

Human Aggregated Aß

For quantitative determination of aggregated $A\beta$ in human tissue homogenates, tissue culture supernatant and buffered solutions

Catalog no. LHB3491

Rev. 1.0 16 October 2009 *PRLHB3491*

User Manual

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Kit Contents and Storage

Storage	All components of the Human Aggregated A β kit are shipped at 2 to 8°C. Upon receipt, store all kit components at 2 to 8°C. Do not freeze .				
Contents	The components and amounts included in the Human Aggregated $A\beta$ kit are listed below.				
	Reagents Provided	100 Test Kit			
Aggregated Aβ Bead (azide)	Concentrate (10X) (contains 7.5 mM sodium	0.25 mL × 1 vial			
Aggregated Aβ Detect 3.3 mM thymol)	$1 \text{ mL} \times 1 \text{ vial}$				
Aggregated Aβ Peptid	2 vials				
Wash Solution Concer	15 mL \times 1 bottle				
Assay Diluent (contain	15 mL \times 1 bottle				
Biotin Diluent (contain	12 mL \times 1 bottle				
Streptavidin-RPE Con azide)	$1 \text{ mL} \times 1 \text{ vial}$				
Streptavidin-RPE Dilu	12 mL \times 1 bottle				
96-well Filter Plate	1×96 -well plate				

Introduction

Overview

Intended Use

Invitrogen's Multiplex Bead Immunoassay Kits are developed to maximize flexibility in experimental design, permitting the measurement of one or multiple proteins in panels designed by the researcher. The Human Aggregated A β kit contains all the reagents that are intended for use with 100^{TM} or 200^{TM} dual laser detection system manufactured by Luminex Corporation and sold by Invitrogen and other vendors. For research use only. Not intended for any animal or human therapeutic or diagnostic use.

Background Information

Several recent reports indicate that an important neurotoxic form of $A\beta$ is an oligomer, composed of approximately 12 subunits, with a composite molecular weight of 54 kDa. This oligomeric form of $A\beta$ can be separated from fibrillar and protofibrillar forms of aggregated $A\beta$ by high speed centrifugation (i.e., 100,000 x g for 1 hour) or by size exclusion methods.

Invitrogen's Multiplex Bead Immunoassays are solid phase protein immunoassays which use spectrally encoded antibody-conjugated beads as the solid support. The spectral beads are suitable for use in singleplex assays or may be mixed for multiplex assays according to the researcher's requirements. Each assay is carefully designed and tested to assure that sensitivity, range, and correlation are maximized. The assay is performed in a 96-well plate format and analyzed with a Luminex[®] 100^{TM} or 200^{TM} instrument which monitors the spectral properties of the capture beads while simultaneously measuring the quantity of associated fluorophore. Standard curves generated with this assay system extend over several orders of magnitudes of concentrations, while the sensitivity and quantitation of the assays are comparable to ELISAs (Enzyme Linked-Immuno-Sorbent Assays). Assay standards are calibrated to NIBSC (National Institute for Biological Standards and Controls) reference preparations, when available, to assure accurate and reliable results.

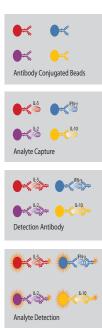
Overview, Continued

Background Information, Continued

Invitrogen's Human Aggregated A β kit is designed for the *in vitro* quantitative determination of Aggregated A β in tissue homogenates that have been clarified by high speed centrifugation, tissue culture supernatant and buffered solutions. This kit may also be combined with other Invitrogen Human Neuroscience Antibody Bead Kits, allowing higher multiplexing of the assay.

Visit the Invitrogen web site for a current listing of available Invitrogen multiplex bead immunoassays and reagents, at www.invitrogen.com/luminex.

Assay Overview



The xMAP[®] technology combines the efficiencies of multiplexing up to 100 different proteins for simultaneous analysis, with reproducibility similar to ELISA. The technology uses 5.6 μ m polystyrene beads which are internally dyed with red and infrared fluorophores of differing intensities. Each bead is given a unique number, or bead region, allowing differentiation of one bead from another.

Beads of defined spectral properties are conjugated to protein-specific capture antibodies and added along with samples (including standards of known protein concentration, control samples, and test samples), into the wells of a filter-bottom microplate and where proteins bind to the capture antibodies over the course of a 2 hour incubation.

After washing the beads, protein-specific biotinylated detector antibodies are added and incubated with the beads for 1 hour. During this incubation, the protein-specific biotinylated detector antibodies bind to the appropriate immobilized proteins.

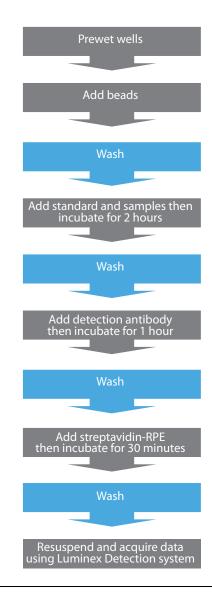
After removal of excess biotinylated detector antibodies, streptavidin conjugated to the fluorescent protein, R-Phycoerythrin (Streptavidin-RPE), is added and allowed to incubate for 30 minutes. The Streptavidin-RPE binds to the biotinylated detector antibodies associated with the immune complexes on the beads, forming a four-member solid phase sandwich.

After washing to remove unbound Streptavidin-RPE, the beads are analyzed with the Luminex detection system. By monitoring the spectral properties of the beads and the amount of associated R-Phycoerythrin (RPE) fluorescence, the concentration of one or more proteins can be determined.

Experimental Overview

Experimental Outline

The experimental outline for using the Human Aggregated A β kit is shown below.



Methods

Before Starting

Materials Required but Not Provided

- Luminex[®] xMAP[®] system with data acquisition and analysis software (Invitrogen, Cat. no. MAP0200), contact Invitrogen for instrument and software placement services, see page 26
- Filtration vacuum manifold for bead washing (Pall, Cat. no. 5017 is recommended)
- Sonicating water bath
- Vortex mixer
- Orbital shaker (small diameter rotation recommended)
- Calibrated, adjustable, precision pipettes, preferably with disposable plastic tips (A manifold multi-channel pipette is desirable)
- Distilled or deionized water
- Glass or polypropylene tubes
- Aluminum foil or opaque 96-well plate cover (Invitrogen, Cat. no. PC10)

Before Starting, Continued



Review the procedural notes below before starting the protocol.

- Before mixing plexes, **check that each protein is represented by a unique bead region**. Up to 10 bead concentrates (singleplexes or pre-mixed multiplexes) can be combined to increase the number of proteins measured.
- **Do not invert the filter plates during the assay**. The filter plates are designed to be used in conjunction with a vacuum manifold (**do not exceed** 5 mm Hg) and emptied from the bottom.
- Do not freeze any component of this kit. Store kit components at 2 to 8°C when not in use. Allow all reagents to warm to room temperature before use (air-warm all reagents at room temperature for at least 30 minutes, or alternatively, in a room-temperature water bath for 20 minutes (except plate and standard vials).
- The fluorescent beads are light-sensitive. Protect the beads from light to avoid photobleaching of the embedded dye. Use aluminum foil to cover test tubes used in the assay. Cover filter plates containing beads with an opaque or aluminum foil-wrapped plate cover. Since the amber vial does not provide full protection, keep the vial covered in the box or drawer when not in use.
- Do not expose beads to organic solvents.
- Do not place filter plates on absorbent paper towels during loading or incubations, as liquid may be lost due to contact wicking. An extra plate cover is a recommended surface to rest the filter plate. Following plate washing, remove excess liquid and blot from the bottom of the plate by pressing the plate on clean paper towels.
- When pipetting reagents, maintain a consistent order of addition from well-to-well to ensure equal incubation times for all wells.
- To prevent filter tearing, avoid touching the filter plate membrane with pipette tips.
- Do not use reagents after kit expiration date.
- It is recommended that in-house controls be included with every assay. If control values fall outside pre-established ranges, the assay may be suspect. Contact Invitrogen Technical Support for product and technical assistance.
- Do not mix or substitute reagents with those from other lots or sources.

Before Starting, Continued



- Handle all blood components and biological materials as potentially hazardous. Follow standard precautions as established by the Centers for Disease Control and Prevention and by the local Occupational Safety and Health Administration when handling and disposing of infectious agents.
- This kit contains materials with small quantities of sodium azide. Sodium azide reacts with lead and copper plumbing to form explosive metal azides. Upon disposal, flush drains with a large volume of water to prevent azide accumulation. Avoid ingestion and contact with eyes, skin and mucous membranes. In case of contact, rinse affected area with plenty of water. Observe all federal, state and local regulations for disposal.

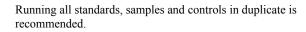
Recommended Plate Plan

It is recommended that a plate plan be designed before starting the assay. A plate plan template is provided on page 30. The following is a suggested plate plan:

	1	2	3	4	5	6	7	8	9	10	11	12
А	В	В										
В	Std 7	Std 7										
С	Std 6	Std 6										
D	Std 5	Std 5										
Е	Std 4	Std 4										
F	Std 3	Std 3										
G	Std 2	Std 2										
Н	Std 1	Std 1										

B= blank (Assay Diluent), Standards 7 through 1, lowest concentration to highest.

The remainder of the plate is available for controls and samples which may be run as a singlet or in duplicate, as desired.





Note

Preparing Reagents

Introduction	Review the information in this section before starting. The Human Aggregated $A\beta$ kit includes both antibody bead reagents and buffer reagents. Prepare components of the Human Aggregated $A\beta$ kit according to instructions below. Note: bring all reagents and samples to room temperature before use.		
Preparing Wash Solution	 Upon storage at 2 to 8°C, a precipitate may form in the 20X Wash Solution Concentrate. If this occurs, warm the 20X Wash Solution Concentrate to 37°C and mix until the precipitate is dissolved. Prepare a 1X Working Wash Solution for use with a 96-well plate by transferring the entire contents of the Wash Solution Concentrate bottle to a 500 mL container (or equivalent) and then add 285 mL of deionized water. Mix well. The 1X Working Wash Solution is stable for up to 2 weeks when stored at 2 to 8°C. Note: To prepare smaller volumes of 1X Working Wash Solution, mix 1 part of 20X concentrate with 19 parts of deionized water. Mix well. 		
Sample Preparation Guidelines	 Aggregated Aβ has been detected in tissue homogenates, tissue culture supernatant, and buffered solution samples using the reagents provided in this kit. Additional sample types may be suitable but have not been thoroughly validated. Collect samples in pyrogen/endotoxin-free tubes. Clarify all samples by centrifugation (1,000 × g for 10 minutes) and/or filter prior to analysis to prevent clogging of the filter plates. 		
	 In the event that the sample concentrations exceed the standard curve, dilute samples and reanalyze. Due to shared epitopes, the following analytes may cause interference with this assay: APP, monomeric Aβ40, monomeric Aβ42, protofibrillar Aβ, and fibrillar Aβ. Sample preparation should therefore be carefully considered when using this assay. Centrifugation at 14,000 x g for 10 minutes has been shown to minimize fibrils in aggregated Aβ-containing samples, while centrifugation at 100,000 x g for one hour at 4°C has been shown to minimize fibrils and protofibrils. Size exclusion methods, such as gel permeation chromatography or ultrafiltration, may also improve assay performance. Aggregated Aβ is sensitive to freeze-thaw cycles. Avoid multiple freeze-thaw cycles of frozen samples. Thaw completely and mix well prior to analysis. 		

Extraction of Aggregated Aβ from tissue

The procedure below provides guidelines for extraction of tissue samples. This procedure may require optimization for each specific application.

Recommended Tissue Extraction Buffer (Tris Buffered Saline)

25 mM Tris-HCl, pH 7.4

150 mM NaCl

This tissue extraction buffer may be stored at 2 to 8°C for up to 1 month. For maximal stability, apportion the tissue extraction buffer into aliquots and store at -20°C.

This buffer should be freshly supplemented with protease inhibitors just prior to use. Suggested protease inhibitors: 1.46 mM pepstatin A, 0.154 mM aprotinin, 2.03 mM leupeptin, 0.5 mM 4-[2-aminoethyl] benzenesulfonyl fluoride hydrochloride [AEBSF], and 0.29 mM PMSF. If desired, this buffer may also be supplemented with phosphatase inhibitors. Suggested phosphatase inhibitors: 0.05 mM fenvalerate, 0.05 mM cantharidin, 1 mM sodium vanadate, 1 mM sodium pyrophosphate, and 50 mM sodium fluoride.

Extraction Procedure

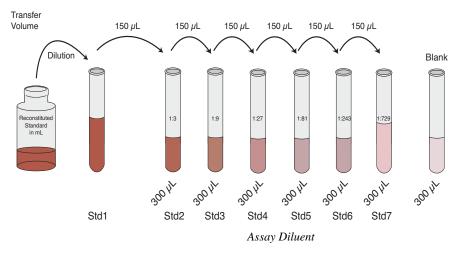
- 1. Determine the mass of the tissue.
- Add 10X volume tissue extraction buffer. Homogenize the tissue thoroughly with a hand-held motor (i.e., Fisher K749540-0000).
- 3. Sonicate the homogenized tissue (2 blasts, 10 seconds each).
- Centrifuge the samples at 100,000 x g for 1 hour at 4°C. Carefully withdraw the clear supernatant fraction, and transfer to a clean polypropylene tube on ice. This supernatant fraction contains TBS-soluble aggregates of Aβ.
- 5. Samples are ready for analysis using the Aggregated A β kit without further dilution.

Guidelines for Standard Curve	•	Each kit comes with 2 complete sets of standard vials, so that 2 runs on the plate can be made with freshly prepared standards.
Preparation	•	Reconstitute the protein standard within 1 hour of performing the assay. All standards are calibrated to NIBSC preparations, when available. Additional standards are available from Invitrogen custom services
	•	Before performing standard mixing and serial dilutions confirm reconstitution volumes on the INFORMATION SHEET, included in the Antibody Bead Kit(s).
	•	The concentrations of the protein components of the standard are indicated on the INFORMATION SHEET.
	•	Perform standard dilutions in glass or polypropylene tubes.
Q Important	pro	tein standards may be analyzed alone, or combined with other tein standards for higher levels of multiplexing. Do not nbine more than 4 vials.
Reconstituting Lyophilized	1.	To the standard vial(s), add the suggested reconstitution
Standards		volume of the appropriate diluent (see table on next page). Do not vortex. When mixing or reconstituting protein solutions, always avoid foaming.
	2.	not vortex. When mixing or reconstituting protein

One vial of standard	Reconstitute the standard vial in the suggested reconstitution volume, usually 1 mL, of appropriate diluent.
Two vials of standards	Reconstitute each vial with 0.5 mL of appropriate diluent. Combine 300 μ L from each vial and mix by gently pipetting up and down 5 to 10 times.
Three vials of standards	Reconstitute each vial with 0.333 mL of appropriate diluent. Combine 200 μ L from each vial and mix by gently pipetting up and down 5 to 10 times.
Four vials of standards	Reconstitute each vial with 0.250 mL of appropriate diluent. Combine 150 μ L from each vial and mix by gently pipetting up and down 5 to 10 times.

Preparing Standard Curve

The standard curve is made by serially diluting the reconstituted standard in Assay Diluent. See below. **Do not vortex.** Mix by gently pipetting up and down 5 to 10 times.



Discard all remaining reconstituted and diluted standards after completing assay. Return the Assay Diluent to the kit.



Preparing 1X Antibody Beads Go to http://www.invitrogen.com/luminex under Multiplex Solution Tools, click Luminex[®] Calculation Worksheet for auto calculation of all assay dilutions.

Determine the number of wells required for the assay.

The Antibody Bead Concentrate is supplied as a **10X concentrate** and must be diluted prior to use. The fluorescent beads are light-sensitive. Protect antibody conjugated beads from light during handling.

- 1. Immediately before dispensing, vortex the 10X Antibody Bead Concentrate for 30 seconds followed by sonication in a sonicating water bath for 30 seconds.
- Prepare 1X Antibody Bead stock by diluting 2.5 μL of 10X beads in 25 μL of Working Wash Solution (page 11) per assay well. Each well requires 25 μL of the diluted beads. See table below for examples of volumes to combine.

Number of Wells	Vol. 10X Antibody Bead Concentrate	Vol. Working Wash Solution
24	0.06 mL	0.6 mL
32	0.08 mL	0.8 mL
40	0.10 mL	1.0 mL
48	0.12 mL	1.2 mL
56	0.14 mL	1.4 mL
64	0.16 mL	1.6 mL
72	0.18 mL	1.8 mL
80	0.20 mL	2.0 mL
88	0.22 mL	2.2 mL
96	0.24 mL	2.4 mL

Note: Dilution factor is 1:11 for extra pipetting volume.

Assay Procedure

Method of Washing

Incomplete washing adversely affects assay results. Perform all wash steps with the Wash Solution supplied with the kit. All phases of the assay, including incubations, washing steps, and loading beads, are performed in the filter bottom plate supplied with the kit. For a demonstration of proper washing techniques, download Luminex video from www.invitrogen.com/luminexinstrument.

- To wash beads, place the filter plate on the vacuum manifold and aspirate the liquid with gentle vacuum (do not exceed 5 mm Hg). Excessive vacuum can cause the membrane to tear, resulting in antibody bead loss. Prevent any vacuum surge by opening and adjusting the vacuum on the manifold before placing the plate on the manifold surface.
- 2. Stop the vacuum pressure as soon as the wells are empty. Do not attempt to pull the plate off the vacuum manifold while the vacuum is still on or filter plate damage may occur. Release the vacuum prior to removing the plate.
- 3. If solution remains in the wells during vacuum aspiration, **do not detach the bottom of the 96 well filter plate.** In some cases, minor clogs in the filter plate may be dislodged by carefully pressing the bottom of the plate under the clogged well with the pointed end of a 15 mL plastic conical tube. Place the filter plate on a clean paper towel and use a gloved thumb or a 1 mL Pasteur pipette bulb to plunge the top of the clogged well. Empty all clogged wells entirely before continuing the washes. **Note**: Do not attempt to repetitively pull vacuum on plates with clogged wells. This can compromise the unclogged wells and bead loss may occur.
- 4. After all wells are empty, lightly tap or press the filter plate onto clean paper towels (hold the plate in the center for tapping) to remove excess fluid from the bottom of the filter plate. **Do not invert plate.**
- 5. Following the last aspiration and plate taps, use a clean absorbent towel to blot the bottom of the filter plate before addition of next liquid phase or data acquisition step.
- 6. Do not leave plate on absorbent surface when adding reagents.



Reverse pipetting recommendation:

To reduce bubbles and loss of reagents due to residual fluid left in pipette tips, use the recommended reverse pipetting technique.

- 1. To reverse pipette, set the pipette to the appropriate volume needed. Note: Do not reverse pipette volumes $<20 \ \mu$ L.
- 2. Press the push-button slowly to the first stop and then press on past it. Note: the amount past the first stop will depend on the volume of liquid available to aspirate from.
- 3. Immerse the tip into the liquid, just below the meniscus.
- 4. Release the push-button slowly and smoothly to the top resting position to aspirate the set volume of liquid.
- 5. Place the end of the tip against the inside wall of the recipient vessel at an angle.
- Press the push button slowly and smoothly to the first stop. Some liquid will remain in the tip, this should not be dispensed.
- 7. Remove the tip, keeping the pipette pressed to the first stop.

Bring all reagents and samples to room temperature before use

Analyte Capture

Note

- 1. Use an adhesive plate cover to seal any unused wells. This will keep the wells dry for future use.
- Pre-wet the designated assay wells by adding 200 μL of Working Wash Solution into designated wells. Incubate plate 15 to 30 seconds at room temperature.
- 3. Aspirate the Working Wash Solution from the wells using the vacuum manifold.
- 4. Vortex the diluted bead solution (prepared on page 15) for 30 seconds, then sonicate for at least 30 seconds immediately prior to use in the assay.
- Pipette 25 μL of the diluted bead solution into each well. Once the beads are added to the plate, keep the plate protected from light.
- Add 200 μL Working Wash Solution to the wells. Allow the beads to soak for 15 to 30 seconds.
- 7. Aspirate the Working Wash Solution from the wells with the vacuum manifold. Repeat this washing step.
- 8. Blot the bottom of the filter plate on clean paper towels to remove any residual liquid.

Note: Place the filter plate on a plate cover or non-absorbent surface before all incubations.

Analyte Capture,	 To wells designated for the standard curve, pipette 100 μL of appropriate standard dilution.
Continued	 To the wells designated for the sample, pipette 50 μL Assay Diluent followed by 50 μL sample to each well or 50 μL in-house controls, if used.
	11. Cover filter plate containing beads with an aluminum foil-wrapped plate cover. Incubate the plate for 2 hours at room temperature on an orbital shaker. Shaking should be sufficient to keep beads suspended during the incubation (500-600 rpm). Larger radius shakers will need a lower speed and smaller radius shakers will typically handle higher speeds without splashing.
	12. Ten to fifteen minutes prior to the end of this incubation, prepare the biotinylated detector antibody, and then proceed to Analyte Detection, Step 1.
Preparing 1) Biotinylated	The Biotinylated Antibody is supplied as a 10X concentrate and must be diluted prior to use.
Antibody	To prepare a 1X Biotinylated Antibody stock, dilute 10 μ L of 10X Biotinylated Antibody in 100 μ L of Biotin Diluent per assay well. Each well requires 100 μ L of the diluted Biotinylated Antibody. See table below for examples of volumes to combine.
	Note: Dilution factor is 1:11 for extra pipetting volume.
Number of Wells	Vol. 10X Biotinylated Antibody Concentrate

Number of Wells	Vol. 10X Biotinylated Antibody Concentrate	Vol. Biotin Diluent
24	0.24 mL	2.4 mL
32	0.32 mL	3.2 mL
40	0.40 mL	4.0 mL
48	0.48 mL	4.8 mL
56	0.56 mL	5.6 mL
64	0.64 mL	6.4 mL
72	0.72 mL	7.2 mL
80	0.80 mL	8.0 mL
88	0.88 mL	8.8 mL
96	0.96 mL	9.6 mL

Analyte Detection	1.	After the 2 hour capture bead incubation, remove the liquid from wells by aspiration with the vacuum manifold.
	2.	Add 200 μ L of Working Wash Solution to the wells. Allow the beads to soak for 15 to 30 seconds, then aspirate with the vacuum manifold. Repeat this washing step.
	3.	Blot the bottom of the filter plate on clean paper towels to remove residual liquid.
	4.	Add 100 μ L of prepared 1X Biotinylated Detector Antibody (page 18) to each well and incubate the plate for 1 hour at room temperature on an orbital shaker. Shaking should be sufficient to keep the beads suspended during incubation (500-600 rpm).
	5.	Prepare the Luminex [®] 100 [™] or 200 [™] instrument during this incubation step. Refer to the Luminex Instrument Quick Reference card provided in kit. Refer to the INFORMATION SHEET for all bead regions and standard concentration values.
	6.	Ten to fifteen minutes prior to the end of the detector incubation step, prepare the Streptavidin-RPE, and then proceed with Assay Reading, Step 1.

Preparing	The Streptavidin-RPE is supplied as a 10X concentrate and must
Streptavidin-RPE	be diluted prior to use. Protect Streptavidin-RPE from light during handling.

To prepare a 1X Streptavidin-RPE stock, dilute 10 μ L of 10X Streptavidin-RPE in 100 of μ L Streptavidin-RPE Diluent per assay well. Each well requires 100 μ L of the diluted Streptavidin-RPE. See table below for examples of volumes to combine.

Number of Wells	Vol. 10X Streptavidin-RPE Concentrate	Vol. Streptavidin-RPE Diluent
24	0.24 mL	2.4 mL
32	0.32 mL	3.2 mL
40	0.40 mL	4.0 mL
48	0.48 mL	4.8 mL
56	0.56 mL	5.6 mL
64	0.64 mL	6.4 mL
72	0.72 mL	7.2 mL
80	0.80 mL	8.0 mL
88	0.88 mL	8.8 mL
96	0.96 mL	9.6 mL

Note: Dilution factor is 1:11 for extra pipetting volume.

Assay Reading		Remove the liquid from wells by aspiration with the vacuum manifold.
-	2.	Add 200 µL Working Wash Solution to the wells. Allow the beads to soak for 15 to 30 seconds, then aspirate with the vacuum manifold. Repeat this washing step.
	3.	Blot the bottom of the filter plate on clean paper towels to remove residual liquid.
	4.	Add 100 μ L of prepared 1X Streptavidin-RPE to each well and incubate the plate for 30 minutes at room temperature on an orbital shaker. Shaking should be sufficient to keep the beads suspended during incubation (500-600 rpm).
	5.	Remove the liquid from wells by aspiration with the vacuum manifold.
	6.	Wash beads by adding 200 µL Working Wash Solution to the

- 6. Wash beads by adding 200 μL Working Wash Solution to the wells; allow the beads to soak for 10 seconds, then aspirate with the vacuum manifold. Repeat this washing step 2 additional times for a total of 3 washes.
- 7. Blot the bottom of the filter plate on clean paper towels to remove residual liquid.
- 8. Add 100 μ L of Working Wash Solution to each well. Shake the plate on an orbital shaker (500-600 rpm) for 2 to 3 minutes to resuspend the beads.

Note: If the plate cannot be read on the day of the assay, cover and store the plate in the dark overnight at 2 to 8°C for reading the following day without significant loss of fluorescent intensity. Aspirate Working Wash Solution from stored plates and add 100 μ L fresh Working Wash Solution. Place the plate on an orbital shaker for 2 to 3 minutes at 500-600 rpm prior to analysis.

- 9. Uncover the plate and insert the plate into the XY platform of the Luminex[®] 100[™] or 200[™] instrument, and analyze the samples.
- 10. Determine the concentration of samples from the standard curve using curve fitting software. It is recommended to use the five parameter algorithm with a weighted function $(1/y^2)$, depending on the software package used.

Performance Characteristics and Limitations of the Procedure

Performance Characteristics	Refer to analyte specific INFORMATION SHEET for performance claims.	
Procedure Limitations	• Do not extrapolate the standard curve beyond the highest or lowest standard point; the dose-response and data collected in these regions may be non-linear and should be considered inaccurate. Note: In some cases, further dilution of the standard beyond 7 points may be possible to extend the low end of the standard curve.	
	• Dilute samples that are greater than the highest standard with Assay Diluent or appropriate matrix diluent; reanalyze these samples and multiply results by the appropriate dilution factor.	
	• Samples are diluted in the assay 1:2 (50 µL of sample and 50 µL of diluent) relative to the standards. Be sure to account for this dilution factor during sample calculations.	
	• The influence of various drugs, aberrant sera (hemolyzed, hyperlipidemic, jaundiced, etc.) and the use of biological fluids in place of serum, plasma and tissue culture supernatant samples have not been thoroughly investigated. The rate of degradation of analytes in various matrices may not have been investigated. The immunoassay literature contains frequent references to aberrant signals seen with some sera, attributed to heterophilic antibodies. Though such samples have not been seen to date, the possibility of this occurrence cannot be excluded.	

Troubleshooting

IntroductionRefer to the table below to troubleshoot your experiments with the
use of Invitrogen's Multiplex Bead Kits on the Luminex®
platform.
To troubleshoot problems with the Luminex® instrument, refer to
the manual supplied with the instrument.
For more troubleshooting solutions, visit

www.invitrogen.com/luminex.

Problem	Cause	Solution
During data analysis, insufficient and/or erratic bead count is observed	Bead aggregation	Make sure to vortex the beads for 30 seconds and then sonicate the beads for at least 30 seconds prior to beginning the assay, to break up any bead aggregates.
		Empty wells and add fresh wash buffer. Shake for 2 to 3 minutes to resuspend the beads.
	Loss of beads due to the filter plate membrane tearing	To prevent membrane tearing, place pipette tips on the side of the well, rather than straight down onto the membrane when dispensing liquid into the wells.
		Turn the vacuum manifold on before placing the filter plate on the top to prevent vacuum surge. When evaluating a new vacuum manifold, adjust the vacuum force so that 3 seconds are required to empty 0.2 mL from the wells of a plate.
	Clog in instrument or probe	Remove probe, sonicate for 5 minutes, rinse the probe and reinstall. Run an unclog protocol. See instrument manual.
	Probe height set incorrectly	Readjust the instrument probe height. If it is too low, it could puncture the well membrane. If it is too high, air could be pulled up with the liquid which may appear as bead fragments to the instrument.

Troubleshooting, Continued

Problem	Cause	Solution
During washing steps, the vacuum manifold does not aspirate the liquid from wells of the filter plate	The filter plate is clogged Lack of a tight seal	Dislodge the clog by gently pushing the pointed end of a 15 mL plastic conical tube into the bottom of the plate under the clogged well. This procedure clears the small opening in the plastic casing.
		Dislodge by placing the filter plate on a clean paper towel and use a gloved thumb or a 1 mL Pasteur pipette bulb to plunge the top of the clogged well.
		To prevent filter plate clogging, clarify samples by centrifugation at $1,000 \times g$ for 10 minutes prior to analysis. Some samples may also require filtration prior to analysis.
		Hold the plate firmly against the vacuum manifold to form a tight seal. If only a partial plate is being analyzed, cover the empty wells with a self-adhesive plate seal.

Problem	Cause	Solution
In-house controls perform differently in subsequent assays	Incorrect concentration entered in data analysis software	The standard proteins included in Invitrogen's Antibody Bead Kits are calibrated to NIBSC preparations, whenever possible. This calibration assures lot-to-lot consistency in performance. However, the concentration of the reconstituted standards may vary with each new lot of standard. Therefore, it is important to check the concentration of the standard listed on the INFORMATION SHEET, and to verify all concentration values entered into the data analysis software.
	Improper reconstitution or dilution of the standard	Check standard reconstitution and dilution as described on page 14.

Troubleshooting, Continued

Problem	Cause	Solution
Leaky plate	Solution remains on the bottom of the wells after vacuum aspiration, causing wicking and leakage of well contents during next incubation	After final wash step and plate taps, use a clean absorbent towel to blot the bottom of the filter plate before addition of next liquid phase or data acquisition step.
	Filter plate membrane tearing	Excessive vacuum can cause the membrane to tear, resulting in antibody bead loss. Prevent any vacuum surge by opening and adjusting the vacuum on the manifold before placing the plate on the manifold surface.

Appendix

Technical Support



Visit the Invitrogen website at www.invitrogen.com for:

- Technical resources, including manuals, INFORMATION SHEET, quick calculation worksheet, application notes, MSDSs, FAQs, formulations, citations, handbooks, and more.
- Complete Technical Support contact information
- Access to the Invitrogen Online Catalog
- Additional product information and special offers

Contact Us

For more information or technical assistance, call, write, fax, or email. Additional international offices are listed on our website (www.invitrogen.com).

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Protocol Summary

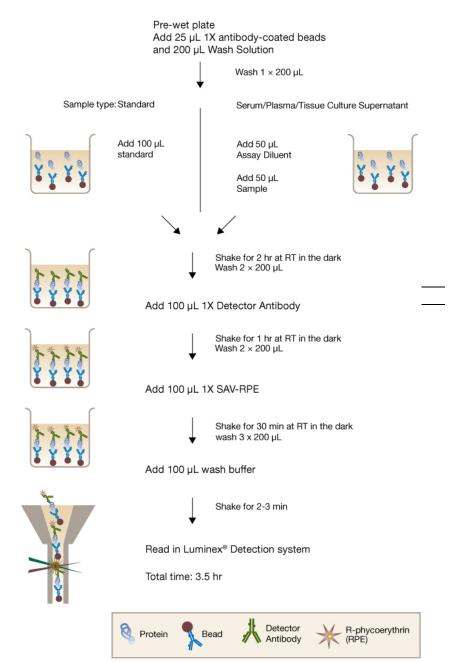
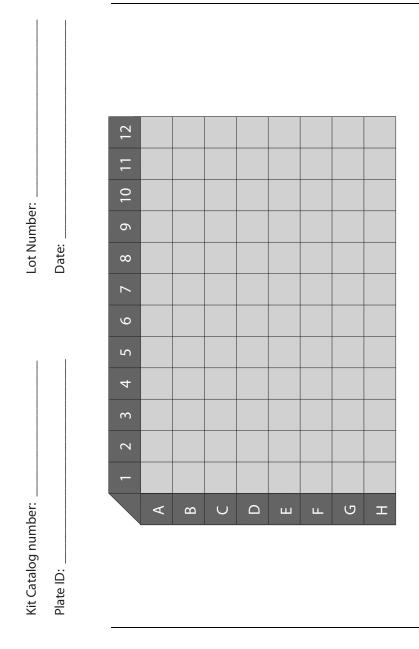


Plate Plan Template



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