

## MOLECULAR MARKERS FOR TRACKING SKELETAL MUSCLE FUNCTION IN DUCHENNE MUSCULAR DYSTROPHY

Camilla Johansson<sup>1\*</sup>, Andreas Hober<sup>1</sup>, Fredrik Edfors<sup>1</sup>, Erik Niks<sup>2</sup>, Pietro Spitali<sup>2</sup>, Cristina Al-Khalili Szigyarto<sup>1</sup>

<sup>1</sup> KTH - Royal Institute of Technology, Stockholm, Sweden. <sup>2</sup> Leiden University Medical Center, Leiden, The Netherlands.

## Summary

The use of sensitive, accurate and specific bloodderived biomarkers can improve assessment of disease progression and therapeutic effects in DMD. Using paralell reaction monitoring mass spectroscopy (PRM-MS) and quantitative immunoassays, we have developed a pipeline to analytically validate blood biomarker candidates for DMD using longitudinally collected serum or plasma. This was illustrated for carbonic anhydrase 3, with high agreement between both methods.

## Conclusions

**1. Orthogonally validated** 4 of 10 biomarker candidates for DMD using PRM-MS.

2. Developed quantitative immunoassay for carbonic anhydrase 3 (CA3) with high correlaton to PRM-MS.

**3.** Proof-of-concept for **analytical validation** of biomarker candidates for DMD

## BACKGROUND



**Duchenne muscular dystrophy (DMD)** is caused by a mutation in the *DMD* gene, wich results in the lack of expression of the muscle protein dystrophin.

The lack of dystrophin expression cause fatal, progressive muscle wasing.

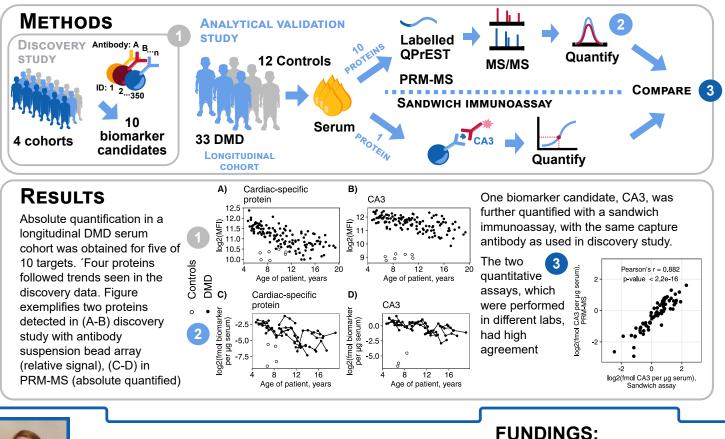
Development of dystrophin-restoring therapies are ongoing. The use of sensitive, accurate and specific blood biomarkers to assess therapeutic outcome could aid in this development.

Proteins from dystrophic tissues leak into the blood stream.

Many blood biomarker candidates have been described in literature, but none clinically validated

AFMTELETHO

**This study focuses on** analytical validation of a subset of blood biomarker candidates. The aim is to define a pipeline for analytical validation ahead of clinical validation.





CONTACT INFORMATION: camjoha@kth.se