The background of the slide is a dark blue field with a fine, light-colored dot pattern. Overlaid on this are several microscopic images of biological structures. In the top right, there is a large, spherical, multi-lobed structure with many thin, radiating filaments extending from its surface. In the bottom left, there are several elongated, rod-shaped structures, some of which appear to have flagella or other appendages. The overall aesthetic is scientific and high-tech.

Identification of *Streptococcus intermedius* Associated with Culture-Negative Brain Abscesses and Meningoencephalitis through CSF mNGS

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Case Studies in Infectious Diseases

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Disclosures

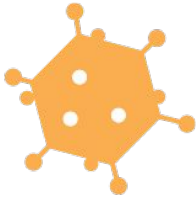
SD, SM: Employee, shareholder, Delve Bio

Meningitis and Encephalitis: Diagnostic and Stewardship Challenges



60%

of cases are due to infectious pathogens



50%

of cases remain undiagnosed



>100

different pathogens linked to ME

Ellul M, *Lancet Infect Dis*, 2022
Glaser CA, *Clin Infect Dis*, 2006
Granerod J, *Lancet Infect Dis*, 2010
Mailles A, *Clin Infect Dis*, 2009
Vora NM, *Clin Infect Dis*, 2017

Empiric antimicrobial use

86% antibiotics

53% antivirals

8% antifungals

Cost per patient

\$64k-260k

ICU level care

22%

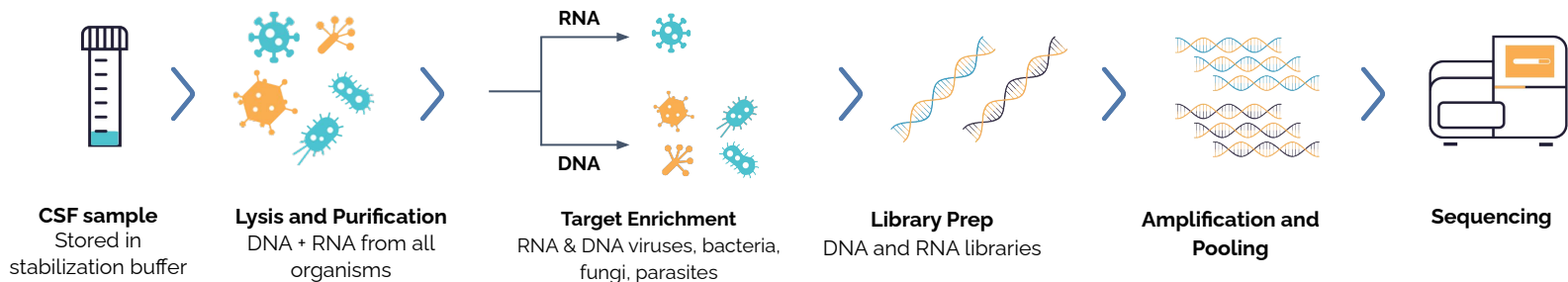
Role of CSF mNGS in CNS infections

- Traditional CSF diagnostics rely on culture, Gram stain, and targeted PCRs
 - May miss fastidious, unexpected, or low burden organisms
- Metagenomic next-generation sequencing (mNGS) provides unbiased detection of all microbial nucleic acids within a sample
- mNGS enables diagnosis of culture-negative CNS infections and supports precision antimicrobial management
- Applications include meningitis, encephalitis, ventriculitis, and brain abscesses with unclear etiology

mNGS Workflow: Laboratory + Bioinformatics

Sample Processing and Sequencing

Optimized for sensitivity, specificity, and contamination-free results



Bioinformatics Pipeline (Delve D-CIDE)

Optimized for speed and removal of non-microbial background



Clinical history

50YO male, previously healthy, limited contact with healthcare

- HPI: 2-week history of progressive altered mental status, ambulatory dysfunction, and two generalized tonic-clonic seizures
- No known immunosuppression or comorbidities
- Social history: Lives in rural Pennsylvania, works in construction, and frequently drinks untreated spring water from a local source
 - Also reports occasional consumption of unpasteurized milk from a nearby farm
- No history of IV drug use, recent dental procedures, or head trauma. No animal exposures or recent travel

Exam and workup

Exam: Afebrile (97.8°F), alert and oriented ×3, mild unsteadiness on gait testing; otherwise normal neurologic and systemic exam

- Labs: WBC $17.8 \times 10^9/L$, normal renal and liver function, mildly low albumin.
- Imaging: MRI revealed multiple peripherally enhancing cystic lesions involving the right frontal and parietal lobes, consistent with brain abscesses
- Microbiology: Stereotactic aspirate showed **Gram-positive cocci**, but no growth on culture after 7 days
- CSF findings: WBC 867 (62% neutrophils, 31% lymphocytes), protein 112 mg/dL, normal glucose; Gram stain and culture negative

Imaging findings

MRI Brain: Multifocal Rim-Enhancing Lesions Consistent with Brain Abscesses

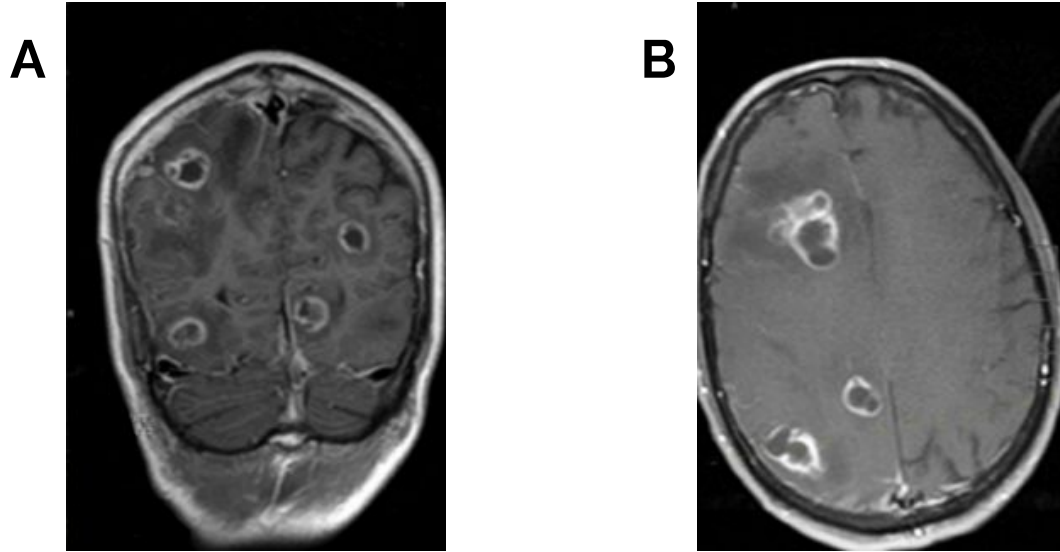


Figure 1. (A) Axial T1 post-contrast MRI showing multiple rim-enhancing cystic lesions involving the right frontal and parietal lobes. (B) Coronal T1 post-contrast MRI demonstrating multifocal abscesses with surrounding vasogenic edema and mass effect.

Workup (continued)

Differential diagnosis:

- Septic emboli
- Nocardiosis
- *Listeria monocytogenes* (history of raw milk/spring water)
- Parasitic CNS infections (cysticercosis, toxoplasmosis)

Broad infectious workup negative: HIV, TB Quantiferon, toxoplasmosis, cysticercosis serologies, and transthoracic echocardiogram (no vegetations)

- Started on broad empiric antibiotics to cover Gram positive, Gram negative, and anaerobic bacteria; anticonvulsants initiated for seizure control

Given persistent culture negativity, CSF was submitted for Delve Detect CSF mNGS testing to investigate fastidious or unculturable pathogens

mNGS results



Table A - Organism Result Summary













| Organism | Total reads | Unique reads | rpM | rpM ratio | Biomass (fg) |
|--|-------------|--------------|------|-----------|-----------------------|
| Streptococcus intermedius (anginosus group) | 21 | 17 | 0.96 | 0.96 | 1.256x10 ³ |

Table B - Sequencing & Host Background Metrics

| Library | Reads PF | % Human | Total biomass (fg) | Control biomass (fg) | Human biomass (fg) |
|------------|------------|---------|-----------------------|-----------------------|-----------------------|
| DNA | 21,863,961 | 99.9456 | 1.308x10 ⁹ | 6.244x10 ⁴ | 1.307x10 ⁹ |
| RNA | 35,213,154 | 99.7029 | 1.118x10 ⁹ | 2.577x10 ⁴ | 1.114x10 ⁹ |

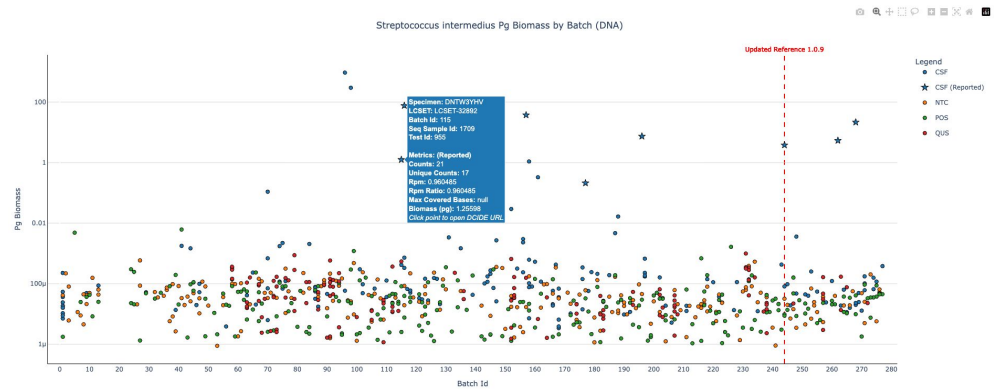
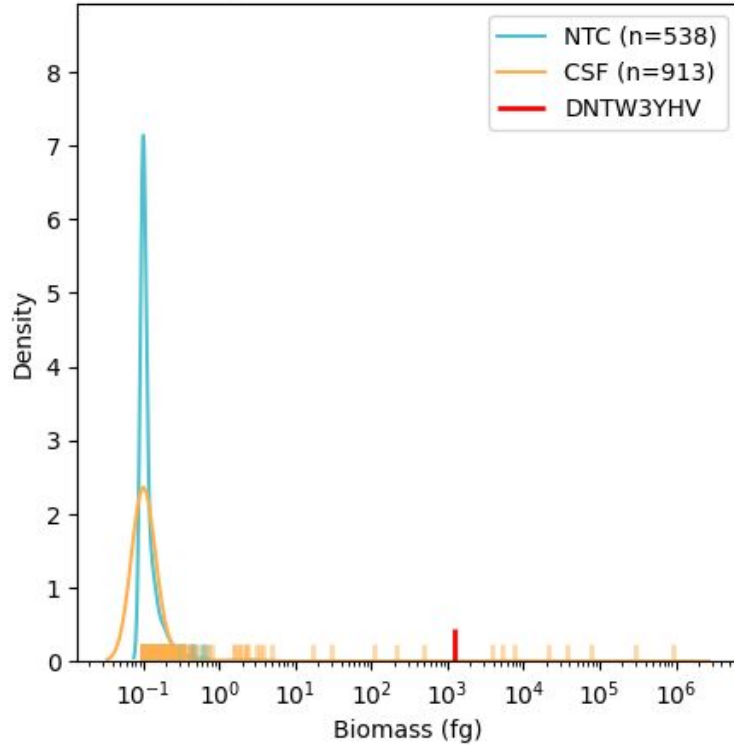
Sequencing metrics (hidden)

⦿ Show raw  

| Total Biomass (pg)   | Control Biomass (pg)   | Human Biomass (pg)   | Organism Biomass (pg)   | Unaligned Biomass (pg)   | Unclassified Biomass (pg)   |
|--|--|--|---|--|---|
| 1,307,650.00 | 62.44 | 1,306,940.00 | 58.37 | 590.07 | 0.84 |
| 11,175.50 | 25.77 | 11,142.30 | 3.31 | 3.04 | 1.08 |

Organism biomass

Streptococcus intermedius - DNA



BLAST summary (top matches)

| Rank | Organism (hit) | Accession | Max score | Query cover | E-value | % identity |
|------|---|------------|-----------|-------------|----------|------------|
| 1 | Streptococcus intermedius B196, complete genome | CP003857.1 | 198 | 100% | 1.00E-46 | 99.09% |
| 2 | Streptococcus intermedius JTH08, complete genome | AP010969.1 | 198 | 100% | 1.00E-46 | 99.09% |
| 3 | Streptococcus intermedius NCTC11324, chromosome 1 | LS483436.1 | 198 | 100% | 1.00E-46 | 99.09% |
| 4 | Streptococcus intermedius FDAARGOS_769, complete genome | CP053999.1 | 198 | 100% | 1.00E-46 | 99.09% |
| 5 | Streptococcus intermedius C270, complete genome | CP003858.1 | 198 | 100% | 1.00E-46 | 99.09% |

Interpretation: Multiple top hits map to *S. intermedius* with 99.09% identity and 1e-46 E-values, supporting species-level identification within the anginosus group

Clinical impact

- mNGS confirmed *Streptococcus intermedius* etiology of culture-negative CNS abscess
 - Supported a monomicrobial etiology without evidence for a mixed infection
- Diagnosis achieved directly from CSF without tissue testing
- Enabled de-escalation to ceftriaxone monotherapy and discontinuation of vancomycin and metronidazole
- Reduced exposure to unnecessary antimicrobials and supported shorter hospitalization

Key takeaways

- CSF mNGS can identify fastidious pathogens in culture-negative CNS infections
 - Interpreting mNGS in high host-background samples requires understanding how relative signal strength reflects organism biomass and confidence of detection
- Provides actionable information for antimicrobial optimization
- Less invasive than brain tissue sequencing; supports precision infectious disease diagnosis
- Integration of mNGS into clinical workflows can reduce morbidity from delayed or indeterminate diagnoses

Thank you!



Delve Detect CSF: Test Overview and Controls

Test Description

- Delve Detect CSF is an mNGS assay for qualitative pathogen detection in CSF from patients with suspected meningitis or encephalitis
- Detects organisms by sequence homology of nucleic acid (DNA & RNA libraries)
- Sequencing on Illumina platform
- Proprietary bioinformatic pipeline for read assignment
- Reports reviewed by board-certified physicians with interpretation and comments
- Results should be considered in the context of clinical history, labs, and imaging

Controls

- Negative control: Synthetic CSF matrix
- Positive control: Synthetic CSF with 7 representative non-pathogenic organisms (DNA & RNA viruses, GN / GP bacteria, fungi, parasites)
- Internal control: DNA and RNA phage spiked into each sample prior to extraction

Clinical use cases for CSF mNGS testing

| | |
|--|---|
| Expedited Diagnosis in High-Risk Patients | <ul style="list-style-type: none">● Critically-ill, ICU● Immunocompromised hosts● Pediatric patients |
| Resolve Atypical Clinical Presentations | <ul style="list-style-type: none">● Suspected atypical or fastidious pathogen● High clinical suspicion with negative conventional testing |
| Optimize Treatment Decisions | <ul style="list-style-type: none">● Antimicrobial de-escalation● Initiation of immunosuppression● Tissue-localized brain infections |
| Public Health and Geographic Medicine | <ul style="list-style-type: none">● Unusual pathogens● Complex exposure history● Outbreak detection and PH investigations |