

Identification and Profiling of Biomarkers in Neurodegeneration

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Introduction

- Novel biomarkers are necessary to further the development of treatments for neurodegenerative disease with advanced and emerging technologies being crucial for their identification and validation in both tissue and fluid samples.
- Tissue samples can be spatially profiled using a range of techniques, including the GeoMx Digital Spatial Profiler to examine transcriptomic or proteomic profiles and mass spectrometry imaging to examine a range of molecules including metabolites and lipids. The outputs from these technologies can be overlaid to examine variations between diseased and control samples.
- Liquid samples like plasma or cerebrospinal fluid can be used to track biomarker profiles longitudinally and can be interrogated using Luminex or Simoa technology alongside mass spectrometry for a wide range of different molecules. Again, the highly multiplexed readouts can be interrogated for a deeper understanding of drug and disease.



Histology Gold standard assavs Mass Spec (IHC/IF& ISH) Lipidomics, metabolomics and Proteomic Imaging Biomarker Identification/ Drug distribution (DESI/ MALDI) • • • **Tissue Profiling** Cell Type Profiling/ arget Engagemen HIA-DE (Confocal/ tiphoton/ Super in vivo Imaging Resolution microscopy) Biodistribution disease phenotyping, efficacy (PET, CT & IVIS ٠ RNA Protein LipidsSmall Molecules Time (Davs)

Deeper understanding of drug and disease – **Circulating**

- Liquid biopsies are just as valuable as they give a real time insight into the patient's disease and response
- Opportunity to utilise longitudinally
- Additional multiplexed readouts can be interrogated/overlaid to support hypothesis observed
- Circulating factors can provide information on the heterogeneity of the disease along with early response or resistance biomarkers, often before clinical responses are observed



md.catapult.org.uk

Technologies and Platforms SlideSeq Platform **Mass Spectrometry Capabilities Sample Acquisition** • We can run analysis on your own samples, assist with acquiring relevant human samples or run in vivo SAMPLE • Spatial transcriptomics method with single cell resolution for use on frozen tissues BIOLOGICA models to acquire animal samples if required. ASSAY DESIGN PREPARATION SAMPLES Human Samples Clustering analysis identifies • A dedicated samples team that has built connections with various sample providers including biobanks with groups of cells present with Liquid 😤 Tissue Unique MS assays Automation global networks – this enables us to access 'hard to find' samples at competitive rates. similar expression profiles. Using novel technologies (acoustic Extensive experience in sourcing a variety of high-quality samples including brain tissue, spinal cord and We cover a wide range of MS analyses: Cell Robust Type We handle most types of biological dispenser, multidrop, 384 well-plate), compound quantification, (un)targeted biofluids from donors with conditions such as Amyotrophic Lateral Sclerosis, Alzheimer's Disease, samples, ranging from tissue to Decomposition (RCTD) uses we handle all the sample preparation, omics, heavy labelled molecules, cell-Parkinson's Disease, Huntington's Disease. plasma. (CSF, sputum, tumour, optimising your samples for MS analysis. single-cell RNA-Seq datasets based and screening assays, etc.. serum, etc...) to identify the cell types • Experience in establishing new bespoke sample collections in a timely manner based on the needs of our • 3 x 3mm coverslip coated with present in each cluster. collaborators. monolayer of 10µm diameter DNA High-Throughput MS barcoded beads. Our samples team is well versed in managing applications for external ethical approvals to ensure DATA We have developed fast & robust screening compliance with HTA guidelines. assay for plasma proteomics. • The spatial position of each cell type can then be mapped. The unique barcode on each bead (LC)-MS We are also specialised in HTS MS-assay. enables the location to be determined Microglia O CA1 Principal Cells Exploitation Targeted MS Samples from in vivo Models Resident Macrophages CA2 Principal Cells ANALYSI • A section of frozen tissue is melted onto Oligodendrocytes CA3 Principal Cells We develop/optimise targeted MS assays, We have a suite of MS software that Alzheimer's Disease (5xFAD model) Dentate Principal Cells Polydendrocyte-1 improve their throughput and data richness.

extensive experience of neurosurgical
techniques we have experience of running
numerous mouse models of disease:

• In addition to validating new models and

 Huntington's Disease (R6/1 model) • Orthotopic Mouse brain tumour model (G261-luc) Acute LPS induced model of acute neuroinflammation • U87-luc subcutaneous xenograft model



GeoMx[®] Digital Spatial Profiler



nanoString

allows for fast processing of the data, as well as a dedicated team of bioinformaticians that perform data analysis and visualisation, in line with your research objectives.

run standard LC-MS/MS compour quantification in any sample type Untargeted MS (-Omics)

We are developing methods for biomarkers

discovery, focusing on small molecules and

proteins using LC-MS/MS approach.

We possess our own dedicated MS laboratory where we access the latest MS technologies (FAIMS, Ion Mobility) providing unique and novel solutions to specific challenges.

Protein Multiplexing (Luminex[®] and Quanterix[™] Simoa [®] Platforms)

• Luminex and Simoa platforms perform bead-based immunoassays, enabling simultaneous multiplex detection of up to 100 analytes



Luminex Bead Spectrum

Pre-validated Assays

- Various commercially pre-validated assays are available and can be run in house on the Luminex MAGPIX instrument
- Catalogues with >600 validated human analytes (includes several multiplex neuroscience/ neurodegeneration panels) • NF-H (but not NF-L), Tau (Total and phospho (pT181 + pT231)), TDP-43

Customised Assays



- Assays to targets not commercially available can be built and validated in house (E.g. NF-L assay validated in house)
- Several neuro focused assays are currently being validated for use with both human and mouse CSF and plasma samples

Pre-validated Assays • GFAP • NF-L • TDP-43 • Total Tau • Aβ40 or Aβ42 • pTau (181 or 231)



Morphology Markers

2. ROI Selection

- Can be selected from a host of pre-validated antibodies (E.g. GFAP (Astrocytes); Iba1 (Microglia), NeuN (Neurons), Amyloid-β (Plaques))
- New markers bespoke to each project can additionally be validated in house

Mass Spectrometry Imaging

- Label free surface analysis of samples (E.g. tissue/ dried blood spots) for a variety of analytes
- Possible to analyse different molecular classes from a single section.
- Relative/semi-quantitation
- Can be correlated and overlaid with microscopy or DSP images (on same or consecutive sections)

MSI Analytes

- Small molecules
- Metabolites
- Lipids Peptides/proteins

What anatomical structures can we see with each technique?



What can we detect with MALDI and DESI? MALDI DESI Daltons > 5000 ~ 100 < 2000 Protein Small molecules Lipids fragments Lipid/metabolite Amino Antibody acids/Peptides fragments fragments Metabolites N-glycoprotein Drugs fragments Glycans

Lipidomic Profiling in Human Spinal Cord





• Amyotrophic Lateral Sclerosis (ALS) is a fatal neurodegenerative disorder characterised by the loss of motor neurons in the brain and spinal cord, with approximately 15% of patients additionally developing a form of Frontotemporal Dementia (FTD)

Spatial Protein Profiling in Neurodegeneration

• The aim of this project was to examine differences in protein profiles in neurodegenerative disease, (specifically in this case, Amyotrophic Lateral Sclerosis (ALS)), and aging.



🔶 ALS

Outer White Matter

Lateral Funiculus

Anterior Horn

Posterior Horn

Luminex Neurofilament-Light Assay



• Neurofilament light is an established biomarker for neuronal damage in a range of neurodegenerative diseases.

• The dysregulation of lipid metabolism is a common hallmark of a range of neurodegenerative diseases, including ALS.

Mass Spectrometry Imaging (MSI)

- A collection of lipids and metabolites previously recognised to be associated with anti-inflammatory pathways was spatially profiled across human spinal cord tissue using MSI.
- The snap frozen tissue was additionally H&E stained in order to define the anterior and posterior orientation.
- One of the lipids, the omega-3 polyunsaturated fatty acid Docosahexaenoic acid (DHA), was found to be increased in the grey matter of the spinal cord in ALS and FTD-ALS patients.



- A panel of 74 proteins was spatially profiled across 6 spinal cord tissues (3 from control donors and 3 from patients suffering from ALS) using the GeoMx DSP platform.
- Data from 4 Regions of each spinal cord were collected, 2 from the grey matter (posterior horn, anterior horn) and 2 from the white matter (lateral funiculus, outer white matter).
- Data from the various regions of each tissue were collected and analysed.



- A Luminex assay was developed and validated to quantify NF-L in a variety of samples, including cerebrospinal fluid, cell lysates and tissue homogenates.
- Multiplexing of this assay with a selection of other proteins has additionally been tested and validated.



