

PROPOSAL TITLE: Targeting Plasma Metagenomic Sequencing to Improve Patient Care and Reduce Waste

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ABSTRACT. Plasma metagenomic sequencing (pmNGS) is an infectious disease diagnostic tool that detects microbial DNA from patient plasma¹. Due to its unbiased nature, pmNGS can identify bacteria, fungi, parasites and DNA viruses, including those clinicians have not considered in their differential diagnosis or that are difficult to identify². The diagnostic power of pmNGS can circumvent invasive, expensive diagnostic procedures³ and potentially shorten patient stays. However, the per-syndrome utility of pmNGS is not well established, and its cost is high compared to other infectious disease diagnostics. Absent institutional or national guidelines for use, UCSF has seen skyrocketing volumes of pmNGS, with an increase from 18 tests in 2018 to 616 in 2024, with an estimated annual cost >\$1 million. Here we propose a multipronged approach to define clinical indications for which pmNGS has highest impact, reduce unnecessary testing costs, analyze hospital days saved, avoid costs related to preventable procedures, and promote equity and excellence. We have created an Infectious Disease/Clinical Microbiology Consensus guidance statement to structure use of pmNGS testing ([Appendices 1-2](#)). We will update the APeX lab order for pmNGS to align with this guidance. Next, we will estimate pmNGS effects on patient antimicrobial management, length of stay, and need for procedures, stratified by clinical syndrome (Appendix 3). This approach will enable assessment of pmNGS utility, improve care, and reduce waste.

TEAM

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PROBLEM: The use of pmNGS has dramatically increased at UCSF over the last few years. Yet, studies of its effectiveness in diagnosis are limited. Single-center retrospective analyses of pmNGS have been small (all <200 patients) and have had extremely disparate estimations of clinical utility, ranging from 7%⁴, to 30.4%⁵, to 46%⁶. This variability is explained at least in part by the fact that pmNGS is ordered on different patient populations and for different clinical syndromes. The few clinical indications which have been studied for pmNGS utility are lower respiratory infection (LRTI) in immunocompromised patients⁷, febrile neutropenia (FN)^{8,9} and sepsis¹⁰. Prospective, real-time assessment of pmNGS impact by clinical syndrome is needed to

understand clinical utility and promote appropriate use. UCSF is poised to lead the nation in this area. Much of the foundational mNGS basic research was performed by UCSF-affiliated scientists, leading to deep scientific and clinical expertise in our institution¹¹⁻¹³. Moreover, UCSF is a marked outlier in use of pmNGS compared to peer institutions, which order this test far less frequently (West Coast Transplant Infectious Disease Society; 10/2/24). Lacking more robust data, it is unclear if practice patterns at UCSF represent an over-use of laboratory resources and significant extra cost outlay; or, conversely, we are under-using this test and creating an equity gap between patients who receive this test, and those who do not.

This is a logical moment to pursue by-syndrome clinical effectiveness analyses. Firstly, the UCSF health center usage, already higher than comparator institutions, is increasing markedly. 164 pmNGS tests were sent in 2022, 255 in 2023, and 614 in 2024 (Figure 1), leading to a total cost in 2024 estimated >\$1 million at approximately \$2,200/test (Figure 1; Appendix 3). 34 tests have been sent in just the first two weeks of 2025, a rate triple that of the equivalent period in 2024. This increase in use suggests increasing clinician awareness of this diagnostic technique, but likely also some percent of overuse. Finally, there is active consideration of implementing pmNGS for diagnosis in common critical illness syndromes, such as sepsis¹⁰. Assessing utility of current clinical use for pmNGS is essential before extending the scope of its use.

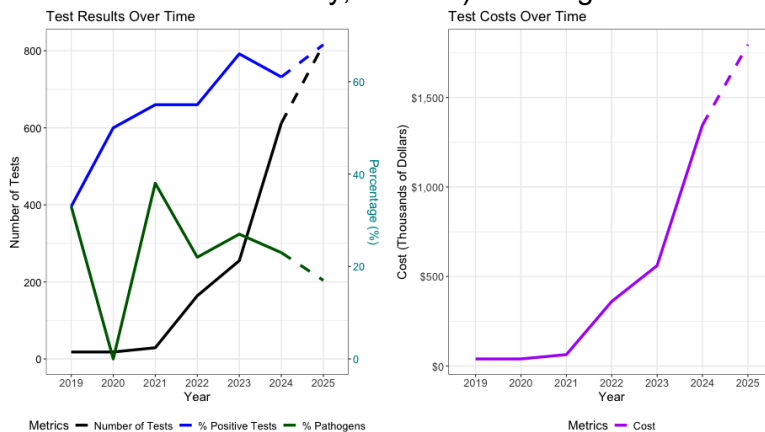


Figure 1. pmNGS volume and cost at UCSF are increasing sharply. Left panel: Volume of pmNGS tests sent at UCSF since the test introduction (black line), % of tests with at least one microbe detected (blue line) and % of tests with high-pathogenicity organisms detected (green). Right panel: estimated cost of pmNGS testing per year. 2025 is projected by # tests sent from 1/1/25-1/17/25 and 2024 per-test cost.

TARGET: The *overarching* goal of this proposal is (1) define the proportion of pmNGS tests that change length of stay and/or procedure need when sent for assessment of each of the syndromes in Appendix 1, and (2) leverage that information to standardize diagnostic use; promote early deployment of test in high-impact scenarios; and reduce use in the clinical scenarios expected to have lowest impact. We *specifically* aim to attenuate the rising rate of testing per year, by avoiding 10-15% of inappropriate tests from 2024 testing levels (5-6 tests per month). Secondly, we aim to ensure that all patients with a given syndrome receive the same guideline-directed care, rather than the current testing landscape in which test use is highly dependent on specific provider practices rather than policy. These efforts are *measurable* by means of clinician surveys coupled with retrospective review (Appendices 2-3). Given high test volume and faculty enthusiasm and engagement, especially from highest-utilizing groups, we expect our goals to be eminently *achievable*. This work is exceptionally *relevant* from a patient care and cost control perspective, given the dramatic yearly increase in pmNGS testing and associated direct costs, as well as the potential anticipated benefit to patient care and cost decrease if this test is deployed to its fullest promise (Appendix 4). This is a *timebound* project, with tracking of outcomes for 1 year, after which the pmNGS use guidelines and APeX order will be iterated to promote care, equity, and cost control. Finally, this project is *equitable* and *inclusive* in its goals, as it seeks to standardize care and ensure all patients are getting timely diagnostic care, while avoiding resource over-use.

GAPS:

Gap 1: Disconnect between stakeholders. The growing capacity for novel diagnostics such as pmNGS is outstripping the robust evaluation of how they should be deployed. Traditional research funds do not support quality improvement/stewardship projects, and personnel with clinical and scientific expertise are siloed from teams focusing on quality improvement and cost control.

Gap 2: Overuse. At UCSF, test volume has increased 20x from 2021 to 2024, with associated increased costs. The utility of this test is not established in different scenarios, with other centers estimating ~50-90% of tests do not change management⁴⁻⁶. Overuse is a gap.

Gap 3. No tracking of current use. At the moment, there is not robust assessment of what pmNGS is used for at UCSF. Therefore, we do not know which services and indications are driving the uptick in test volume and associated costs.

Gap 4: No coordinated guidance. Absent coordinated guidance on when to use this test, test use at UCSF is dependent on the diagnostic familiarity or particular prior experiences of an individual provider, leading to inequities and inconsistencies in which patients receive this test and at what points in their diagnostic journey.

INTERVENTIONS

- Alter the order format for pmNGS to integrate the indication options in the APeX order with the Infectious Disease/Clinical Microbiology Consensus Molecular Testing Guidance ([Appendix 1](#)) such that clinicians must indicate the guideline-concordant reason they are ordering this test.
- Leverage the daily Micro Reports of pmNGS and survey clinicians about patient outcomes (antimicrobial changes, days of stay changes, and procedures avoided or pursued) by clinical syndrome.
 - Publish internal quarterly reports summarizing results by syndrome and ordering service, and disseminate to key stakeholders.
- Promote effective test use by:
 - Iterating test order format in APeX to restrict to higher-yield clinical scenarios.
 - Establish and maintain a weekly clinical microbial sequencing board¹² to review pmNGS cases in real time between microbiologists and clinicians, and promote effective use by regular discussions with high-use clinical services.

PRACTICE SETTING AND TARGET POPULATION: This project will focus specifically on the inpatient use of pmNGS testing with a focus on the infectious disease and immunocompromised transplant services that guide either the use or interpretation of the majority of pmNGS tests at this time. However, we anticipate that lessons learned will be applicable to all clinicians using pmNGS within UCSF, and will also directly impact practices of other peer institutions.

BARRIERS: Key to this project is the iterative assessment of test utility by the ordering or recommending clinicians. This requires short-term follow up of results, which in the first year of this project will be labor-intensive. Thus, we have requested dedicated time for a clinical research coordinator (Hannah Teal) and for a supervising clinician (Dr. Spottiswoode).

ADVERSE EVENTS: pmNGS results have the potential to drastically improve patient care, but false negatives or false positives have the potential to cause harm by causing under- or over-treatment. The iterative nature of this project, in which indications for future years will be based on this funded year, will help to minimize these potential unwanted consequences. Moreover, the identification of low-yield clinical scenarios for which pmNGS should be avoided will not only help to reduce overall costs, but also reduce adverse events.

RETURN ON INVESTMENT (ROI) – We estimated the ROI of this proposed project as between **\$172,074-245,385**. [Appendix 4](#) contains all baseline costs and per-month estimations.

- **Estimated costs saved by avoiding testing in low-yield scenarios (est: \$132,000-\$158,400)** Other centers have estimated 54-94% pmNGS tests did not change management.

Sources Cited

1. Chiu CY, Miller SA. Clinical metagenomics. *Nat Rev Genet* 2019;20(6):341-355. DOI: 10.1038/s41576-019-0113-7.
2. Wilson MR, Shanbhag NM, Reid MJ, et al. Diagnosing Balamuthia mandrillaris Encephalitis With Metagenomic Deep Sequencing. *Ann Neurol* 2015;78(5):722-30. DOI: 10.1002/ana.24499.
3. Madut DB, Chemaly RF, Dadwal SS, et al. Clinical Utility of Plasma Microbial Cell-Free DNA Sequencing Among Immunocompromised Patients With Pneumonia. *Open Forum Infect Dis* 2024;11(8):ofae425. DOI: 10.1093/ofid/ofae425.
4. Hogan CA, Yang S, Garner OB, et al. Clinical Impact of Metagenomic Next-Generation Sequencing of Plasma Cell-Free DNA for the Diagnosis of Infectious Diseases: A Multicenter Retrospective Cohort Study. *Clin Infect Dis* 2021;72(2):239-245. DOI: 10.1093/cid/ciaa035.
5. Vinh Dong H, Saleh T, Kaur I, Yang S. Elucidating the Clinical Interpretation and Impact of a Positive Plasma Cell-Free DNA Metagenomics Test Result-A Single Center Retrospective Study. *J Appl Lab Med* 2024;9(1):14-27. DOI: 10.1093/jalm/jfad083.
6. Shishido AA, Noe M, Saharia K, Luethy P. Clinical impact of a metagenomic microbial plasma cell-free DNA next-generation sequencing assay on treatment decisions: a single-center retrospective study. *BMC Infect Dis* 2022;22(1):372. DOI: 10.1186/s12879-022-07357-8.
7. Bergin SP, Chemaly RF, Dadwal SS, et al. Plasma Microbial Cell-Free DNA Sequencing in Immunocompromised Patients with Pneumonia: A Prospective Observational Study. *Clin Infect Dis* 2023. DOI: 10.1093/cid/ciad599.
8. Benamu E, Gajurel K, Anderson JN, et al. Plasma Microbial Cell-free DNA Next-generation Sequencing in the Diagnosis and Management of Febrile Neutropenia. *Clin Infect Dis* 2022;74(9):1659-1668. DOI: 10.1093/cid/ciab324.
9. Schulz E, Grumaz S, Hatzl S, et al. Pathogen Detection by Metagenomic Next-Generation Sequencing During Neutropenic Fever in Patients With Hematological Malignancies. *Open Forum Infect Dis* 2022;9(8):ofac393. DOI: 10.1093/ofid/ofac393.
10. Kalantar KL, Neyton L, Abdelghany M, et al. Integrated host-microbe plasma metagenomics for sepsis diagnosis in a prospective cohort of critically ill adults. *Nat Microbiol* 2022. DOI: 10.1038/s41564-022-01237-2.
11. Langelier C, Kalantar KL, Moazed F, et al. Integrating host response and unbiased microbe detection for lower respiratory tract infection diagnosis in critically ill adults. *Proc Natl Acad Sci U S A* 2018;115(52):E12353-E12362. DOI: 10.1073/pnas.1809700115.
12. Wilson MR, Sample HA, Zorn KC, et al. Clinical Metagenomic Sequencing for Diagnosis of Meningitis and Encephalitis. *N Engl J Med* 2019;380(24):2327-2340. DOI: 10.1056/NEJMoa1803396.
13. Gu W, Deng X, Lee M, et al. Rapid pathogen detection by metagenomic next-generation sequencing of infected body fluids. *Nat Med* 2021;27(1):115-124. DOI: 10.1038/s41591-020-1105-z.