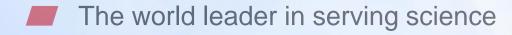


Current Trends in Clinical Diagnosis with Orbitrap High-Resolution Mass Spectrometry-based Techniques

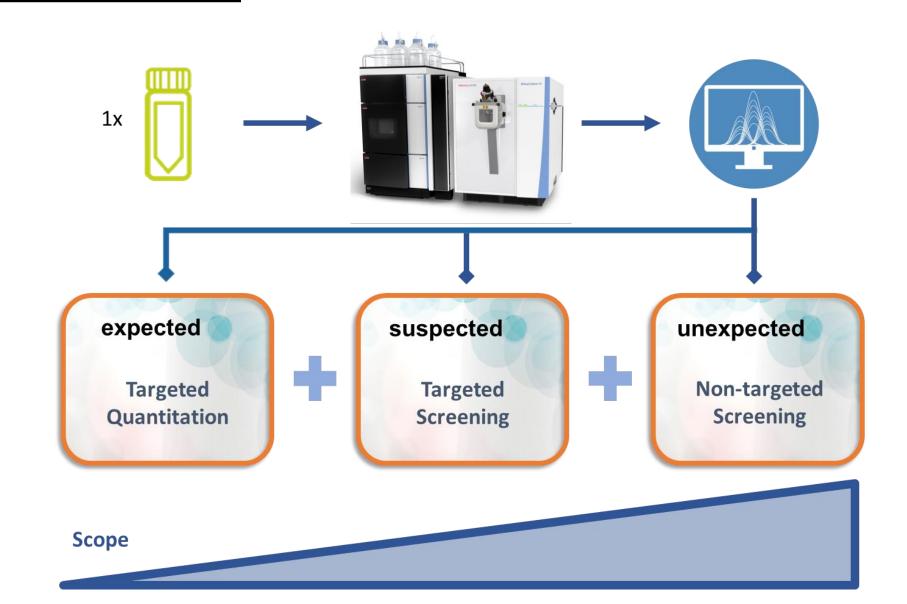
Jitnapa Voranitikul

Product Specialist, SciSpec Co., Ltd.





System Main Workflows





Thermo Scientific

Orbitrap HRAMS System





Mass Analyzer: Orbitrap™ Technology

Year	Orbitrap	TOF
2010	61	165
2011	87	189
2012	112	219
2013	188	255
2014	219	361
2015	380	516
2016	625	734
2017	794	830
2018	755	813
2019	934	934
2020	1042	971
2021	1176	1042
Total	6373	7029



By the end of 2021, the publication ratios in each years, compared to those of 2010, show superiority of the publication growth from Orbitrap technology 3 times higher than that of TOF.

Latest Update: June 13, 2022



Agenda Application

Application

- Endocrinology IGF-1
- Metabolic syndromes IEM hemoglobinopathy
- Precision medicine & advanced clinical diagnostics
 - Infectious disease (Ucalgary, Nanopin, Fleury), pandemic responsiveness
 - Alzheimer's disease C2N, Reproductive Health NX Prenatal



Quest Diagnostics: IGF-1 LCMS assay



Quantitation and identifying variants of IGF-1 using high resolution mass spectrometry.

Quantification of insulin-like growth factor 1 in human serum by Vanquish UHPLC with Q Exactive high-resolution accurate-mass mass spectrometry for clinical research

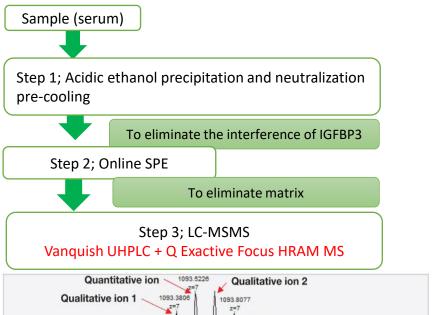
LLOQ

10 ng/mL

3.96

Identification of IGF-1 variants helps support physician's decisions when treating growth-related patients.

- Development of a robust protein assay by "top-down" measurement of IGF-1 in human serum
- Improve assay performance using HRAM MS
- Identify IGF-1 variants and understanding its clinical significance



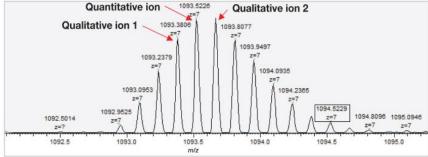
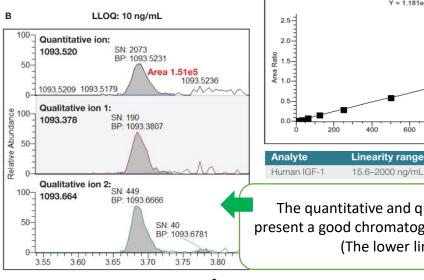


Figure 4. Chromatography mass spectra for insulin-like growth factor quantitative, qualitative, and internal standard ions



The quantitative and qualitative ions at the LLOQ (10 ng/mL) still present a good chromatographic peak shape and an excellent S/N ratio. (The lower limit of the ref range is 14 ng/mL)

r2 linear fit, 1/x weight

0.9996

Outstanding of Benefits;

Established reference range comparable to gold standard immunoassay method

Calibration Curve

- ✓ Better sensitivity, specificity vs immunoassay approach (the main way to measure IGF-1)
- ✓ Developed an effective and high-throughput method to monitor IGF-1 and its variants using high resolution LC-MS coupled with online extraction in a clinical laboratory
- ✓ High resolution MS revealed the complexity of IGF-1 variants



Case Study: Archimed Life (2016 – 2019)

HRAM/MS method fits ideal requirements for Clinical

- · Collaborator: David Kasper (Founder), Thomas Wiesinger
- Implementation of novel HRAM/MS methods for Newborn screening on Q Exactive Focus MS
 - Prospective study of ~5000 de-identified newborn samples
 (UKE Hamburg) –Screening of hemoglobinopathy and thalassemia
 - Applicability of HRAM/MS in untargeted screening protocol for rare mutations

Thomas Wiesinger, Thomas Mechtler, Markus Schwarz, Xiaolei Xie, Regine Grosse, Paulina Nieves Cobos, David Kasper* and Zoltan Lukacs*

Investigating the suitability of high-resolution mass spectrometry for newborn screening: identification of hemoglobinopathies and β-thalassemias in dried blood spots

https://doi.org/10.1515/cclm-2019-0832 Received August 8, 2019; accepted December 23, 2019; previously published online February 6, 2020 Introduction

identification of clinically relevant mutations and differentiation between hetero, homozygous form"



The major challenge for a screening assay

"The method is suitable for the

☐ Identification and differentiation between carrier- and sickle cell disease-positive samples

UKE Hamburg

Eppendorf

☐ Single mutation of clinical relevance in the beta-chain (e.g. Hb C, D, E), these Hb variants differ only by 1 Da for the entire protein.



- ✓ Minimal sample preparation and handling
- ✓ High throughput (2 min/sample)
- ✓ Inexpensive (6–7 € Euro/sample)
- ✓ Robust system (no maintenance only one time source cleaning during the study)
- ✓ Automated clinical assessment

A B C D 3.2 mm DBS punch Tryptic protein fragments 1 DBS punch 1 Extraction buffer 2 Info: intact protein & tryptic frag. Automated data processing

Figure 1: Illustration of the general workflow.

(A) Single DBS punch extracted with tryptic buffer (40 mM NH_4HCO_3 , 9% MeCN, 1 mM $CaCl_3$, 5 mg/mL TPCK-treated trypsin from bovine pancreas) for 30 min; (B) mixture of the intact protein (α , β and γ) and tryptic fragments (e.g. C, D, E and S); (C) flow injection (duplex mode) is performed after quenching (2 min/sample); (D) results are processed with TraceFinderTM.

Archimedlife

Science GmbH



Automated Dried Spot Analysis –

Comprehensive workflows in one platform

The world leader in serving science



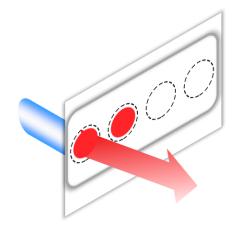
What is Flow-Through Desorption?

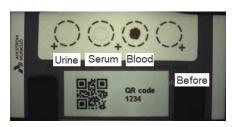






- Whatman DMPK
- PerkinElmer 226
- Ahlstrom Autocollect
- HemaXis DB10









- No disc-punching Direct elution of DBS from cards
- Leak-tight clamp heads
- Highly efficient desorption



Dried Matrix Sent Directly to LC/MS

- Flow injection
- Analytical Flow
- Turboflow

^{*} Patented



Tuberculosis (TB) by NanoPin's Technology

NanoPin uses an antibody enrichment approach to capture species-specific biomarker peptides.



May 21, 2020

Desert Platforms MDC Rahul Rao Project Manager 275 N. Gateway Dr. Phoenix, Arizona 85034

Re: O191789/S001 Trade/Device Name: NanoDetect-TB Received: February 26, 2020

Dear Rahul Rao

received the above submission requesting designation as a Breakthrough Device. The proposed indications for use includes "NanoDetect-TB is indicated for use in the detection of active TB disease. It is an in-vitro diagnostic test for the direct detection of target peptides derived from proteins secreted by Mtb bacilli into the circulation, which are then assayed in serum or plasma samples collected by standard intravenous blood draws. Antibody-conjugated nanoparticles are employed to enrich the target peptides from trypsin digested serum or plasma samples, which are then analyzed by MS to report a positive or negative result for active TB disease. The intended use population are symptomatic patients suspected to have active TB disease and highrisk individuals." We are pleased to inform you that your device and proposed indication for use meet the criteria and have been granted designation as a Breakthrough Device. Please refer to the FDA guidance document entitled "Breakthrough Devices Program", for more information regarding the program, available at https://www.fda.gov/media/108135/download.

We recommend you review the FDA guidance document for the Breakthrough Devices Program referenced above for the available mechanisms for obtaining feedback from the Agency on device development for designated breakthrough devices. When submitting any new requests, please reference Q191789/S001. Any new submission should be provided as an eCopy, it should include the FDA reference number for this submission, and should be submitted to the following address:

U.S. Food and Drug Administration Center for Devices and Radiological Health IDE Document Control Center - WO66-G609 10903 New Hampshire Avenue Silver Spring, MD 20993-0002

U.S. Food & Drug Administration 10903 New Hampshire Avenue Silver Spring, MD 20993



Prof. Tony Hu Tulane University | Co-Founder of NanoPin Technologies, Inc.

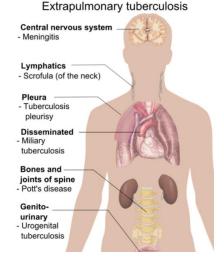
The Center for Devices and Radiological Health (CDRH) of the Food and Drug Administration (FDA) has

New Assay (NanoPin) Current method Tissue biopsy Serum

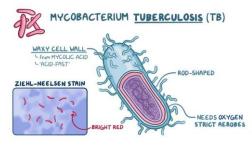


Pulmonary and ExtraPulmonary Detection





Main sites of



Selection of target peptide from the untargeted analysis.

HRMS (Orbitrap)



target peptide

TSQ (Altis)



Tuberculosis (TB) by NanoPin's Technology

- Collaboration press release (<u>link</u>)
- Support upcoming clinical trial strategies in US and China
- Infectious disease diagnostics \$26B with TB Segment >\$3 billion by 2024 and HIV Segment \$5 billion by 2023







Challenge:

- Low concentration in the bloodstream (Especially early in infection)
- Disturb by abundant serum protein
- Conserved protein by closely related pathogens



Benefits from digestion:

- Remove the masking effect
- Increase the amount of target peptide
- Distinguish highly conserved proteins by closely related pathogens

Conclusion

 Antibody-mediated enrichment of target peptides prior to their analysis by LC-MS/MS

Workflow

- R&D discovery on Thermo Fisher Orbitrap and triple quadrupole MS
- Successfully transitioned and implemented NanoPin TB assay on Vanquish MD + TSQ Altis MD

Advantages of the alternative method

- High increase in both sensitivity and specificity
- ✓ No tissue biopsy
- Quick response (LC-MS/MS)





SARS-CoV-2

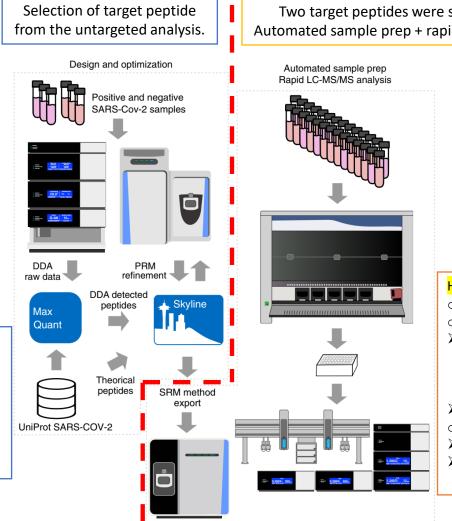
Establishing a mass spectrometry-based system for rapid detection of SARS-CoV-2 in large clinical sample cohorts.



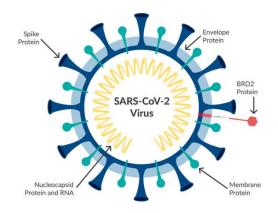
Dr. Valdemir Carvalho Fleury Group, Brazil

Develop Phase

- o LC: Ultimate 3000 Nano
- o MS: Q-Exactive HF-X Mass spectrometer
- o Column
- > PepMap 100 C18, 5 um, 0.3 x 5 mm; sample trapping precolumn
- PepMap RSLC C18, 2 um, 150 um x 15 cm; analytical column



Two target peptides were selected. Automated sample prep + rapid LC-MS/MS



High throughput method; Routine Lab

- o Automated sample prep: Haliton
- o Rapid LC-MS/MS
- LC: Transcend TLX-4
 - + four Dionex UltiMate 3000 quaternary pumps
 - + four Dionex UltiMate 3000 binary pumps
 - + VIM+ one CTC PAL autosampler
- ➤ MS: TSQ Altis (ESI) Mass Spectrometer
- o Column
- > Cyclone-P HPLC 0.5 x 50 mm column; TurboFlow column
- > Acquity UPLC BEH C18, 1.7 um, 2.1 mm x 50 mm column; Analytical column

Thermo Fisher

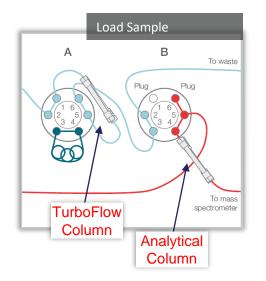
Transcend II TLX-1 System

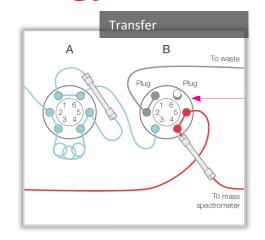
- Flow injection (FIA)
- LX Mode Analytical Flow
- TLX Mode TurboFlow
- Control and timing controlled by Aria MX Software
 - Pressure 1000 bar
 - Viper Fingertight Fittings
 - Features a biocompatible 2 channel binary highpressure gradient mixing pump with 2 x 3 solvent channels. Select among 3 solvents per piston assembly (A1, A2, A3 & B1, B2, B3)

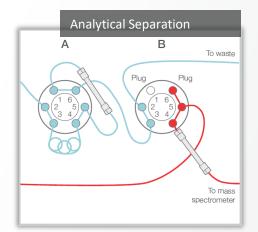


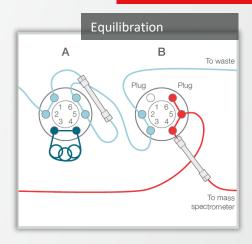


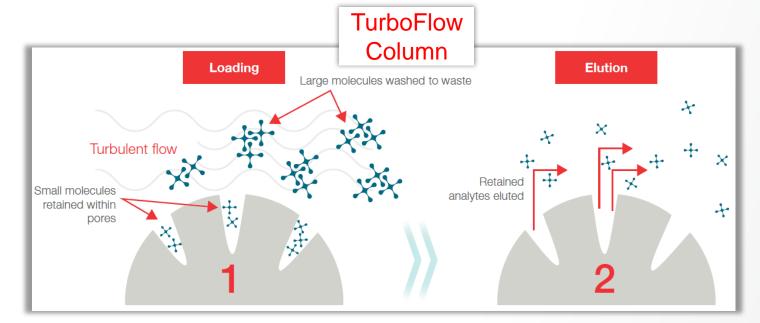
TurboFlow Technology







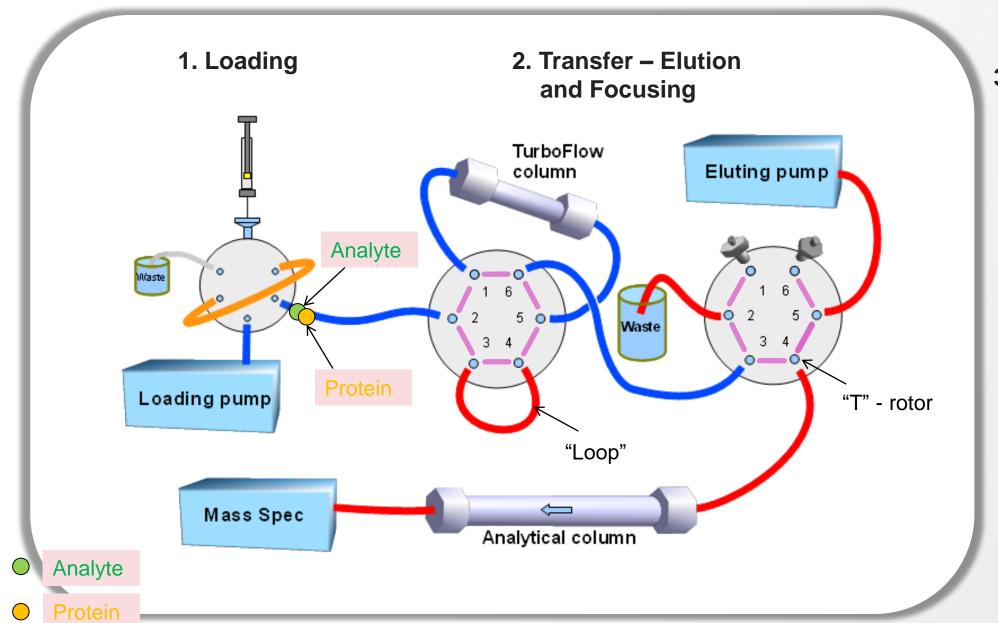




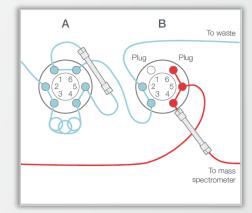
- Small molecules diffuse into porous particles faster than large molecules
- Sample components of interest Analytes are well retained
- Less retained components (e.g., salts & sugars) are rinsed away

Thermo Fisher SCIENTIFIC

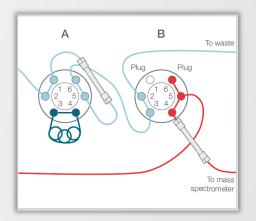
TurboFlow Focus Mode Plumbing



3. TX column cleaning LC column gradient



4. Re-conditioning





SARS-CoV-2

The concept application of automated sample preparation and multiplexing TFC coupled to triple quadrupole MS as a feasible alternative for detecting SARS-CoV-2 in clinical respiratory tract samples on a large scale at a population, level was proved.

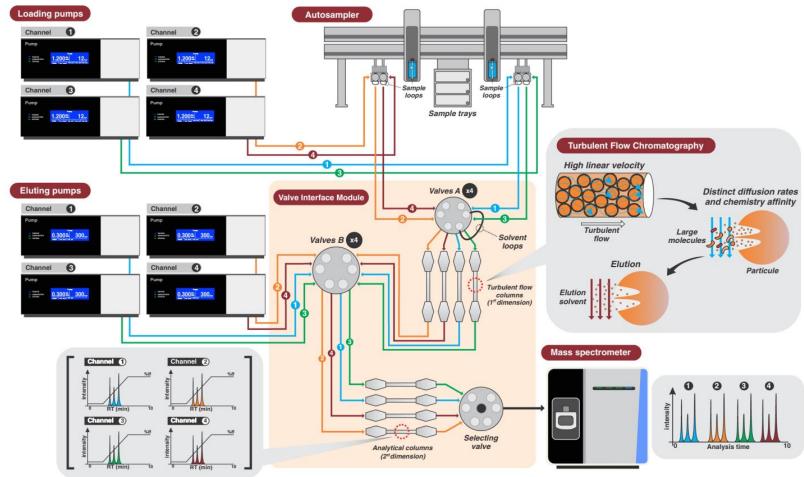


Fig. Schematic illustration of turbulent flow chromatography (TFC) setup in Transcend TLX-4 system coupled to TSQ Altis triple quadrupole mass spectrometer

Conclusion

 ✓ High throughput targeted proteomics assay i.e., LCMS approach as an alternative testing strategy for large populations

Outstanding of Benefits

- Analysis resulted in a 2.5-min acquisition time per sample (96 sample/4h ~500 samples/day)
- ✓ Circumvent reagent shortages
- Better specificity vs. immunoassay approach (standard method)
- Automated sample prep
- o Rapid LC-MS/MS
- LC: Transcend TLX-4
 - + four Dionex UltiMate 3000 quaternary pumps
 - + four Dionex UltiMate 3000 binary pumps
 - + VIM+ one CTC PAL autosampler
- MS: TSQ Altis (ESI) Mass Spectrometer
- o Column
- > Cyclone-P HPLC 0.5 x 50 mm column; TurboFlow column
- Acquity UPLC BEH C18, 1.7 um, 2.1 mm x 50 mm column; Analytical column



C₂N Diagnostics



1st precise clinical Alzheimer's blood test on Thermo Fisher Orbitrap Lumos Tribrid MS

Objectives

- Develop a LCMS approach to better diagnose and treat Alzheimer's and other neurodegenerative diseases
- Non-invasive, less expensive and scalable
- Need for amyloid tests as **companion diagnostics** for AD treatment

Deliverables & Impact

- Thermo Fisher-enabled platform for early brain pathology detection and early treatment
- Address public health burden, \$600B blood-based AD diagnostics, health and wellness market and companion diagnostics for AD treatment
- **Deployment of IVD kits** to qualified reference labs globally

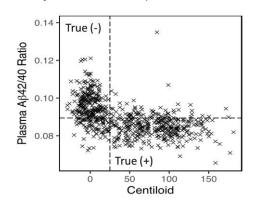
C₂N Diagnostics Introduces the PrecivityAD2™ Blood Test

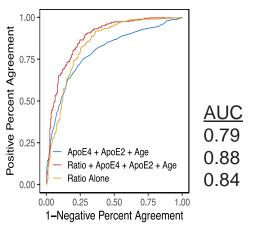
Next Generation Blood Test Aims to Establish New Standard in Alzheimer's Disease Diagnosis with Combined Measures of Amyloid Beta and Tau Protein



PrecivityAD was granted designation as a Breakthrough Device by FDA in 2018

- PrecivityAD™ Performance High Correlation to PET scan results
- Positioning PrecivityAD™ as a Rule In/Rule Out Amyloid test for ~80% of the intended use patient population
- Comparison between PrecivityAD™ test and amyloid PET scan results (+ PET defined by centiloid >25)



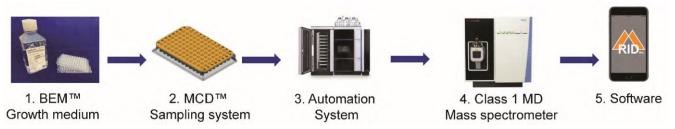




Rapid infections diagnostics: University of Calgary

• Thermo Fisher-enabled platform for rapid, metabolomics-based diagnostics approach for infectious disease

Hopper™ | Rapid diagnostic platform for identification and antibiotic susceptibility testing of pathogens





	Sample preparation (incubated period)	Method
Current method	2-5 days	MALDI-TOF (DI), AST
New method	4 h (MPA)	LC-MS/MS

MPA = Metabolic preference assay; metabolic fluxes observed in ex-vivo)





Ian Lewis, PhD; Associate Professor
Al Translational Health Chair
University of Calgary

Lewis Research Group, UCalgary MS lab showcase at the ACAD grand opening

Conclusion

- Diagnostic platform for implementing high-volume diagnostic tests for bloodstream
- Rapidly identify microorganisms and measuring their antibiotic susceptibility profiles (AST) using a LCMS approach

Workflow

- R & D discovery on Thermo Fisher Orbitrap (Q Exactive HF MS (negative)-MAVEN (SW))
- Successfully established the use of Vanquish Flex+ TSQ Altis MS-TraceFinder software for microbial identification and antimicrobial susceptibility testing (AST)



Potential implementation into **DynaLIFE** – **multiple reference sites** across **Alberta and Canada**

Outstanding of Benefits

- Replaces standard tests performed by MALDI-TOF (ID) and VITEK (AST)
- ✓ Reduces current microbiology testing workflow form 2-5 days to 4 hours



NX Prenatal, US

Blood-based diagnostics for preeclampsia and preterm-birth risk stratification

Addressing a critical clinical unmet need in women's health

NX Prenatal Announces Publication of Peer-Reviewed Study in the American Journal of Obstetrics & Gynecology for Validation of First Trimester Preterm Birth Risk Biomarkers

The findings represent the first reported multi-site validation of blood-based biomarker panels at 10-12 weeks gestation to stratify pregnant patients for the risk of preterm birth

Feb 28, 2019, 12:18 ET

LOUISVILLE, Ky. and HOUSTON, Feb. 28, 2019 /PRNewswire/ -- NX Prenatal Inc., focused on development of proprietary blood-based, early warning molecular diagnostic tests for adverse pregnancy outcomes, announced today the publication of a peer-reviewed study, "Circulating Microparticle Proteins Obtained in the Late First Trimester Predict Spontaneous Preterm Birth at Less than 35 Weeks Gestation: A Panel Validation with Specific Characterization by Parity" in the American Journal of Obstetrics & Gynecology. The publication is available online: https://doi.org/10.1016/j.ajog.2019.01.220

Objectives

 Exosome isolation analytical approach for targeted biomarker panel development

Conclusion

Blood test for assessing the risk of pre-term birth as easily as 10-12 weeks (compatible with NIPT testing)

Workflow

- R & D discovery on Thermo Fisher Orbitrap Fusion Tribrid
- Successfully transitioned and implemented targeted peptide biomarker panel on LC-MS/MS Vanquish Horizon UHPLC + TSQ Altis MS

scientific reports Microparticle protein Check for update Late first trimester circulating microparticle proteins predict Size Exclusion the risk of preeclampsia < 35 weeks Chromatography and suggest phenotypic differences among affected cases Thomas F. McElrath¹⁵⁶, David E. Cantonwine¹, Kathryn J. Gray¹, Hooman Mirzakhani², Robert C. Doss³, Najmuddin Khaja³, Malik Khalid³, Gail Page³, Brian Brohman³, Zhen Zhang⁴, (Microparticles) TurboFlow + Analytical Column Size: 100 - 1000 nm Markers: Integrins, glycoprotein-1b, P-+ Fusion Lumos Orbitrap selectin, VCAMP3, ARF6 Annexin V Size: 40 - 100 nm Isolation speed: 10,000 Markers: Alix, TSG101, 60,000g Rab, chaperones, CD63, Isolation speed: 100,000-200,000g Peptide spectrum library Transfer Method

→ LC (Vanguish Horizon)

+ TSQ (Altis)

Target peptide biomarker panel

Outstanding of Benefits

- Quick access to prophylaxis and therapy
- Reduce NICU dependency
- Enabled development of new pipeline product (Placenta Accreta Study)



Summary

		Benefit of using HRMS
•	Endocrinology – IGF-1	 High sensitivity and selectivity vs immunoassay approach Reveal complex of IGF-1 variants High throughput
•	Metabolic syndromes – IEM hemoglobinopathy	 Identification and differentiation of carrier and sickle cell disease-positive in beta chain Single mutation of clinical relevance

Precision medicine & advanced clinical diagnostics
 Infectious disease (Ucalgary, Nanopin, Fleury), pandemic responsiveness
 Alzheimer's disease C2N, Reproductive Health NX Prenatal

R & D discovery on Thermo Fisher Orbitrap (HRMS)

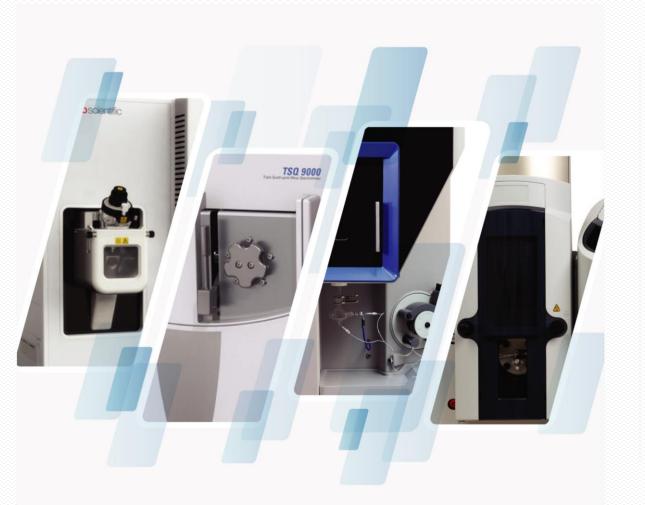
Selection of target peptide (Biomarker) by untargeted peptide analysis



Routine Lab on Thermo Fisher Triple Quadrupole

- ✓ Adding some sample preparation technology ex. Nanopin technology for TB assay
- ✓ Adding some automated system ex. Online SPE, Transcended TLX 4 which supports high throughput targeted peptide (Biomarker) assay

ติดตามกิจกรรมของบริษัทได้ที่













@scispec

















