### PERKINELMER, INC.



# AlphaLISA<sup>®</sup> Assay Development Guide

For Laboratory Use Only Research Chemicals for Research Purposes Only

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### **Precautions**

- Alpha streptavidin-coated Donor beads (referred in this guide as streptavidin Donor beads) are light-sensitive. All assays using the streptavidin Donor beads should be performed under subdued laboratory lighting of less than 100 lux. Alternatively, green filters (e.g., Roscolux filters #389 from Rosco Laboratories, Inc., or equivalent) can be applied to light fixtures. Any incubation of streptavidin Donor beads should be performed in the dark.
- Due to the small volumes used in the assay, it is recommended to cover microplates with TopSeal-A<sup>TM</sup> adhesive sealing film to reduce evaporation during incubation periods (PerkinElmer, Inc., Cat. No. 6005185). The microplates can be read with the TopSeal-A film in place, except if condensation is present.
- AlphaLISA Acceptor beads in the stock solution may slightly sediment over several days. This is normal. It is advised to vortex the beads prior to use.
- Streptavidin Donor beads and AlphaLISA Acceptor beads should be stored in the dark at 4°C.
- AlphaLISA is intended for research purposes only and is not for use in diagnostic procedures.

### I. BEFORE STARTING

### Receiving the AlphaLISA Acceptor Beads

Upon receiving the AlphaLISA Acceptor beads, ensure that they are on blue ice and that the ice packs are not completely thawed.

### **Reagents Provided**

The following sizes are available:

1 mg (catalog number 6772001)

5 mg (catalog number 6772002)

50 mg (catalog number 6772003)

Bulk format (catalog number 6772004)

Table I: Reagents provided

COMPONENT	1 mg	5 mg	50 mg
	6772001	6772002	6772003
AlphaLISA Acceptor beads stored in 0.1M Tris-HCl pH 8.0, 0.05% Proclin-300	50 μL (20 mg/mL)	250 μL (20 mg/mL)	5 x 500 μL (20 mg/mL)

### Notes before use

- For maximum recovery of contents, briefly spin the vials prior to removing the caps and resuspend the beads by vortexing.
- AlphaLISA Acceptor beads should be stored at +2 to +8°C.

### **Recommended Additional Reagents and Materials**

**Table II**: Recommended reagents and materials

Item	Suggested source	Catalog #
AlphaScreen Streptavidin-coated Donor beads	PerkinElmer, Inc.	6760002S (1 mg) 6760002 (5 mg) 6760002B (50 mg)
Analyte of choice	N/A	N/A
Pair of anti-analyte antibodies	N/A	N/A
Microplate	PerkinElmer, Inc.	see Table III
TopSeal-A Adhesive Sealing Film	PerkinElmer, Inc.	6005185 (pack size of 100 pieces)
Single-channel pipettors <sup>§</sup>	N/A	N/A
NHS activated biotinylating reagent (ChromaLink)	SoluLink Inc.	B1001-105
Carboxymethylamine hemihydrochloride (CMO)	Sigma-Aldrich Co.	C13408
Sodium cyanoborohydride	Sigma-Aldrich Co.	152159
Zeba desalting columns	Pierce (ThermoFisher Scientific Inc.)	89882 (0.5 mL) 89889 (2 mL) 89891 (5 mL) 89893 (10 mL)
Proclin-300	Sigma-Aldrich Co.	48912-U
Tween-20 (Surfact-Amps 20)	Pierce (ThermoFisher Scientific Inc.)	28320
Dextran 500 MW ~500000	Sigma-Aldrich Co.	D1037
Casein 5% Alkali-soluble solution	Novagen (EMD Chemicals Inc.)	70955
Triton-X100 (Surfact Amps X100)	Pierce (ThermoFisher Scientific Inc.)	28314
Streptavidin-Sepharose beads	GE Healthcare, Inc.	17-5113-01

 $<sup>^\</sup>S$  For lower volume additions (< 10  $\mu$ L), we recommend a pipettor precision of  $\leq$  2%. For higher volume additions (25-1000  $\mu$ L), a pipettor precision of  $\leq$  1% is recommended.

Table III: Microplates

Assay Format	Microplate	Catalog # §	Total well volume	Recommendations
	OptiPlate™-96	6005290	400 μL	ldeal for assay volume ≥ 50 μL
96	½ AreaPlate-96	6005560	180 µL	Ideal for assay volume of 50 µL in 96-well format
	ProxiPlate™-96	6006290	100 μL	Not recommended
	OptiPlate-384	6007290	105 µL	Good for assays of 25-50 µL in 384-well format
384	AlphaPlate™-384	6005350	105 µL	Ideal for high-count assays of 25-50 µL in 384-well format (lower cross-talk than OptiPlate-384)
	ProxiPlate-384	6008280	30 µL	Good for assays of 10-25 µL in 384-well format
1536	AlphaPlate-1536	6004350	12 µL	Ideal for assays of 5-10 µL in 1536-well format (lower cross-talk than OptiPlate-1536)
	OptiPlate-1536	6004290	12 μL	Not recommended for AlphaLISA due to higher cross-talk

<sup>§</sup> For 50 plates; refer to PerkinElmer, Inc. website for other quantities

The protocols included in this manual are for OptiPlate-384 microplates. Other plate types listed in Table III may be used with the appropriate assay volume.

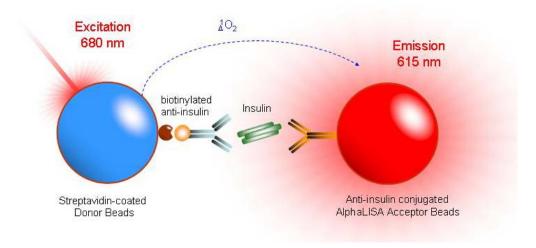
Assays must be read using an AlphaScreen-compatible reader such as all  $EnVision^{\mathbb{R}}$  multilabel plate readers with an AlphaScreen module, the Fusion-Alpha $^{\mathbb{T}}$  multilabel reader or the AlphaQuest $^{\mathbb{R}}$  original AlphaScreen reader (PerkinElmer, Inc.). The normal AlphaScreen settings should be used.

### II. INTENDED USE

The AlphaLISA Acceptor beads are intended to perform biomarker detection (referred in this guide as analyte), protein-protein interaction, intracellular target detection assays or any other biological detection assay. AlphaLISA is for research purposes only and not for use in any diagnostic procedures.

### III. PRINCIPLE OF THE ASSAY

The AlphaLISA technology allows the detection of molecules of interest in serum, plasma, cell culture supernatants or cell lysates in a very sensitive, quantitative, reproducible, and user-friendly assay. Almost, any sandwich assay can be developed to detect an analyte of interest. As shown in Figure 1, an anti-analyte antibody which is biotinylated binds the streptavidin Donor bead while another anti-analyte antibody is conjugated to AlphaLISA Acceptor beads. In the presence of the analyte, the beads come into close proximity. The excitation of the Donor beads provokes the release of singlet oxygen molecules that triggers a cascade of energy transfer in the Acceptor beads, resulting in light emission. For more details about the AlphaLISA technology, please refer to the guide entitled 'A Practical Guide to Working with AlphaScreen'.



**Figure 1**: Illustration of the detection of an analyte (insulin in this example) using AlphaLISA Acceptor beads

### IV. ASSAY DEVELOPMENT

A few simple steps are needed to develop an AlphaLISA analyte assay. This generic protocol is the starting point for the development of all AlphaLISA assays. Some assays may require modified protocols based on the specific nature of the analyte under investigation.

### A. Selection of antibody pairs

To develop an AlphaLISA analyte detection assay, an antibody pair that recognizes the targeted analyte is required. It is always preferable to find an antibody pair that has already been tested in a sandwich assay. Most of the identified pairs should perform well in AlphaLISA. If no identified pairs are available, different antibodies need to be tested to find a pair that will perform optimally in AlphaLISA. All antibodies must be selected according to the following guidelines:

- Each antibody should recognize a different epitope on the analyte.
- The two antibodies must be specific to the analyte.
- Monoclonal or purified polyclonal antibodies will perform best; avoid the use of anti-sera.
- The antibodies must not be in any amine-based buffer, including Tris, glycine, bicine, tricine, etc. If buffer exchange is necessary, the buffer should be replaced with a neutral to slightly alkaline buffer, such as PBS or carbonate buffer pH 8.
- Antibody solutions must not contain any protein or peptide-based stabilizers (such as BSA or gelatin).
- Optimal results will be obtained with antibody concentrations above 0.5 mg/mL.

One of the two antibodies will have to be biotinylated and the other one should be conjugated to AlphaLISA Acceptor beads. It is recommended to test the two possible combinations as the sensitivity and the level of counts may differ dramatically depending on the set-up. The two combinations are as follows:

- 1. biotinylated antibody A + antibody B-AlphaLISA Acceptor beads
- 2. biotinylated antibody B + antibody A-AlphaLISA Acceptor beads

Therefore, it is recommended to biotinylate both antibodies of the pair as well as conjugating both to AlphaLISA acceptor beads to eventually select the optimal assay configuration.

(Note: Alternatively, the antibodies can be captured in an indirect assay set-up using anti-species IgG AlphaLISA Acceptor beads and/or biotinylated anti-species IgG. See AlphaLISA products on p. 24.)

### B. Biotinylation of the antibody

The following protocol is used for biotinylation of antibodies:

- a. A preliminary check of the antibody to be biotinylated is mandatory. The User must check for the following:
  - The biotinylation will perform best when the antibody concentration is at least 0.5 mg/mL.
  - The antibodies must not be in any amine-based buffer, including Tris, glycine, bicine, tricine, etc. If buffer exchange is necessary, the buffer should be replaced with a neutral to slightly alkaline buffer, such as PBS or carbonate buffer pH 8. Optimal performance will be obtained in sodium azide-free conditions.
  - Antibody solutions must not contain any protein or peptide-based stabilizers (such as BSA or gelatin).
  - The antibody will be labeled at slightly alkaline pH values (7.0-8.0) in aqueous buffer. Ensure that the antibody is soluble in these conditions.
- b. Prepare antibody solution. If the antibody is provided in a powder form, resuspend at 5 mg/mL in PBS. If already in solution at neutral to slightly alkaline pH (pH ≥ 7.0), use as provided.
- c. On the day of the assay, prepare a fresh solution of N-hydroxysuccinimido-ChromaLink-biotin (NHS-ChromaLink-biotin) at 2 mg/mL in PBS. Alternatively, other NHS reagents such as NHSbiotin, NHS-LC-biotin or NHS-LC-LC-biotin can also be used.
- d. Add NHS-ChromaLink-biotin to the antibody solution. A 30:1 molar ratio of biotin to antibody is recommended. This represents using 7.6  $\mu$ L of a 2 mg/mL NHS-ChromaLink-biotin solution for 100  $\mu$ g of antibody. Adjust the volume to 200  $\mu$ L with phosphate buffer pH 7.4.
- e. Incubate for 2 hours at 21-23°C.
- f. When using NHS-ChromaLink-biotin, a purification step using Zeba columns is required to remove free biotin. To evaluate biotinylation efficiency, refer to SoluLink's website (http://solulink.com).

### C. AlphaLISA acceptor bead conjugation with the antibody

### C1. Preliminary Notes

A preliminary check of the material to be conjugated is mandatory. The User must check for the followings:

 The conjugation will perform best when the antibody concentration is at least 1 mg/mL (when conjugating 1-2 mg of beads) or 0.53 mg/mL (when conjugating 2.5 mg of beads or higher amounts). Lower concentrations of antibody yield lower coupling efficiency.

- The antibodies must not be in any amine-based buffer, including Tris, glycine, bicine, tricine, etc. If buffer exchange is necessary, the buffer should be replaced with a neutral to slightly alkaline buffer, such as PBS or carbonate buffer pH 8. (Note: Although both buffers can be utilized, for clarity purposes, phosphate buffer will be used in the protocol below).
- Ideally, antibody solutions should not contain any protein or peptide-based stabilizers (such as BSA or gelatin). However in the presence of such substances, the conjugation process can still be performed, but may result in lower coupling efficiency in some cases.
- Glycerol will significantly impact coupling efficiency (10% glycerol final in the conjugation mix will cause approx. 50% signal reduction). Dialysis of the antibody is recommended prior to coupling.

The ratio of antibody to mg of beads is an **important parameter** for a successful assay development. Typical coupling weight ratios (amount of beads to amount of antibody) vary from 10:1 to 50:1. When preparing low amounts of beads (1-2 mg), a 10:1 ratio is recommended (i.e. 1 mg of Acceptor beads to 0.1 mg of antibody), while a ratio of 50:1 is used with bead amounts equal to or higher than 2.5 mg to minimize the antibody consumption (i.e. 5 mg of Acceptor beads to 0.1 mg of antibody).

# C2. Protocol for conjugating <u>1 mg</u> AlphaLISA Acceptor beads (10:1 coupling ratio)

This procedure is appropriate for a solution of antibody ≥ 1 mg/mL; if the concentration is below 1 mg/mL, the antibody solution must be concentrated using an iCON Concentrator (Pierce Cat# 89886), Microcon or Centricon (Ultracell YM-30, Millipore, cat# 4208 or 42409).

### **Washing**

 Wash AlphaLISA Acceptor beads (50 μL at 20 mg/mL) once with 50 μL PBS: centrifuge at 16,000 × g or maximum speed for 15 min and then discard the supernatant.

### Conjugation

In an eppendorf tube, add the following reagents:

1 mg of AlphaLISA Acceptor bead pellet (prepared as described above)

- 0.1 mg of antibody
- the appropriate volume of 0.13 M phosphate buffer pH 8.0 to obtain a final reaction volume of 200 µL
- 1.25 μL of 10% Tween-20
- 10 μL of a 25 mg/mL solution of NaBH<sub>3</sub>CN in water (prepare fresh as required).

Incubate for 18-24 hours at 37°C. Longer incubation times up to 48 hours might be used, which could result in higher conjugation yields.

### **Blocking**

- Prepare a fresh 65 mg/mL solution of carboxy-methoxylamine (CMO) in a 0.8 M NaOH solution.
- Add 10 μL of CMO solution to the reaction (to block unreacted sites).
- Incubate for 1 hour at 37°C.

### **Purification**

- Centrifuge for 15 minutes at 16,000 × g (or maximum speed).
- Remove the supernatant with a micropipette and resuspend the bead pellet in 200 μL of 0.1 M Tris-HCl pH 8.0.
- Centrifuge for 15 minutes at 16,000 × g (or maximum speed), then remove the supernatant.
- Repeat the washing steps (resuspend the pellet and centrifuge) another time.
- After the last centrifugation, resuspend the beads at 5 mg/mL in storage buffer (200 μL of PBS + 0.05% Proclin-300 as a preservative).
- Vortex, briefly spin down and sonicate the bead solution (20 short pulses of 1 second using a probe sonicator).

### Storage

• Store the conjugated Acceptor bead solution at 4°C. <u>Important note</u>: always vortex conjugated AlphaLISA Acceptor beads before use, as beads tend to settle with time.

# C3. Protocol for conjugating <u>5 mg</u> AlphaLISA Acceptor beads (50:1 coupling ratio)

This procedure is appropriate for a solution of antibody ≥ 0.53 mg/mL; if the concentration is below 0.53 mg/mL the antibody solution must be concentrated using an iCON Concentrator (Pierce Cat# 89886), Microcon or Centricon (Ultracell YM-30, Millipore, cat# 4208 or 42409).

### **Washing**

 Wash AlphaLISA Acceptor beads (250 μL at 20 mg/mL) once with 250 μL PBS: centrifuge at 16,000 × g or maximum speed for 15 min and then discard the supernatant.

### Conjugation

In an eppendorf tube, add the following reagents:

- 0.1 mg of antibody to 5 mg of bead pellet (prepared as described above)
- the appropriate volume of 0.13 M phosphate buffer pH 8.0 to obtain a final reaction volume of 200 µL
- 1.25 µL of 10 % Tween-20
- 10 μL of a 25 mg/mL solution of NaBH<sub>3</sub>CN in water (prepare fresh as required).

Incubate for 18-24 hours at 37°C.

### **Blocking**

- Prepare a fresh 65 mg/mL solution of carboxy-methoxylamine (CMO) in a 0.8 M NaOH solution.
- Add 10 μL of CMO solution to the reaction (to block unreacted sites).
- Incubate for 1 hour at 37°C.

### **Purification**

- Centrifuge for 15 minutes at 16,000 × g (or maximum speed).
- Remove the supernatant with a micropipette and resuspend the bead pellet in 1 mL of 0.1 M Tris-HCl pH 8.0 (200 μg per mg of beads).
- Centrifuge for 15 minutes at 16,000 × g (or maximum speed), then remove the supernatant.
- Repeat the washing steps (resuspend the pellet and centrifuge) another time.
- After the last centrifugation, resuspend the beads at 5 mg/mL in storage buffer (1 mL of PBS + 0.05% Proclin-300 as a preservative).
- Vortex, briefly spin down and sonicate the bead solution (20 short pulses of 1 second using a probe sonicator).

### Storage

Store the conjugated Acceptor bead solution at 4°C.

<u>Important note</u>: always vortex conjugated AlphaLISA Acceptor beads before use, as beads tend to settle with time.

### D. Determination of the best antibody pair

One of the first development steps is the selection of the antibody combination. As previously mentioned, both combinations of antibody set-up must be tested:

- 1. biotinylated antibody A + antibody B-AlphaLISA Acceptor beads
- 2. biotinylated antibody B + antibody A-AlphaLISA Acceptor beads

This section describes the method to be used to perform this selection.

### D1. Selection of optimal antibody pair(s) using a matrix assay

A matrix assay should be performed using fixed concentrations of antibodies and 2-3 different concentrations of analyte (within the usual working range) and a negative control (no analyte).

It is important to note that this method is a **guideline**; it will need to be optimized for every analyte studied.

These preliminary conditions could be used for this experiment:

### Typical assay buffer:

- 25 mM HEPES pH 7.4
- 0.5% Triton X-100
- 0.1% Casein
- 1 mg/mL Dextran 500
- Adjust pH to 7.4

This buffer is available from PerkinElmer, Inc. as a 10X solution:

AlphaLISA ImmunoAssay Buffer (10X) (Cat No. AL000). Dilute 10-fold with water prior to use.

In an OptiPlate-384 microplate add:

- 5 μL of the analyte diluted in assay buffer (use various concentrations that are in the working range for the target detection)
- 10 μL of fixed amount of the biotinylated antibody (Use a final concentration between 0.3 nM and 3 nM of biotinantibody. Often the optimal biotinylated antibody

concentration is 1 nM.)

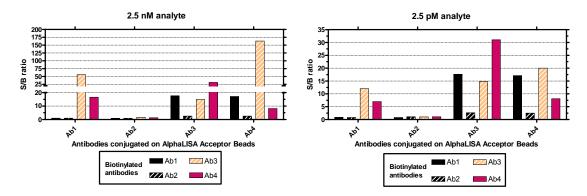
 10 μL of antibody-coupled AlphaLISA Acceptor beads at 50 μg/mL (10 μg/mL final concentration in each well)

Incubate at 23°C for 1 h then add:

 25 μL of streptavidin Donor beads at 80 μg/mL prepared under subdued light conditions (40 μg/mL final concentration in each well)

Incubate at 23°C in the dark for 30 min and read on an AlphaScreen reader.

Select the pair(s) of antibodies providing the highest signal-to-background (S/B) ratio and the best sensitivity as defined as the highest S/B ratio for the condition with lowest concentration of analyte. Once selected, determination of optimal biotinylated antibody concentration can be performed.



**Figure 2**: Determination of the best antibody pair. S/B ratio obtained for each antibody pair (each permutation possible), with 2.5 nM and 2.5 pM of analyte (S/B ratio is calculated as AlphaLISA signal obtained for 2.5 nM or 2.5 pM analyte ÷ AlphaLISA signal obtained without analyte (background)). In this example, Antibody-Ab3-conjugated AlphaLISA Acceptor Beads and Biotinylated-Antibody-Ab4 were selected.

## D2. Determination of optimal biotinylated antibody concentration

An antibody titration curve should be performed using a fixed concentration of analyte (within the usual working range).

The following preliminary conditions may be used for this titration.

It is important to note that the method is a **guideline**; it will need to be optimized for every analyte studied.

In an OptiPlate-384 microplate add:

- 5 µL of the analyte diluted in assay buffer (use a fixed concentration that is in the working range for the target detection)
- 10 μL of increasing amounts of the biotinylated antibody (as a starting point, use concentrations between 0.1 nM to 100 nM final concentration of biotin-antibody)
- 10 μL of antibody-coupled AlphaLISA Acceptor beads at 50 μg/mL (10 μg/mL final concentration in the well)

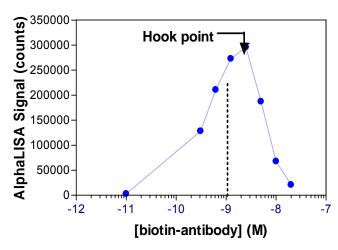
Incubate at 23°C for 1 h then add:

 25 μL of streptavidin-Donor bead solution at 80 μg/mL prepared under subdued light conditions (40 μg/mL final concentration in each well)

Incubate at 23°C in the dark for 30 min and read on an AlphaScreen reader.

A bell-shaped curve should be obtained (Figure 3). The highest signal obtained indicates the hook point (highest signal before saturation of the bead binding capacity) of the biotinylated anti-analyte antibody. A sub-hooking concentration should be used for the next optimization steps. A lack of specific signal likely indicates that the selected antibodies are not able to capture the analyte simultaneously in this specific configuration.

For the selection of the best combination of antibody pair, the usual criterion is the highest S/B signal. However, in order to confirm the choice of the antibody pair, a calibration curve of the analyte should be performed. This will allow the measurement of the analyte detection limit and the dynamic range of the specific pair of antibody in the AlphaLISA assay.



**Figure 3**: Titration curve for an insulin detection assay, prepared in assay buffer: in this particular example, the optimal antibody concentration was determined to be 1 nM.

### D3. Calibration curve of analyte

The most important parameters to determine the best combination of antibodies are the detection limit, the assay window (S/B) and the assay dynamic range.

The following conditions may be used for the analyte calibration curve.

In an OptiPlate-384 microplate add:

- 5 µL of the analyte diluted in assay buffer (use dilutions that cover the working range for the target detection plus few points above and below the range of target detection as well as negative controls (background) consisting of 3 independent conditions without analyte in triplicate for a total of 9 points)
- 20 µL of the antibody mix:

biotinylated antibody (as a starting point, use the concentration determined by titration, Section D2)

### AND

antibody-conjugated AlphaLISA Acceptor beads at 25 μg/mL (10 μg/mL final concentration in each well)

Incubate at 23°C for 1 h then add:

 25 μL of streptavidin-Donor bead solution at 80 μg/mL prepared under subdued light conditions (40 μg/mL final concentration in the well)

Incubate at 23°C in the dark for 30 min and read on an

AlphaScreen reader.

The lower detection limit (LDL) is usually calculated as follows:

- Average the 9 background values and calculate the standard deviation (SD).
- Multiply the standard deviation (SD) by 3
- Add the 3XSD value calculated to the average background signal.
- On the graph, extrapolate the value obtained (AlphaLISA signal counts) to determine the corresponding analyte concentration. This concentration is the lowest the assay can detect and corresponds to the lower detection limit (LDL).

The assay dynamic range corresponds to the concentration window in the standard curve running from the lower detection limit to the maximum concentration up to (but excluding) the hook point.

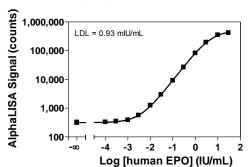
Data can be analyzed using linear or nonlinear regression analysis. However, wider dynamic ranges are usually interpreted using nonlinear regression Eq.1. Four-Parameter Logistic Model also described as Sigmoidal Dose-Response (variable slope) where the four parameters to be estimated are Top, Bottom,  $EC_{50}$  and Slope. For more details refer to the NIH Chemical Genomics Center manual

(http://ncgc.nih.gov/guidance/manual toc.html).

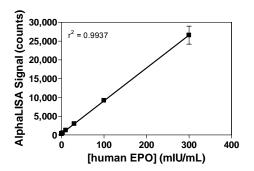
Re 
$$sponse = Top + \frac{(Bottom - Top)}{1 + (\frac{concentration}{EC_{50}})^{Slope}}$$
 Eq. (1)

Top' refers to the top asymptote, 'Bottom' refers to the bottom asymptote, and ' $EC_{50}$ ' refers to the concentration at which the response is halfway between Top and Bottom.





Linear-Linear scale (low analyte concentrations):



**Figure 4**: Calibration curve for a human EPO detection assay prepared in assay buffer. The graph can be analyzed using nonlinear (A) or linear (B) regression analysis. Note that nonlinear representations are more suitable for a wide range of analyses. The lower concentrations of the calibration curve (up to 300 mlU/mL) are shown as a linear plot (B). Reported concentrations are those in the samples.

### E. Assay buffer optimization

The assay buffer (our recommended AlphaLISA Immunoassay buffer is described on page 11) may be optimized for maximum response. The following parameters can be tested:

- Buffer type: Tris, HEPES, at different pH values.
- The presence of Dextran 500: for some serum or plasma samples, it is important to add Dextran 500 at 1 mg/mL in order to prevent nonspecific bead aggregation.
- The presence of detergent, like Tween-20, CHAPS, and Triton X-100, at a concentration between 0.01 to 1%.
- The presence of protein blockers, such as casein or BSA, at a concentration between 0.01 to 1%.

# F. Optimization of the assay volume, incubation time and order of addition

The assay may be further optimized by changing the following parameters.

### F1. Assay volume

Most analyte assays will perform best in 50-75  $\mu$ L final volume in 96-well plates, 25-50  $\mu$ L in 384-well plates and 5-10  $\mu$ L in 1536-well plates. However, the volume can be adjusted to fulfill specific user requirements. If working with serum or plasma samples, it is strongly recommended that the sample volume represents no more than 10% of the total volume in the well in order to reduce interference. Less interference will be observed using the lowest sample volume possible. It is always possible to pre-dilute the sample in the matrix solution if the concentration of the analyte in the sample permits.

### F2. Incubation time

A time course could be performed following each addition step. A time course could also be performed after the addition of Donor beads. Usually the signal will reach a plateau after a few hours.

### F3. Order of addition

For the majority of analyte assays, the order of addition described in Section D3 performs best. However, the addition steps can be changed in order to optimize the assay sensitivity. It could be beneficial to add the biotin-antibody and the Acceptor beads separately. In this case, an incubation time may be necessary between each addition step.

From our experience, pre-mixing the biotinylated antibody with the Donor beads lowers the sensitivity of the assay and should in most cases be avoided.

# G. Determination of the level of analyte in unknown samples by interpolation from the calibration curve

In order to determine the concentration of an analyte in a serum or plasma sample, a calibration curve with known concentrations of the analyte should be performed first. The calibrators (solutions containing a known concentration of analyte) should be prepared in a "matrix

solution". We recommend using a serum or plasma (similar to the samples) first depleted of the analyte of interest as matrix solution then spiked with known concentrations of analyte.

### G1. Analyte depletion of serum

It is possible to prepare analyte-depleted sera by treatment with streptavidin-Sepharose beads and the biotinylated antibody.

- Evaluate the amount of biotinylated-antibody and streptavidin-Sepharose beads required for the depletion. Those conditions will need to be adjusted and we suggest using as starting condition a molar ratio of analyte/antibody of 1/100 and a molar ratio of antibody/streptavidin of 1/20.
- Wash the Sepharose beads with PBS buffer prior to use.
- Mix biotinylated antibody with Sepharose beads for 2 h at 4°C.
- Centrifuge at 1000 × g for 5 min, remove the supernatant and add serum to pelleted beads and incubate overnight at 4°C.
- Remove beads from serum by centrifuging at 16,000 × g (or maximum speed) for 5 min. Transfer supernatant to a new tube and centrifuge again at 16,000 × g (or maximum speed) for 5 min.
- Collect the supernatant and store depleted serum in aliquots at -20°C.

Alternatively, a charcoal treatment may be used to remove the analyte of interest. Several protocols are described in the literature (see references page 20).

An artificial matrix solution could also be prepared. Such a matrix should behave the same way in the AlphaLISA assay as the real samples (same level of signal interference). In such instance, high concentrations of BSA or any other signal quencher could be used for this purpose. Finally, the concentrations of analyte used to perform the calibration curve must cover the range expected to be found in the samples.

### G2. Calibration curve

A general protocol to generate a calibration curve in the matrix solution and determine the analyte concentration of unknown samples is presented below. As mentioned earlier, it might be necessary to optimize the assay conditions to meet the specific assay requirements.

For calibration curve:

Add 1-5 µL of calibrators diluted in matrix solution

For unknown samples:

Add 1-5 μL of unknown samples

Then to calibrator or unknown sample,

 Add 20-24 µL of antibody-conjugated AlphaLISA Acceptor beads + biotinylated antibody solution

Incubate for 30 min at 23°C.

Add 25 μL of streptavidin-Donor bead solution

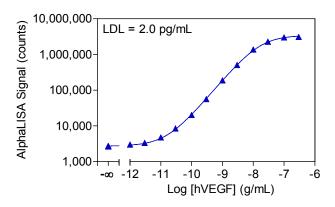
Incubate for 60 min at 23°C, in the dark and read plate on an AlphaScreen reader.

It is advised to prepare three to five replicates of each calibrator concentration.

The calibrator serial dilutions are prepared in the matrix solution, while the solutions of antibody-conjugated AlphaLISA Acceptor beads and of biotinylated antibody are prepared in the assay buffer (see page 11).

It is also possible to predilute the calibrators in the matrix solution, aliquot the solutions and store the vials at -80°C for future use.

On the calibration curve (Figure 5), the increasing concentrations of analyte are on the x-axis and the AlphaLISA signal on the y-axis.



**Figure 5**: Calibration curve for a human VEGF detection assay, prepared in matrix solution. The biotinylated antibody was used at a final concentration of 1 nM in the wells. The antibody-conjugated AlphaLISA Acceptor beads and the streptavidin-Donor beads were used at a final concentration of 10 μg/mL and 40 μg/mL, respectively.

The level of analyte in unknown samples is determined by interpolation from the calibration curve generated in the matrix solution (see previous section). Indeed, the AlphaLISA signals obtained from the unknown samples are used to interpolate the concentration of analyte from the curve.

### V. REFERENCES

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### Appendix I: Summary of AlphaLISA Assay Development Steps

### A. Identify optimal antibody pair

# Biotinylate all antibodies (see: IV. B) Conjugate all antibodies to AlphaLISA Acceptor beads (see: IV. C) Test all possible antibody combinations: Titrate the biotinylated antibody in AlphaLISA (see: IV. D)

# B. Optimize assay conditions & Determine assay characteristics

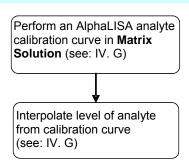
Determine optimal:

Assay volume (see: IV. F1) Incubation time (see: IV. F2) Order of addition (see: IV. F3) Assay buffer (see: IV. E)

by generating analyte calibration curves in **Assay Buffer** 

Determine assay characteristics from calibration curves performed in **Assay Buffer** and in **Matrix Solution**: detection limit, dynamic range, reproducibility (see: IV. D3)

# C. Determine analyte level in unknown samples



### **Appendix II: Serum Sample Preparation**

- a. To prepare serum samples, whole blood is directly drawn into a Vacutainer® serum tube that contains no anticoagulant. Let blood clot at room temperature for 30 minutes.
- b. Promptly centrifuge the clotted blood at  $2,000 3,000 \times g$  for 15 minutes at 4 + /- 2°C.
- c. Transfer and store serum samples in separate tubes. Date and identify each sample.
- d. Use freshly prepared serum or store samples in aliquots at ≤ -20°C for later use. Avoid freeze/thaw cycles.

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- AlphaLISA anti-FLAG Acceptor beads (Cat No. AL112)
- AlphaLISA anti-DIG Acceptor beads (Cat No. AL113)
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- AlphaScreen Glutathione Donor beads (Cat No. 6765300)
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<u>Note</u>: AlphaLISA is for research purposes only and not for use in diagnostic procedures.

<sup>\*</sup> Visit our Web site for the different formats available:



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