

# Diagnosis by mass spectrometry

## Problem

Mass spectrometry emerged as a breakthrough technology for differentiated diagnostics. Some routine examples are diagnostics of pathogenic microorganisms, anti-doping tests, and aminoacidopathies. However, the solutions that exist in state-of-the-art laboratories cannot, for example, differentiate between bacteria that are or are not resistant to antibiotics, point out the agents that cause immediate sepsis, or even, in more unusual cases, point out the origin of poisoning.

## Solution

Methodology that includes a differentiated form of mass spectrometry analysis (LC/MS/MS) in conjunction with an artificial intelligence platform that solves and overcomes the above bottlenecks. Our computing environment is able to "learn" to diagnose virtually any pathology with protein changes.

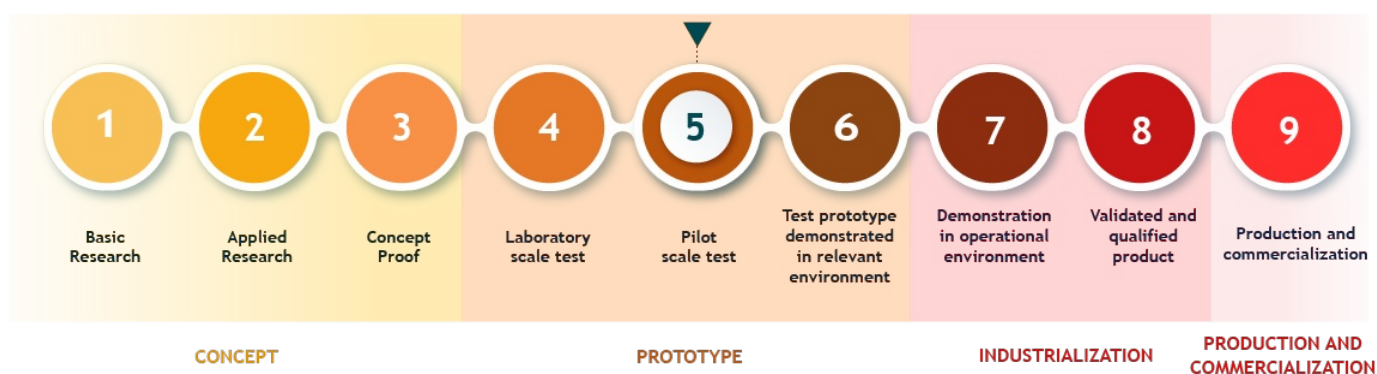
### Differential

Specificity

Artificial intelligence

Time reduction

### Development stage



### What we are searching for

Commercialization of the methodologies together with the software.

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# Inventors

Paulo Costa Carvalho



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Campus Fiocruz Maré - Av. Brasil, 4036 - Maré, Rio de Janeiro  
- RJ

CEP: 21040-361



[portfolio@fiocruz.br](mailto:portfolio@fiocruz.br)



+55 (21) 3282-9080