

RADx-rad: Comparative Analysis of Novel Diagnostic Methods for SARS-CoV-2 Detection and Surveillance

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Comparative Analysis of Novel Diagnostic Methods for SARS-CoV-2 Detection and Surveillance

The RADxSM Radical program (RADx-rad) is advancing non-traditional diagnostics to improve SARS-CoV-2 detection, surveillance, and response. This study compares novel diagnostic methods for detecting SARS-CoV-2 and its variants, including electronic noses (VOCs in breath), chemosensory testing (at-home olfactory assessment), and automatic detection technologies (microfluidic saliva assays for viral and antibody detection). It also evaluates technologies, such as electrochemical sensors, touch-based detection, mechanical detection, and aerosolized virus monitoring, alongside biosensing platforms like smart masks.

Methods are assessed based on analyte targets (spike protein, protease, antibodies), specimen types (saliva, blood, breath), and biophysical techniques. Key performance metrics include sensitivity, specificity, area under the curve (AUC), and limit of detection (LoD), with validation across diverse populations and SARS-CoV-2 variants. Cross-reactivity with common respiratory viruses such as influenza and human coronaviruses is also examined to ensure diagnostic specificity.

This comparative analysis will guide the selection of optimal diagnostics for point-of-care and at-home-testing, improving pandemic response and preparedness for COVID-19 and future infectious diseases. The innovative and complementary diagnostic methods developed by the RADx-rad program offer practical solutions for rapid deployment and scalable impact in real-world infectious disease surveillance.

RADx-rad Research Areas

RADx-rad research areas are diverse and can be grouped into three major research areas:

- Wastewater-based surveillance (7)
- Diagnosis of multisystem inflammatory syndrome in children (MIS-C) (8)
- **Novel diagnostic method development (35)**

- Evaluate the use the NIH RADx Data Hub to access and analyze RADx-rad dataset
- Review the data elements used to describe RADx-rad datasets
- Characterize the diagnostics methods developed by RADx-rad
 - Technology metadata
 - Samples and specimens used for validation
 - Performance metrics
- Demonstrate the diversity of diagnostic methods

Motivation: Traditional Testing Methods have Limitations

- **PCR (Polymerase Chain Reaction)**

- ✓ High sensitivity, detects asymptomatic cases

- ✗ Long turnaround time (24-48 h), lab required, labor-intensive, expensive

- 🩺 Use: Confirm active infection

- **Antigen (Rapid) Test**

- ✓ Fast (15-30 min), easy access, inexpensive

- ✗ Less accurate, false negatives possible

- 🩺 Use: Quick screening

- **Antibody Test**

- ✓ Shows past infection, research tool

- ✗ Not for current diagnosis

- 🩺 Population studies

Novel Diagnostic Method Development 1



Research Topics

- **Electronic Noses:** Developing portable breath analysis and electronic nose technology for COVID-19 detection through volatile organic compounds (VOCs)
- **Chemosensory Testing:** Implementing rapid, self-administered, and at-home chemosensory tests to detect and assess olfactory function and chemosensory loss in COVID-19 patients
- **Exosome-based Technologies:** Utilizing saliva-based and microfluidic technologies to detect SARS-CoV-2, antibodies, and respiratory pathogens, and to predict outcomes in COVID-19 patients through the analysis of exosomes

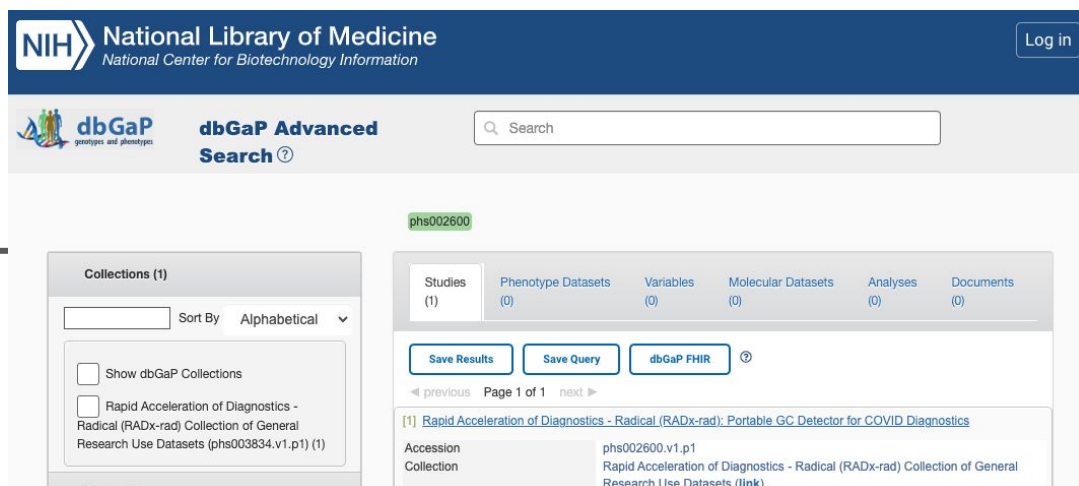
Novel Diagnostic Method Development 2



Research Topics

- **Novel Biosensing:** Developing novel technologies, including home-based diagnostics, and breathalyzers for rapid and sensitive detection of SARS-CoV-2
- **Automatic Detection and Tracing:** Developing technologies for SARS-CoV-2, including privacy-protected contact tracing, electrochemical sensors, touch-based detection, mechanical detection, and aerosolized virus detection
- **Multimodal Surveillance:** Utilizing multimodal data sources and smart technologies for early detection and surveillance of COVID-19 in various populations, including dialysis facilities and through smart masks

1. Select studies in dbGaP or RADx Data Hub



NIH National Library of Medicine
National Center for Biotechnology Information

dbGaP dbGaP Advanced Search

Search

phs002600

Collections (1)

Sort By Alphabetical

Show dbGaP Collections

Rapid Acceleration of Diagnostics - Radical (RADx-rad) Collection of General Research Use Datasets (phs003834.v1.p1) (1)

Studies (1) Phenotype Datasets (0) Variables (0) Molecular Datasets (0) Analyses (0) Documents (0)

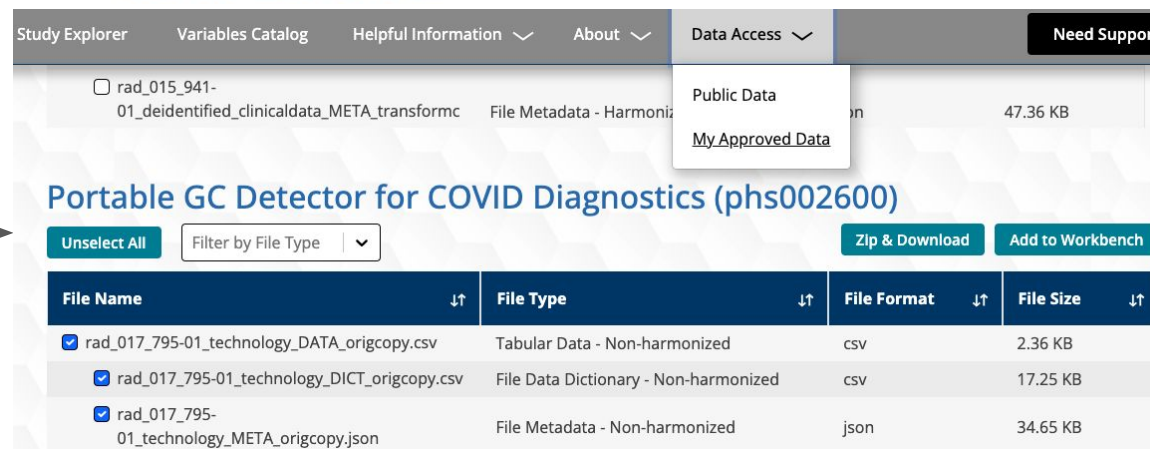
Save Results Save Query dbGaP FHIR

previous Page 1 of 1 next

[1] Rapid Acceleration of Diagnostics - Radical (RADx-rad): Portable GC Detector for COVID Diagnostics

Accession Collection phs002600.v1.p1
Rapid Acceleration of Diagnostics - Radical (RADx-rad) Collection of General Research Use Datasets ([link](#))

2. Apply for data access in dbGaP

3. Select approved datasets & add to Workbench
RADx[®] Data Hub

Study Explorer Variables Catalog Helpful Information About Data Access Need Support?

rad_015_941-01_deidentified_clinicaldata_META_transformc File Metadata - Harmonized 47.36 KB

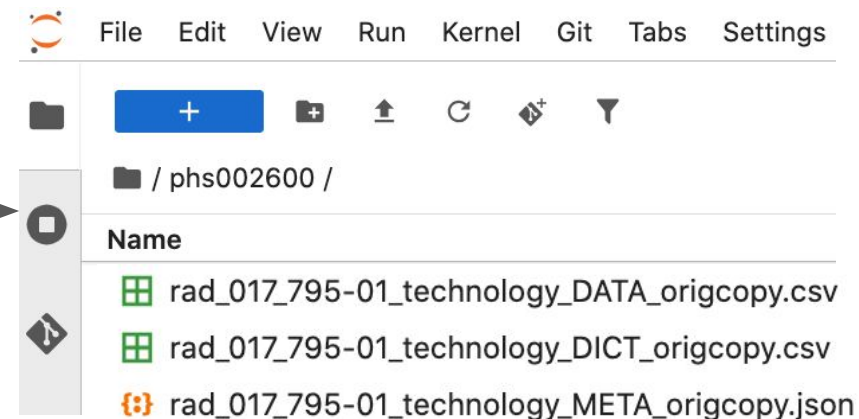
Public Data
My Approved Data

Portable GC Detector for COVID Diagnostics (phs002600)

Unselect All Filter by File Type Zip & Download Add to Workbench

File Name	File Type	File Format	File Size
rad_017_795-01_technology_DATA_origcopy.csv	Tabular Data - Non-harmonized	csv	2.36 KB
rad_017_795-01_technology_DICT_origcopy.csv	File Data Dictionary - Non-harmonized	csv	17.25 KB
rad_017_795-01_technology_META_origcopy.json	File Metadata - Non-harmonized	json	34.65 KB

4. Access datasets in Jupyter Lab



File Edit View Run Kernel Git Tabs Settings

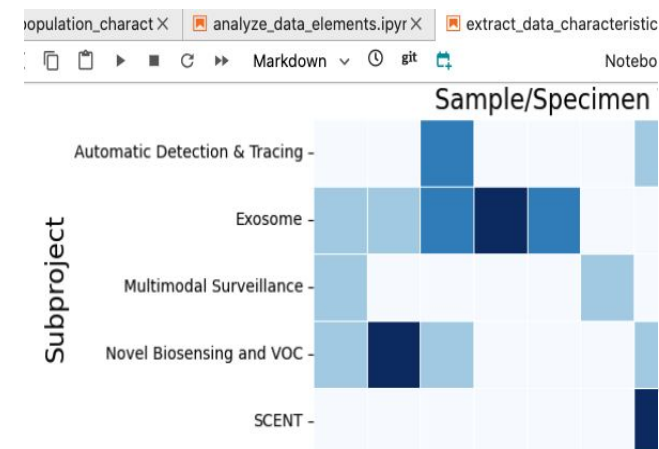
+ + + + +

/ phs002600 /

Name

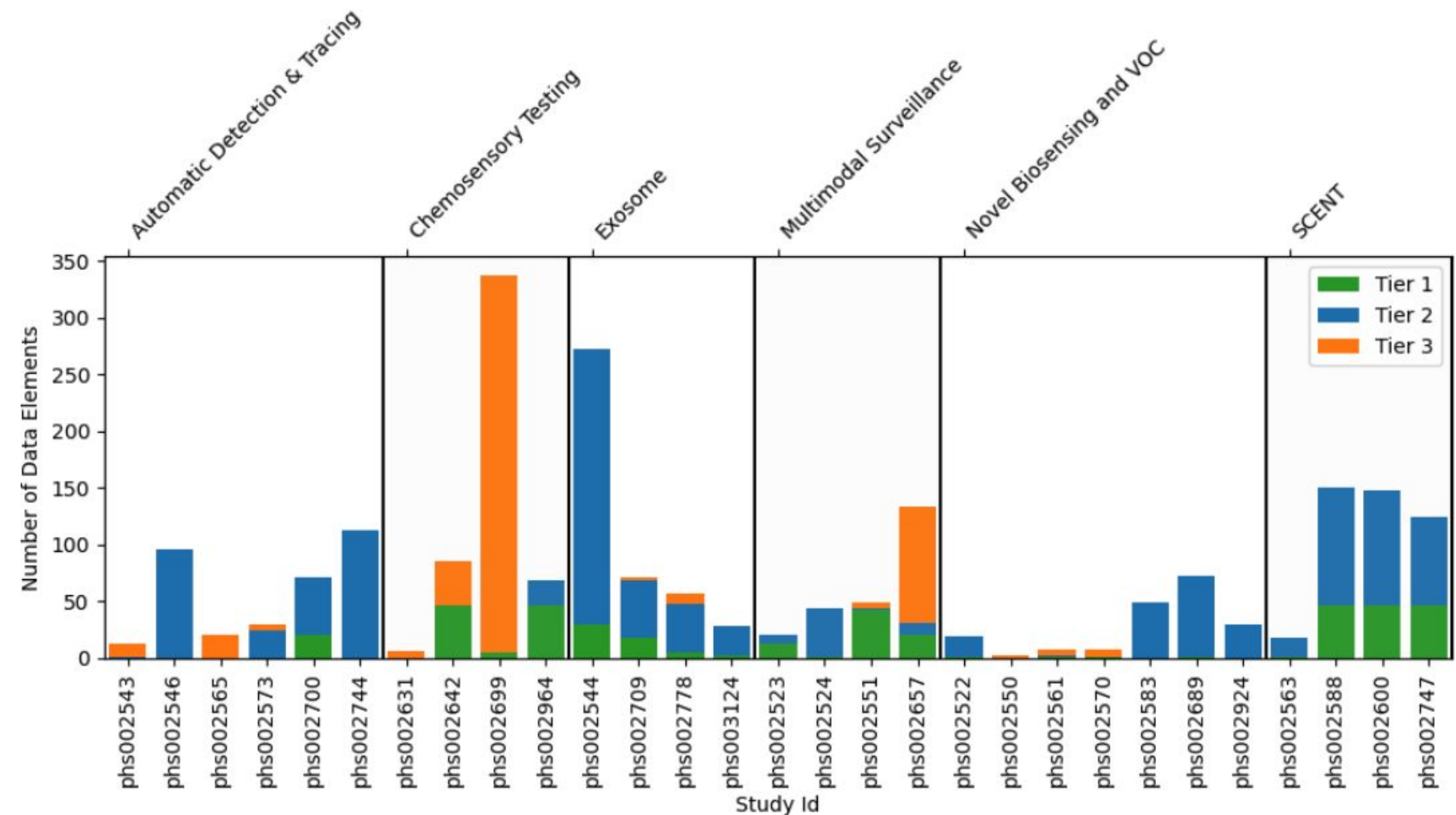
- rad_017_795-01_technology_DATA_origcopy.csv
- rad_017_795-01_technology_DICT_origcopy.csv
- rad_017_795-01_technology_META_origcopy.json

5. Perform Analysis



- Added all RADx-rad datasets to the workspace
- Used the META data files to select the six technology development project-related studies
 - SCENT
 - Chemosensory Testing
 - Exosome
 - Novel Biosensing and VOC
 - Automated Detection & Tracing
 - Multimodal Surveillance

Data Element Harmonization Status

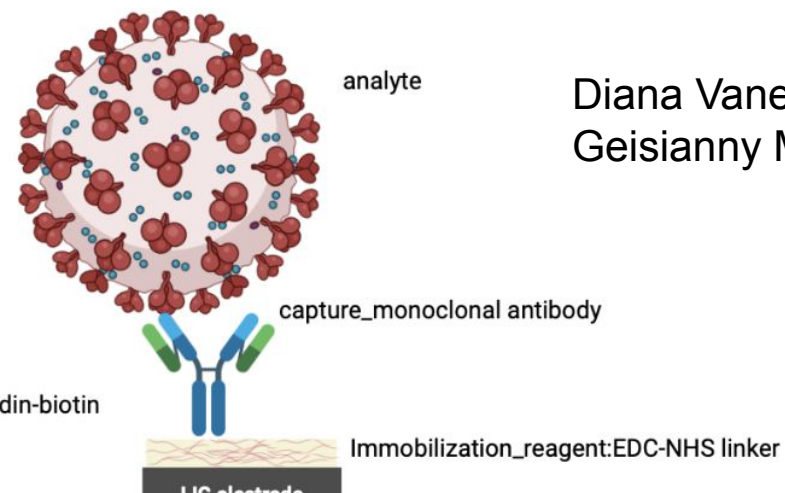
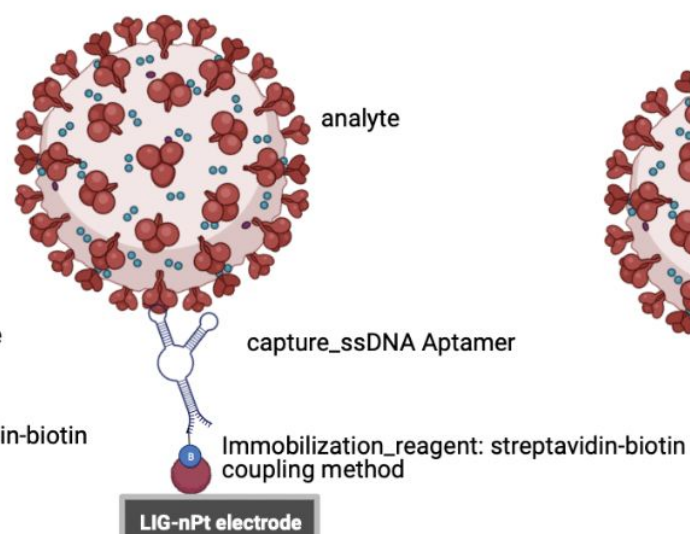
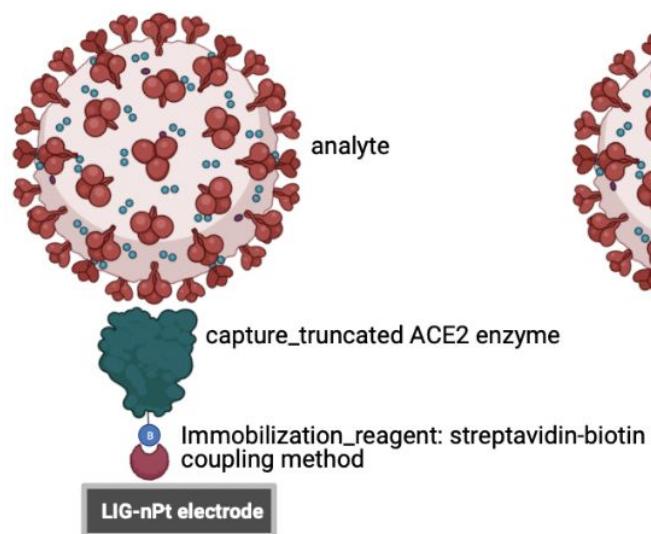


Data Element Tiers

Tier 1: RADx harmonized demographic and medical history data elements

Tier 2: RADx-rad-specific harmonized data elements (used in this study)

Tier 3: RADx-rad study-specific data elements



Diana Vanegas,
Geisianny Moreira

capture_receptor_source
capture_receptor_source_name
capture_receptor_target_organism
...

capture_aptamer_source
capture_aptamer_source_name
capture_aptamer_target_organism
...

capture_antibody_source
capture_antibody_source_name
capture_antibody_target_organism
...

generalization

capture_receptor_region
capture_receptor_region_start
capture_receptor_region_end
...

capture_aptamer_sequence
capture_aptamer_Kd
...

capture_antibody_host_organism
capture_antibody_clonality
...

specialization

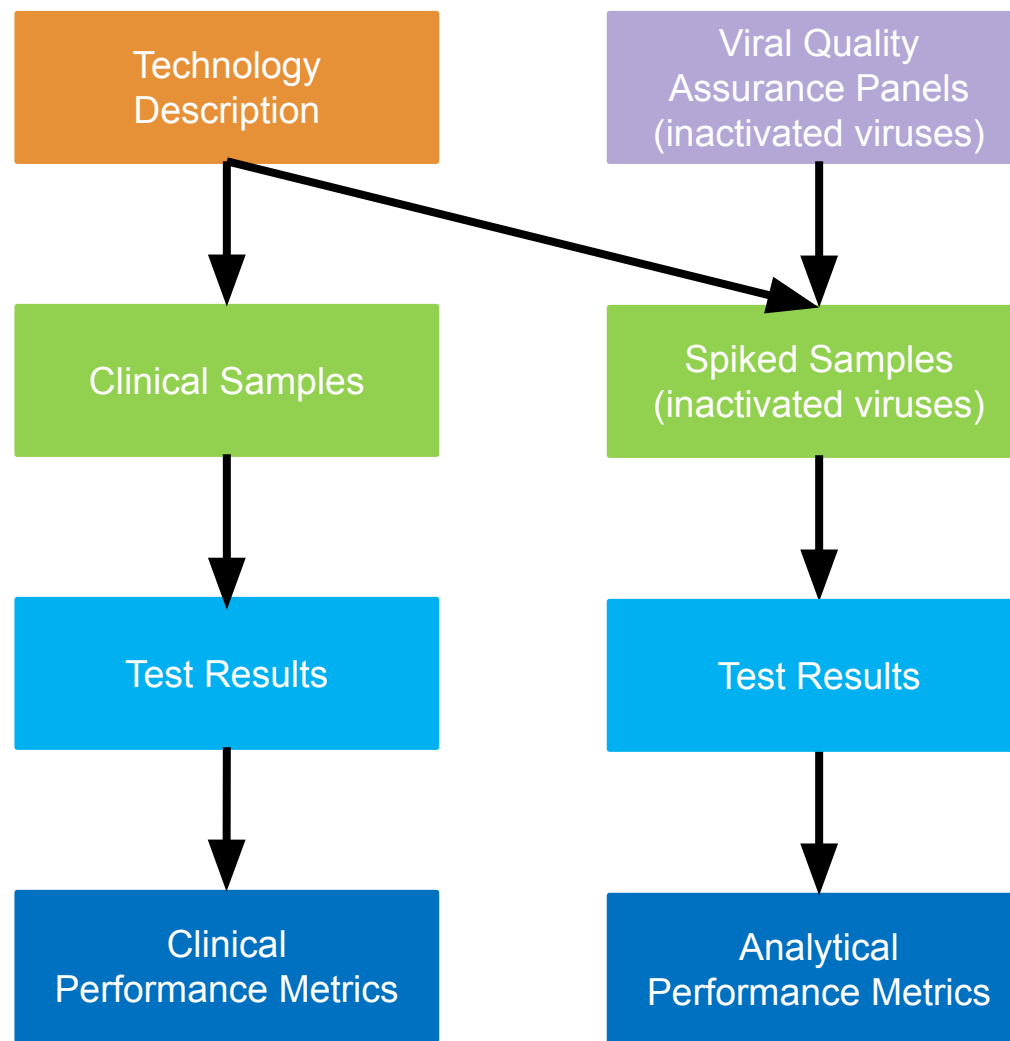
Method Characteristics

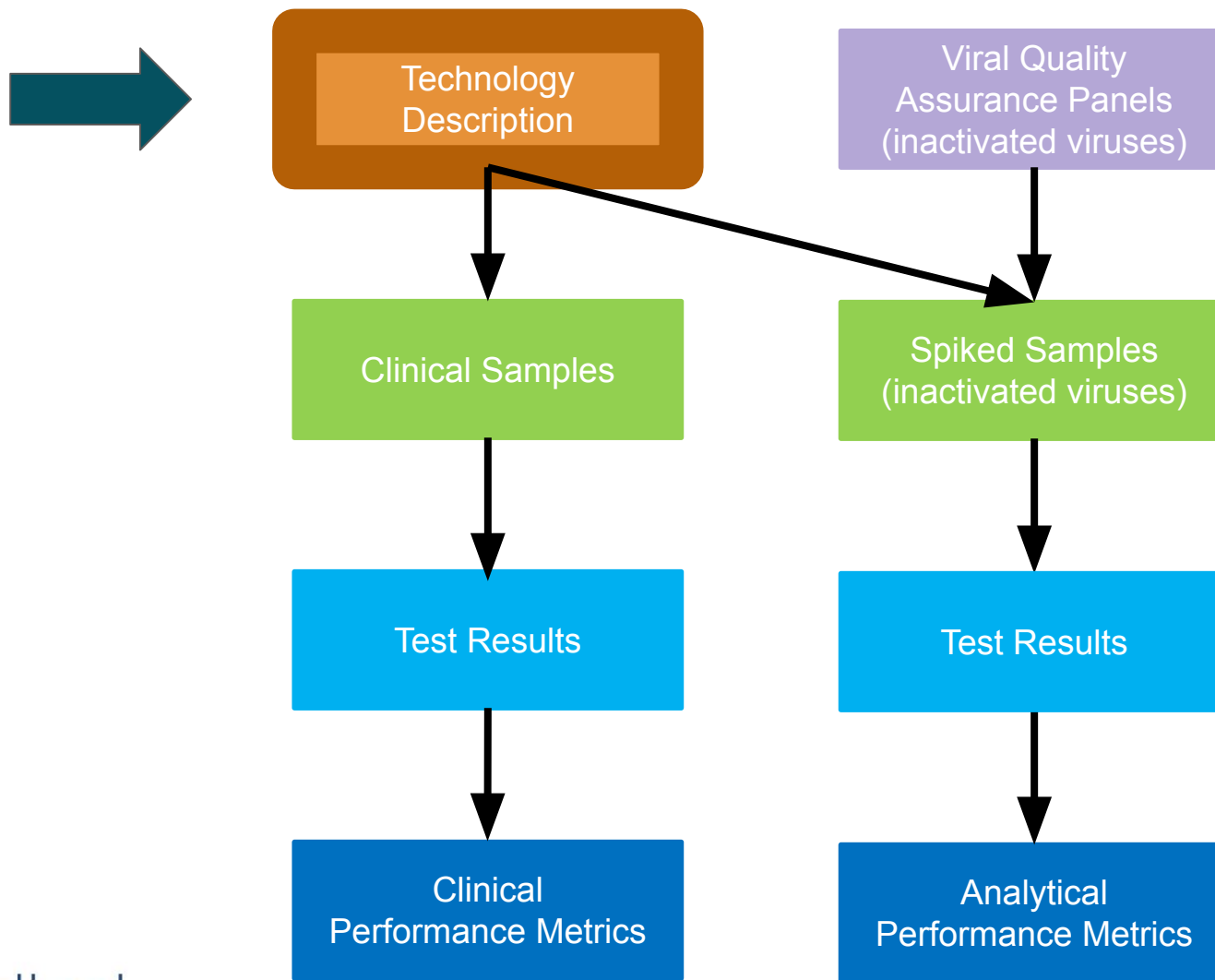
- **Analyte:** Spike protein, protease, antibodies
- **Biophysical Techniques:** SARS-CoV-2/COVID-19 detection methods
- **Specimens Used:** Saliva, breath, odor

Method Validation

- **Variants Tested:** SARS-CoV-2 variants
- **Cross-Reactivity:** Influenza, HCoVs
- **Clinical Testing:** Study demographics
- **Performance Metrics:** Sensitivity, specificity, AUC, limit of detection

Data Organization for Diagnostic Method Development Projects





Technology Description

Aptamer-Antibody Sandwich ELISA

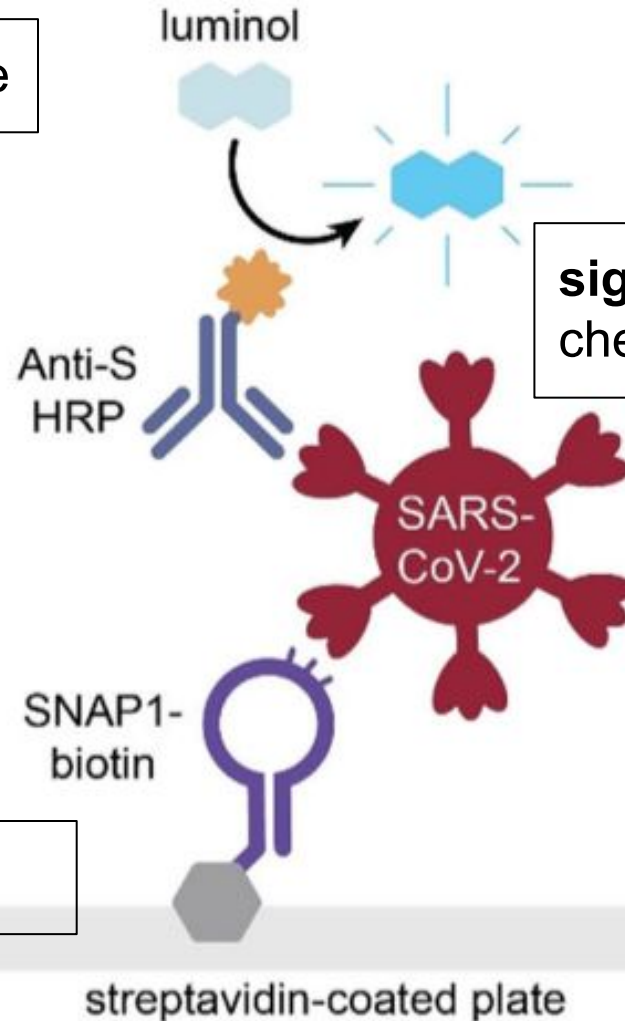
**biorecognition
elements:**

substrate

detector_antibody

capture_apramer

surface: streptavidin-coated plate



technology_platform: sandwich ELISA

technology_description: sandwich
enzyme-linked immunosorbent assay

signal_detection:
chemiluminescent

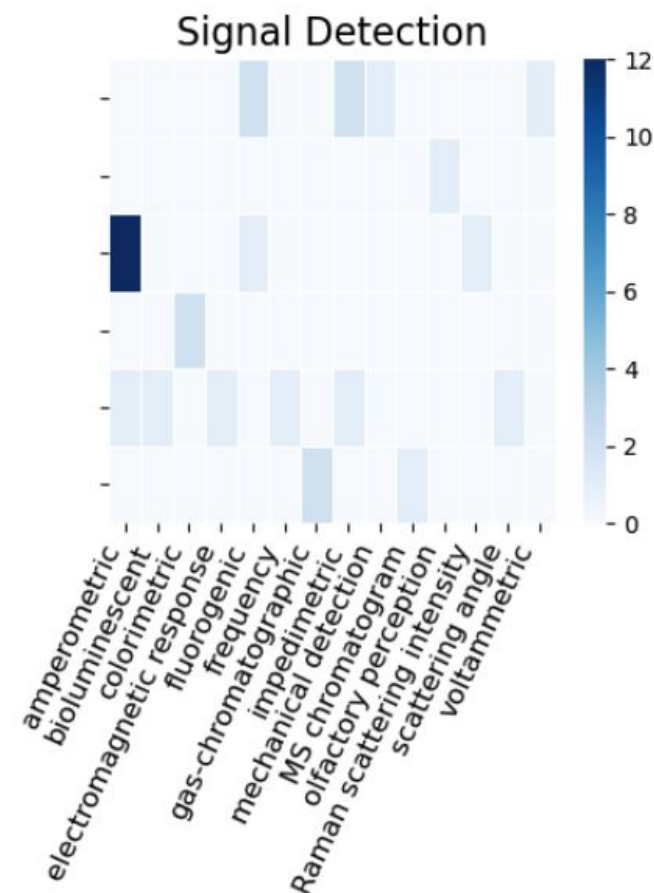
analyte_type: virus

technology_reference:
<https://doi.org/10.1002/anie.202107730>

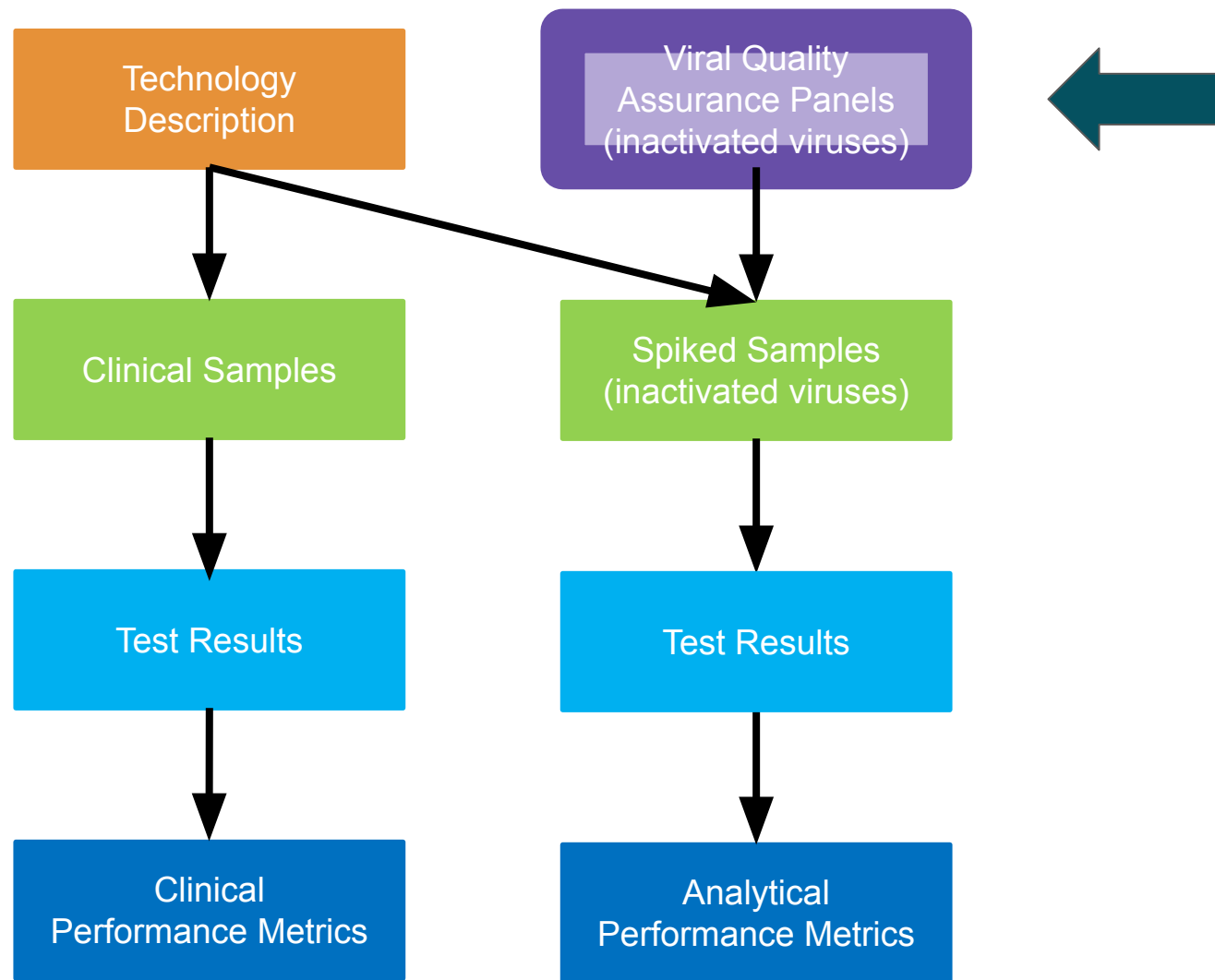
Subproject



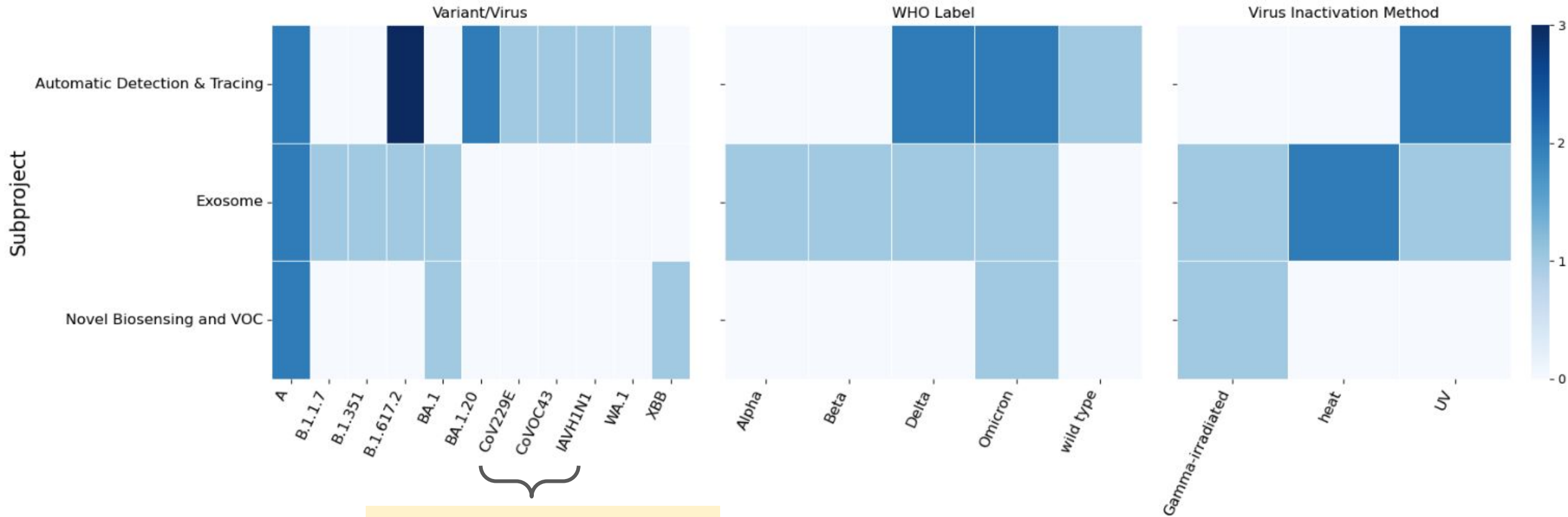
The molecular mechanism or component used to specifically bind to the analyte.



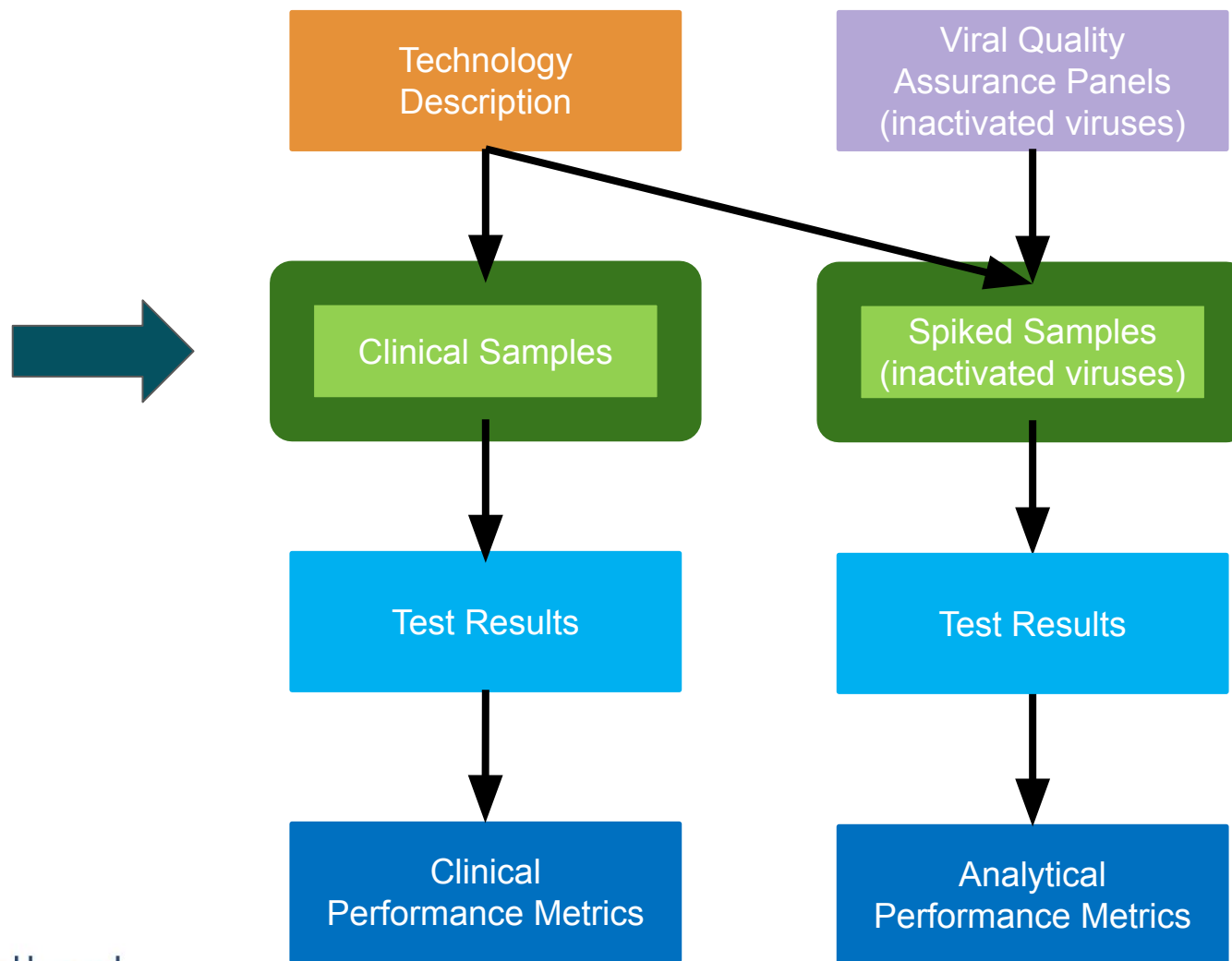
The physical or chemical technique used to quantify the recognition event between analyte and biorecognition element.



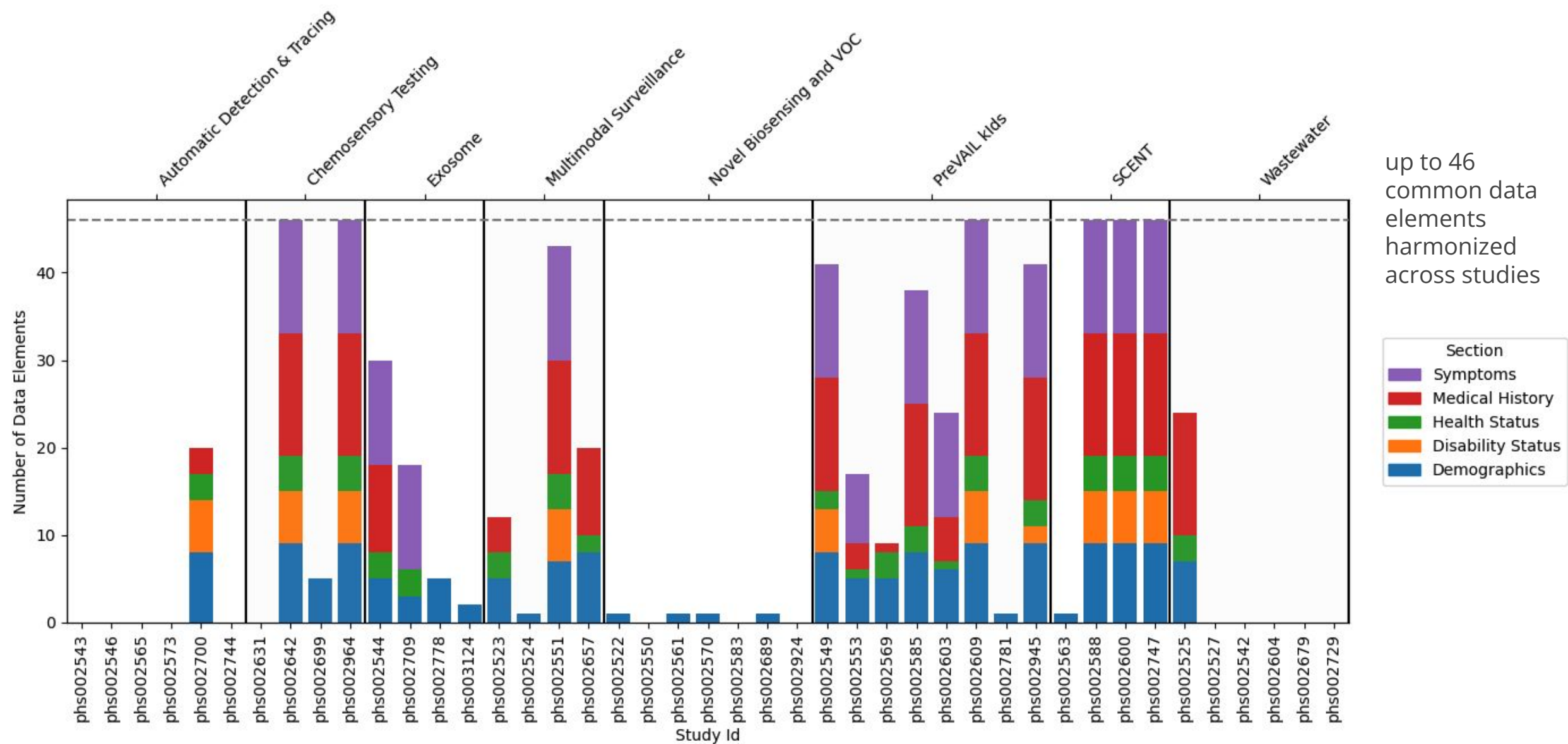
Diverse Variants for Method Validation - Inactivated Viruses



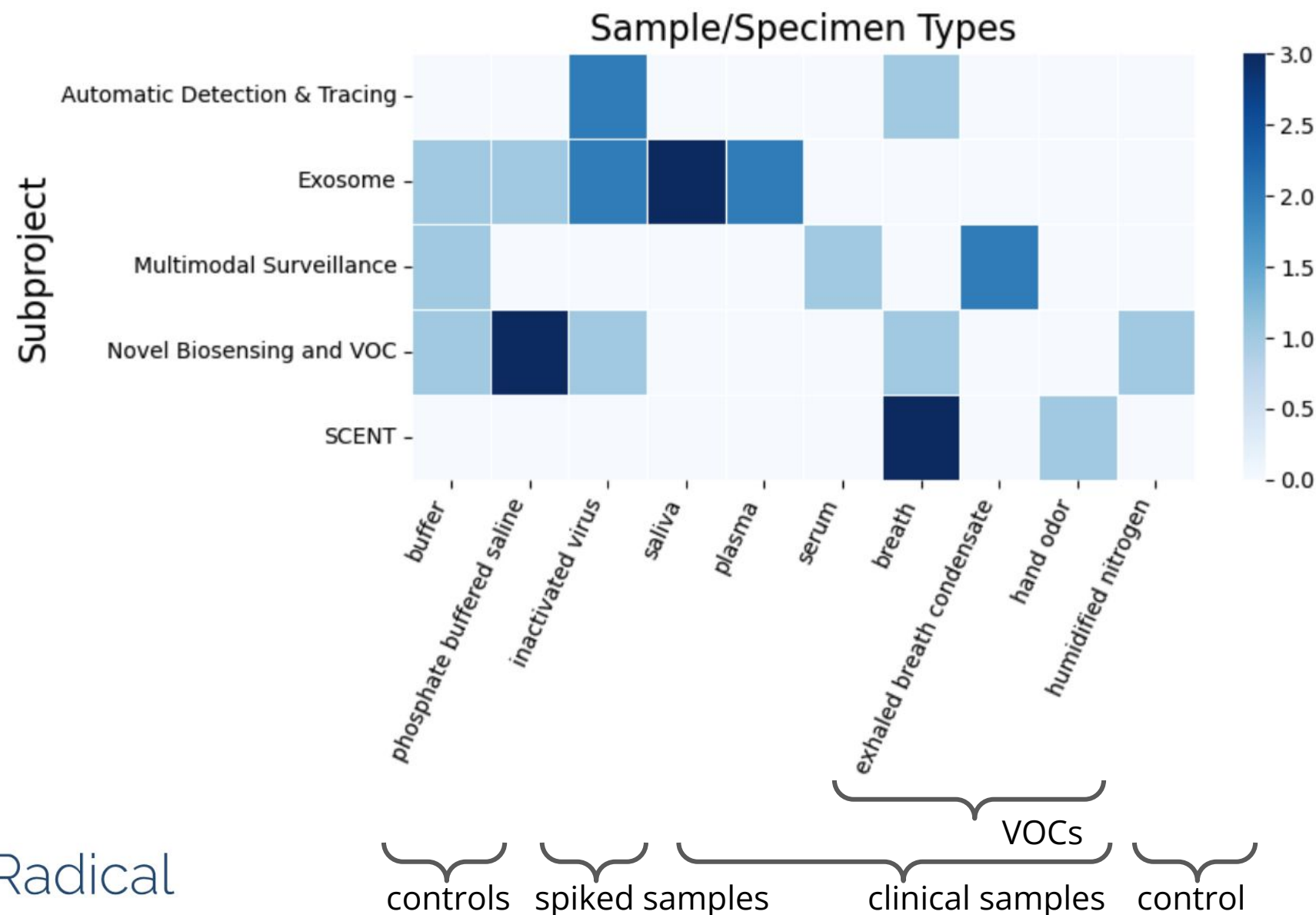
Evaluated cross-reactivity against common respiratory viruses (human coronaviruses, influenza)

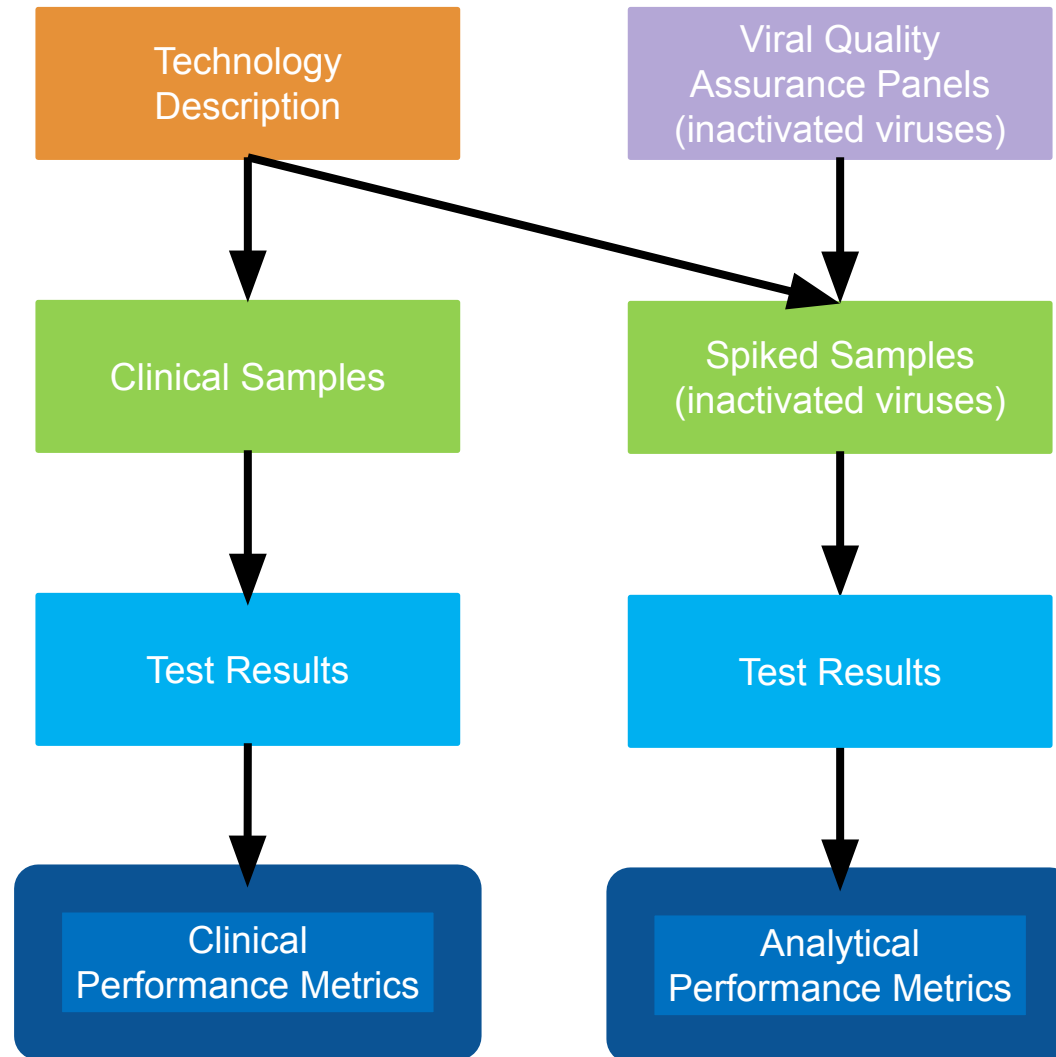


Distribution of Tier 1 Data Elements (clinical samples)

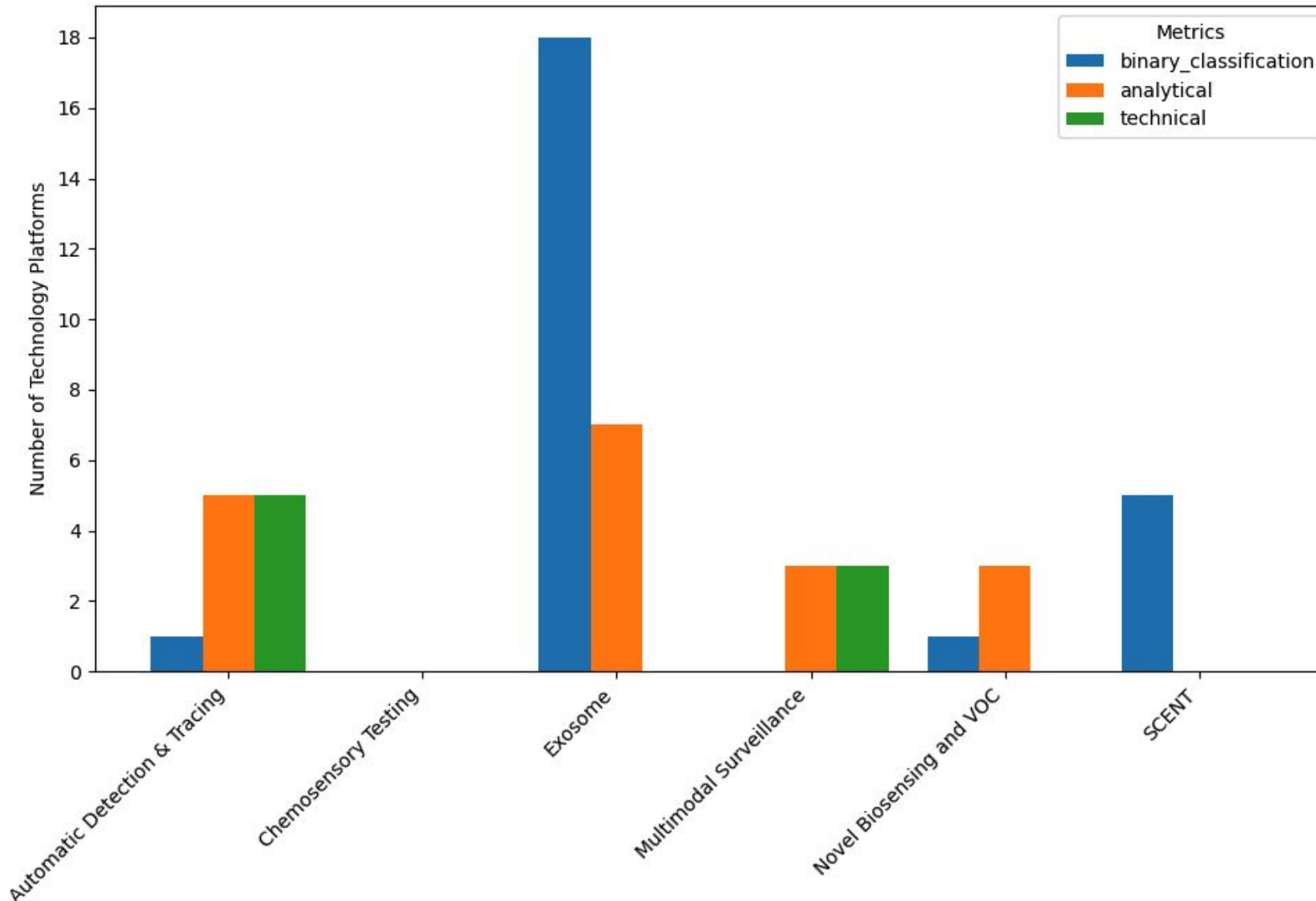


Diverse Sample/Specimen Types for Method Validation





Performance Metrics for Diagnostic Methods



binary classification metrics

sensitivity
specificity
predictive values
percent agreements
AUC
ROC curve

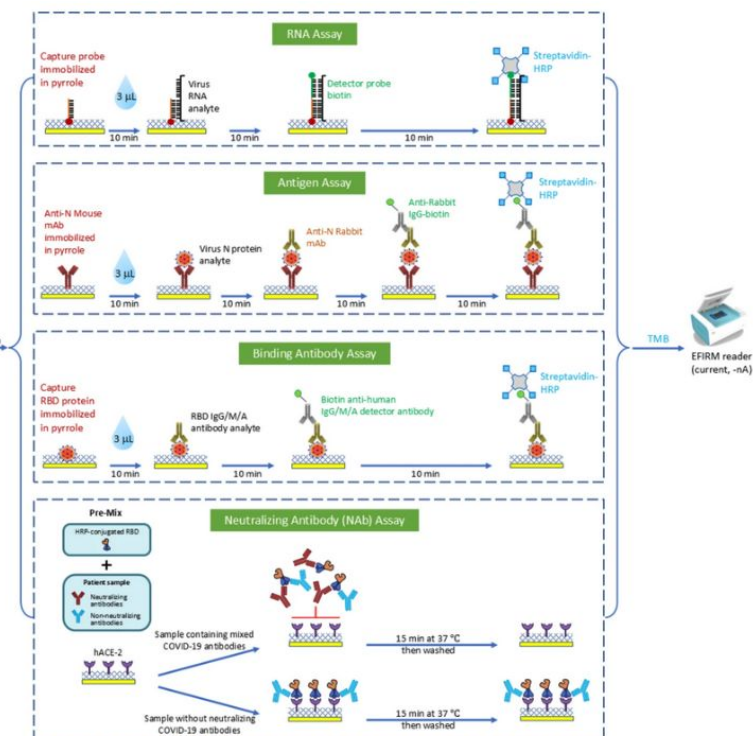
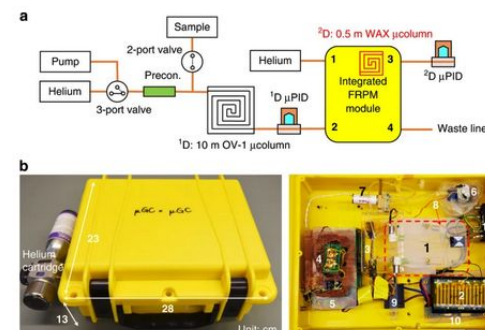
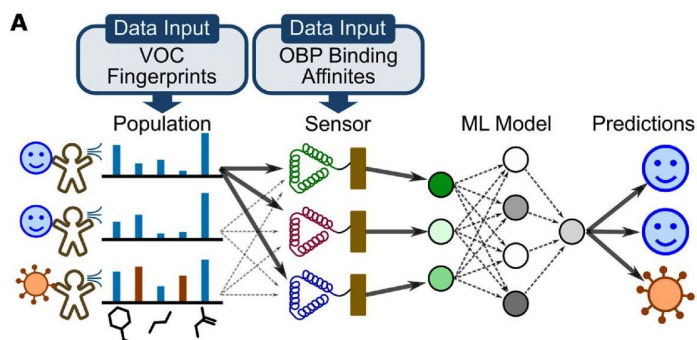
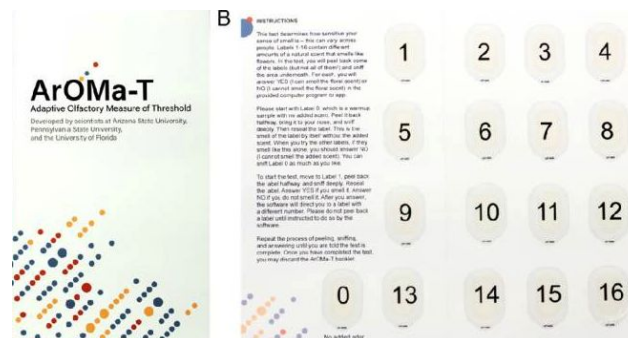
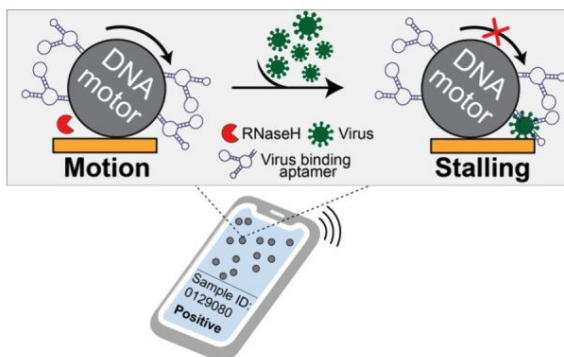
analytical metrics

analytical sensitivity
limit of detection
limit of quantitation
limit of blank

technical metrics

price per test
turnaround time

Examples of Novel Diagnostic Methods



Multimodal Surveillance
doi: 10.1021/acs.analchem.1c02399

- The RADx-rad program delivers cutting-edge diagnostic methods, validated across diverse samples and virus variants.
- Harmonized datasets from the NIH Data Hub support real-world evaluation and AI-driven insights.
- The RADx Data Hub provides tools for querying, analyzing, and machine learning in a secure environment.
- Novel diagnostic technologies from RADx-rad are transforming how we can detect infectious diseases—quickly, accurately, and at scale.
- The program's harmonized datasets provide a foundation for validating and deploying future-ready diagnostic and surveillance systems.

The source code used to analyze the RADx-rad data and instructions how to setup and run the code in the NIH RADx Data Hub are here:

<https://github.com/radxrads/radx-analyzer>