



## Short Communication

# Comparative clinical evaluation of the IsoAmp<sup>®</sup> HSV Assay with ELVIS<sup>®</sup> HSV culture/ID/typing test system for the detection of herpes simplex virus in genital and oral lesions

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## ABSTRACT

**Background:** The novel IsoAmp<sup>®</sup> HSV Assay employs isothermal helicase-dependent nucleic acid amplification and a user-friendly disposable test device to achieve rapid (<1.5 h), on-demand qualitative detection of herpes simplex virus (HSV) types 1 and 2 in oral and genital lesions.

**Objectives:** To compare performance of the IsoAmp<sup>®</sup> HSV Assay with the ELVIS<sup>®</sup> HSV ID/typing (shell-vial culture and DFA) test system for clinical specimens collected from oral and genital lesions in symptomatic patients.

**Study design:** A total of 994 specimens from male and female genital and oral lesions were obtained and evaluated at five study sites in the United States. Results from the IsoAmp<sup>®</sup> HSV Assay were compared to those from the ELVIS<sup>®</sup> system. Separate reproducibility studies were performed at 3 sites using a blinded and randomized study panel. Discrepant specimens were resolved by bidirectional sequencing analysis. **Results:** After discrepant analysis, overall agreement of IsoAmp<sup>®</sup> with ELVIS<sup>®</sup> was 98.8% with 37.0% overall prevalence (all study sites). Reproducibility rates were well within expectations.

**Conclusion:** The IsoAmp<sup>®</sup> HSV Assay showed excellent performance for clinical use for detection of HSV in genital and oral specimens. In contrast to ELVIS<sup>®</sup>, IsoAmp<sup>®</sup> HSV offers excellent sensitivity plus rapid on-demand testing and simpler specimen preparation.

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## 1. Background

Herpes simplex viruses (HSV), serotype 1 (HSV-1) and 2 (HSV-2), are the etiologic agents responsible for a spectrum of human diseases with local, or severe or fatal disseminated presentations including: skin and genital infections, viral meningitis, meningo-encephalitis, and neonatal herpes.<sup>1–4</sup> Currently there is no therapeutic cure for HSV latent infection; treatment aims

to reduce symptoms, viral shedding, frequency of recurrence, and transmission during antiviral administration. Diagnosis of genital herpes solely by clinical presentation is insensitive and nonspecific.<sup>5</sup> Timely and accurate diagnosis is necessary to assist antiviral therapeutic management and counseling for primary infection, intrapartum delivery, and suppressive therapy.

Optimized culture methods are widely available and often preferred for in vitro detection of HSV in mucocutaneous, genital and ocular lesions. The Enzyme-Linked Virus Inducible System (ELVIS<sup>®</sup>, Diagnostics Hybrids, Inc., Athens, OH) detects HSV using transgenic shell vial culture followed by typing of HSV-1 or HSV-2 by fluorescein-labeled monoclonal antisera.<sup>6–8</sup> The detection limit for ELVIS<sup>®</sup> culture is estimated to be between 0.65- and 8.5-TCID<sub>50</sub> for HSV-1 and 0.1- and 8.0-TCID<sub>50</sub> for HSV-2 depending on the strain.<sup>9</sup> Compared to conventional roll tube culture, ELVIS<sup>®</sup> reduces the maximum time to detection from days to 24 h while maintaining adequate sensitivity and eliminating subjective detection of cytopathic effect, but both methods require a cell culture

**Abbreviations:** HSV, herpes simplex virus; FDA, Food and Drug Administration; HIV, human immunodeficiency virus; ELVIS<sup>®</sup>, enzyme linked virus inducible system; ID, identification; DFA, direct fluorescent antibody; PCR, polymerase chain reaction; TCID<sub>50</sub>, 50% tissue culture infective dose; DNA, deoxyribose nucleic acid; HDA, helicase-dependent amplification; VTM, viral transport medium.

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facility and associated technical expertise. Polymerase chain reaction (PCR)-based molecular assays detect and subtype HSV with greater sensitivity than shell vial and ELVIS® methods.<sup>7,10,11</sup> Currently, only a few PCR-based assays are FDA-cleared as in vitro diagnostic tests for HSV: The MultiCode® HSV-1&2 assay (EraGen Biosciences, Inc., Madison, WI) uses novel probe-free PCR-based technology for qualitative detection and differentiation of HSV-1 and HSV-2 in vaginal lesions<sup>12</sup>; and, the BD ProbeTec™ HSV Q<sup>X</sup> Amplified DNA Assays (BD Diagnostics, Sparks, MD) employ automated strand-displacement amplification technology for detection and typing of HSV in external anogenital specimens. However, these assays require specialized and relatively expensive instruments. In comparison, the IsoAmp® HSV Assay (BioHelix Corporation, Beverly, MA) offers a facile, user-friendly approach for rapid instrument-free detection of HSV that maintains the performance benefits of PCR.

## 2. Objectives

To compare performance of the IsoAmp® HSV Assay (IsoAmp®) with the ELVIS® HSV ID/typing (shell-vial and DFA) test system (ELVIS®) for clinical specimens collected from oral and genital lesions in symptomatic patients suspected of having herpes infection.

## 3. Study design

### 3.1. Comparative clinical evaluation

A total of 994 clinical specimens (962 prospective, 32 retrospective) from male and female genital and oral lesions were obtained and evaluated at five study sites in the United States: Boston Medical Center, Boston, MA; Cleveland Clinic, Cleveland, OH; University of Virginia Health System, Charlottesville, VA; Vanderbilt University Medical Center, Nashville, TN; Laboratory Alliance of Central New York, Liverpool, NY. Institutional Review Board approval was obtained at each site as needed prior to the study. Clinical lesional specimens were collected at each site using Dacron swabs, transported in viral transport medium (VTM) to the respective microbiology laboratory, and processed for HSV detection using ELVIS®. An aliquot of left over eluate from each sample was tested by IsoAmp®. Specimens were de-identified and linked only by a unique study number. Results from IsoAmp® were compared to ELVIS® at the end of the study.

### 3.2. Reproducibility

Reproducibility studies were performed at 2 clinical sites and at BioHelix using a blinded and randomized 7-member study panel.<sup>13</sup> HSV viral stock was diluted in HSV Negative Matrix (pooled clinical HSV negative samples) to make the different panel concentration levels measured as TCID<sub>50</sub>/mL with a negativity (neg %) or positivity rate (pos %) as follows: HSV-1 High negative =  $1.75 \times 10^4$  (neg 30%); HSV-2 high negative =  $2.2 \times 10^3$  (neg 30%); HSV-1 and HSV-2 low positives =  $1.1 \times 10^5$  and  $1.1 \times 10^4$ , respectively (pos 95% each). HSV-1 and HSV-2 moderate positives =  $3.3 \times 10^5$  and  $3.3 \times 10^4$ , respectively (pos 100%). HSV negative matrix was used as the HSV Negative (pos 0%). Panels and controls from one lot were tested twice daily for five days by two operators at each site. Lot-to-lot reproducibility was performed by one site using three lots of the panel.

### 3.3. IsoAmp® HSV Assay procedure

Specimens were tested on IsoAmp® according to the manufacturer's Research Use Only kit (BioHelix Corp.) as detailed previously.<sup>13</sup> Briefly, specimen in VTM was diluted with buffer and transferred to an amplification tube to which was added master mix, then mineral oil. Tubes were placed in a 64°C heat block for 60 min to amplify target DNA, then placed in a Type II BEST™ Cassette (BioHelix Corp.) for amplicon detection.<sup>14,15</sup> IsoAmp® uses helicase-dependent amplification (HDA) to achieve isothermal PCR-based amplification of the target HSVglycoproteinB (gB) gene.<sup>16,17</sup> Fluorescein- and digoxigenin-labeled target amplicon and an internal control amplicon are captured and visualized as colored lines on a vertical flow strip within the disposable cassette. Results were read visually after 15 min and scored as HSV-present or HSV-absent based on the presence or absence of a test line plus control. This version of the BEST™ Cassette is not designed to discriminate between HSV-1 and HSV-2. The specifics of HDA technology and preliminary evaluation of IsoAmp® HSV were previously reported.<sup>13,16,17</sup> Analytical sensitivity of the assay was estimated at 5.5 and 34.1 copies/reaction for HSV-1 and HSV-2 respectively with excellent specificity.<sup>13</sup>

### 3.4. ELVIS viral cultures

Viral cultures for detecting HSV were performed using the Enzyme-Linked Virus Inducible System (ELVIS®, Diagnostics Hybrids, Inc., Athens, OH) shell vial assay according to the package insert.<sup>9</sup>

### 3.5. Discrepant analysis

Residual aliquots of discrepant specimens were analyzed by bidirectional sequencing performed at the GMP sequencing facility of Beckman Coulter Genomics (Morrisville, NC) after sample-processing and purification of PCR products at BioHelix. Primers were designed to have M13 tails at the 5' ends for bidirectional sequencing; the PCR reaction amplified the 399-base pair target sequence of HSV that encompasses the target sequence of the IsoAmp® HSV Assay. Raw sequence files were imported using DNASTAR Lasergene 8, SeqMan Pro Version 8.0.2 (16), 402. Consensus sequences were analyzed using DNASTAR Lasergene 8, MegAlign Version 8.0.2 (13), 402; Jotun Hein method was used for alignment.

## 4. Results

### 4.1. Comparative clinical evaluation

A total of 962 prospective specimens (803 genital, 159 oral) and 32 retrospective specimens (15 genital, 17 oral) were comparatively evaluated by ELVIS® and IsoAmp®. Retrospective specimens (all specimens, all sites), showed 100% agreement between IsoAmp® and ELVIS® (data not shown). Agreement was obtained for 902 prospective specimens (93.8%) with 309 specimens positive and 593 specimens negative by both methods (Table 1). Using ELVIS® as the reference, there were 60 discrepant results for IsoAmp® (49 false-positive and 11 false-negative). After bidirectional sequencing of these discrepant specimens determined that 42 of 49 alleged false-positive IsoAmp® specimens were true positives and 6 of 11 of alleged false-negative IsoAmp® specimens were true negatives, there was 98.8% total agreement with ELVIS® (Table 1). The false negative and false positive samples that were HSV-1 or HSV-2 positive by sequencing were compared to see if discrepancies

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