

Subject:	Identification of Microorganisms using Nucleic Acid Probes		
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I. Policy Description

Nucleic acid hybridization technologies utilize complementary properties of the DNA double-helix structures to anneal together DNA fragments from different sources. These techniques are utilized in polymerase chain reaction (PCR) and fluorescent resonance energy transfer (FRET) techniques to identify microorganisms.¹

A discussion of every infectious agent that might be detected with a probe technique is beyond the scope of this policy. Many probes have been combined into panels of tests. For the purposes of this policy, only individual probes are reviewed.

For guidance on nucleic acid identification of *Candida* in vaginitis, please refer to AHS-M2057-Diagnosis of Vaginitis.

Indications and/or Limitations of Coverage

Application of coverage criteria is dependent upon an individual's benefit coverage at the time of the request. Specifications pertaining to Medicare and Medicaid can be found in the "Applicable State and Federal Regulations" section of this policy document.

1. The coverage status of nucleic acid identification using direct probe, amplified probe, or quantification for the microorganism's procedure codes is summarized in Table 1 below. "MCC" in the table below indicates that the test **MEETS COVERAGE CRITERIA**; while "DNMCC" tests indicates that the test **DOES NOT MEET COVERAGE CRITERIA**.

Microorganism	Direct Probe	Amplified Probe	Quantification
<i>Bartonella henselae</i> or <i>quintana</i>		87471 (MCC)	87472 (DNMCC)
<i>Chlamydia pneumoniae</i>	87485	87486 (MCC)	87487 (DNMCC)
<i>Clostridium difficile</i>		87493 (MCC)	
<i>Cytomegalovirus</i>	87495	87496 (MCC)	87497 (MCC)

Microorganism	Direct Probe	Amplified Probe	Quantification
<i>Enterococcus</i> , Vancomycin-resistant (e.g., <i>enterococcus vanA</i> , <i>vanB</i>)		87500 (MCC)	
<i>Enterovirus</i>		87498 (MCC)	
Hepatitis G	87525 (DNMCC)	87526 (DNMCC)	87527 (DNMCC)
Herpes virus-6	87531 (DNMCC)	87532 (DNMCC)	87533 (MCC)
<i>Legionella pneumophila</i>	87540	87541 (MCC)	87542 (DNMCC)
Orthopoxvirus		87593 (MCC)	
<i>Mycoplasma pneumoniae</i>	87580	87581 (MCC)	87582 (DNMCC)
Respiratory syncytial virus		87634 (MCC)	
<i>Staphylococcus aureus</i>		87640 (MCC)	
<i>Staphylococcus aureus</i> , methicillin resistant		87641 (MCC)	

2. Simultaneous ordering of amplified probe and quantification for the same organism in a single encounter **DOES NOT MEET COVERAGE CRITERIA.**

Scientific Background

Nucleic acid hybridization technologies, including polymerase chain reaction (PCR), ligase- or helicase-dependent amplification, and transcription-mediated amplification, are beneficial tools for pathogen detection in blood culture and other clinical specimens due to high specificity and sensitivity.¹ The use of nucleic acid-based methods to detect bacterial pathogens in a clinical laboratory setting offers “increased sensitivity and specificity over traditional microbiological techniques” due to its specificity, sensitivity, reduction in time, and high-throughput capability; however, “contamination potential, lack of standardization or validation for some assays, complex interpretation of results, and increased cost are possible limitations of these tests.”²

Guidelines and Recommendations

World Health Organization (WHO)

For detection of mpox, the WHO recommends “detection of viral DNA by polymerase chain reaction (PCR)” as the preferred laboratory test and recommends that any individual with a suspected case should be offered testing. They note that the best specimens for diagnosis are taken directly from the rash. Antigen and antibody detection may not be able to distinguish between orthopoxviruses.³

Infectious Diseases Society of America (IDSA)

Specific guidelines for testing of many organisms listed within the policy coverage criteria is found in the updated 2018 Infectious Diseases Society of America (IDSA) guidelines and recommendations titled, “A Guide to Utilization of the Microbiology Laboratory for Diagnosis of Infectious Diseases: 2018 Update by the Infectious Diseases Society of America and the American Society for Microbiology.”⁴ “This document is organized by body system, although many organisms are capable of causing disease in >1 body system. There may be a redundant mention of some organisms because of their propensity to infect multiple sites. One of the unique features of this document is its ability to assist clinicians who have specific suspicions regarding possible etiologic agents causing a specific type of

disease. When the term “clinician” is used throughout the document, it also includes other licensed, advanced practice providers. Another unique feature is that in most chapters, there are targeted recommendations and precautions regarding selecting and collecting specimens for analysis for a disease process. It is very easy to access critical information about a specific body site just by consulting the table of contents. Within each chapter, there is a table describing the specimen needs regarding a variety of etiologic agents that one may suspect as causing the illness. The test methods in the tables are listed in priority order according to the recommendations of the authors and reviewers.”⁴

The IDSA, in conjunction with the American Society for Microbiology (ASM,) released a 2024 update to their Guide to Utilization of the Microbiology Laboratory for Diagnosis of Infectious Diseases. “The current 2024 version provides new knowledge, discusses new infections, and suggests new laboratory procedures to assist in confirming the causes of infectious diseases.”⁵ Like the 2018 version, this guide covers key point for the laboratory diagnosis for various infections and symptoms broken down by body system. Overall, this guide highlights NAATs and PCR tests as important testing tools for modern infectious disease diagnostics due to rapid direction of pathogens with a high level of sensitivity and specificity in the clinical setting.

Centers of Disease Control and Prevention (CDC)

Candida Auris (C. auris)

The CDC writes that “Molecular methods based on sequencing the D1-D2 region of the 28s rDNA or the Internal Transcribed Region (ITS) of rDNA can identify *C. auris*.” The CDC further notes that various PCR methods have been developed for identifying *C. auris*.

Chlamydia Pneumoniae (C. pneumoniae)

The CDC writes that RT-PCR is the “preferred” method of detecting an acute *C. pneumoniae* infection.

Clostridioides difficile (C. diff)

The CDC writes that “Nucleic acid amplification tests (NAAT), including real-time polymerase chain reaction (PCR), are the preferred method of diagnostic testing for acute *C. pneumoniae* infection. . . . Molecular tests, including NAATs, offer high sensitivity and specificity and provide timely results for treatment decisions. These tests can also determine antibiotic susceptibilities.”⁷

Clostridioides difficile (C. diff)

The CDC states that there are four laboratory tests that can be used to diagnose *Clostridioides difficile* infection (CDI). “FDA-approved PCR assays are same-day tests that are highly sensitive and specific for the presence of a toxin-producing *C. diff* organism.” The CDC does note that “molecular assays can be positive for *C. diff* in asymptomatic individuals and those who do not have an infection” and

“when using multi-pathogen (multiplex) molecular methods, read the results with caution as the pre-test probability of *C. diff* infection might be less.”⁸

Cytomegalovirus (CMV)

The CDC states that “the standard laboratory test for diagnosing congenital CMV infection is a PCR on saliva, with a confirmatory test on urine. . . The reason for the confirmatory test on urine is that most CMV seropositive mothers shed CMV in their breast milk. This can cause a false-positive CMV result on saliva collected shortly after the baby has breast fed.”⁹

Mpox Virus

The CDC defines a suspect case of Mpox as a “new characteristic rash or meets one of the epidemiologic criteria and has a high clinical suspicion for mpox.” A probable case is defined as “no suspicion of other recent *Orthopoxvirus* exposure (e.g., Vaccinia virus in ACAM2000 vaccination) AND demonstration of the presence of *Orthopoxvirus* DNA by polymerase chain reaction of a clinical specimen OR *Orthopoxvirus* using immunohistochemical or electron microscopy testing methods OR Demonstration of detectable levels of anti-orthopoxvirus IgM antibody during the period of 4 to 56 days after rash onset.” A confirmed case of Mpox is defined as “demonstration of the presence of Mpox virus DNA by polymerase chain reaction testing or Next-Generation sequencing of a clinical specimen OR isolation of Mpox virus in culture from a clinical specimen.”¹⁰

The CDC states that “Mpox is diagnosed using real time PCR tests” and further notes “clinicians should collect two swabs from each lesion (generally from 2-3 lesions) in case additional testing, such as clade-specific testing, is needed for these patients.”¹¹

MRSA

The CDC remarks that “Providers can test some patients to see if they carry MRSA in their nose or on their skin. This test involves rubbing a cotton-tipped swab in the patient’s nostrils or on the skin. The only way to know if MRSA is the cause of an infection is to test for the bacteria in a laboratory.” The CDC further states “There are many methods laboratorians can use to test for MRSA” and lists that “Phenotypic methods recommended for the detection of MRSA include: cefoxitin broth microdilution, oxacillin broth microdilution, and cefoxitin disk diffusion testing.” The CDC includes additional methods including “Nucleic acid amplification tests, such as the polymerase chain reaction (PCR), to detect the *mecA* gene, which mediates oxacillin resistance in staphylococci” but notes “*mecA* PCR tests will not detect novel resistance mechanisms or uncommon phenotypes (e.g., *mecC* or borderline-resistant oxacillin resistance).”¹²

Mycoplasma pneumoniae

The states that “(NAATs) are the preferred method of diagnostic testing for *M. pneumoniae* infections. Molecular tests such as nucleic acid amplification tests (NAATs) offer high sensitivity and specificity

and provide timely results for treatment decisions. These tests can also predict antibiotic susceptibilities.”¹³

Non-Polio Enterovirus

The CDC remarks that their laboratories “routinely” perform qualitative testing for enteroviruses, parechoviruses, and uncommon picornaviruses and states that “CDC and some health departments test with molecular sequencing methods, or a real-time reverse transcription polymerase chain reaction (rRT-PCR) lab test.”¹⁴

Respiratory Syncytial Virus (RSV)

The CDC writes that “PCR tests can be used to diagnose anyone for RSV. Antigen tests are only effective when testing infants and young children.”¹⁵

Miscellaneous

The CDC does not mention the need to quantify [through PCR] *Bartonella*, *Legionella pneumophila*, or *Mycoplasma pneumoniae*. However, PCR can be performed for both *Bartonella* and *Legionella pneumophila* specimen.^{16,17} No guidance was found on Hepatitis G.

Committee on Infectious Diseases, American Academy of Pediatrics, 31st Edition (2018-2021, Red Book)

The Committee on Infectious Diseases released joint guidelines with the American Academy of Pediatrics. In it, they note that “the presumptive diagnosis of mucocutaneous candidiasis or thrush usually can be made clinically.” They also state that FISH probes may rapidly detect *Candida* species from positive blood culture samples, although PCR assays have also been developed for this purpose.¹⁸

European Centre for Disease Prevention and Control (ECDC)

On May 23, 2022, the ECDC released a rapid risk assessment of the Mpox multi-country outbreak. They recommend that patients with probable cases should be tested with a “Mpox virus specific PCR or an orthopoxvirus specific PCR assay which is then confirmed through sequencing.”¹⁹

On June 2, 2022, ECDC released interim advice on risk communication and community engagement during the 2022 Mpox outbreak in Europe. This is a joint report with the WHO regional office for Europe. They recommend speaking to your doctor about getting tested for Mpox if you develop a rash with a fever or feeling of discomfort or illness.²⁰

United Kingdom Health Security Agency (UKHSA)

The UKHSA states that “Mpox is diagnosed by PCR test for the Mpox virus (MPXV) on a viral swab taken from one or more vesicles or ulcers.” Specifically, it is recommended that healthcare workers “Take a viral swab in viral culture medium or viral transport medium (for example Virocult®) from an open sore or from the surface of a vesicle. If other wounds are present, ensure that the sample is

definitely taken from a vesicle, an ulcer or a crusted vesicle. Rub the swab over the lesion and place the swab in the collection tube. If there are pharyngeal lesions, a throat swab should also be taken.”²¹ UKHSA also suggests that “A viral throat swab can be taken for high-risk contacts of a confirmed or highly probable case who have developed systemic symptoms but do not have a rash or lesions that can be sampled. Please note that even if the throat swab is negative, the individual must continue with monitoring and isolation as instructed by their local health protection team and should be reassessed and sampled if further symptoms develop.” Lastly, “If follow-up testing is required from a confirmed or highly probable case, either because of clinical deterioration or to inform discharge from isolation to an inpatient setting, additional samples should be taken and should include the following:

- a lesion swab and throat swab in viral transport medium
- a blood sample in an EDTA tube
- a urine sample in a universal sterile container.”²¹

The UKHSA states that “Following the identification of a cluster of sexually transmitted HCID Clade I mpox in 2023, there is an increased risk of mpox HCID infection circulating unrecognized on the background of Clade II infections.” They therefore recommend “All diagnostic samples from all individuals testing positive for mpox should now be subject to clade confirmation. Positive mpox samples should be sent to RIPL for clade specific testing if clade differentiation is not available through local mpox testing services.”²¹

The UKHSA states that mpox DNA viruses can be detected in semen up to 11 days after acute infection, and recommends that: “Following the initial 12 weeks and up to 6 months after recovery from infection, UKHSA recommends performing MPXV PCR on semen samples (and where necessary, oropharyngeal and/or rectal swabs) if the patient:

- is undergoing fertility treatment or planning pregnancy
- is undergoing planned semen storage (for example prior to chemotherapy)
- has an immunocompromised sexual partner (including a pregnant partner)
- is concerned about transmission to sexual partner or partners for any other reason and requests a test from their clinician.”²¹

HHV-6 Foundation

The human herpesvirus 6 (HHV-6) foundation also states that “a negative finding in the plasma does not rule out a localized active infection in an organ (e.g. uterus, brain, thyroid, liver). Persistent HHV-6 infections have been found in the liver, brain, lungs, heart tissue and uterus, with no trace of HHV-6 DNA in the plasma. Quantitative testing on blood and tissues is preferred because it can differentiate between the very low levels occasionally found in healthy controls and high levels found in diseased tissues.”²²

The HHV-6 foundation states that *qualitative* PCR DNA tests on whole blood are “useless for differentiating active from latent infection” but notes that the test may be useful for differentiating between herpes virus-6A and herpes virus-6B. The HHV-6 foundation states that *quantitative* PCR DNA tests on whole blood can differentiate active from latent infection “If the viral load is >200 copies per ml or 20 copies per microgram of DNA then this is an active infection.”²²

Food and Drug Administration (FDA)

Many labs have developed specific tests that they must validate and perform in house. These laboratory-developed tests (LDTs) are regulated by the Centers for Medicare and Medicaid (CMS) as high-complexity tests under the Clinical Laboratory Improvement Amendments of 1988 (CLIA '88). LDTs are not approved or cleared by the U. S. Food and Drug Administration; however, FDA clearance or approval is not currently required for clinical use.

A list of current U.S. Food and Drug Administration²³ approved or cleared nucleic acid-based microbial tests is available at: <https://www.fda.gov/medical-devices/vitro-diagnostics/nucleic-acid-based-tests>.

II. Applicable Codes

Code	Description	Comment
87471	Infectious agent detection by nucleic acid (DNA or RNA); Bartonella henselae and Bartonella quintana, amplified probe technique	
87472	Infectious agent detection by nucleic acid (DNA or RNA); Bartonella henselae and Bartonella quintana, quantification	
87485	Infectious agent detection by nucleic acid (DNA or RNA); Chlamydia pneumoniae, direct probe technique	
87486	Infectious agent detection by nucleic acid (DNA or RNA); Chlamydia pneumoniae, amplified probe technique	
87487	Infectious agent detection by nucleic acid (DNA or RNA); Chlamydia pneumoniae, quantification	
87493	Infectious agent detection by nucleic acid (DNA or RNA); Clostridium difficile, toxin gene(s), amplified probe technique	
87495	Infectious agent detection by nucleic acid (DNA or RNA); cytomegalovirus, direct probe technique	
87496	Infectious agent detection by nucleic acid (DNA or RNA); cytomegalovirus, amplified probe technique	
87497	Infectious agent detection by nucleic acid (DNA or RNA); cytomegalovirus, quantification	
87498	Infectious agent detection by nucleic acid (DNA or RNA); enterovirus, amplified probe technique, includes reverse transcription when performed	

87500	Infectious agent detection by nucleic acid (DNA or RNA); vancomycin resistance (eg, enterococcus species van A, van B), amplified probe technique	
87525	Infectious agent detection by nucleic acid (DNA or RNA); hepatitis G, direct probe technique	
87526	Infectious agent detection by nucleic acid (DNA or RNA); hepatitis G, amplified probe technique	
87527	Infectious agent detection by nucleic acid (DNA or RNA); hepatitis G, quantification	
87531	Infectious agent detection by nucleic acid (DNA or RNA); Herpes virus-6, direct probe technique	
87532	Infectious agent detection by nucleic acid (DNA or RNA); Herpes virus-6, amplified probe technique	
87533	Infectious agent detection by nucleic acid (DNA or RNA); Herpes virus-6, quantification	
87540	Infectious agent detection by nucleic acid (DNA or RNA); Legionella pneumophila, direct probe technique	
87541	Infectious agent detection by nucleic acid (DNA or RNA); Legionella pneumophila, amplified probe technique	
87542	Infectious agent detection by nucleic acid (DNA or RNA); Legionella pneumophila, quantification	
87580	Infectious agent detection by nucleic acid (DNA or RNA); Mycoplasma pneumoniae, direct probe technique	
87581	Infectious agent detection by nucleic acid (DNA or RNA); Mycoplasma pneumoniae, amplified probe technique	
87582	Infectious agent detection by nucleic acid (DNA or RNA); Mycoplasma pneumoniae, quantification	
87593	Infectious agent detection by nucleic acid (DNA or RNA); Ortho poxvirus (eg, monkeypox virus, cowpox virus, vaccinia virus), amplified probe technique, each	
87634	Infectious agent detection by nucleic acid (DNA or RNA); respiratory syncytial virus, amplified probe technique	
87640	Infectious agent detection by nucleic acid (DNA or RNA);	

	Staphylococcus aureus, amplified probe technique	
87641	Infectious agent detection by nucleic acid (DNA or RNA); Staphylococcus aureus, methicillin resistant, amplified probe technique	

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Procedure codes appearing in Medical Policy documents are included only as a general reference tool for each policy. They may not be all-inclusive.

III. Definitions

Term	Meaning

IV. Related Policies

Policy Number	Policy Description
PO-RE-050	Hepatitis Testing
PO-RE-011	Lyme Disease
PO-RE-012	Pathogen Panel Testing
PO-RE-014	Diagnostic Testing of Common Sexually Transmitted Infections
PO-RE-015	Testing for Vector-Borne Infections
PO-RE-018	Diagnosis of Vaginitis

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Procedure codes appearing in Reimbursement Policy documents are included only as a general reference tool for each policy. They may not be all-inclusive.

V. Reference Materials

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HHV-6 Foundation. Overview on Testing for HHV-6 infection. 2024;
FDA. Nucleic Acid Based Tests. Updated March 05. https://www.fda.gov/medical-devices/vitro-diagnostics/nucleic-acid-based-tests

VI. Revision History

Revision Date	Summary of Changes
06/04/2025	<p>Reviewed and Updated: Updated background, guidelines, and evidence-based scientific references. Literature review necessitated the following changes in coverage criteria:</p> <p>Removed “Non-vaginal Candida species” and associated codes from the table, as the codes for all Candida species are the same and appropriate ordering scenarios for NAAT testing for Candida are addressed in M2057 and M2172.</p> <p>Direct probe testing for Chlamydia pneumoniae, Cytomegalovirus, Legionella pneumophila, and Mycoplasma pneumoniae all changed from "MCC" to "DNMCC". All direct probes in policy do not meet coverage criteria.</p> <p>Change of direct probe management in CC1 results in removal of direct probe from CC2 and reorganization of the criteria to describe the exclusion of same day ordering. Now reads: “2) Simultaneous ordering of amplified probe and quantification for the same organism in a single encounter DOES NOT MEET COVERAGE CRITERIA.”</p> <p>Removed CPT code 87480, 87481, 87482</p>
09/04/2024	<p>Reviewed and Updated: Updated background, guidelines, and evidence-based scientific references. Literature review necessitated the following changes in coverage criteria:</p> <p>Removed Mycoplasma genitalium from this table, as management of testing for M. genitalium is now contained within G2157- Diagnostic Testing of Common Sexually Transmitted Infections</p> <p>Changed direct probe for HHV6 (87531) from MCC to DNMCC, as quantitative testing for HHV6 can differentiate between active and latent infection, whereas qualitative does not and direct and amplified probe coverage should match for HHV6.</p> <p>Removed CPT code 87563</p>

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