

# SPIFE® Nexus Split Beta SPE Procedure in Combination with Template Urine Application

Cat. No. 2420, 2421, 2422

## INTENDED USE

The SPIFE Nexus Split Beta SPE method is intended to quantitatively determine the presence of normal and abnormal serum proteins and qualitatively determine the presence of urine proteins by agarose electrophoresis using the SPIFE Nexus System.

For *In Vitro* Diagnostic Use Only.

Rx Only

## SUMMARY

Serum contains over one hundred individual proteins, with specific functions and various concentrations under different pathologic conditions.<sup>1</sup> Since introduction of moving-boundary electrophoresis by Tiselius<sup>2</sup> and the subsequent use of zone electrophoresis, serum proteins have been fractionated on the basis of their electrical charge at a particular pH into five classical fractions: albumin, alpha<sub>1</sub>, alpha<sub>2</sub>, beta and gamma proteins. Each of these classical electrophoretic zones, with the exception of albumin, normally contains two or more components. Knowing the relative proportions of these fractions has proven useful in the diagnosis and prognosis of certain disease states.<sup>3,4,5</sup>

## PRINCIPLE

Proteins are large molecules composed of covalently linked amino acids. Depending on electron distributions resulting from covalent or ionic bonding of structural subgroups, proteins can be either polar or nonpolar at a given pH. Proteins are separated according to their respective electrical charges on agarose gel using both the electrophoretic and electroendosmotic forces present in the system. The proteins are then stained with a visible stain.

## REAGENTS

### 1. SPIFE Split Beta SPE Gel

**Ingredients:** Each gel contains agarose in a tris-barbital/MOPS buffer with calcium lactate, a stabilizer, and a preservative.

**WARNING: FOR IN-VITRO DIAGNOSTIC USE ONLY.** The gel contains barbital which, in sufficient quantity, can be toxic.

**Preparation for Use:** The gels are ready for use as packaged.

**Storage and Stability:** The gels should be stored at room temperature (15 to 30°C) and are stable until the expiration date indicated on the package. The gels must be stored horizontally in the protective packaging in which they are shipped. **DO NOT REFRIGERATE OR FREEZE THE GELS.** Avoid storage close to a window or heat source, and avoid temperature variation during storage.

**Signs of Deterioration:** Any of the following conditions may indicate deterioration of the gel: (1) crystalline appearance indicating the agarose has been frozen, (2) cracking and peeling indicating drying of the agarose, (3) bacterial growth indicating contamination, (4) thinning of the gel blocks.

### 2. SPIFE Nexus Blue

**Ingredients:** The stain contains 0.5% (w/v) acid blue stain, 5% acetic acid, and surfactant.

**WARNING: FOR IN-VITRO DIAGNOSTIC USE ONLY. DO NOT INGEST.**

**Preparation for Use:** The stain is ready for use as packaged.

**Storage and Stability:** The stain solution is stable for one year when stored at 15 to 30°C in a closed container.

**Signs of Deterioration:** The prepared stain should be a homogeneous mixture free of precipitate. Discard if precipitate forms.

### 3. Citric Acid Destain

**Ingredients:** After dissolution, destain contains 0.3% (w/v) citric acid.

**WARNING: FOR IN-VITRO DIAGNOSTIC USE. DO NOT INGEST - IRRITANT.**

**Preparation for Use:** Pour 11 L of deionized water into Destain vat. Add full package of destain and mix until completely dissolved.

**Storage and Stability:** Store the destain at 15 to 30°C. It is stable until the expiration date on the package.

**Signs of Deterioration:** Discard if solution becomes cloudy.

## INSTRUMENT

A SPIFE Nexus analyzer must be used to apply serum samples, electrophorese, stain, destain, dry and then scan the gels. The gels may also be scanned on a separate densitometer such as the QuickScan Touch Plus (Cat. No. 1640). Refer to the Operator's Manuals for detailed instructions.

## SPECIMEN COLLECTION AND HANDLING

**Specimen:** Fresh serum or urine is the specimen of choice. Use of plasma will cause a fibrinogen band to appear as a distinct narrow band between the beta and gamma fractions.

**Storage and Stability:** If storage is necessary, store serum samples covered at 15 to 30°C for 4 days, 2 to 8°C for 2 weeks or -20°C for 6 months.<sup>6</sup> Urine samples may be stored covered at 2 to 8°C for up to 72 hours or at -20°C for 1 month.

## Interfering Factors:

1. Hemolysis may cause false elevation in the alpha<sub>2</sub> and beta fractions.
2. Uncovered specimens may yield inaccurate results due to evaporation.

## PROCEDURE

**Materials provided:** The following materials needed for the procedure are contained in the SPIFE Nexus Split Beta SPE Kit. Individual items are not available.

Test Size	Cat. No.
60 Samples	2420
40 Samples	2421
20 Samples	2422

**Cat. No. 2420, 2421, 2422**

SPIFE Split Beta SPE Gels (10)	
SPIFE Nexus Blue (1 vial)	
SPIFE Blotter C (10)	
Citric Acid Destain (1 pkg)	
Serrated Blade Applicator Kit, 20 Sample (10/20/30)	

## Materials provided but not contained in the kit:

Item	Cat. No.
SPIFE Nexus Analyzer	1650
QuickScan Touch Plus	1640
Gel Block Remover	1115
SPE Normal Control	3424
SPE Abnormal Control	3425
SPIFE Dispo Sample Cups, Deep Well	3360
SPIFE 20,40,60 Dispo Cup Tray	3370
SPIFE Nexus Cassette	2580
SPIFE Nexus Applicator Templates	2570
SPIFE Nexus Applicator Blade Weights	2572
SPIFE Nexus Dispo Stain Cups	2575
Pos ID Barcode Labels for Touch and SPIFE Nexus Systems	1696
REP Prep	3100
SPIFE Nexus Reagent Roller	2583
SPIFE Nexus Ready Run Kit	2582
SPIFE Nexus Carbon Electrode Insert	2576
SPIFE Urine Alignment Guide	3380
SPIFE Urine Protein Accessories	3427

## Materials needed but not provided:

0.85% saline

## STEP-BY-STEP METHOD

### I. Sample Preparation

To allow serum and urine samples to be run in combination on the same gel, the SPIFE Nexus analyzer pipettes the serum samples

into the sample tray. During serum sample processing, the gel is prepared with urine samples manually applied to the bottom rows of the gel by template application and then placed onboard. The test continues with blade application of the serum samples.

**Serum:** No specimen preparation is necessary for serum. Serum samples will be automatically pipetted into sample cups at a volume of 80 µL per sample.

**Urine:** Due to differences in the running parameters, urine specimens cannot be processed on the SPIFE Nexus in combination with serum samples unless urine specimens are applied using template application. Urine samples should be concentrated if a higher sensitivity is desired. A total protein range of 2,000-3,000 mg/dL is generally sufficient for optimum sensitivity. The sensitivity is 15-30 mg/dL for a single band.

Shake samples to homogenize. Centrifuge desired volume at 2000 x g for 5 minutes. Remove supernatant and concentrate to lab specifications per laboratory protocol.

## II. SPIFE Nexus Preparation

A. Fill designated bottles with 0.85% saline, deionized water, and destain.

B. Turn on the SPIFE Nexus. Click on the SPIFE Nexus icon to initialize.

C. If this is the first test of the day, prime the instrument according to the instructions in the SPIFE Nexus Operator's Manual.

D. Serum samples are placed onboard the instrument for sample processing. Load the correct number of uncapped patient serum sample test tubes into test tube racks and place racks within the tube transport area. **NOTE: Urine samples are not loaded into the tube transport area. During gel preparation, urine samples will be manually applied to the bottom rows of the gel by template application.**

E. Open the main door of the instrument and prepare the items onboard the instrument.

1. Ensure that the following items are in their respective onboard storage locations: **Platen Cover** with the Carbon Electrode Insert and **Dryer Cover** with the red sticker toward the back of the instrument.

### 2. Sample Cup Tray

a. Prepare the sample cup tray with the appropriate disposable deep well sample cups. Slide the Disposable Sample Cups into the cup tray that correspond to the loaded serum samples. Use only the top row for 20 or fewer samples, top and middle rows for up to 40 samples.

b. Place the cup tray onto the sample tray platform.

### 3. Stain/Reagent Dispenser

a. Fill three Stain Cups each with 700 µL of SPIFE Nexus Blue stain and place a Stain Cup in each slot of the Stain/Reagent Dispenser.

b. Place a clean Reagent Roller bar between the hooks on the Stain/Reagent Dispenser.

### 4. Consumables Tray

a. Slide the Consumables Tray forward from its home position.

b. Prepare the Applicator Holder

(1) Place a Split Beta (60) Applicator Template on top of the Applicator Holder. Place Applicator Blades in the designated slots corresponding to the sample cups loaded within the sample tray. **NOTE: The Applicator Blades will only fit into the slots in the Applicator Holder one way; do not try to force the Applicator Blades into the slots.**

(2) Place the Applicator Blade Weights on top of the Applicator Blades with the thick side facing the front of the instrument.

c. Slide the Consumables Tray into position in the back of the instrument.

### 5. Gel Cassette

a. Place an empty Gel Cassette on the electrophoresis platen. Make sure that the 2D barcode is located in the upper right corner of the cassette.

F. Close the main door of the instrument.

## III. Sample Loading

A. Click the Start button on the menu bar. Select the **SPIFE Split Beta Serum Proteins with Urine Template 60 (Acid Blue)** test name from the drop down menu. Ensure the toggles for all Run Processes are set to "Yes" and click the Start Run button. The analyzer will load serum samples.

B. While the SPIFE Nexus automatically loads serum samples, proceed with template application of the urine specimens as follows.

1. Remove the gel from the protective packaging and discard the overlay. Carefully place the gel on the SPIFE Urine Alignment Guide.

2. Using a SPIFE Blotter C, gently blot the entire gel. Discard the blotter.

3. The Urine Templates have been marked with a hole in one corner. Hold the template so that the marked corner is in the lower left position. Align the template pin holes with the pins on the sides of the Alignment Guide and carefully slide the template over the alignment pins until the template contacts the gel surface. Apply slight fingertip pressure across the template, making sure there are no air bubbles between the gel and the template.

4. Pipette 3 µL of each urine sample onto an individual slit of the Urine Template. After pipetting the final sample, use the top half of the clam shell to carefully cover the gel in the Alignment Guide.

5. Allow urine samples to absorb for 5 minutes.

6. After 5 minutes, uncover the gel and gently blot the unabsorbed urine from the Urine Template with a Blotter A-Plus.

7. Carefully remove the blotter and templates and discard as biohazardous waste.

8. Cover the gel until the SPIFE Nexus has completed the serum sample processing.

C. When the SPIFE Nexus has completed serum sample processing, the alarm will sound, the door will unlock, and the following message will be displayed: **Place gel with template-applied urine samples in the gel cassette. Close the door and click 'OK' to continue.**

D. Open the main door of the SPIFE Nexus and place the gel with template-applied urines into the cassette as follows:

1. Remove the top half of the Gel Cassette from the electrophoresis platen.

2. Dispense 2 mL of REP Prep on the platen.

3. Place the left edge of the gel into the bottom of the cassette fitting the round hole over the upper pin and the obround hole over the lower pin. Gently lay the gel down over the REP Prep making sure no bubbles remain under the gel.

4. Place the top half of the Gel Cassette over the gel. Make sure the 2D barcode is located in the upper right corner of the cassette. **Note: Do not use a Positive ID Barcode Label on the gel for this test type.**

E. Close the main door of the instrument.

## IV. Automated Gel Electrophoresis and Scanning

A. Click the OK button on the onscreen message. The analyzer will apply serum samples, electrophorese, stain, destain, dry and scan the serum samples. For details of Automated Gel Electrophoresis parameters, contact Technical Services.

B. After scanning the serum samples, the Gel Cassette with the finished gel will be located in the scanner port on the front side of the instrument.

1. On the finished gel, record the Gel Identifier number beginning with T000 displayed on the open Worklist. This will serve as the serum scan record locator.

2. If scanning of the urine samples is desired, perform a second scan of the gel as follows:

a. Clean any residual stain from the electrophoresis platen, and ensure that the platen and the back of the finished gel are clean and dry.

- b. Lay the cassette containing the finished gel on the platen. Make sure the 2D barcode is located in the upper right corner of the cassette.
- c. Close the main door of the instrument.
- d. Click the Start button on the menu bar. Select the **SPIFE Urine Proteins Scanning for Template Application 60 (Acid Blue)** test name from the drop down menu. Ensure that only the toggle for the Scan process is set to "Yes" and click the Start Run button.
- e. The SPIFE Nexus will move the finished gel from the platen to the scanner. After scanning the urine samples, the Gel Cassette with the finished gel will be located in the scanner port on the front side of the instrument.
- f. Record on the finished gel the Gel Identifier number beginning with T000 displayed on the open Worklist. This will serve as the urine scan record locator.

- C. If gel storage is required, remove and discard the two gel blocks.
- D. After every test: discard the used blotters, Applicator Blades, Stain Cups and sample cups as biohazardous waste. Clean any residual stain from the electrophoresis platen, Gel Cassette and the Reagent Roller bar. For daily, weekly, and monthly maintenance, reference the SPIFE Nexus Operator's Manual.

### Evaluation of the Protein Bands

**Quantitative Evaluation of Serum:** The SPIFE Nexus Split Beta SPE Gel will be automatically scanned. An aperture size of 5 with the acid blue setting is recommended. Refer to the QuickScan Touch Plus Operator's Manual for scanning parameters.

**Qualitative Evaluation of Urine:** The urine samples run on the SPIFE Nexus Split Beta SPE Gel can only be visually inspected for the presence of the bands.

**Stability of End Product:** The completed, dried SPIFE Split Beta SPE Gel is stable for an indefinite period of time.

**Quality Control:** SPE Normal Control (Cat. No. 3424) and SPE Abnormal Control (Cat. No. 3425) may be used to verify all phases of the procedure and should be used on each gel run. If desired, a control or patient sample may be diluted 1:7 with 0.85% saline (1 part sample + 6 parts saline) and run with urines for qualitative comparison. Refer to the package insert provided with the control for assay values.

### REFERENCE VALUES

The reference ranges presented were established with the Split Beta SPE System on 38 normal specimens using the SPIFE Nexus. These values are presented as a guideline.

Protein Fraction	% of Total Mean ± 2 S.D.
Albumin	46.2 - 63.0
Alpha <sub>1</sub>	1.7 - 4.4
Alpha <sub>2</sub>	6.8 - 13.7
Beta	12.5 - 18.5
Gamma	10.6 - 22.7

Each laboratory should perform its own normal range study.

### Variations of Expected Values<sup>5</sup>

Studies show that values are the same for both males and nonpregnant females (Some differences are seen in pregnant females at term and in women on oral contraceptives).

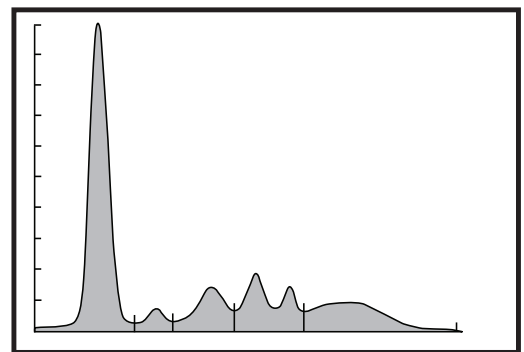
Age has some effect on normal levels. Cord blood has decreased total protein, albumin, alpha<sub>2</sub> and beta fractions, slightly increased alpha<sub>1</sub> and normal or increased gamma fractions (largely of maternal origin). The gamma globulins drop rapidly until about three months of age, while the other fractions have reached adult levels by this time. Adult levels of the gamma globulins are not reached until 16 years of age. The albumin decreases and beta globulin increases after the age of 40.

### RESULTS

Figure 1 illustrates the electrophoretic mobilities of the albumin, alpha<sub>1</sub>, alpha<sub>2</sub>, beta and gamma protein bands on a SPIFE Split Beta SPE Gel. The fastest moving band, and normally the most prominent, is the albumin band found closest to the anodic edge of the gel. The faint band next to this is alpha<sub>1</sub>, followed by alpha<sub>2</sub> globulin, split beta and gamma globulins.



**Figure 1:** A SPIFE Split Beta SPE-60 Gel showing relative position of the bands.



**Figure 2:** A scan of a SPIFE Split Beta SPE pattern.

### Calculations of the Unknown

The SPIFE Nexus scanner will automatically calculate and print the relative percent and the absolute value of each band when the total protein is entered. Refer to the SPIFE Nexus and QuickScan Touch Plus Operator's Manuals provided with the instrument.

### INTERPRETATION OF RESULTS<sup>5,6</sup>

Results on normal individuals will cover age and sex-related variations and day-to-day biologic variations. Disease states in which abnormal patterns are observed include inflammatory response, rheumatic disease, liver diseases, protein-loss disorders, plasma cell dyscrasias, infectious disorders, renal disorders, pregnancy, and genetic deficiencies. Patients with high levels of IgG4 can produce a relatively restricted band cathodic in the beta gamma region or beta gamma bridging.<sup>7</sup> Proteins migrating in the alpha<sub>2</sub> and beta region may show slight variation in migration under a variety of circumstances.<sup>8</sup>

### Further Testing Required

The serum or urine protein electropherogram, or densitometric tracing, should be evaluated for abnormalities. If abnormalities are observed, appropriate follow-up studies should be initiated.<sup>9</sup> Not all clinically significant monoclonal gammopathies will display a distinct band detectable by protein electrophoresis.<sup>10</sup> Further studies may be indicated based on clinical context. These may include immunofixation, quantitation of immunoglobulins, bone marrow examination and other appropriate tests.

## LIMITATIONS

1. Since all electrophoretic procedures are nonlinear, it is critical to fill the wells with the recommended volume of serum or urine to obtain optimal resolution and reproducible results. Noncompliance with the recommended procedure may affect the results.
2. Therapeutic monoclonal antibodies may be used in the treatment of multiple myeloma as well as various other malignancies or medical conditions. If present in sufficient concentration, these agents may be indistinguishable from a pathologic monoclonal protein on serum protein electrophoresis.<sup>11,12</sup>
3. Use of plasma will cause a fibrinogen band to appear as a distinct narrow band between the beta and gamma fractions. A fibrinogen band may also be present in patients on heparin therapy.
4. Hemolyzed samples should be avoided as the changes in hemoglobin-haptoglobin may affect the alpha<sub>2</sub> and beta migration.<sup>13,14</sup>
5. The mobility of beta lipoprotein (low density lipoprotein) can vary considerably and may migrate under normal conditions anywhere between the alpha<sub>2</sub> and beta region. Beta lipoproteins can be recognized by their characteristic appearance as a thin, irregular line, regardless of its migration location.
6. An artifact may be present at the point of application, particularly with the use of frozen samples, older samples, or samples containing debris.<sup>13</sup> An application artifact may appear as fine clear line (negative space) that may be visible to a faint degree across the entire gel in the beta region. This can on occasion cause the edge of a normal blush to appear slightly blunted.

## SPECIFIC PERFORMANCE CHARACTERISTICS

Reproducibility was assessed over a 5 day period. Normal and abnormal serum controls were tested on two gels per day on each of three SPIFE Nexus instruments. Three hundred sixty determinations per protein fraction in total were collected for the normal and abnormal serum protein controls respectively.

Normal Control Fraction	N	Mean %	Within Day		Between Day		Between Instrument		Total	
			SD	CV%	SD	CV%	SD	CV%	SD	CV%
Albumin	360	60.3	1.94	3.21	0.26	0.43	1.95	3.24	2.06	3.42
Alpha <sub>1</sub>	360	3.2	0.40	12.53	0.10	3.05	0.41	12.9	0.43	13.53
Alpha <sub>2</sub>	360	9.8	0.99	10.15	1.00	10.21	1.41	14.4	1.47	14.95
Beta	360	14.0	1.02	7.26	0.87	6.24	1.34	9.57	1.34	9.58
Gamma	360	12.7	1.39	10.95	0.53	4.15	1.49	11.71	1.50	11.84

Normal Control Fraction	N	Mean %	Within Day		Between Day		Between Instrument		Total	
			SD	CV%	SD	CV%	SD	CV%	SD	CV%
Albumin	360	57.7	1.63	2.82	0.24	0.42	1.64	2.85	1.75	3.04
Alpha <sub>1</sub>	360	2.6	0.31	12.07	0.02	0.91	0.31	12.10	0.34	13.00
Alpha <sub>2</sub>	360	7.8	0.56	7.21	0.36	4.62	0.67	8.56	0.67	8.61
Beta	360	10.4	0.64	6.18	0.22	2.10	0.68	6.52	0.76	7.33
Gamma	360	21.4	1.34	6.24	0.10	0.49	1.33	6.22	1.35	6.32

## SENSITIVITY

A pathological serum sample with a monoclonal protein at 1.76 g/dL (1760 mg/dL) was serially diluted and the dilutions electrophoresed on the SPIFE Split Beta SPE gel on the SPIFE Nexus. After visual inspection and densitometric analysis of the gel, the lowest detectable concentration of a monoclonal protein was between 0.014 and 0.028 g/L (14 and 28 mg/dL).

Serial dilutions of a pathological urine sample containing a monoclonal were analyzed on the SPIFE Nexus using template application on the SPIFE Split Beta SPE gel. The sensitivity was determined to be between 0.015 and 0.030 g/L (15 and 30 mg/dL) for a single band.

**NOTE:** The migration position of the monoclonal protein and the presence of a polyclonal background in the gamma zone may affect the detection limit.

## CORRELATION

Normal and abnormal serum samples were analyzed using the SPIFE Touch Split Beta SPE system and the SPIFE Nexus Split Beta SPE system. Deming regression with 95% confidence intervals and Pearson correlation coefficient are presented below.

n = 30

Slope: 1.046 (1.026 to 1.066)

Intercept: -0.93 (-1.48 to -0.38)

R = 0.99

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