Proteomics & Metabolomics analysis of COVID-19 patients to identify biomarkers



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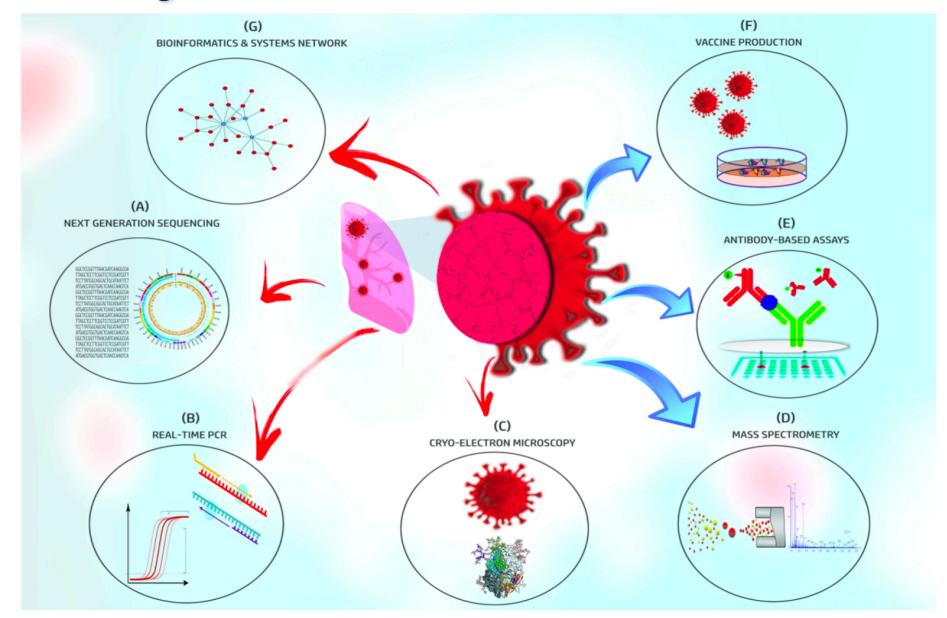
Guest Editor:

Proteomics, J of Proteomics, Nature India Editorial board, JPP

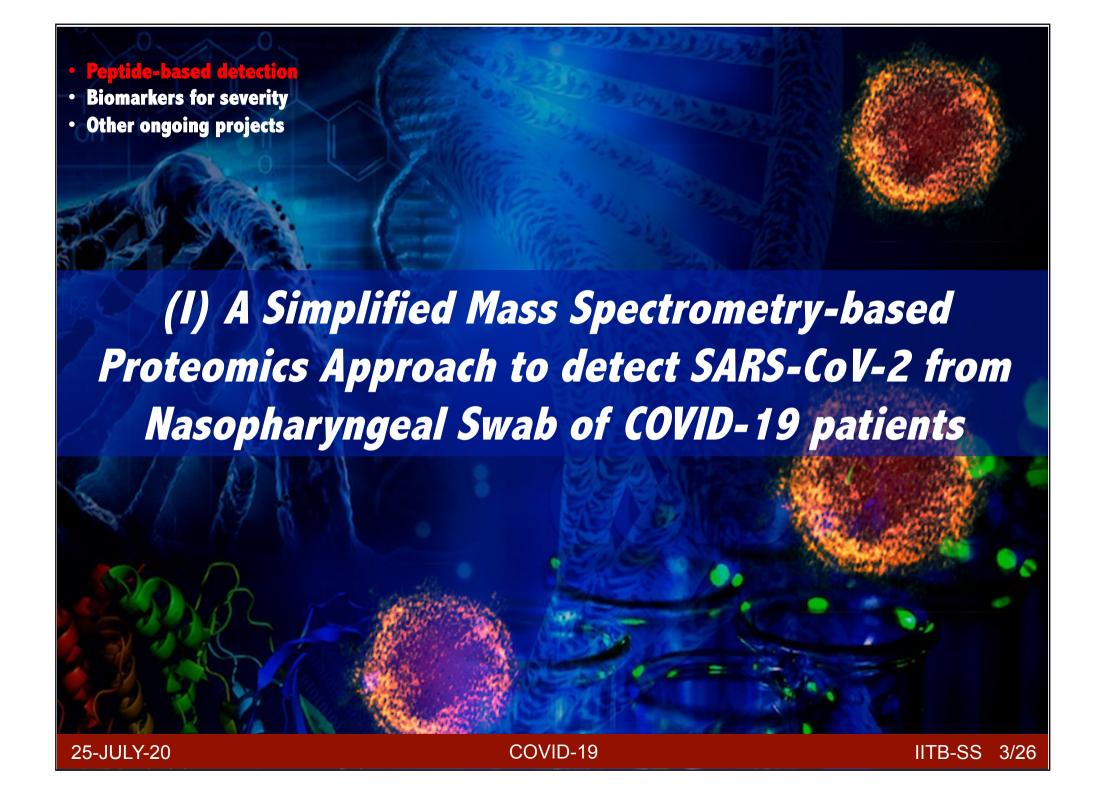
Outline

- Introduction
- Peptide based detection
- Biomarkers for severity prognosis
- Other ongoing projects
- Summary

COVID-19: Proteomics and Multi-omics Research

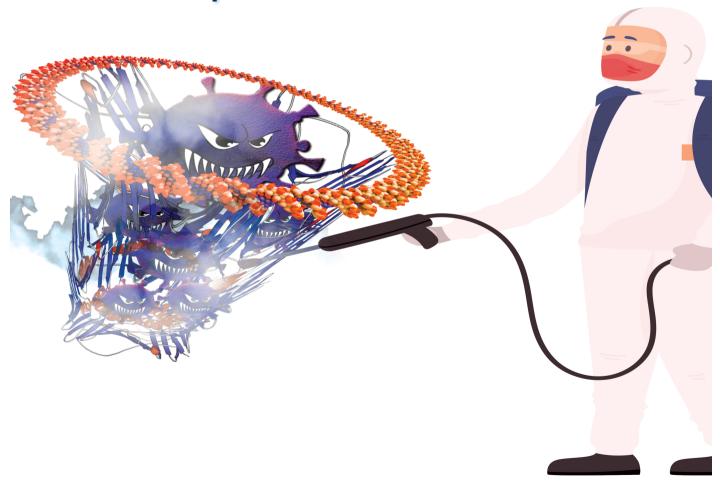


Ray S, Srivastava S. OMICS A Journal of Integrative Biology, 2020

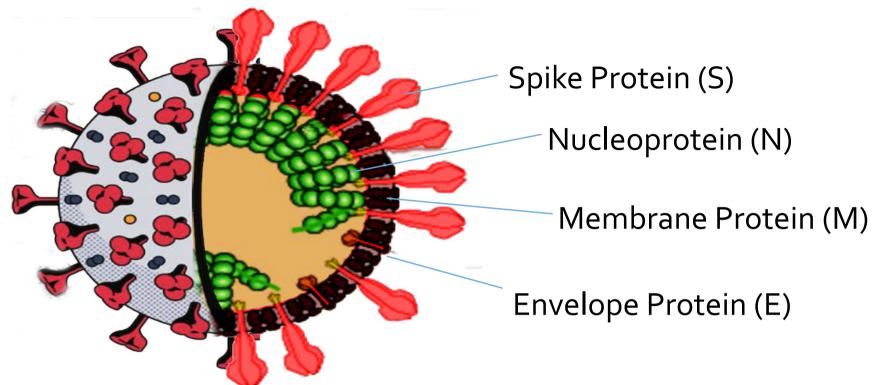




How do we capture SARS-CoV-2 Peptides/ Proteins?



SARS-CoV-2 Proteins



- Viral envelope is made of (S, E, M); N-protein encapsulates & protects RNA genome
- S (less copies); E, M (Short & tightly membrane bound) difficult to identify
- N-Protein is main target for detection

Proteomics National Facility

High Resolution Mass Spectrometry based Proteomics Research and Training Facility at Indian Institute of Technology Bombay



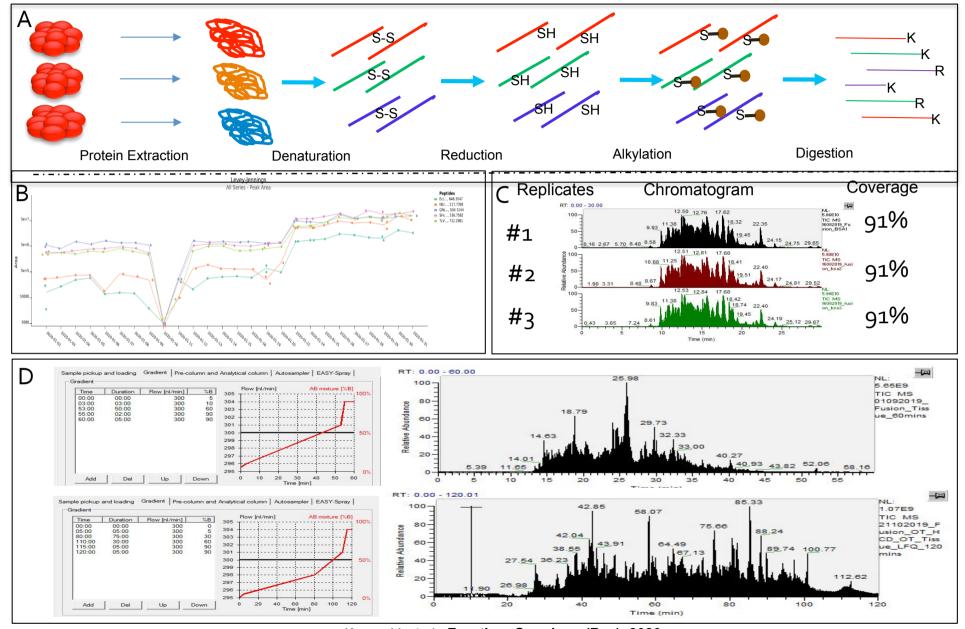
URL: http://www.bio.iitb.ac.in/~sanjeeva/massfiitb/



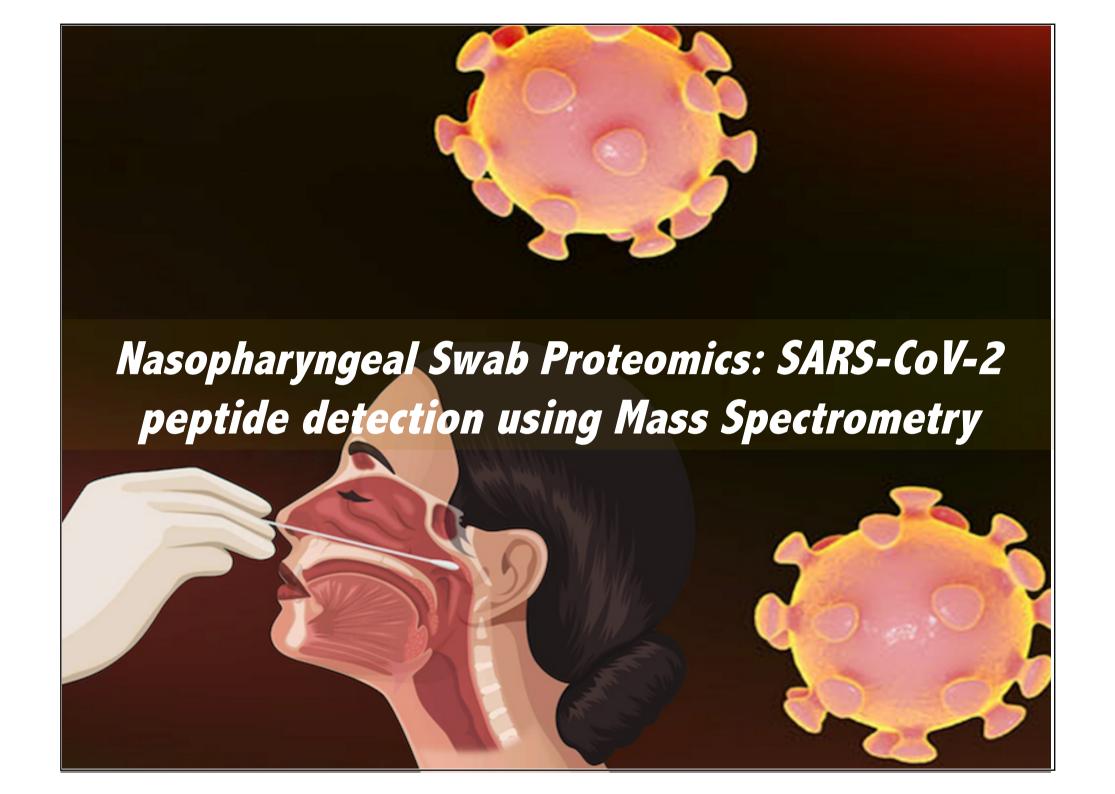




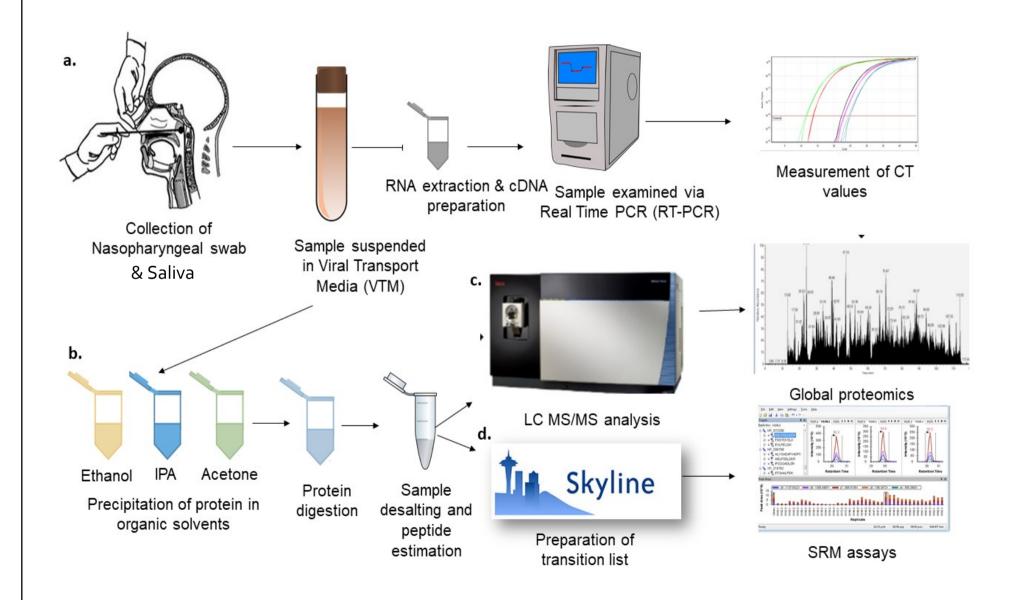
MS-based Proteomics Workflow



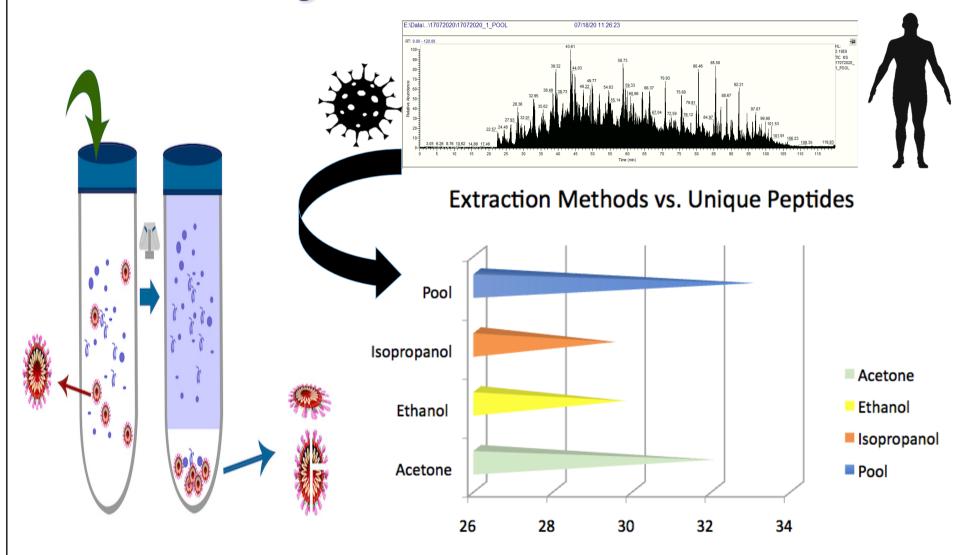
Kumar V. et al.. Frontiers Oncology (Rev), 2020



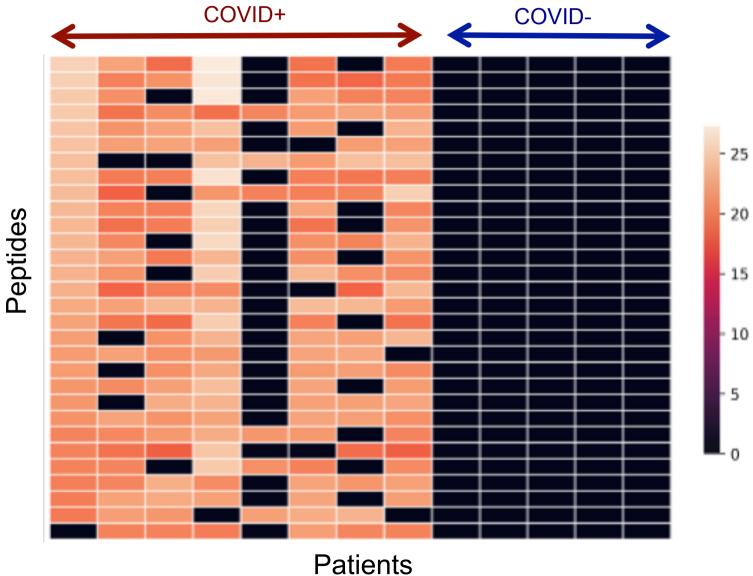
Proteomics Workflow for SARS-CoV-2 Detection From NP Swab



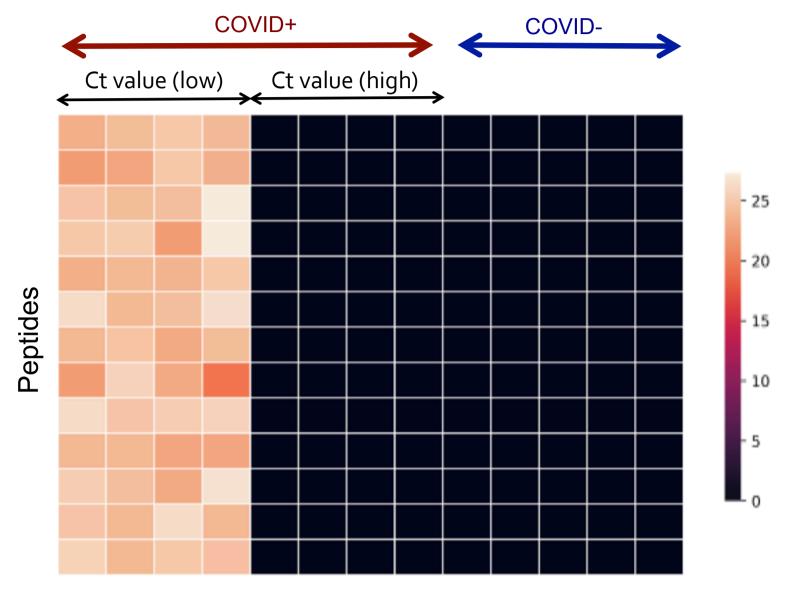
How to capture SARS-CoV-2 Peptides/ Proteins: Need to investigate various extraction methods!



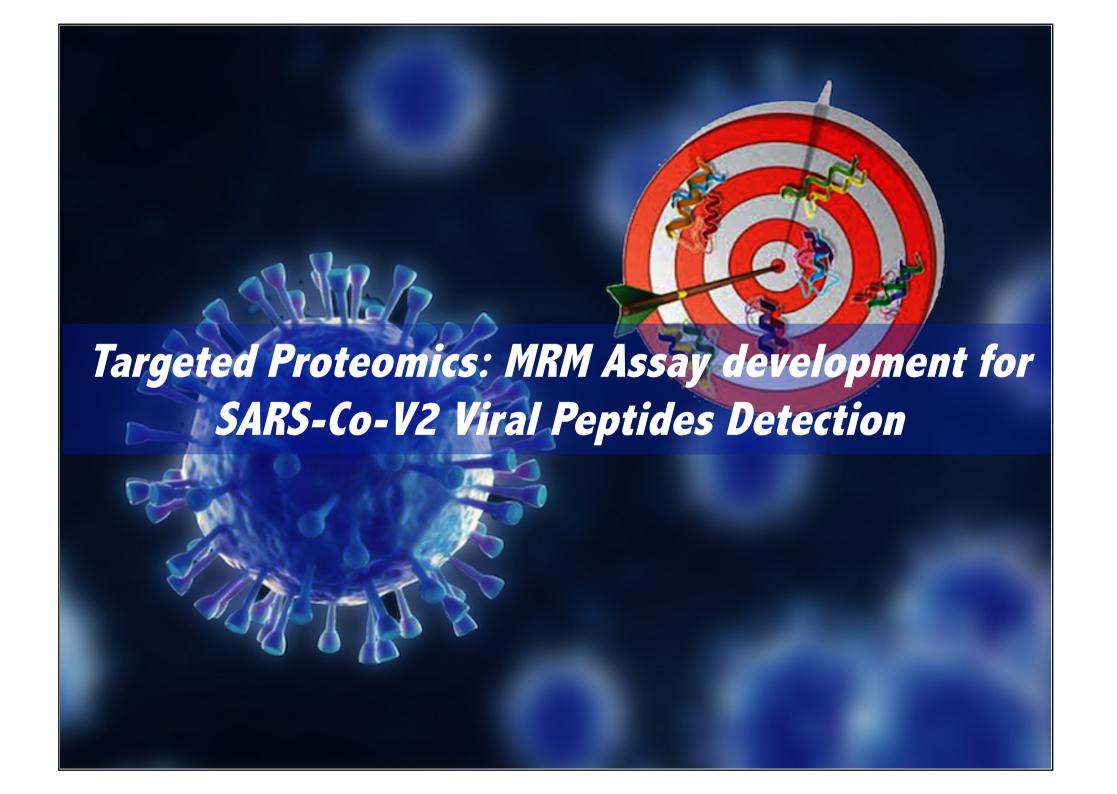
Viral peptides identified from respiratory specimen of COVID-19 patients



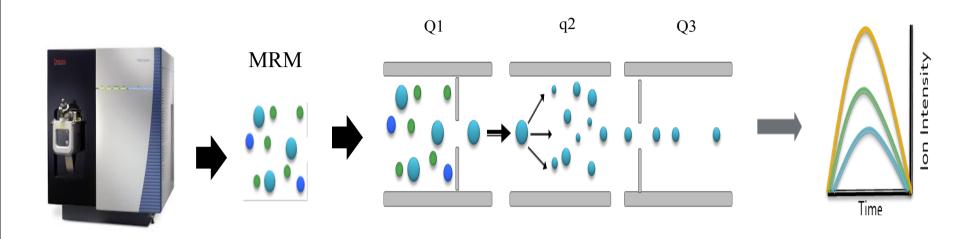
Viral associated peptides correlation with Ct values

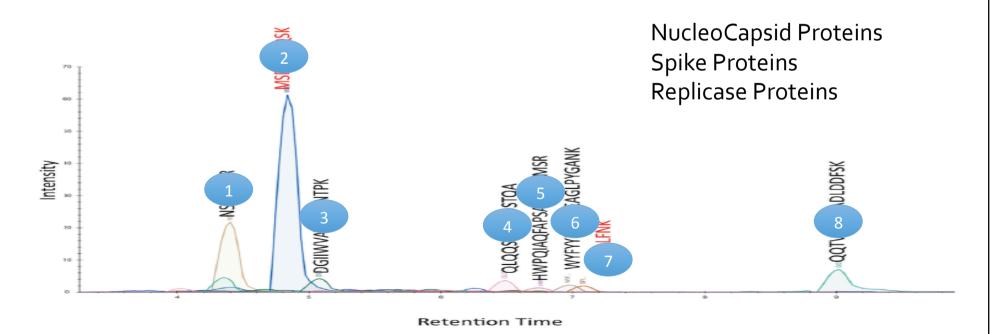


Patients



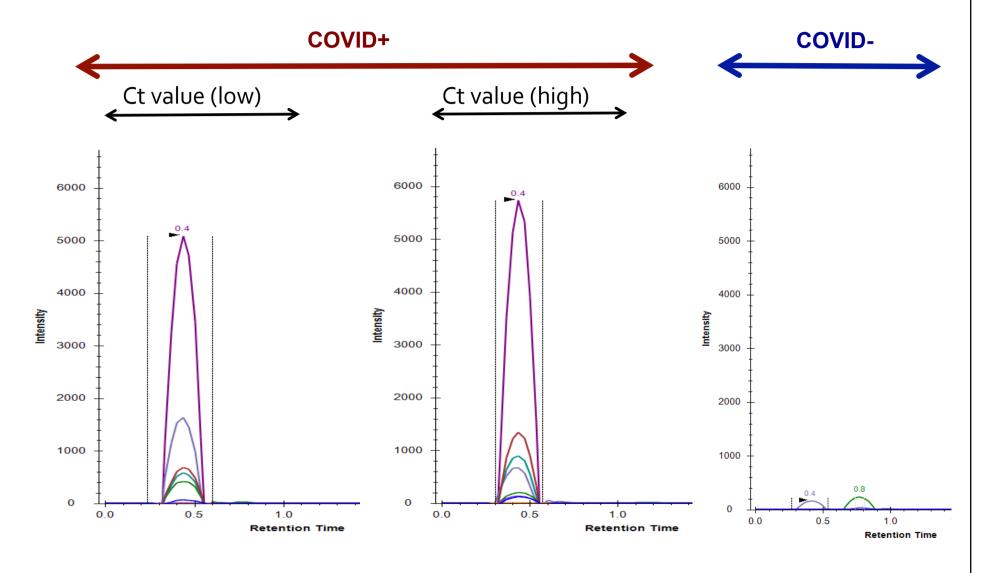
Detection of Viral Peptides Using MRM Assay





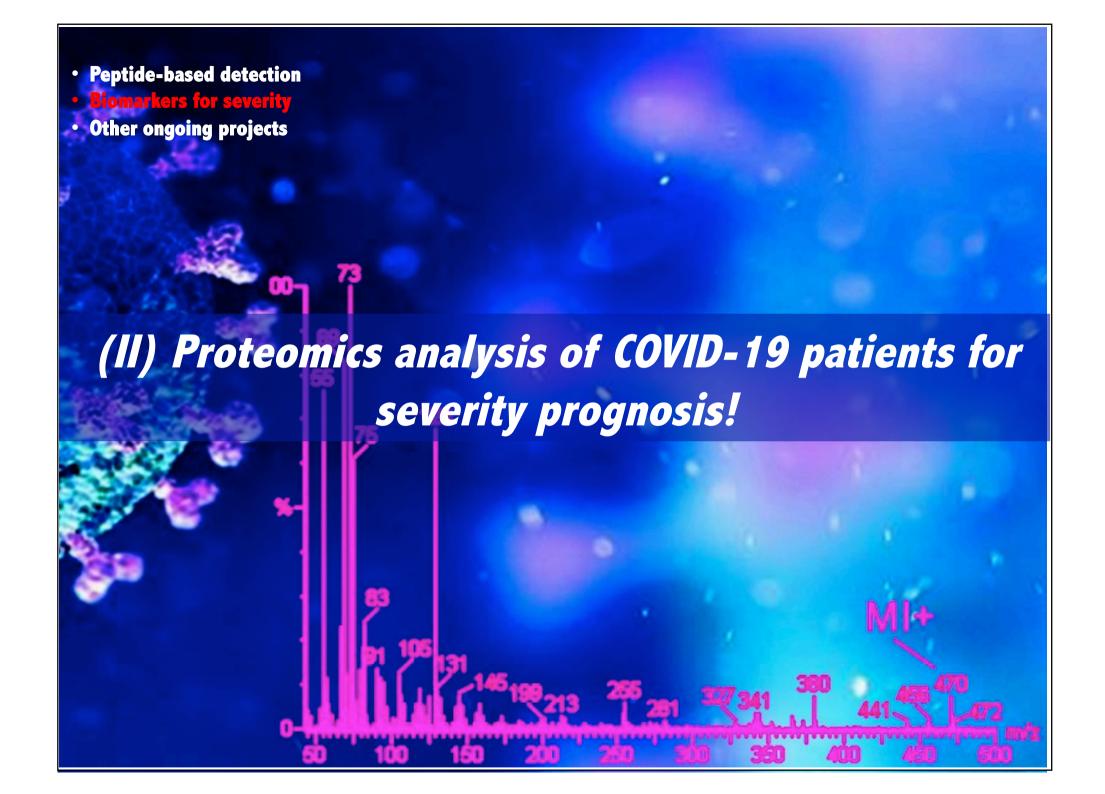
25-JULY-20 COVID-19 IITB-SS 15/26

Development of MRM Assays from NP Swab Samples

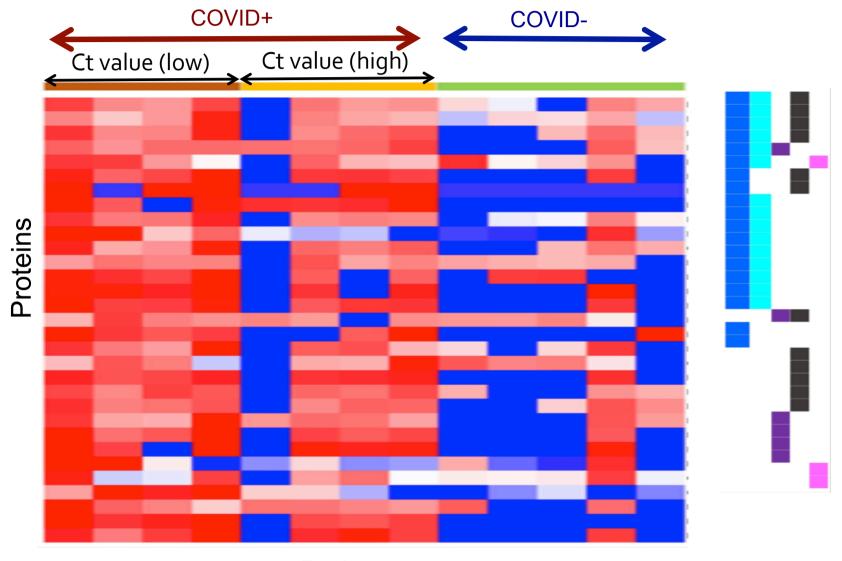


IDF: PAT/BS/SS-9/20-21

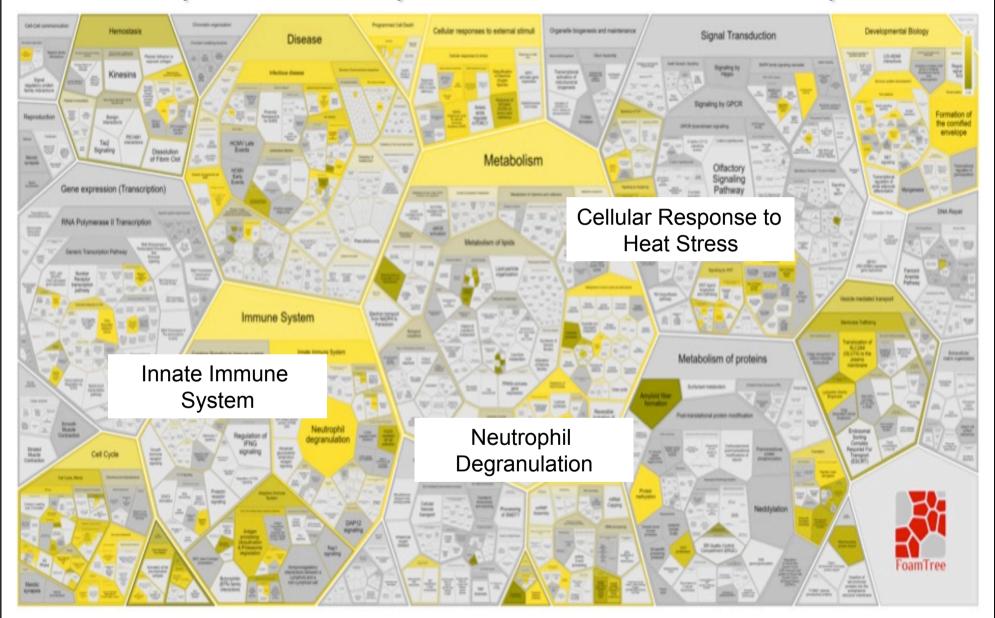


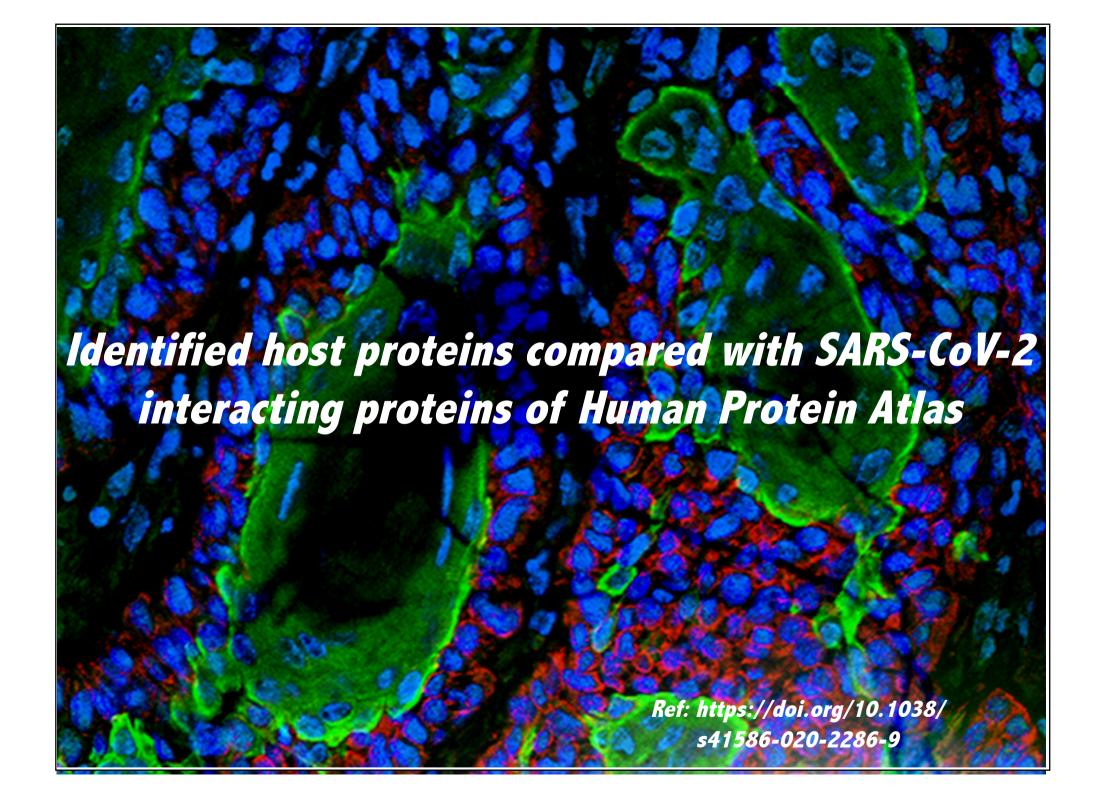




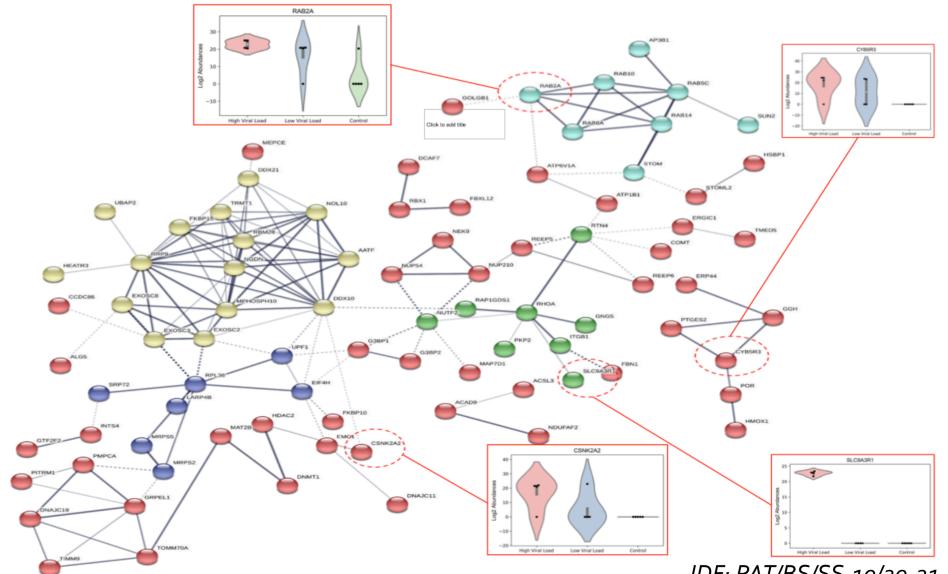


A landscape of Pathways modulated in COVID-19 patients

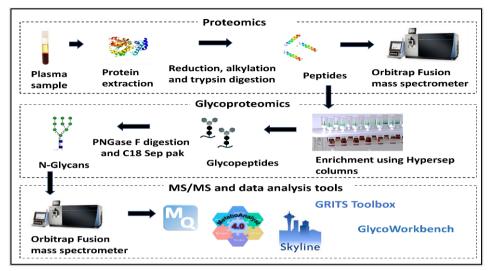




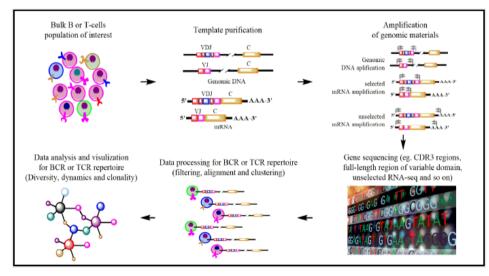
Host-COVID-19 Interaction Map & Biomarkers for **Severity Prognosis**



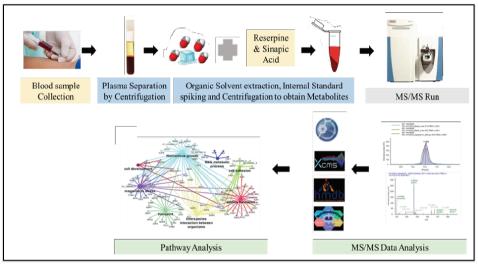
Other initiatives on COVID-19!



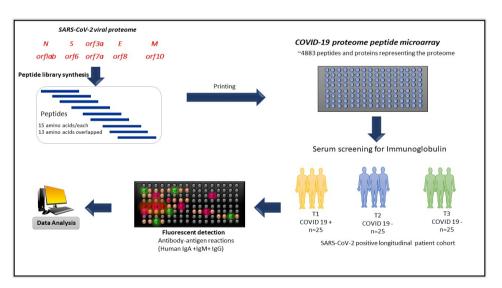
Proteomics investigation – targets for diagnosis



NGS - Immune system repertoire



Metabolomics investigation – therapeutics



Microarrays – epitope mapping & antibody response

Summary:

- Identification of peptides/ proteins that can aid diagnosis of SARS-CoV-2 from the nasal swab (& potentially Saliva) from COVID-19 patients using HT-MS platforms
- Implementation of global & targeted proteomics workflow to identify most significantly altered host proteins for severity prediction and prognostic value
- Other projects...epitope mapping, identification of targets for therapeutics

Thanks to Clinicians & Team of COVID-19 Project



Acknowledgement

Science and Engineering Research Board







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