



510(k) Summary

K001825

**BD ProbeTec™ *Neisseria gonorrhoeae* (GC) Q<sup>s</sup>  
Amplified DNA Assay**

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|                                       |   |                    |
|---------------------------------------|---|--------------------|
| <b>Applicant</b>                      | BD Diagnostic Systems<br>7 Loveton Circle<br>Sparks, MD 21152   | <b>DEC 11 2008</b> |
| <b>Establishment Registration No.</b> | 1119779   |                    |
| <b>Contact Person</b>                 | Kathryn Babka Carr, RAC<br>tel. 410-316-4260<br>fax. 410-316-4041<br><a href="mailto:Kathy_Carr@bd.com">Kathy_Carr@bd.com</a> |                    |
| <b>Summary Date</b>                   | December 1, 2008  |                    |
| <b>Proprietary Name</b>               | BD ProbeTec™ <i>Neisseria gonorrhoeae</i> (GC) Q <sup>s</sup> Amplified DNA Assay   |                    |
| <b>Generic Name</b>                   | DNA probe, nucleic acid amplification, <i>Neisseria</i>   |                    |
| <b>Classification</b>                 | Class II  |                    |
| <b>Classification Name</b>            | <i>Neisseria</i> spp. direct serological test reagents  |                    |
| <b>Regulation Number</b>              | 866.3390  |                    |
| <b>Product Code</b>                   | LSL   |                    |
| <b>Predicate Devices</b>              | BD ProbeTec ET CT/GC Amplified DNA Assay (K984631),<br>APTIMA Combo 2 Assay (K003395)   |                    |

**Device Description**

The BD ProbeTec GC Q<sup>s</sup> Amplified DNA Assay is based on the simultaneous amplification and detection of target DNA using amplification primers and a fluorescently-labeled detector probe. The reagents for SDA are dried in two separate disposable microwells: the Priming Microwell contains the amplification primers, fluorescently-labeled detector probe, nucleotides and other reagents necessary for amplification, while the Amplification Microwell contains the two enzymes (a DNA polymerase and a restriction endonuclease) that are required for SDA. The BD Viper™ System pipettes a portion of the purified DNA solution from each Extraction Tube into a Priming Microwell to rehydrate the contents. After a brief incubation, the reaction mixture is transferred to a corresponding, pre-warmed Amplification Microwell which is sealed to prevent contamination and then incubated in one of the two thermally-controlled fluorescent readers. The presence or absence of *N. gonorrhoeae* DNA is determined by calculating the peak fluorescence (Maximum Relative Fluorescent Units (MaxRFU)) over the course of the amplification process and by comparing this measurement to a predetermined threshold value.

In addition to the fluorescent probe used to detect amplified *N. gonorrhoeae* target DNA, a second labeled oligonucleotide is incorporated in each reaction. The Extraction Control (EC) oligonucleotide is labeled with a different dye than that used for detection of the *N. gonorrhoeae*-specific target and is used to confirm the validity of the extraction process. The EC is dried in the Extraction Tubes and is rehydrated



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upon addition of the specimen and extraction reagents. At the end of the extraction process, the EC fluorescence is monitored by the **BD Viper** System and an automated algorithm is applied to both the EC and *N. gonorrhoeae*-specific signals to report results as positive, negative, or EC failure.

#### **Intended Use**

The **BD ProbeTec™ *Neisseria gonorrhoeae* (GC) Q<sup>x</sup> Amplified DNA Assay**, when tested with the **BD Viper™ System** in Extracted Mode, uses Strand Displacement Amplification technology for the direct, qualitative detection of *Neisseria gonorrhoeae* DNA in clinician-collected female endocervical and male urethral swab specimens, patient-collected vaginal swab specimens (in a clinical setting), and male and female urine specimens (both UPT and Neat). The assay is indicated for use with asymptomatic and symptomatic females and symptomatic males to aid in the diagnosis of gonococcal urogenital disease.

#### **Summary and Principles of Operation**

When used with the **BD Viper** System, the **BD ProbeTec GC Q<sup>x</sup> Amplified DNA Assay (GC Q<sup>x</sup> Assay)** involves automated extraction of DNA from clinical specimens through the chemical lysis of cells, followed by binding of DNA to para-magnetic particles, washing of the bound nucleic acid and elution in an amplification-compatible buffer. When present, *N. gonorrhoeae* DNA is then detected by Strand Displacement Amplification (SDA) of a specific target sequence in the presence of a fluorescently labeled detector probe.

#### **Analytical Performance Characteristics**

##### **Limit of Detection (Analytical Sensitivity)**

The Limits of Detection (LODs) for the GC Q<sup>x</sup> Assay with *Neisseria gonorrhoeae* strain ATCC 19424 in urine and swab specimens when extracted on the **BD Viper** System were determined to be  $\leq 50$  cells per mL for neat and UPT treated urine and  $\leq 100$  GC cells per mL for expressed vaginal and endocervical swab specimens. The GC Q<sup>x</sup> Assay on the **BD Viper** System in extracted mode was able to detect 17 GC strains with  $\geq 95\%$  proportion positive at a concentration of 50 cells per mL in Q<sup>x</sup> Swab Diluent.

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### Analytical Specificity

The 141 organisms listed in **Table 1** were tested with the **BD ProbeTec GC Q<sup>x</sup> Amplified DNA Assay** on the **BD Viper System**. All potential cross-reactive species were tested at  $\geq 1 \times 10^8$  cells/mL except where noted. Two *N. cinerea* and two *N. lactamica* strains were shown to cross-react with the GC Q<sup>x</sup> assay.

|                                    |  |   |   |
|------------------------------------|--|---|---|
| <i>Acinetobacter calcoaceticus</i> | Epstein Barr Virus ***                     | <i>Peptostreptococcus productus</i>             | <i>Neisseria elongata</i> subsp. <i>nitroreducens</i> (2) |
| <i>Acinetobacter lwoffii</i>       | <i>Escherichia coli</i>                    | <i>Plesiomonas shigelloides</i>                 | <i>Neisseria elongata</i>                                 |
| <i>Actinomyces israelii</i>        | <i>Flavobacterium meningosepticum</i>      | <i>Propionibacterium acnes</i>                  | <i>Neisseria flava</i> (4)                                |
| Adenovirus***                      | <i>Gardnerella vaginalis</i>               | <i>Providencia stuartii</i>                     | <i>Neisseria flavescens</i> (4)                           |
| <i>Aeromonas hydrophilia</i>       | <i>Gemella haemolysans</i>                 | <i>Pseudomonas aeruginosa</i>                   | <i>Neisseria gonorrhoeae</i>                              |
| <i>Alcaligenes faecalis</i> *      | <i>Haemophilus influenzae</i>              | <i>Salmonella minnesota</i>                     | <i>Neisseria lactamica</i> (7)                            |
| <i>Bacillus subtilis</i> *         | Herpes Simplex Virus **                    | <i>Salmonella typhimurium</i>                   | <i>Neisseria meningitidis</i> (12)                        |
| <i>Bacteroides fragilis</i>        | Human papillomavirus (16 and 18)***        | <i>Staphylococcus aureus</i>                    | <i>Neisseria mucosa</i> (5)                               |
| <i>Candida albicans</i> *          | <i>Kingella kingae</i>                     | <i>Staphylococcus epidermidis</i>               | <i>Neisseria perflava</i> (8)                             |
| <i>Candida glabrata</i> *          | <i>Klebsiella pneumoniae</i>               | <i>Streptococcus agalactiae</i>                 | <i>Neisseria polysaccharea</i> (2)                        |
| <i>Candida tropicalis</i> *        | <i>Lactobacillus acidophilus</i> *         | <i>Streptococcus mitis</i>                      | <i>Neisseria sicca</i> (5)                                |
| <i>Chlamydia pneumoniae</i> ****   | <i>Lactobacillus brevis</i>                | <i>Streptococcus mutans</i>                     | <i>Neisseria subflava</i> (15)                            |
| <i>Chlamydia psittaci</i> *        | <i>Lactobacillus jensenii</i> *            | <i>Streptococcus pneumoniae</i> *               | <i>Neisseria weaverii</i> (3)                             |
| <i>Citrobacter freundii</i>        | <i>Listeria monocytogenes</i>              | <i>Streptococcus pyogenes</i>                   |   |
| <i>Clostridium perfringens</i>     | <i>Mobiluncus mulieris</i>                 | <i>Streptomyces griseus</i> **                  |   |
| <i>Corynebacterium renale</i>      | <i>Moraxella lacunata</i> *                | <i>Trichomonas vaginalis</i> **                 |   |
| <i>Cryptococcus neoformans</i> *   | <i>Moraxella osloensis</i>                 | <i>Veillonella parvula</i>                      |   |
| Cytomegalovirus**                  | <i>Morganella morganii</i>                 | <i>Vibrio parahaemolyticus</i>                  |   |
| <i>Edwardsiella tarda</i>          | <i>Mycobacterium gordonae</i>              | <i>Yersinia enterocolitica</i>                  |   |
| <i>Enterobacter cloacae</i>        | <i>Mycobacterium smegmatis</i>             | <i>Branhamella catarrhalis</i> (5)              |   |
| <i>Enterococcus faecalis</i>       | <i>Peptostreptococcus anaerobius</i>       | <i>Neisseria cinerea</i> (2)                    |   |
| <i>Enterococcus faecium</i>        | <i>Peptostreptococcus asaccharolyticus</i> | <i>Neisseria elongata</i> ss <i>glycolytica</i> |   |

(n) number of strains tested in the **BD ProbeTec GC Q<sup>x</sup> Assay**

\* Tested at  $>1 \times 10^7$  cells/mL; \*\*Tested at  $>1 \times 10^6$  cells or viral particles per mL; \*\*\*Tested at  $\geq 1 \times 10^8$  genomic equivalents per mL;

\*\*\*\* tested at  $\geq 1 \times 10^2$  TCID<sub>50</sub>/mL



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### Interfering Substances

Potential interfering substances which may be encountered in swab and/or urine specimens were extracted from urine and vaginal swab matrix in the absence and presence of GC target (150 GC cells/mL for urine and 300 GC cells/mL for swabs) and tested with the **BD ProbeTec GC Q<sup>x</sup> Amplified DNA Assay** on the **BD Viper System**. Results are summarized in **Table 2**.

**Table 2: Interfering Substances**

| Interpretation                             | Swab  | Urine  |
|--|---|--|
| No Interference Observed                   | Blood ( $\leq 60\%$ )<br>Seminal Fluid<br>Mucus<br>Over The Counter vaginal products and contraceptives<br>Hemorrhoidal cream<br>Prescription vaginal treatments<br>Leukocytes ( $1 \times 10^6$ cells/mL)<br>$1 \times 10^6$ cells/mL <i>Chlamydia trachomatis</i> | Blood (1%)<br>Seminal fluid<br>Mucus<br>Antibiotics<br>Analgesics<br>Over The Counter deodorant sprays and powders<br>Hormones<br>Leukocytes<br>Albumin $<1$ mg/mL<br>Glucose<br>Acidic urine (pH 4.0)<br>Alkaline urine (pH 9.0)<br>Bilirubin<br>Organisms associated with Urinary Tract Infections |
| May cause extraction control (EC) failures | Blood ( $> 60\%$ )  | Not observed   |

### Clinical Performance Characteristics

Clinician-collected endocervical and male urethral swab specimens, patient-collected vaginal swab specimens (in a clinical setting), and male and female Q<sup>x</sup> UPT and neat urine specimens were collected from 1059 female subjects and 479 male subjects attending OB/GYN, sexually transmitted disease (STD) and family planning clinics at seven geographically diverse clinical sites in North America. Subjects were classified as symptomatic if they reported symptoms such as dysuria, urethral discharge, coital pain/difficulty/bleeding, testicular or scrotum pain/swelling, abnormal vaginal discharge, or pelvic/uterine/adnexal pain. Subjects were classified as asymptomatic if they did not report symptoms. Sixty five female subjects and 7 male subjects were excluded from the data analysis due to age requirement violations, antibiotic treatment in the last 21 days, opting to withdraw from the study after initially consenting, failure to obtain paired swab and urine specimens, urine quantity less than 20 mL, or



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transport and storage errors related to specimen collection. Therefore, the final data analysis included 994 compliant female subjects and 472 compliant male subjects.

Five specimens were collected from each of the 994 eligible female subjects. A urine specimen was collected and split into Q<sup>x</sup> UPT, neat urine and the two reference urine specimen collection devices followed by a vaginal swab specimen and three randomized endocervical swab specimens. Up to four specimens were collected from each of the 472 eligible male subjects. Up to three randomized urethral swab specimens were collected followed by a urine specimen that was split into Q<sup>x</sup> UPT, neat urine and the two reference urine specimen collection devices. **BD ProbeTec GC Q<sup>x</sup>** assay results were generated from the Q<sup>x</sup> UPT and neat urine specimens, the vaginal swab specimen, one endocervical swab specimen and one male urethral swab specimen. The remaining two endocervical swab specimens, up to two male urethral swab specimens, and the two reference urine specimens for each male and female subject were tested using two reference methods: the **BD ProbeTec ET GC/AC** assay and another commercially available NAAT (Nucleic Acid Amplification Test). Specimen testing was conducted either at the site of collection or at a designated **BD Viper** testing site.

All performance calculations were based on the total number of **BD ProbeTec GC Q<sup>x</sup>** assays results for endocervical, vaginal and male urethral swab specimens, and male and female Q<sup>x</sup> UPT and neat urine specimens compared to a patient infected status (PIS) algorithm for each gender. In the algorithm, the designation of a subject as being infected with GC or not was based on endocervical swab and urine specimen results from the commercially available **BD ProbeTec ET GC/AC** assay and the other commercially available NAAT. Subjects were considered infected with GC if two of the four endocervical swab and urine specimens (or two of the three or four urethral swab and urine specimens) tested positive in the **BD ProbeTec ET GC/AC** assay and the other reference NAAT (one specimen testing positive in each NAAT). Subjects were considered non-infected if less than two reference NAAT results were positive. A total of 5387 **BD ProbeTec GC Q<sup>x</sup>** assay results was used to calculate sensitivity and specificity. Sensitivity and specificity by specimen type and symptomatic status are presented in **Table 3**.



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**Table 3: GC Q<sup>x</sup> Assay Performance Compared to Patient Infected Status  
(by specimen type and symptomatic status)**

| Specimen Type | Symptomatic | N    | Sensitivity      | 95% C.I.         | Specificity       | 95% C.I.         | PPV%  | NPV%  | Error Initial/Final |
|---------------|-------------|------|------------------|------------------|-------------------|------------------|-------|-------|---------------------|
| FS            | N           | 450  | 96.3% (26/27)    | (81.0% - 99.9%)  | 99.5% (421/423)   | (98.3% - 99.9%)  | 92.9  | 99.8  | 3/0                 |
|               | Y           | 542  | 100.0% (38/38)   | (90.7% - 100.0%) | 99.8% (503/504)   | (98.9% - 100.0%) | 97.4  | 100.0 | 2/2                 |
|               | Total       | 992  | 98.5% (64/65)    | (91.7% - 100.0%) | 99.7% (924/927)   | (99.1% - 99.9%)  | 95.5  | 99.9  | 5/2                 |
| FV            | N           | 449  | 100.0% (27/27)   | (87.2% - 100.0%) | 98.6% (416/422)   | (96.9% - 99.5%)  | 81.8  | 100.0 | 0/0                 |
|               | Y           | 544  | 100.0% (38/38)   | (90.7% - 100.0%) | 99.6% (504/506)   | (98.6% - 100.0%) | 95.0  | 100.0 | 0/0                 |
|               | Total       | 993  | 100.0% (65/65)   | (94.5% - 100.0%) | 99.1% (920/928)   | (98.3% - 99.6%)  | 89.0  | 100.0 | 0/0                 |
| FN            | N           | 450  | 96.3% (26/27)    | (81.0% - 99.9%)  | 99.3% (420/423)   | (97.9% - 99.9%)  | 89.7  | 99.8  | 0/0                 |
|               | Y           | 543  | 97.4% (37/38)    | (86.2% - 99.9%)  | 99.6% (503/505)   | (98.6% - 100.0%) | 94.9  | 99.8  | 0/0                 |
|               | Total       | 993  | 96.9% (63/65)    | (89.3% - 99.6%)  | 99.5% (923/928)   | (98.7% - 99.8%)  | 92.6  | 99.8  | 0/0                 |
| FUPT          | N           | 450  | 100.0% (27/27)   | (87.2% - 100.0%) | 99.5% (421/423)   | (98.3% - 99.9%)  | 93.1  | 100.0 | 0/0                 |
|               | Y           | 543  | 97.4% (37/38)    | (86.2% - 99.9%)  | 99.8% (504/505)   | (98.9% - 100.0%) | 97.4  | 99.8  | 0/0                 |
|               | Total       | 993  | 98.5% (64/65)    | (91.7% - 100.0%) | 99.7% (925/928)   | (99.1% - 99.9%)  | 95.5  | 99.9  | 0/0                 |
| MS            | N           | 215  | 100.0% (7/7)     | (59.0% - 100.0%) | 100.0% (208/208)  | (98.2% - 100.0%) | 100.0 | 100.0 | 0/0                 |
|               | Y           | 257  | 100.0% (100/100) | (96.4% - 100.0%) | 98.7% (155/157)   | (95.5% - 99.8%)  | 98.0  | 100.0 | 1/0                 |
|               | Total       | 472  | 100.0% (107/107) | (96.6% - 100.0%) | 99.5% (363/365)   | (98.0% - 99.9%)  | 98.2  | 100.0 | 1/0                 |
| MNU           | N           | 215  | 100.0% (7/7)     | (59.0% - 100.0%) | 100.0% (208/208)  | (98.2% - 100.0%) | 100.0 | 100.0 | 0/0                 |
|               | Y           | 257  | 100.0% (100/100) | (96.4% - 100.0%) | 98.1% (154/157)   | (94.5% - 99.6%)  | 97.1  | 100.0 | 0/0                 |
|               | Total       | 472  | 100.0% (107/107) | (96.6% - 100.0%) | 99.2% (362/365)   | (97.6% - 99.8%)  | 97.3  | 100.0 | 0/0                 |
| MUPT          | N           | 215  | 100.0% (7/7)     | (59.0% - 100.0%) | 99.5% (207/208)   | (97.4% - 100.0%) | 87.5  | 100.0 | 0/0                 |
|               | Y           | 257  | 100.0% (100/100) | (96.4% - 100.0%) | 98.7% (155/157)   | (95.5% - 99.8%)  | 98.0  | 100.0 | 0/0                 |
|               | Total       | 472  | 100.0% (107/107) | (96.6% - 100.0%) | 99.2% (362/365)   | (97.6% - 99.8%)  | 97.3  | 100.0 | 0/0                 |
| Total         |             | 5387 | 99.3% (577/581)  | (98.2% - 99.8%)  | 99.4% (4779/4806) | (99.2% - 99.6%)  | 95.5  | 99.9  | 6/2                 |

A Asymptomatic  
 CI Confidence Interval  
 FNU Female Neat Urine  
 FS Female endocervical swab  
 FUPT Female urine in Q<sup>x</sup> UPT  
 FV Female vaginal swab  
 MNU Male Neat Urine  
 MS Male urethral swab  
 MUPT Male urine in Q<sup>x</sup> UPT  
 n number  
 S Symptomatic

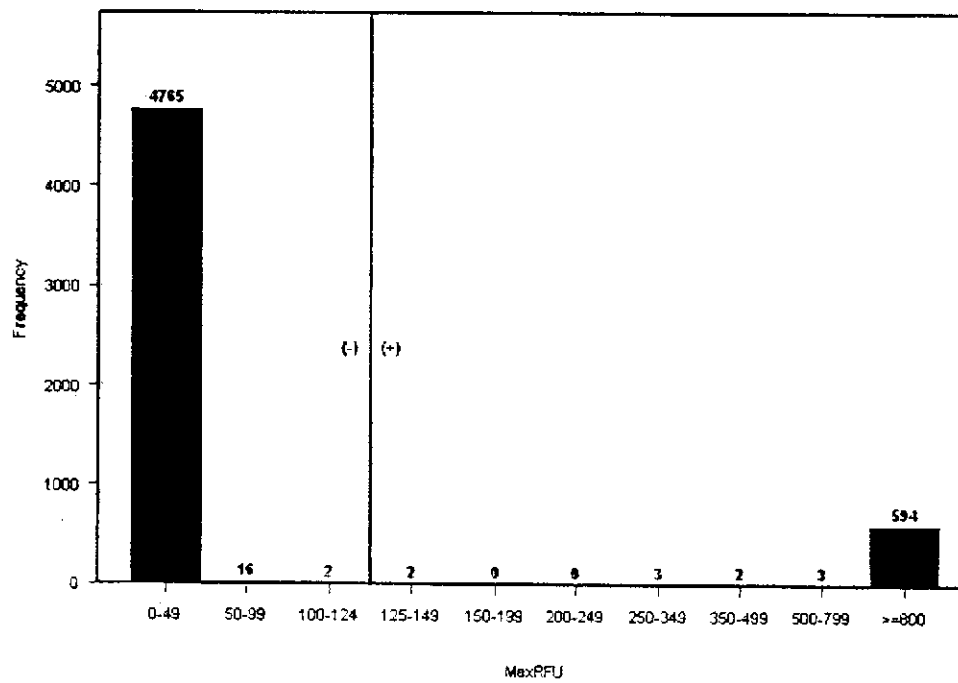


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A total of 5387 GC Q<sup>x</sup> Assay results was evaluated at seven geographically diverse clinical sites. A frequency distribution of the initial MaxRFU values for the GC Q<sup>x</sup> Assay with an assay cutoff of 125 MaxRFU is shown in **Figure A**.

**Figure A: Frequency Distribution of MaxRFU for the GC Q<sup>x</sup> Assay**





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#### Reproducibility

Reproducibility of the **BD Viper System** using the **BD ProbeTec GC Q<sup>x</sup> Assay** was evaluated at three clinical sites on one **BD Viper System** per site. A panel of simulated specimens was tested that comprised CT and GC organisms seeded into swab diluent for the **BD ProbeTec GC Q<sup>x</sup> Assay**. Simulated endocervical and urethral specimens contained a clean endocervical swab whereas the simulated urine and vaginal swab specimens did not. Uninoculated swab diluent for the **BD ProbeTec GC Q<sup>x</sup> Assay** was used for the GC negative samples. Nine replicates of each panel member were tested every day for five days on each **BD Viper System**. The data are summarized in **Table 4**.

**Table 4: Summary of Reproducibility Data on the BD Viper System for the GC Q<sup>x</sup> Assay**

| Specimen Type             | OT, CFU/ml | GC Cells/ml | Sensitivity         |                    | Mean   |       | SD     |      | CV   |      |       |
|---------------------------|------------|-------------|---------------------|--------------------|--------|-------|--------|------|------|------|-------|
|                           |            |             | %                   | (95% CI)           | Mean   | SD    | %      | CV   |      |      |       |
| Endocervical/<br>Urethral | 0          | 0           | 99.3%<br>(134/135)  | (95.9%,<br>100.0%) | 13.8   | 151.3 | 1096.3 | 0.0  | 0.0  | 0.6  | 4.3   |
|                           | 30         | 0           | 98.5%<br>(133/135)  | (94.8%,<br>99.8%)  | 28.1   | 220.7 | 785.3  | 0.0  | 0.0  | 33.8 | 120.3 |
|                           | 0          | 100         | 100.0%<br>(135/135) | (97.3%,<br>100.0%) | 1859.5 | 94.1  | 5.1    | 0.0  | 0.0  | 19.2 | 1.0   |
|                           | 30         | 250         | 100.0%<br>(135/135) | (97.3%,<br>100.0%) | 1847.3 | 117.6 | 6.4    | 0.0  | 0.0  | 25.9 | 1.4   |
|                           | 75         | 100         | 100.0%<br>(135/135) | (97.3%,<br>100.0%) | 1855.9 | 119.4 | 6.4    | 0.0  | 0.0  | 42.2 | 2.3   |
| Urine/Vaginal             | 0          | 0           | 99.3%<br>(134/135)  | (95.9%,<br>100.0%) | 15.7   | 162.3 | 1031.1 | 0.0  | 0.0  | 0.0  | 0.0   |
|                           | 30         | 0           | 100.0%<br>(135/135) | (97.3%,<br>100.0%) | 1.1    | 3.1   | 295.8  | 0.7  | 69.7 | 0.5  | 48.3  |
|                           | 0          | 100         | 100.0%<br>(135/135) | (97.3%,<br>100.0%) | 1899.0 | 86.1  | 4.5    | 22.8 | 1.2  | 0.0  | 0.0   |
|                           | 30         | 250         | 100.0%<br>(135/135) | (97.3%,<br>100.0%) | 1884.2 | 94.0  | 5.0    | 13.8 | 0.7  | 0.0  | 0.0   |
|                           | 75         | 100         | 100.0%<br>(135/135) | (97.3%,<br>100.0%) | 1867.2 | 87.7  | 4.7    | 0.0  | 0.0  | 19.2 | 1.0   |

A second study was conducted internally to characterize the reproducibility of test results (i.e., proportion positive or negative) at target levels below the analytical Limit of Detection (LOD) of the **BD ProbeTec GC Q<sup>x</sup> Assay**. A panel of simulated specimens was tested that comprised GC and CT organisms seeded into Q<sup>x</sup> swab diluent at two different levels each of which was below the respective analytical LOD for the organisms (1:10, 1:100). These levels were selected to fall within the dynamic range of the analytical LOD curve of the assay. Fifteen replicates of each panel member were tested every day for five days across three **BD Viper Systems**. The data are summarized in **Table 5**.





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**Table 15: Characterization of System Reproducibility at Target Levels below the Analytical Limit of Detection for the GC Q<sup>x</sup> Assay**

| Sample Type           | Dilution | Mean              | 95% CI       | Median | 95% CI            | 95% CI       | Median |
|-----------------------|----------|-------------------|--------------|--------|-------------------|--------------|--------|
| Endocervical/Urethral | 1:10     | 92.9<br>(209/225) | (88.7, 95.9) | 1324.6 | 7.1<br>(16/225)   | (4.1, 11.3)  | 41.4   |
| Endocervical/Urethral | 1:100    | 30.7<br>(69/225)  | (24.7, 37.1) | 835.9  | 69.3<br>(156/225) | (62.9, 75.3) | 7.2    |
| Urine/Vaginal         | 1:10     | 90.7<br>(204/225) | (86.1, 94.1) | 1165.9 | 9.3<br>(21/225)   | (5.9, 13.9)  | 34.2   |
| Urine/Vaginal         | 1:100    | 22.7<br>(51/225)  | (17.4, 28.7) | 872.7  | 77.3<br>(174/225) | (71.3, 82.6) | 7.8    |

### Conclusions

The analytical and clinical study results for the **BD ProbeTec** *Neisseria gonorrhoeae* (GC) Q<sup>x</sup> Amplified DNA Assay support the determination of substantial equivalence in accordance with the intended use as stated in the product labeling.



DEPARTMENT OF HEALTH & HUMAN SERVICES

Public Health Service

Food and Drug Administration  
2098 Gaither Road  
Rockville MD 20850

Ms. Kathryn Babka Carr  
Regulatory Affairs Specialist  
BD Diagnostics Systems  
Becton, Dickinson and Company  
7 Loveton Circle  
Sparks, MD 21152

DEC 11 2008

Re: K081825  
Trade/Device Name: BD ProbeTec™ *Neisseria gonorrhoeae* (GC) Q<sup>x</sup> Amplified DNA Assay  
Regulation Number: 21 CFR 866.3390  
Regulation Name: *Neisseria* spp. direct serological test reagents  
Regulatory Class: Class II  
Product Code: LSL  
Dated: December 5, 2008  
Received: December 9, 2008

Dear Ms. Babka Carr:

We have reviewed your Section 510(k) premarket notification of intent to market the device referenced above and have determined the device is substantially equivalent (for the indications for use stated in the enclosure) to legally marketed predicate devices marketed in interstate commerce prior to May 28, 1976, the enactment date of the Medical Device Amendments, or to devices that have been reclassified in accordance with the provisions of the Federal Food, Drug, and Cosmetic Act (Act) that do not require approval of a premarket approval application (PMA). You may, therefore, market the device, subject to the general controls provisions of the Act. The general controls provisions of the Act include requirements for annual registration, listing of devices, good manufacturing practice, labeling, and prohibitions against misbranding and adulteration.

If your device is classified (see above) into either class II (Special Controls) or class III (PMA), it may be subject to such additional controls. Existing major regulations affecting your device can be found in Title 21, Code of Federal Regulations (CFR), Parts 800 to 895. In addition, FDA may publish further announcements concerning your device in the Federal Register.

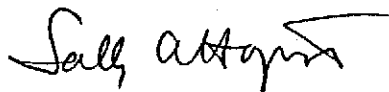
Please be advised that FDA's issuance of a substantial equivalence determination does not mean that FDA has made a determination that your device complies with other requirements of the Act or any Federal statutes and regulations administered by other Federal agencies. You must comply with all the Act's requirements, including, but not limited to: registration and listing (21 CFR Part 807); labeling (21 CFR Parts 801 and 809); and good manufacturing practice requirements as set forth in the quality systems (QS) regulation (21 CFR Part 820).

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This letter will allow you to begin marketing your device as described in your Section 510(k) premarket notification. The FDA finding of substantial equivalence of your device to a legally marketed predicate device results in a classification for your device and thus, permits your device to proceed to the market.

If you desire specific advice for your device on our labeling regulation (21 CFR Part 801), please contact the Office of In Vitro Diagnostic Device Evaluation and Safety at 240-276-0450. Also, please note the regulation entitled, "Misbranding by reference to premarket notification" (21CFR Part 807.97). For questions regarding postmarket surveillance, please contact CDRH's Office of Surveillance and Biometric's (OSB's) Division of Postmarket Surveillance at 240-276-3474. For questions regarding the reporting of device adverse events (Medical Device Reporting (MDR)), please contact the Division of Surveillance Systems at 240-276-3464. You may obtain other general information on your responsibilities under the Act from the Division of Small Manufacturers, International and Consumer Assistance at its toll-free number (800) 638-2041 or (240) 276-3150 or at its Internet address <http://www.fda.gov/cdrh/industry/support/index.html>.

Sincerely yours,



Sally A. Hojvat, M.Sc., Ph.D.  
Director  
Division of Microbiology Devices  
Office of *In Vitro* Diagnostic Device  
Evaluation and Safety  
Center for Devices and  
Radiological Health

Enclosure

