



# OPTIMISATION OF MULTI-OMIC SPATIAL ANALYSES OF ENDOMETRIOSIS FFPE TISSUES USING MALDI MSI

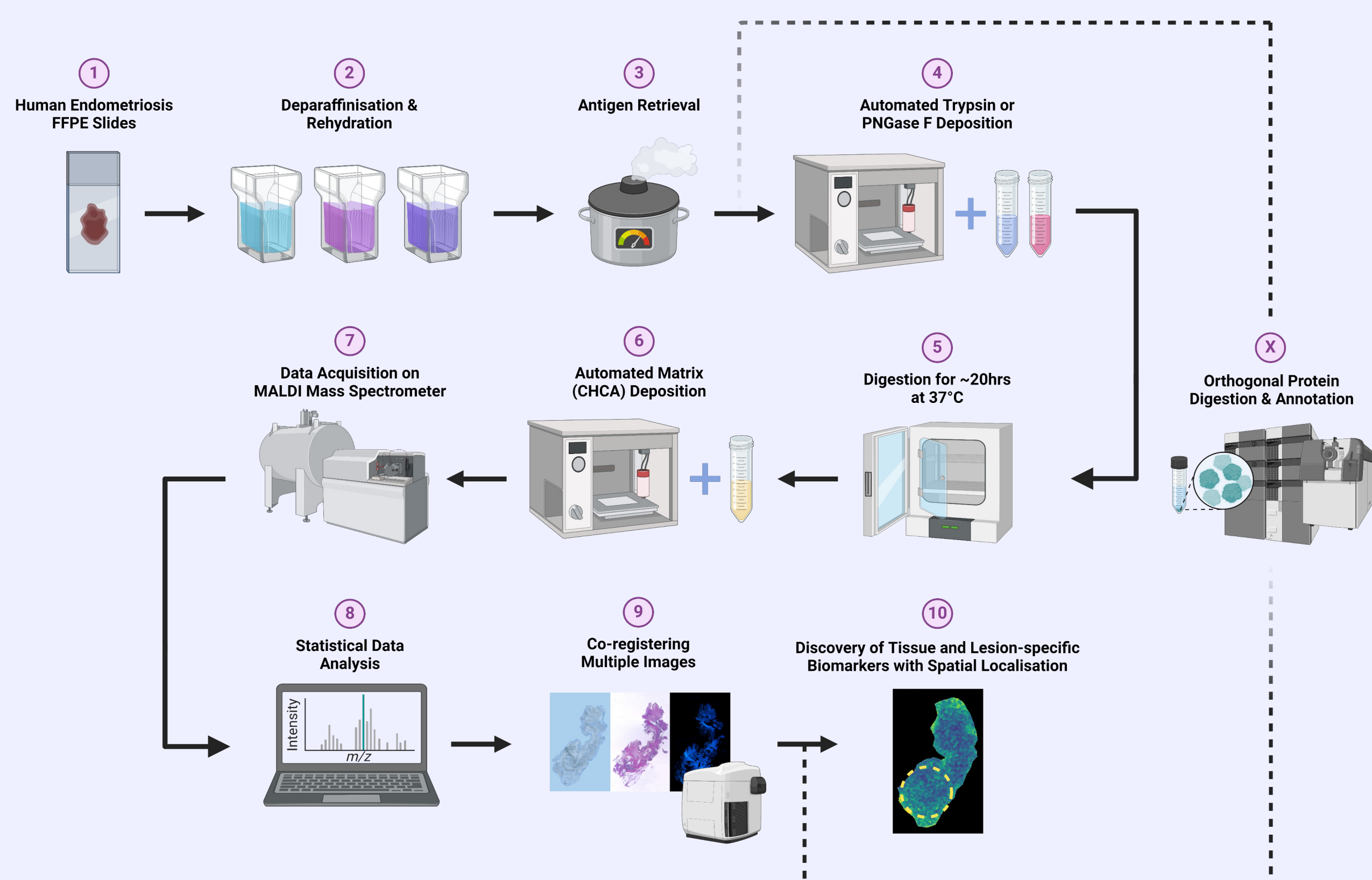
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## INTRODUCTION

- **Endometriosis** is a complex, chronic disease which causes **lesion growth** of endometrial-like tissues outside of the uterine cavity.
  - Affects **~11.4%** of Australian women of reproductive age.<sup>1</sup>
  - Symptoms include **chronic pain, painful menstruation and infertility**.
  - Diagnosis only currently possible with invasive surgery.
    - = Discovery of **new disease biomarkers urgently required**.
- Both **proteins and glycans** show much promise as disease biomarkers in both endometriosis and other diseases.<sup>2</sup>
- **Mass Spectrometry (MS)** is an advanced method for the measurement and **identification** of biomolecules **including proteins and glycans**.
- **Mass Spectrometry Imaging (MSI)** allows for the analysis of whole FFPE (formalin-fixed paraffin-embedded) tissues allowing for **molecular localisation and measurement of spatial distribution**.
- MSI permits accurate analysis of **endometriotic lesion-specific molecules** which can be **compared to healthy tissue to identify disease biomarkers**.

## METHOD

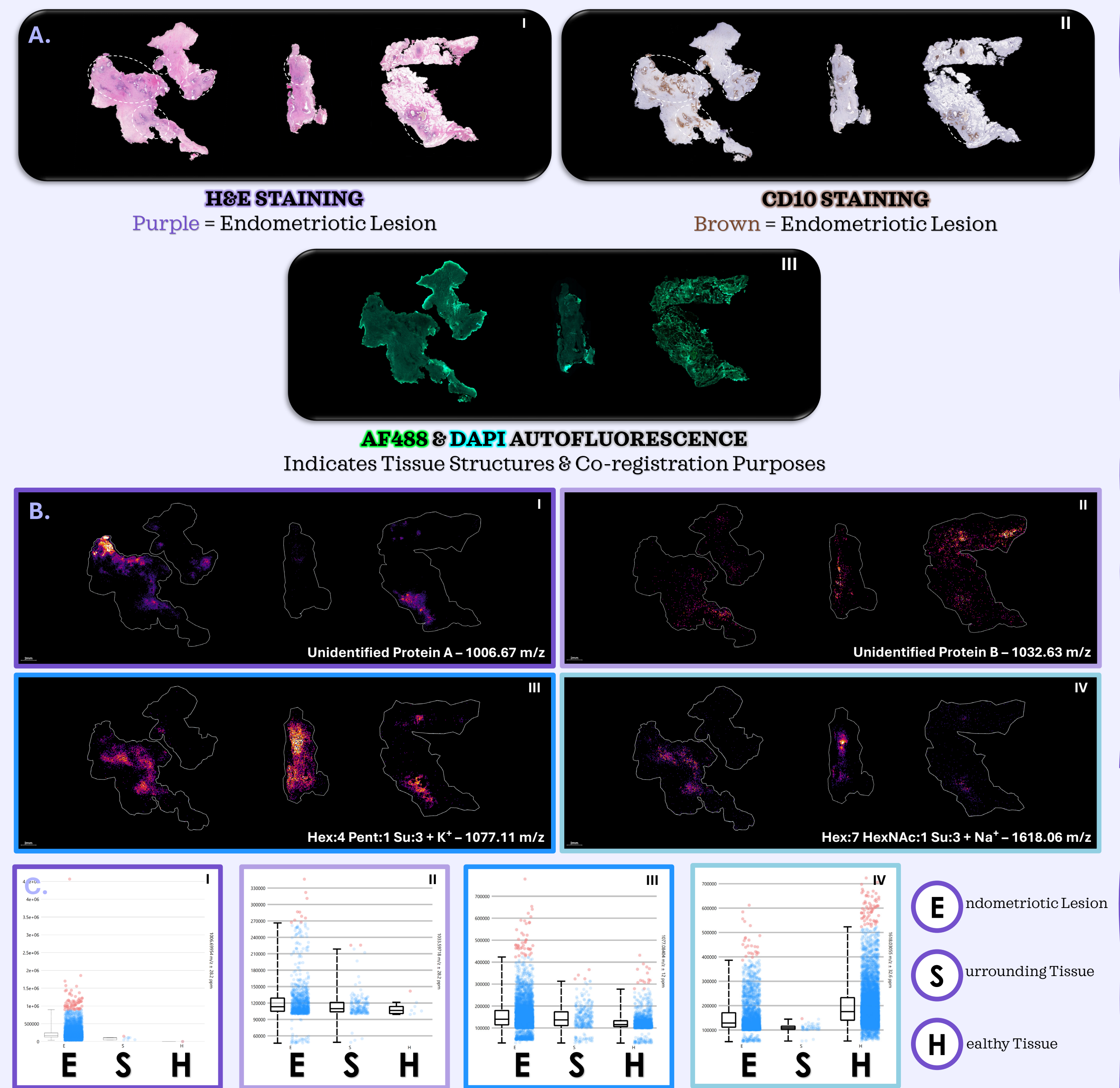


**Figure 1** - Diagram illustrating the process of MALDI-MSI sample preparation for FFPE (formalin-fixed paraffin-embedded) tissues and the identification and analysis of protein peptides and N-linked Glycans. Instruments shown include (3) Aptom Retriever 2100, (4/6) HTX TM-Sprayer M3, (7) Bruker Solarix 7T 2XR hybrid ESI/MALDI-FT-ICR MS, (9) Zeiss AxioScan Z.1 Digital Slide Scanner and (X) Shimadzu Prominence Nano HPLC coupled with a Sciex 5600 TripleTOF MS (created with BioRender.com)

- **10 confirmed endometriosis patients** with **≥ 1 deep and ≥ 2 superficial lesions** (= 44 sections) AND full-thickness uterine tissues AND a **healthy endometrium tissue microarray**.
  - Endometriosis and full-thickness uterine tissues sourced from The Royal Women's Hospital Tissue Bank under ethical approval (Project 10/43).
- **Method optimisation on full-thickness uterine sections** and the healthy endometrium microarray (sourced from CHTN) to define a healthy endometrium, then progress to using diseased endometriosis tissues.<sup>3-5</sup>
- Assess **spatial proteomic and glycomic profile differences** of deep and superficial endometriotic lesions as well as healthy endometrium samples from MSI data.
- **Statistically assess** using -
  - Principal Component Analysis (PCA)
  - Linear Discriminant Analysis (LDA) models
  - Two-sided Independent t-test (Student's or Welch's)
  - Receiver Operative Characteristics (ROC) curves
  - Area Under the Curve (AUC) analysis.

- Define tissue types and regions  
- Build classification models  
- Determine discriminant features
- **Annotate disease biomarkers** via orthogonal protein digestion and LC-MS/MS for proteins and METASPACE annotations for N-glycans.

## RESULTS & DISCUSSION



**Figure 2** - Graphs illustrating (A) the (I) H&E and (II) CD10 staining of three selected endometriotic FFPE sections with either purple or brown staining reflective of endometriotic lesions (circled in white), respectively; and (I) shows the autofluorescence of said sections using merged AF488 (excitation = 493 nm, emission = 517 nm) and DAPI (excitation = 353 nm, emission = 465 nm) filters collected on a ZEISS AxioScan Z1 digital slide scanner. (B) MALDI MSI ion images of discriminatory ions co-localised to the endometriotic lesions with I-II demonstrating unidentified protein peptides at 1006.67 and 1032.63 m/z and III-IV demonstrating tentative N-glycan annotations at 1077.11 and 1618.06 m/z. (C) Comparative Box Plots of the above MALDI MSI ions (B - I-IV) between different tissues and sites - endometriotic lesions, surrounding tissue and healthy tissue.

- From the assessed data we have shown **different proteomic and glycomic molecular signatures between gynaecological tissue regions**.
  - Different lesion sites express **presence or absence** of varying molecules at **higher or lower intensities** from surrounding and healthy tissues.
  - Some lesions appear to have **stronger proteomic or glycomic involvement**.
  - **Lesions appear heterogenous** - displaying **different molecular signatures** between both lesion sites within individuals and between different patients.
  - **Endometrial cycle phase** appears to have a notable effect on tissue and lesion molecular signatures (data not shown).
- MSI can therefore allow for the **accurate analysis and localisation** of differential molecular profiles between endometriotic lesions, surrounding tissues and healthy tissues.
- Next steps include:
  - **Further annotation** of data to identify affected proteins and glycans.
  - Cross-validation with different sample types.

## CONCLUSION

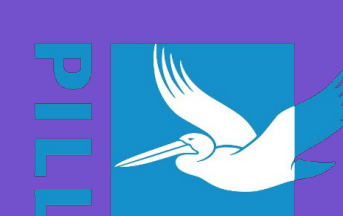
- MSI has the potential to be a **vital technology** in the discovery of biomarkers and understanding of pathophysiology in diseases such as endometriosis.
  - Tissue biomarkers can then be tested in more **non-invasive mediums** (blood and urine) to generate **diagnostic tests**.
  - The heterogeneity of endometriotic lesion sites, however complicates the discovery and assessment of biomarkers.
- Endometriosis needs further research to identify biomarkers and **assist in earlier diagnosis, thus improving quality of life** of those affected.

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## ACKNOWLEDGEMENTS:

- MRF Grant - 2019 Endometriosis Research GNT1199715: (Improving Diagnosis and Treatment of Endometriosis)
- Proteomics International - Orthogonal Protein Identification
- Cooperative Human Tissue Network (CHTN), US Dept. of Health and Human Services, National Institute of Health, National Cancer Institute, USA - Provision of Normal Endometrial Cycle Tissue Microarray
- Western Australian Government - Subsidised Analyses



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