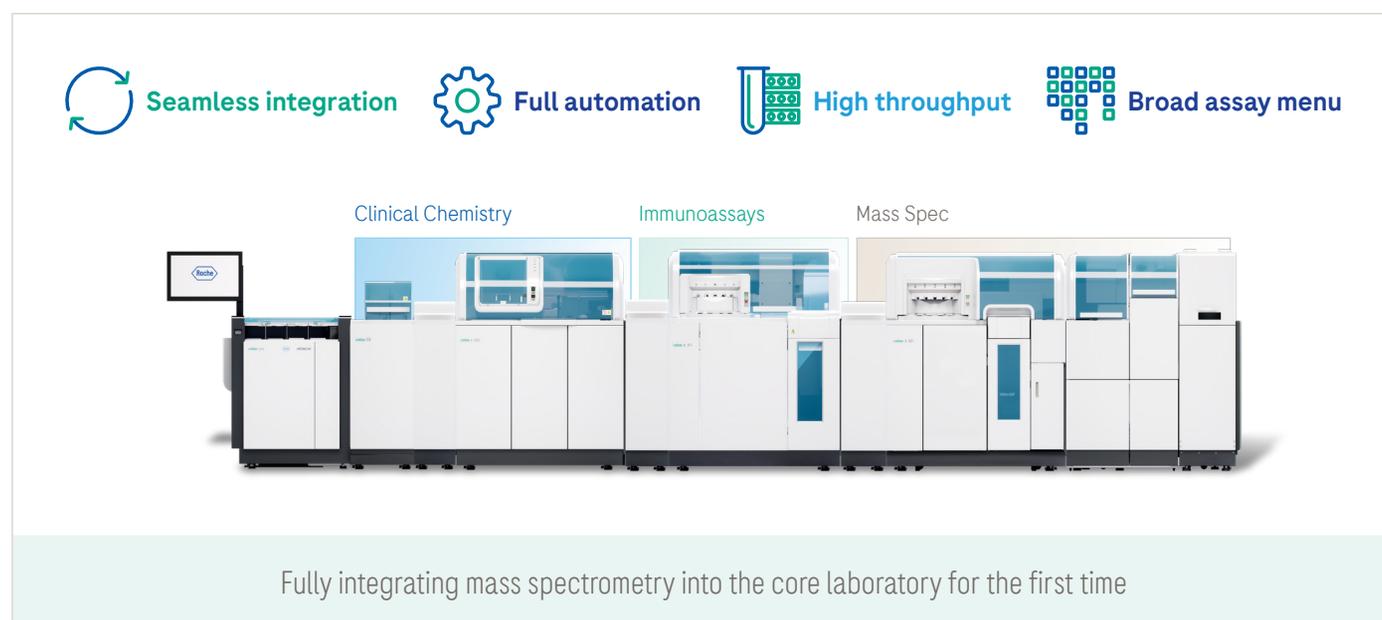


Chapter 5 Summary

In summary, the primary barriers to adoption of mass spec technology for routine clinical testing are high operational complexity, lack of automation, standardization, and integration, and the need for skilled technical staff.

The cobas® Mass Spec solution fully automates mass spec based testing - making it as easy to use as current clinical chemistry or immunochemistry systems. The system's design addresses the needs of a routine clinical laboratory, such as throughput and random access testing. It seamlessly integrates mass spec testing into the core lab. See Figure 6.

Figure 6: The future of mass spectrometry



At launch, **cobas**® Mass Spec will be in a stand-alone configuration only. All other configurations, including clinical chemistry and immunochemistry units, will be available at a later date.

References

1. Greaves et al. Key questions about the future of laboratory medicine in the next decade of the 21st century. *Clin Chim Acta*. 2019;S0009-8981(19)31882-0.
2. Seger, et al. Assuring the proper analytical performance of measurement procedures for immunosuppressive drug concentrations in clinical practice: recommendations of the International Association of Therapeutic Drug Monitoring and Clinical Toxicology Immunosuppressive Drug Scientific Committee. *Ther Drug Monit*. 2016;38:170-89.
3. Seger C, Vogeser M. Immunosuppressant drug monitoring – a routine undertaking? *J Lab Med*. 2010;34(3):1-11
4. Keevil BG. LC-MS/MS analysis of steroids in the clinical laboratory. *Clin Biochem*. 2016 Sep;49(13-14):989-97. doi: 10.1016/j.clinbiochem.2016.04.009)
5. Vogeser M, Zhang YV. Understanding the strategic landscape surrounding the implementation of mass spectrometry in the clinical laboratory: A SWOT analysis. *Clinical Mass Spectrometry*. 2018;9:1-6
6. Zhang YV , Rockwood A. Impact of automation on mass spectrometry. *Clinica Chimica Acta*. 2015;450:298-303
7. Stone JA, Fitzgerald RL. Liquid Chromatography-Mass Spectrometry Education for Clinical Laboratory Scientists. *Clin Lab Med*. 2018;38(3):527-537.
8. CLSI C62-A Liquid Chromatography-Mass Spectrometry Methods; Approved Guideline (2014).
9. Vogeser, et al. A proposal to standardize the description of LC-MS-based measurement methods in laboratory medicine. *Clin. Mass Spectrom*. 2019;13:36-38
10. Ouweland & Kema. The role of liquid chromatography-tandem mass spectrometry in the clinical Laboratory. *J Chromatogr B Analyt Technol Biomed Life Sci*. 2012;883-884:18-32
11. Seger, et al. Establishing metrological traceability in laboratory medicine. *Clin Chem Lab Med*. 2023; doi: 10.1515/cclm-2022-0995



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