Infectious Diseases of the Central Nervous System

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Disclosures

• I have no relevant financial relationships to disclose
Learning Objectives

• Identify appropriate molecular testing assays for pathogen(s) of interest
• Interpret molecular testing results in context of histological and clinical findings
• Discuss benefits and limitations of unbiased sequencing assays
Diagnosis of CNS Infections

- Clinical history
- Radiology
- Blood
- CSF
- Brain tissue

Neuropathological Evaluation

- Gross examination
- Intraoperative frozen section and smear
- Routine H&E
- Special stains
- Immunohistochemistry
- *In situ* hybridization
- Electron microscopy
- Molecular testing
Molecular Testing for Infectious Diseases

• Generation of rapid, clinically actionable information
• Expensive to perform compared to conventional microbiology and anatomic pathology laboratory testing
• Selection of appropriate samples and molecular assays is critical to patient care and resource utilization

Image Courtesy of CDC PHIL
Selection of Specimen Types

- Culture isolate
  - High rate of positive testing
  - May take days to weeks for growth
- Fresh/frozen tissue/fluid
  - Can be performed immediately; no fixation interference
  - Prone to contamination
- Formalin-fixed paraffin-embedded (FFPE) tissue
  - Can be screened for organisms/inflammation
  - Decreased sensitivity due to nucleic acid cross-linking
Selection of Molecular Assay

• Single pathogen (e.g. *Toxoplasma gondii*)
  – High sensitivity; automatable with rapid turnaround
  – Requires high degree of suspicion to select correct test

• Targeted panel (e.g. Meningitis/Encephalitis)
  – Conserves specimen volume by testing for most common pathogens associated with specific symptoms
  – Lower sensitivity/specificity; will miss unusual organisms

Rand *et al.* (2011) *J Clin Micro.* 49(7) 2449-2453
Selection of Molecular Assay

• Broad spectrum (e.g. Bacterial 16S rRNA gene sequencing)
  – Provides genus and often species identification for bacteria or fungus
  – Requires interpretation and limited by sequencing databases
• Unbiased (e.g. metagenomic next-generation sequencing)
  – Detects any non-human nucleic acids including novel pathogens
  – High sensitivity results in false positives; more expensive
Case 1

- 32-year-old woman with acute myelogenous leukemia; neutropenia
- Presented with fever, headache, photophobia, blurred vision, right lower extremity pain
- MRI: 2.8 x 1.9 cm ring enhancing lesion with moderate vasogenic edema
PATHOLOGIC DIAGNOSIS:

A. SPECIMEN DESIGNATED "PARIETAL OCCIPITAL LESION" :

Necroinflammatory debris (abscess contents) with Gram-positive bacilli.

NOTE: Organisms highlighted by GMS and PAS-D stains.
Nocardia (mAFB) stain negative.

<table>
<thead>
<tr>
<th>Culture:</th>
<th>Negative</th>
</tr>
</thead>
<tbody>
<tr>
<td>16S rRNA (FFPE):</td>
<td>Bacillus cereus species</td>
</tr>
<tr>
<td>Bacillus spp. IHC (FFPE):</td>
<td>Positive</td>
</tr>
</tbody>
</table>
16S rRNA gene sequencing

- 16S rRNA gene (1500 bp) contains multiple conserved and hypervariable regions
  - V1: nucleotides 69-99 sufficient to identify *Staphylococcus* spp. and *Streptococcus* spp.
  - V2: nucleotides 137-242 sufficient to identify many other species (110+)
- Targeting full length gene vs V1/V2 determined by specimen (culture isolate, fresh/frozen tissue, FFPE tissue) and sequencing method (Sanger vs NGS)
### Histological Findings

<table>
<thead>
<tr>
<th>Group No.</th>
<th>Gram</th>
<th>GMS</th>
<th>AI</th>
<th>WS</th>
</tr>
</thead>
<tbody>
<tr>
<td>1 (n=43)</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>nd</td>
</tr>
<tr>
<td>2 (n=15)</td>
<td>X</td>
<td>+</td>
<td>+</td>
<td>nd</td>
</tr>
<tr>
<td>3 (n=10)</td>
<td>X</td>
<td>X</td>
<td>+</td>
<td>?</td>
</tr>
<tr>
<td>4 (n=10)</td>
<td>X</td>
<td>X</td>
<td>X</td>
<td>?</td>
</tr>
</tbody>
</table>

#### Table 1

<table>
<thead>
<tr>
<th>Group No.</th>
<th>Gram</th>
<th>GMS</th>
<th>AI</th>
<th>WS</th>
<th>16S Positive, No. (%)</th>
<th>Decalcification</th>
<th>Antibiotic Treatment Effect</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td>Total Positive Cases</td>
<td>Yes</td>
<td>No</td>
</tr>
<tr>
<td>1 (n=43)</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>nd</td>
<td>28/43 (65)</td>
<td>4/11 (36)</td>
<td>24/32 (75)</td>
</tr>
<tr>
<td>2 (n=15)</td>
<td>–</td>
<td>+</td>
<td></td>
<td>nd</td>
<td>5/15 (33)†</td>
<td>0/2 (0)</td>
<td>5/13 (38)</td>
</tr>
<tr>
<td>3 (n=10)</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td></td>
<td>0/10 (0)</td>
<td>0/2 (0)</td>
<td>0/8 (0)</td>
</tr>
<tr>
<td>4 (n=10)</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td></td>
<td>0/10 (0)</td>
<td>0/3 (0)</td>
<td>0/7 (0)</td>
</tr>
</tbody>
</table>

†Includes three cases of Gram-negative bacilli (Cardiobacterium hominis, Haemophilus parainfluenzae, and Streptobacillus moniliformis).

AI, acute inflammation; GMS, Grocott methenamine silver; +, positive; -, negative.

Minimum Number of Organisms

**NO (0) ORGANISMS**
0/10 (0%) positive

**RARE (1+) ORGANISMS**
2/3 (66%) positive
1 negative case decalcified
Failed at 1:2 dilution

**ABUNDANT (2+) ORGANISMS**
31/55 (56%) positive overall
Detected to 1:16 dilution
Environmental Contamination

33/78 (42%) samples showed evidence of environmental contamination (including 9/10 negative controls!)

Common Contaminants

- *Meiothermus silvanus*
- *Geobacillus jurassicus*
- *Acinetobacter radioresistens*
- *Sphingomonas hankookensis*
Summary – 16S rRNA sequencing

• Useful for detecting bacteria directly from primary samples including common surgical pathology specimens (e.g. brain abscess, endocarditis)
• Diagnostic yield markedly increased by the presence of organisms on histological review
• Submit frozen tissue (if saved) or FFPE
• Optimal timing usually after routine aerobic cultures are finalized (5 days)
Case 2

- 36-year-old woman with no significant past medical history
- Presented with 5 weeks of headaches unresponsive to triptans
- Lived in Philippines for 6 years during 20s to attend school
- MRI: Right frontal dural based mass, along right frontal convexity with associated edema (no images available)
PATHOLOGIC DIAGNOSIS:

A. SPECIMEN DESIGNATED "LEFT FRONTAL BRAIN MASS" :

Necrotizing granulomas containing AFP-positive organisms involving dura mater consistent with TUBERCULOUS PACHYMENINGITIS

Note: Acid fast bacilli identified by ZN AFB stain.
Gram, PAS-D, and MSS stains were negative.

Culture: Mycobacterium tuberculosis complex
Mycobacterial Molecular Testing

- 16S rRNA gene
  - Single copy, highly conserved
  - Can distinguish most *Mycobacterium* spp.
- hsp65
  - Shorter sequence (134 bp), less conserved
  - Less susceptible to interference from formalin-fixation
- IS6110
  - Multicopy gene (70 bp) present in MTB complex
  - Increased sensitivity compared to non-tuberculous mycobacteria
- Human beta-globin (DNA quality indicator)
AFB (Ziehl-Neelsen) vs Fite-Farcaoo (mAFB) vs Mycobacteria IHC

AFB (Ziehl-Neelsen) vs Fite-Farcao (mAFB) vs Mycobacteria IHC

<table>
<thead>
<tr>
<th>Characteristic</th>
<th>TP, No.</th>
<th>FP, No.</th>
<th>FN, No.</th>
<th>TN, No.</th>
<th>Total, No.</th>
<th>Sensitivity (95% CI), %</th>
<th>Specificity (95% CI), %</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mycobacterial IHC vs culture/PCR</td>
<td>29</td>
<td>25</td>
<td>27</td>
<td>99</td>
<td>180</td>
<td>52 (38-65)</td>
<td>80 (71-86)</td>
</tr>
<tr>
<td>AFB vs culture/PCR</td>
<td>9</td>
<td>8</td>
<td>33</td>
<td>90</td>
<td>140</td>
<td>21 (11-37)</td>
<td>92 (84-96)</td>
</tr>
<tr>
<td>mAFB vs culture/PCR</td>
<td>23</td>
<td>15</td>
<td>15</td>
<td>76</td>
<td>129</td>
<td>61 (43-76)</td>
<td>84 (74-90)</td>
</tr>
<tr>
<td>Mycobacterial IHC vs culture/PCR/AFB/mAFB</td>
<td>46</td>
<td>12</td>
<td>37</td>
<td>172</td>
<td>267</td>
<td>55 (44-66)</td>
<td>93 (89-96)</td>
</tr>
</tbody>
</table>

AFB, acid-fast bacilli; CI, confidence interval; FN, false negative; FP, false positive; IHC, immunohistochemistry; mAFB, modified acid-fast bacilli; PCR, polymerase chain reaction; TM, true negative; TP, true positive.

Mycobacterial Molecular Testing Algorithm

Organisms on mAFB/IHC?  No → Clinical suspicion for MTB?  No → Necrotizing granulomas or AI?

Yes → Yes → Yes → Cultures negative at 1 week?

Yes → Send FFPE for PCR

No → PCR not indicated
Summary – Mycobacterial Testing

- Multi-target sequencing approach can identify MTB and NTM at low concentrations
- Identification of unusual organisms should raise concern for environmental contamination
- Presence of bacilli on mycobacteria IHC, mAFB, or AFB stains correlates with higher yield
- Optimal timing after ~1 week to exclude rapid growers
Case 3

- 74-year-old man with chronic lymphocytic leukemia (on ibrutinib) and AFib presented with impaired speech, nausea/vomiting, and possible seizures
- MRI: irregularly enhancing 2.4 x 2.5 x 2.9 cm lesion in left frontal centrum semiovale with central restricted diffusion and surrounding vasogenic edema
- Also found to have lung nodules (possibly post infectious, recent history of pneumonia)
Review Virtual Slide

https://pathpresenter.net/#/public/presentation/display?token=00b87721
PATHOLOGIC DIAGNOSIS:

B. SPECIMEN DESIGNATED "LEFT FRONTAL MASS" :

Fungal abscess

NOTE: MSS and PAS-D highlight abundant fungal forms involving blood vessels and brain parenchyma, including narrow hyphae with abundant septations and irregular branching, numerous chlamydoconidia, and occasional pyriform conidia. Gram and Fite (mAFB) stains are negative. These findings are morphologically compatible with *Scedosporium* or *Lomentospora* sp., while *Fusarium, Candida, Aspergillus*, and other hyaline molds remain lower on the differential.

Culture: Positive for mold
Sequencing (TUB, CAL genes): *Scedosporium boydii* (member of *Scedosporium apiospermum* complex)
Laboratory Testing

• Serology
  – Antigen/antibody detection
    • *Cryptococcus*
    • *Histoplasma*
    • *Blastomyces*
    • *Coccidioides*
  – Fungal wall components
    • Galactomannan
    • (1,3)-β-D-glucan

Filamentous Fungi

- Broad ribbon-like hyphae, pauciseptate, 90-degree angle branching
  - Mucorales order: *Mucor, Rhizopus, Absidia, Cunninghamella*
    - Galactomannan – (1,3)-β-D-glucan –
  - Galactomannan + (1,3)-β-D-glucan +

- Uniform septate hyphae that branch at acute angles
  - *Aspergillus*
  - *Scedosporium, Fusarium, etc.*
    - Galactomannan – (1,3)-β-D-glucan–
    - Galactomannan + (1,3)-β-D-glucan +
    - Galactomannan – (1,3)-β-D-glucan + *(Cultures/Molecular)*

- Hyphae, pseudohyphae, Gram-positive yeast
  - *Candida*
  - Galactomannan – (1,3)-β-D-glucan–
  - Galactomannan + (1,3)-β-D-glucan +

- Pigmented hyphae, yeast (Fontana Masson-positive)
  - Phaeohyphomycosis: *Alternaria, Bipolaris, Curvularia, Exophiala, etc.*
Yeast Forms in Tissue

Small (2 to 4 μM)
- *Candida* (Gram positive)
- Chromoblastomycosis (pigmented)
- *Paracoccidioides* (Ship’s wheel)
- Lobomycosis (chains of yeast)
- *Penicillium* (“drug capsule”)
- Sporotrichosis (asteroid bodies)

Narrow budding
- *Histoplasma*
- *Pneumocystis* ((1,3)-β-D-glucan)
- *Coccidioides* endospores
- Bacteria post-treatment

Non-budding
- *Cryptococcus* (Mucicarmine, Fontana-Masson)

Large (4 to 20 μM)
- Narrow budding

Broad Based budding
- *Blastomyces*

Spherules w/ endospores
- *Coccidioides*

Very Large (20 to 100+ μM)
- *Other: Rhinosporidium Chrysosporium*

Other:
- *Histoplasma*
- *Coccidioides* endospores
- Bacteria post-treatment
Fungal Sequencing

- Species directed PCR
  - e.g. Cryptococcus spp.
  - Higher sensitivity; requires high clinical suspicion
- Broad spectrum
  - 18S
  - ITS
  - D1/D2

http://sites.biology.duke.edu/fungi/mycolab/primers.htm
Summary – Fungal Sequencing

• Very useful for identification of slow growing molds or species difficult to identify based on morphology
• Identification of unusual organisms should raise concern for environmental contamination, but may also represent disease in immunocompromised patients
• Tissue specimens without observable organisms are usually false positives
• Timing depends on clinical urgency; typically after 1-2 weeks if cultures negative or unable to further classify
Case 4

- 65-year-old man with history of grade 2 follicular lymphoma (on rituximab) and occasional tick bites
- Presented with fever and subsequently developed slurred speech followed by neck stiffness
- LP: Normal glucose, elevated protein, mix of lymphocytes and neutrophils
- MRI: Cerebellum with diffuse swelling, obstructive hydrocephalus, leptomeningeal enhancement; additional signal abnormality within midbrain and thalami
POWASSAN ENCEPHALITIS

mNGS (CSF): POWV (DTV, lineage II)
IgM (blood): negative
IgM (CSF): negative
PCR (serum): POSITIVE
PCR (CSF): POSITIVE
PCR (FFPE brain): POSITIVE
POWV IHC (FFPE brain): POSITIVE
CNS Viral Infections

• Morphological Diagnosis of Viral Infections
  • Adenovirus
  • Herpes Simplex Virus 1 and 2 (HSV-1, HSV-2)
  • Varicella Zoster Virus (VZV; HHV-3)
  • Epstein-Barr Virus (EBV; HHV-4)
  • Cytomegalovirus (CMV; HHV-5)
  • Polymavirus (i.e. JC and BK)
  • Measles Virus
  • Rabies Virus

• Viral Infections Requiring Ancillary Tests
  • Mumps Virus
  • Poliovirus
  • Rubella
  • Enterovirus
  • Human T-Lymphotropic Virus Type 1 (HTLV-1)
  • Human Immunodeficiency Virus (HIV)
  • Influenza Virus
  • Parainfluenza
  • Arboviruses (i.e. West Nile, Eastern Equine, Powassan, Zika)
  • Hendra virus and Nipah virus
  • Lymphocytic Choriomeningitis Virus (LCMV)
  • HHV-6 and HHV-7
Metagenomic next-generation sequencing (mNGS)

• Advantages
  – Single test
  – Unbiased
  – Moderate turn around time

• Limitations
  – Relatively expensive
  – Low sensitivity
  – Requires sufficient quantity of pathogen nucleic acid

Summary – mNGS

• Powerful tools for identification of pathogens in cases with broad infectious differential (e.g. viral meningoencephalitis)
• Can identify novel pathogens, but must distinguish from environmental contamination (varies with laboratory)
• Relatively expensive but could provide cost savings if ordered instead of large panel of molecular tests
• Turn around time improving with better informatics pipelines; interpretation still required
Case 5

- 56-year-old man with untreated HCV, alcoholic liver disease, interstitial lung disease
- Present with progressive cognitive decline over several days without fever or meningismus
- MRI: Multiple large ring enhancing lesions (up to 3 cm) with extensive vasogenic edema and local mass effect
PATHOLOGIC DIAGNOSIS:

A. SPECIMEN DESIGNATED "RIGHT FRONTAL TUMOR":

Abscess with TOXOPLASMA GONDII organisms identified.

Toxoplasma IGG EIA: 5.57 POSITIVE (Suggestive of infection at undetermined time)
HIV ELISA test positive
Parasitic Infections

Table 5.2 Principal Protozoa Responsible for Human Infections

1. Amoebiasis
   a. Entamoeba histolytica: cerebral amoebic abscesses
   b. Primary amoebic encephalitis
      i. Primary amoebic meningoencephalitis (Naegleria fowleri)
      ii. Granulomatous amoebic encephalitis (Acanthamoeba spp. and Leptomyxid)
      iii. Acanthamoeba keratitis

2. Cerebral malaria (Plasmodium falciparum infection)
3. Toxoplasmosis (Toxoplasma gondii infection)
4. Trypanosomiasis
   a. African trypanosomiasis (Trypanosoma brucei spp.)

Table 5.3 Major Helminthic Infections of the CNS

1. Cestodes
   a. Neurocysticercosis (Taenia solium)
   b. Hydatid cyst (Echinococcus granulosus)
   c. Coenurosis (Taenia multiceps)
   d. Sparganosis (Spirometra)

2. Trematodes
   a. Paragonimiasis (Paragonimus westermani)
   b. Schistosomiasis (Schistosoma mansoni, japonicum, haematobium, mekongi)
   c. Other trematode infections

3. Nematodes
   a. Eosinophilic meningoencephalitis
      i. Angiostrongylus cantonensis
      ii. Gnathostoma spinigerum
   b. Toxocariasis (visceral larva migrans)
      Other forms of larva migrans
      Trichinella spiralis
   c. Human filariasis
      i. Loa-loa
      ii. Dracunculus medinensis
      iii. Onchocerca volvulus
   d. Nematodes and immunosuppression
      Strongyloides stercoralis
Summary - Parasitic Infections

- Morphological diagnosis sufficient in many cases
- Often unable to identify to species, but can specify helminth vs protozoan, nematode vs trematode vs cestode, amoeba vs toxoplasma
- Consultation available from reference centers
- Serology (and exposure history) can narrow differential
- Targeted assays (e.g. *Naegleria fowleri* vs *Acanthamoeba* spp.) and mNGS (CSF) are clinically available
Case 6

- 56-year-old woman with no significant past medical history
- Presented with multiple neurological complaints (headaches, gait abnormalities, slowed speech, trouble swallowing, difficulty with driving, memory, and calculations), and insidious decline in functioning and ADLs over 2-3 months
- MRI: Cortical restricted diffusion identified in occipital, temporal, frontal and parietal lobes; T2/FLAIR hyperintensity in bilateral basal ganglia and thalami
FINAL NEUROPATHOLOGICAL DIAGNOSIS:

PRION DISEASE, most likely CREUTZFELDT-JAKOB DISEASE

NOTE:
Western blot and immunohistochemistry (3F4) for PrPSc were performed at the National Prion Disease Pathology Surveillance Center, which revealed granular deposits as seen in prion disease.
Human Prion Diseases

- MRI
- CSF (14-3-3, tau, RT-QuiC)
- Histology: Spongiform degeneration
- Confirm by WB, IHC
- Sequence PRNP gene for mutations

2018 CDC Diagnostic Criteria for CJD

1. Sporadic CJD

**Definite:**
- Diagnosed by standard neuropathological techniques; and/or immunocytochemically; and/or Western blot confirmed protease-resistant PrP; and/or presence of scrapie-associated fibrils.

**Probable:**
- Neuropsychiatric disorder **plus** positive RT-QuIC in cerebrospinal fluid (CSF) or other tissues
- OR
  - Rapidly progressive dementia; and at least two out of the following four clinical features:
    1. Myoclonus
    2. Visual or cerebellar signs
    3. Pyramidal/extrapyramidal signs
    4. Akinesis/mutism
- AND a positive result on at least one of the following laboratory tests
  - a typical EEG (periodic sharp wave complexes) during an illness of any duration
  - a positive 14-3-3 CSF assay in patients with a disease duration of less than 2 years
  - High signal in caudate/putamen on magnetic resonance imaging (MRI) brain scan or at least two cortical regions (temporal, parietal, occipital) either on diffusion-weighted imaging (DWI) or fluid attenuated inversion recovery (FLAIR)
- AND without routine investigations indicating an alternative diagnosis.

**Possible:**
- Progressive dementia; and at least two out of the following four clinical features:
  1. Myoclonus
  2. Visual or cerebellar signs
  3. Pyramidal/extrapyramidal signs
  4. Akinesis/mutism
- AND the absence of a positive result for any of the four tests above that would classify a case as "probable"
- AND duration of illness less than two years
- AND without routine investigations indicating an alternative diagnosis.

2. Iatrogenic CJD

Progressive cerebellar syndrome in a recipient of human cadaveric-derived pituitary hormone; or sporadic CJD with a recognized exposure risk, e.g., antecedent neurosurgery with dura mater implantation.

3. Familial CJD

Define or probable CJD **plus** definite or probable CJD in a first degree relative; and/or Neuropsychiatric disorder **plus** disease-specific PrP gene mutation.
Overall Summary

• CNS infections can be effectively diagnosed by gross and histological findings with appropriate ancillary testing

• Targeted assays useful when a specific pathogen is suspected on clinical or histological grounds

• Consider panel or unbiased testing if a broad differential exists

• Molecular testing results should be correlated with histological findings for a final integrated diagnosis
References

