BD Cytometric Bead Array (CBA) Mouse Inflammation Kit



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History

Revision	Date	Change made
23-12720-00 Rev. 01	1/2011	New document
23-12720-01 Rev. 01	2/2012	Updated image
Rev. 2	4/2015	Update hazard statements

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About this kit

This section covers the following topics:

- Purpose of this kit (page 6)
- Limitations (page 8)
- Kit contents (page 9)
- Storage and handling (page 11)

Purpose of this kit

Use of the kit

The BDTM CBA Mouse Inflammation Kit can be used to quantitatively measure Interleukin-6 (IL-6), Interleukin-10 (IL-10), Monocyte Chemoattractant Protein-1 (MCP-1), Interferon-γ (IFN-γ), Tumor Necrosis Factor (TNF), and Interleukin-12p70 (IL-12p70) protein levels in a single sample. The kit performance has been optimized for analysis of specific proteins in tissue culture supernatants, EDTA plasma, and serum samples. The kit provides sufficient reagents for 80 tests.

Principle of CBA assays

BD CBA assays provide a method of capturing a soluble analyte or set of analytes with beads of known size and fluorescence, making it possible to detect analytes using flow cytometry.

Each capture bead in the kit has been conjugated with a specific antibody. The detection reagent provided in the kit is a mixture of phycoerythrin (PE)-conjugated antibodies, which provides a fluorescent signal in proportion to the amount of bound analyte.

When the capture beads and detector reagent are incubated with an unknown sample containing recognized analytes, sandwich complexes (capture bead + analyte + detection reagent) are formed. These complexes can be measured using flow cytometry to identify particles with fluorescence characteristics of both the bead and the detector.

Principle of the test

Six bead populations with distinct fluorescence intensities have been coated with capture antibodies specific for IL-6, IL-10, MCP-1, IFN-γ, TNF, and IL-12p70 proteins. The six bead populations are mixed together to form the bead array, which is resolved in a red channel of a flow cytometer (see Figure 1).

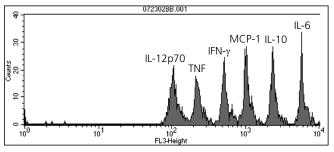


Figure 1

During the assay procedure, you will mix the chemokine capture beads with the recombinant standards or unknown samples and incubate them with the PEconjugated detection antibodies to form sandwich complexes. The intensity of PE fluorescence of each sandwich complex reveals the concentration of that cytokine. After acquiring samples on a flow cytometer, use FCAP ArrayTM software to generate results in graphical and tabular format.

Advantages over ELISA

The broad dynamic range of fluorescence detection via flow cytometry and the efficient capturing of analytes via suspended particles enable BD CBA assays to measure the concentration of an unknown in substantially less time and using fewer sample dilutions compared to conventional ELISA methodology.

- The required sample volume is approximately onesixth the quantity necessary for conventional ELISA assays due to the detection of six analytes in a single sample.
- A single set of diluted standards is used to generate a standard curve for each analyte.
- A BD CBA experiment takes less time than a single ELISA and provides results that would normally require six conventional ELISAs.

Limitations

Assay limitations

The theoretical limit of detection of the BD CBA Mouse inflammation Kit is comparable to conventional ELISA, but due to the complexity and kinetics of this multianalyte assay, the actual limit of detection in a given experiment may vary slightly. Note the reduced sensitivity of the Mouse MCP-1 assay. See Theoretical limit of detection (page 30) and Precision (page 35).

The BD CBA assay is not recommended for use on stream-in-air instruments for which signal intensities may be reduced, adversely affecting assay sensitivity. Stream-in-air instruments include the BD FACStarTM Plus, BD FACSVantageTM, and BD InfluxTM flow cytometers (BD Biosciences).

Serum spike recoveries for IL-10 and TNF are lower than for the other proteins in this assay. This variation is due to assay conditions and serum proteins. It may affect quantitation of these proteins in serum samples.

Quantitative results or protein levels for the same sample or recombinant protein run in ELISA and BD CBA assays may differ. A spike recovery assay can be performed using an ELISA standard followed by BD CBA analysis to assess possible differences in quantitation.

This kit is designed to be used as an integral unit Do not mix components from different batches or kits.

Kit contents

Contents

This kit contains the following components sufficient for 80 tests.

Vial label	Reagent	Quantity
A1	Mouse IL-6 Capture Beads	1 vial, 0.8 mL
A2	Mouse IL-10 Capture Beads	1 vial, 0.8 mL
A3	Mouse MCP-1 Capture Beads:	1 vial, 0.8 mL
A4	Mouse IFN-γ Capture Beads	1 vial, 0.8 mL
A5	Mouse TNF Capture Beads	1 vial, 0.8 mL
A6	Mouse IL-12p70 Capture Beads	1 vial, 0.8 mL
В	Mouse Inflammation PE Detection Reagent	1 vial, 4 mL
С	Mouse Inflammation Standards	2 vials lyophilized
D	Cytometer Setup Beads	1 vial, 1.5 mL

Vial label	Reagent	Quantity
E1	PE Positive Control Detector	1 vial, 0.5 mL
E2	FITC Positive Control Detector	1 vial, 0.5 mL
F	Wash Buffer	1 bottle, 130 mL
G	Assay Diluent	1 bottle, 30 mL

Bead reagents

Mouse Inflammation Capture Beads (A1–A6): An 80-test vial of each specific capture bead (A1–A6). The specific capture beads, having discrete fluorescence intensity characteristics, are distributed from brightest (A1) to dimmest (A6).

Cytometer Setup Beads (D): A 30-test vial of setup beads for setting the initial instrument PMT voltages and compensation settings is sufficient for 10 instrument setup procedures. The Cytometer Setup Beads are formulated for use at 50 µL per test.

Antibody and standard reagents

Mouse Inflammation PE Detection Reagent (B): An 80-test vial of PE-conjugated anti-mouse IL-6, IL-10, MCP-1, IFN-γ, TNF, and IL-12p70 antibodies, formulated for use at 50 μL per test.

Mouse Inflammation Standards (C): Two vials containing lyophilized recombinant mouse proteins. Each vial should be reconstituted in 2 mL of Assay Diluent to prepare the top standard.

PE Positive Control Detector (E1): A 10-test vial of PE-conjugated antibody control that is formulated for use at 50 μL per test. This reagent is used with the Cytometer Setup Beads to set the initial instrument compensation settings.

FITC Positive Control Detector (E2): A 10-test vial of FITC-conjugated antibody control that is formulated for use at $50~\mu L$ per test. This reagent is used with the Cytometer Setup Beads to set the initial instrument compensation settings.

Buffer reagents

Wash Buffer (F): A 130-mL bottle of phosphate buffered saline (PBS) solution (1X), containing protein and detergent, used for wash steps and to resuspend the washed beads for analysis.

Assay Diluent (G): A 30-mL bottle of a buffered protein solution (1X) used to reconstitute and dilute the Mouse Inflammation Standards and to dilute test samples.

Note: Source of all serum proteins is from USDA inspected abattoirs located in the United States.

Storage and handling

Storage

Store all kit components at 2 to 8°C. Do not freeze.

Warning

Components A1–A6, B, D, E1, E2, F, and G contain sodium azide. Sodium azide yields a highly toxic hydrazoic acid under acidic conditions. Dilute azide compounds in running water before discharging to avoid accumulation of potentially explosive deposits in plumbing.

Mouse Inflammation Standards (component 51-9000148) contains 0.02% (w/w) of a CMIT/MIT mixture (3:1), which is a mixture of: 5-chloro-2-methyl-4-isothiazolin-3-one [EC No 247-500-7] and 2-methyl-4-isothiazolin-3-one [EC No 220-239-6] (3:1).

Hazard statements

May cause an allergic skin reaction.

Precautionary statements

Wear protective gloves / eye protection.

Wear protective clothing.

Avoid breathing mist/vapours/spray.

If skin irritation or rash occurs: Get medical advice/ attention.

IF ON SKIN: Wash with plenty of water.

Dispose of contents/container in accordance with local/regional/national/international regulations.

Before you begin

This section covers the following topics:

- Workflow overview (page 14)
- Required materials (page 15)

Workflow overview

Workflow

The overall workflow consists of the following steps.

Step	Description
1	Preparing Mouse Inflammation Standards (page 18)
2	Mixing Mouse Inflammation Capture Beads (page 20)
3	Diluting samples (page 21)
4	Performing instrument setup with Cytometer Setup Beads (instructions can be found at bdbiosciences.com/cbasetup) Can be performed during the incubation in step 5.
5	Performing the Mouse Inflammation Assay (page 24)
6	Acquiring samples (instructions can be found at bdbiosciences.com/cbasetup)
7	Data analysis (page 27)

Incubation times To help you plan your work, the incubation times are listed in the following table.

Procedure	Incubation time
Preparing standards	15 minutes
Preparing Cytometer Setup Beads	30 minutes
Performing the assay	2 hours

Required materials

Materials required but not provided

In addition to the reagents provided in the BD CBA Mouse inflammation Kit, the following items are also required:

• A dual-laser flow cytometer equipped with a 488-nm or 532-nm and a 633-nm or 635-nm laser capable of distinguishing 576-nm, 660-nm, and >680-nm fluorescence. The following table lists examples of compatible instrument platforms.

Flow cytometer	Reporter channel	Bead channels	
BD FACSArray TM	Yellow	Red	
BD FACSCanto TM platform BD TM LSR platform BD FACSAria TM platform	PE	APC	
BD FACSCalibur TM (single laser) BD FACSCalibur (dual laser)	FL2	FL3 FL4	
Note: Visit bdbiosciences.com/cbasetup for setup protocols.			

- Falcon® 12 × 75-mm sample acquisition tubes for a flow cytometer (Catalog No. 352008)
- 15-mL conical, polypropylene tubes (Falcon, Catalog No. 352097), or equivalent

Note: FCAP Array software (Catalog No. 641488 [PC] or 645447 [Mac®])

Materials required for plate loaderequipped flow cytometers

- Millipore MultiScreen_{HTS}-BV 1.2-µm Clear nonsterile filter plates [Catalog No. MSBVN1210 (10 pack) or MSBVN1250 (50 pack)]
- Millipore MultiScreen_{HTS} Vacuum Manifold (Catalog No. MSVMHTS00)
- MTS 2/4 Digital Stirrer, IKA Works, VWR (Catalog No. 82006-096)
- Vacuum source
- Vacuum gauge and regulator (if not using the recommended manifold)

Assay preparation

This section covers the following topics:

- Preparing Mouse Inflammation Standards (page 18)
- Mixing Mouse Inflammation Capture Beads (page 20)
- Diluting samples (page 21)

Preparing Mouse Inflammation Standards

Purpose of this procedure

The Mouse Inflammation Standards are lyophilized and should be reconstituted and serially diluted immediately before mixing with the Capture Beads and the PE Detection Reagent.

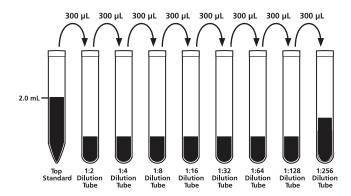
You must prepare fresh cytokine standards to run with each experiment. Do not store or reuse reconstituted or diluted standards.

Procedure

To reconstitute and serially dilute the standards:

- 1. Open one vial of lyophilized Mouse Inflammation Standards. Transfer the standard spheres to a 15-mL conical, polypropylene tube. Label the tube "Top Standard."
- Reconstitute the standards with 2.0 mL of Assay Diluent.
 - a. Allow the reconstituted standard to equilibrate for at least 15 minutes at room temperature.
 - b. Gently mix the reconstituted protein by pipet only. Do not vortex or mix vigorously.
- 3. Label eight 12 x 75-mm tubes and arrange them in the following order: 1:2, 1:4, 1:8, 1:16, 1:32, 1:64, 1:128, and 1:256.
- 4. Pipette 300 μL of Assay Diluent into each of the remaining tubes.
- 5. Perform a serial dilution:
 - a. Transfer 300 μL from the Top Standard to the 1:2 dilution tube and mix thoroughly by pipet only. Do not vortex.

- b. Continue making serial dilutions by transferring $300 \, \mu L$ from the 1:2 tube to the 1:4 tube and so on to the 1:256 tube.
- c. Mix thoroughly by pipet only. Do not vortex.



Prepare one 12 x 75-mm tube containing Assay
Diluent to serve as the 0-pg/mL negative control.

Concentration of standards

See Performing the Mouse Inflammation Assay (page 24) for a list of the concentration (pg/mL) of all six recombinant proteins in each standard dilution.

Next step

Proceed to Mixing Mouse Inflammation Capture Beads (page 20).

Mixing Mouse Inflammation Capture Beads

Purpose of this procedure

The Capture Beads are bottled individually (A1–A6). You must pool all six bead reagents immediately before using them in the assay.

Procedure

To mix the Capture Beads:

- 1. Determine the number of assay tubes (including standards and controls) that are required for the experiment (for example, 8 unknowns + 9 standard dilutions + 1 negative control = 18 assay tubes).
- 2. Vigorously vortex each Capture Bead suspension for 3 to 5 seconds before mixing.

Note: The antibody-conjugated beads will settle out of suspension over time. Vortex the vial immediately before taking a bead-suspension aliquot.

- 3. Add a 10-μL aliquot of each Capture Bead, for each assay tube to be analyzed, into a single tube labeled "Mixed Capture Beads" (for example, 10 μL of IL-6 Capture Beads × 18 assay tubes = 180 μL of IL-6 Capture Beads required).
- 4. Vortex the bead mixture thoroughly.

Next step

The mixed Capture Beads are now ready to be transferred to the assay tubes. Discard excess mixed Capture Beads. Do not store after mixing.

To begin the assay, proceed to Performing the Mouse Inflammation Assay (page 24). If you need to dilute samples having high-protein concentration, proceed to Diluting samples (page 21).

Diluting samples

Purpose of this procedure

The standard curve for each protein covers a defined set of concentrations from 20 to 5,000 pg/mL. It might be necessary to dilute test samples to ensure that their median fluorescence values fall within the range of the generated standard curve. For best results, dilute samples that are known or assumed to contain high levels of a given protein.

Procedure

To dilute samples with a known high-protein concentration:

- 1. Dilute the sample by the desired dilution factor (ie, 1:2, 1:10, or 1:100) using the appropriate volume of Assay Diluent.
- 2. Mix sample dilutions thoroughly.

Next step

Perform instrument setup using the Cytometer Setup Beads. For details, go to bdbiosciences.com/cbasetup and select the appropriate flow cytometer under CBA Kits: Instrument Setup.

Of, if you wish to begin staining your samples for the assay, proceed to Performing the Mouse Inflammation Assay (page 24), and you can perform instrument setup during the 2-hour staining incubation.

Assay procedure

This section covers the following topics:

- Performing the Mouse Inflammation Assay (page 24)
- Data analysis (page 27)

Performing the Mouse Inflammation Assay

Purpose of this procedure

- Prepare the standards as described in Preparing Mouse Inflammation Standards (page 18).
- Mix the Capture Beads as described in Mixing Mouse Inflammation Capture Beads (page 20).
- If necessary, dilute the unknown samples. See Diluting samples (page 21)

Procedure for tubes

To perform the assay:

- 1. Vortex the mixed Capture Beads and add 50 μL to the all assay tubes.
- 2. Add 50 μ L of the Mouse Inflammation Standard dilutions to the control assay tubes as listed in the following table.

Tube label	Concentration (pg/mL)	Standard dilution
1	0 (negative control)	no standard dilution (Assay Diluent only)
2	20	1:256
3	40	1:128
4	80	1:64
5	156	1:32
6	312	1:16
7	625	1:8
8	1,250	1:4
9	2,500	1:2
10	5,000	Top Standard

3. Add 50 μL of each unknown sample to the appropriately labeled sample assay tubes.

- 4. Add 50 μL of the Mouse Inflammation PE Detection Reagent to all assay tubes.
- 5. Incubate the assay tubes for 2 hours at room temperature, protected from light.

Note: If you have not yet performed cytometer setup, you may wish to do so during this incubation.

- 6. Add 1 mL of Wash Buffer to each assay tube and centrifuge at 200*g* for 5 minutes.
- 7. Carefully aspirate and discard the supernatant from each assay tube.
- 8. Add 300 μ L of Wash Buffer to each assay tube to resuspend the bead pellet.

Procedure for filter plates

To perform the assay:

- 1. Wet the plate by adding 100 μL of wash buffer to each well.
- 2. Place the plate on the vacuum manifold.
- 3. Aspirate for 2 to 10 seconds until the wells are drained.
- 4. Remove the plate from the manifold, then blot the bottom of the plate on paper towels.
- 5. Add 50 μL of each of the following to the wells in the filter plate:
 - Capture Beads (vortex before adding)
 - Standard or sample (add standards from the lowest concentration to the highest, followed by samples)
 - Mouse Inflammation PE Detection Reagent
- 6. Cover the plate and shake it for 5 minutes at 1,100 rpm on a plate shaker.

7. Incubate the plate for 2 hours at room temperature on a non-absorbent, dry surface.

Note: Place the plate on a non-absorbent, dry surface during incubation. Absorbent or wet surfaces can cause the contents of the wells to leak.

- 8. Remove the cover from the plate and apply the plate to the vacuum manifold.
- 9. Vacuum aspirate for 2 to 10 seconds until the wells are drained.
- 10. Remove the plate from the manifold, then blot the bottom of the plate on paper towels after aspiration.
- 11. Add 120 µL of wash buffer to each well to resuspend the beads.
- 12. Cover the plate and shake it for 2 minutes at 1,100 rpm before you begin sample acquisition.

Next step

Acquire the samples on the flow cytometer. For details, go to bdbiosciences.com/cbasetup and select the appropriate flow cytometer under CBA Kits: Instrument Setup.

CBA samples must be acquired on the same day they are prepared. Prolonged storage of samples, once the assay is complete, can lead to increased background and reduced sensitivity.

To facilitate the analysis of samples using the FCAP Array software, we recommend the following guidelines:

- Acquire standards from lowest (0 pg/mL) to highest (Top Standard) concentration, followed by the test samples.
- If running sample dilutions, acquire sequentially starting with the most concentrated sample.

 Store all FCS files (standards and samples) in a single folder.

When you are finished acquiring samples, proceed to Data analysis (page 27).

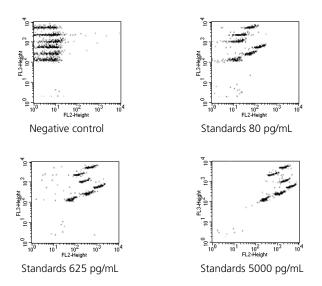
Data analysis

How to analyze

Analyze BD CBA Mouse inflammation Kit data using FCAP Array software. For instructions on analysis, go to bdbiosciences.com/cbasetup and see the *Guide to Analyzing Data from BD CBA Kits Using FCAP Array Software*.

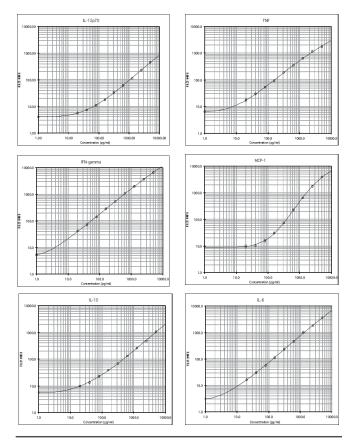
Typical data

The following data, acquired using BD CellQuestTM software, shows standards and detectors alone.



Standard curve examples

The following graphs represent standard curves from the BD CBA Mouse Inflammation Standards.



Performance

This section covers the following topics:

- Theoretical limit of detection (page 30)
- Recovery (page 31)
- Linearity (page 32)
- Specificity (page 33)
- Precision (page 35)

Theoretical limit of detection

Experiment details

The individual standard curve range for a given protein defines the minimum and maximum quantifiable levels using the BD CBA Mouse Inflammation Kit (ie, 20 pg/mL and 5,000 pg/mL). By applying the 4-parameter curve fit option, it is possible to extrapolate values for sample intensities not falling within the limits of the standard curve. It is up to the researcher to decide the best method for calculating values for unknown samples using this assay. The theoretical limit of detection for each protein using the BD CBA Mouse Inflammation Kit is defined as the corresponding concentration at two standard deviations above the median fluorescence of 20 replicates of the negative control (0 pg/mL).

Limit of detection data

Cytokine	Median fluorescence	Standard deviation	Limit of detection (pg/mL)
IL-6	3.3	0.3	5
IL-10	5.9	0.5	17.5
MCP-1	9.3	0.4	52.7
IFN-γ	5.4	0.4	2.5
TNF	6.5	0.3	7.3
IL-12p70	4.1	0.4	10.7

Recovery

Experiment details

Individual protein was spiked into various matrices at three different levels within the assay range. The cell culture media used in these experiments were not diluted before addition of the protein. The pooled mouse serum samples in these experiments were diluted 1:4 in Assay Diluent before addition of the cytokine protein. Results are compared with the same concentrations of the proteins spiked in the Assay Diluent.

Recovery data

Protein	Matrix	Standard spike conc. (pg/mL)	Observed in given matrix (pg/mL)	% Recovery
IL-6	Pooled mouse sera (1:4 dilution)	2,500 625 80	1,990.3 458.4 62.3	80% 73% 78%
	Cell culture media	2,500 625 80	2,476.7 594.3 95.8	99% 95% 120%
IL-10	Pooled mouse sera (1:4 dilution)	2,500 625 80	918.5 208.2 11.2	37% 33% 14%
	Cell culture media	2,500 625 80	2,326.7 581.9 99.7	93% 93% 125%
MCP-1	Pooled mouse sera (1:4 dilution)	2,500 625 80	1,954.5 448.5 35.8	78% 72% 45%
	Cell culture media	2,500 625 80	2,474.2 579.2 82.7	99% 93% 103%

Protein	Matrix	Standard spike conc. (pg/mL)	Observed in given matrix (pg/mL)	% Recovery
IFN-γ	Pooled mouse sera (1:4 dilution)	2,500 625 80	1,386.8 323.9 46.6	55% 52% 58%
	Cell culture media	2,500 625 80	2,423.9 584.8 79.7	97% 94% 100%
TNF	Pooled mouse sera (1:4 dilution)	2,500 625 80	779.8 181.0 26.8	31% 29% 34%
	Cell culture media	2,500 625 80	2,651.7 587.5 83.3	106% 94% 104%
IL-12p70	Pooled mouse sera (1:4 dilution)	2,500 625 80	1,468.5 350.5 23.5	59% 56% 29%
	Cell culture media	2,500 625 80	2,349.0 538.2 73.9	94% 86% 92%

Linearity

Experiment details

In two experiments, the following matrices were spiked with IL-6, IL-10, MCP-1, IFN-γ, TNF, and IL-12p70 and were then serially diluted with Assay Diluent.

Linearity data

		Observed (pg/mL)					
Matrix	Dilution	IL-6	IL-10	MCP-1	IFN-γ	TNF	IL-12p70
Pooled	1:4	4,041.8	1,935.9	4,052.3	2,877.2	1,601.9	3,372.1
mouse	1:8	2,128.9	1,244.7	2,227.9	1,588.3	1,001.2	1,675.3
sera	1:16	1,086.9	797.3	1,183.0	958.8	630.0	913.5
(1:4	1:32	529.3	438.6	619.2	520.4	370.7	491.7
starting	1:64	277.1	238.7	291.8	255.4	211.5	242.8
dilution)	1:128	136.9	136.7	160.5	140.1	122.0	142.2
	1:256	70.6	76.7	77.8	69.9	73.4	82.7
	1:512	33.2	29.9	22.3	34.2	38.0	34.4
	1:1,024	14.5	14.2	0.0	18.1	17.5	12.9
	Slope	1.006	0.882	1.029	0.922	0.801	0.961
Cell	Neat	4,347.7	4,870.6	4,946.1	4,518.4	4,407.9	4,615.7
culture	1:2	2,370.3	2,373.0	2,379.3	2,392.4	2,576.8	2,426.8
media	1:4	1,207.6	1,271.9	1,202.2	1,255.9	1,330.1	1,203.6
	1:8	572.5	588.9	593.1	640.7	616.1	588.7
	1:16	295.0	312.8	296.4	303.3	321.2	328.2
	1:32	132.3	151.0	135.7	145.3	146.3	130.6
	1:64	68.9	72.9	57.8	72.2	75.8	63.0
	1:128	39.8	44.2	34.6	38.6	43.2	44.5
	1:256	17.5	21.6	18.1	20.9	21.6	19.1
	Slope	0.998	0.979	1.026	0.999	0.979	1.113

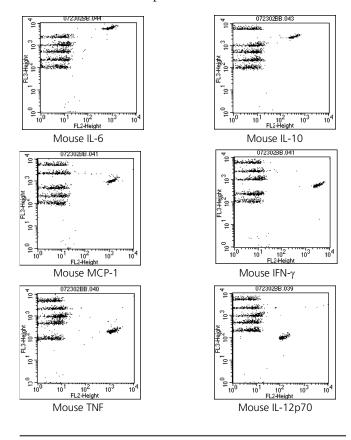
Specificity

Experiment details

The antibody pairs used in the BD CBA Mouse Inflammation assay have been screened for specific reactivity with their specific proteins. Analysis of samples containing only a single recombinant protein found no cross-reactivity or background detection of protein in other Capture Bead populations using this assay.

Specificity data

The following plots show BD CellQuest data for the detection of individual proteins.



Precision

Intra-assay precision

Ten replicates of each of three different levels of IL-6, IL-10, MCP-1, IFN- γ , TNF, and IL-12p70 were tested.

Protein	Actual mean conc. (pg/mL)	SD	%CV
IL-6	76	4	5%
	551	20	4%
	2,107	94	5%
IL-10	78	7	9%
	573	19	3%
	1,987	104	5%
MCP-1	66	6	10%
	614	9	2%
	2,300	46	2%
IFN-γ	73	3	4%
	633	19	3%
	2,239	103	5%
TNF	69	2	3%
	597	21	4%
	2,358	103	4%
IL-12p70	66	10	15%
	600	23	4%
	2,238	103	5%

Inter-assay precision

Inter-assay: Three different levels of IL-6, IL-10, MCP-1, IFN- γ , TNF, and IL-12p70 (80, 625, and 2,500 pg/mL) were tested in four experiments.

Note: Eight replicates (total number of assay tubes) were tested for each concentration of protein.

Protein	Actual mean conc. (pg/mL)	SD	% CV
IL-6	79	8	10%
	557	62	11%
	2,426	198	8%
IL-10	71	10	14%
	581	45	8%
	2,329	162	7%
MCP-1	58	11	19%
	598	33	5%
	2,431	51	2%
IFN-γ	75	6	8%
	631	40	6%
	2,425	120	5%
TNF	75	8	11%
	592	40	7%
	2,697	155	6%
IL-12p70	72	7	9%
_	590	30	5%
	2,430	70	3%

Reference

This section covers the following topics:

- Troubleshooting (page 38)
- References (page 40)

Troubleshooting

Recommended actions

These are the actions we recommend taking if you encounter the following problems.

Note: For best performance, vortex samples immediately before analyzing on a flow cytometer.

Note: The BD CBA Mouse Inflammation assay has been shown to detect TNF produced by the activation of cells from the rat model. Direct quantitation of proteins from the rat model has not been validated using this kit and results may vary.

Problem	Recommended action
Variation between duplicate samples	Vortex Capture Beads before pipetting. Beads can aggregate.
Low bead number in	• Avoid aspiration of beads during wash step.
samples	 Do not wash or resuspend beads in volumes higher than recommended volumes.
High background	• Test various sample dilutions. The sample may be too concentrated.
	 Remove excess Mouse Inflammation PE Detection Reagent by increasing the number of wash steps, since the background may be due to non-specific binding.
Little or no detection of protein in sample	Sample may be too dilute. Try various sample dilutions.
Less than six bead populations are	• Ensure that equal volumes of beads were added to each assay tube.
observed during analysis, or distribution is unequal	 Vortex Capture Bead vials before taking aliquots. Once Capture Beads are mixed, vortex to ensure that the beads are distributed evenly throughout the solution.

Problem	Recommended action		
Debris (FSC/SSC) during sample	• Increase FSC threshold or further dilute samples.		
acquisition	• Increase the number of wash steps, if necessary.		
	• Make a tighter FSC/SSC region gate around the bead population.		
Overlap of bead population fluorescence (FL3) during acquisition	This may occur in samples with very high protein concentration. Ensure that instrument settings have been optimized using the Cytometer Setup Beads.		
Standards assay tubes show low fluorescence	Ensure that all components are properly prepared and stored.		
or poor standard curve	Use a new vial of standard with each experiment, and once reconstituted, do not use after 12 hours.		
	Ensure that incubation times were appropriate.		
All samples are positive or above the high standard median fluorescence value	Dilute the samples further. The samples may be too concentrated.		
Biohazardous samples	It is possible to treat samples briefly with 1% paraformaldehyde before analyzing on the flow cytometer. However, this may affect assay performance and should be validated.		

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Notes	

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