

BD™ Cytometric Bead Array
(CBA) Human Th1/Th2
Cytokine Kit II
Instruction Manual

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Regulatory information

BD cytometers are class 1 laser products.

For Research Use Only. Not for use in diagnostic or therapeutic procedures.

History

| Revision | Date | Change made |
|---------------------|---------|----------------------------|
| 23-12495-00 Rev. 01 | 11/2010 | Initial release |
| 23-12495-01 | 2/2013 | Updated patent information |

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About this kit

This section covers the following topics:

- [Purpose of the kit \(page 6\)](#)
- [Limitations \(page 8\)](#)
- [Kit contents \(page 9\)](#)
- [Storage and handling \(page 11\)](#)

Purpose of the kit

Use of the kit The BD™ CBA Human Th1/Th2 Cytokine Kit II (Catalog No. 551809) can be used to quantitatively measure Interleukin-2 (IL-2), Interleukin-4 (IL-4), Interleukin-6 (IL-6), Interleukin-10 (IL-10), Tumor Necrosis Factor (TNF), and Interferon- γ (IFN- γ) protein levels in a single sample. The kit performance has been optimized for analysis of specific cytokines in tissue culture supernatants, EDTA plasma, and serum samples. The kit provides sufficient reagents for the quantitative analysis of 80 samples.

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 a BD CBA 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 assay

The BD CBA Human Th1/Th2 Cytokine Kit II uses bead array technology to simultaneously detect multiple cytokine proteins in research samples. Six bead populations with distinct fluorescence intensities have been coated with capture antibodies specific for IL-2, IL-4, IL-6, IL-10, TNF, and IFN- γ proteins. The six bead populations are mixed together to form the bead array, which is resolved in a red channel of a flow cytometer.

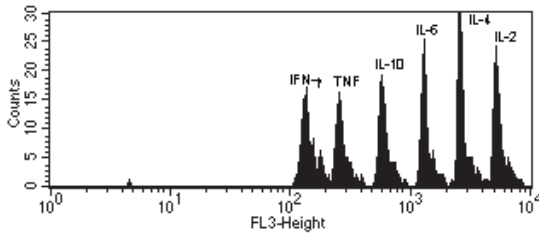


Figure 1

During the assay procedure, you will mix the cytokine capture beads with recombinant standards or unknown samples and incubate them with the PE-conjugated 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 Array™ 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 the BD CBA assay 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 one-sixth 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 Human Th1/Th2 Cytokine Kit II is comparable to conventional ELISA, but due to the complexity and kinetics of this multi-analyte assay, the actual limit of detection on a given experiment may vary slightly. See [Theoretical limit of detection \(page 32\)](#) and [Precision \(page 38\)](#).

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 FACStar™ Plus, BD Influx™, and BD FACSVantage™ flow cytometers (BD Biosciences, San Jose, CA).

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 BD CBA Human Th1/Th2 Cytokine Kit II contains the following components sufficient for 80 tests.

| Vial label | Reagent | Quantity |
|------------|---|-----------------------------|
| A1 | Human IL-2 Capture Beads | 1 vial, 0.8 mL |
| A2 | Human IL-4 Capture Beads | 1 vial, 0.8 mL |
| A3 | Human IL-6 Capture Beads | 1 vial, 0.8 mL |
| A4 | Human IL-10 Capture Beads | 1 vial, 0.8 mL |
| A5 | Human TNF Capture Beads | 1 vial, 0.8 mL |
| A6 | Human IFN- γ Capture Beads | 1 vial, 0.8 mL |
| B | Human Th1/Th2 - II PE Detection Reagent | 1 vial, 4 mL |
| C | Human Th1/Th2 Cytokine Standards | 2 vials, 0.2 mL lyophilized |
| D | Cytometer Setup Beads | 1 vial, 1.5 mL |
| E1 | PE Positive Control Detector | 1 vial, 0.5 mL |
| E2 | FITC Positive Control Detector | 1 vial, 0.5 mL |

| Vial label | Reagent | Quantity |
|------------|--------------------------|------------------|
| F | Wash Buffer | 1 bottle, 130 mL |
| G | Assay Diluent | 1 bottle, 30 mL |
| H | Serum Enhancement Buffer | 1 bottle, 10 mL |

Bead reagents

Human Cytokine 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 $\mu\text{L}/\text{test}$.

Antibody and standard reagents

Human Th1/Th2 - II PE Detection Reagent (B): An 80-test vial of mixed PE-conjugated anti-human IL-2, IL-4, IL-6, IL-10, TNF, and IFN- γ antibodies formulated for use at 50 $\mu\text{L}/\text{test}$.

Human Th1/Th2 Cytokine Standards (C): Two vials containing lyophilized recombinant human cytokine proteins. Each vial should be reconstituted in 2.0 mL of Assay Diluent to prepare the top standard.

PE Positive Control Detector (E1): A 10-test vial of PE-conjugated antibody control formulated for use at 50 $\mu\text{L}/\text{test}$. 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 formulated for use at 50 $\mu\text{L}/\text{test}$. 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 Human Th1/Th2 Cytokine Standards and to dilute test samples.

Serum Enhancement Buffer (H): A 10-mL bottle of a buffered protein solution (1X) used to dilute mixed Capture Beads when testing serum or plasma 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, G, and H 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.

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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 | Preparation of Human Th1/Th2 Cytokine Standards (page 18) |
| 2 | Mixing Human Th1/Th2 Cytokine Capture Beads (page 20) |
| 3 | Diluting samples (page 22) |
| 4 | Performing instrument setup with Cytometer Setup Beads (instructions can be found at bdbiosciences.com/cbasetup) Note: Can be performed during the incubation in step 5. |
| 5 | Performing the Human Th1/Th2 Cytokine II Assay (page 24) |
| 6 | Acquiring samples (instructions can be found at bdbiosciences.com/cbasetup) |
| 7 | Data analysis (page 28) |

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 Capture Beads | 30 minutes (for serum/ plasma samples only) |
| Preparing Cytometer Setup Beads | 30 minutes |
| Performing the assay | 3 hours |

Required materials

Materials required but not provided

In addition to the reagents provided in the BD CBA Human Th1/Th2 Cytokine Kit II, 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 FACSArry™ | Yellow | Red |
| BD FACSCanto™ platform BD™ LSR platform BD FACSAria™ platform | PE | APC |
| BD FACSCalibur™ (single laser) BD FACSCalibur (dual laser) | FL2 | FL3 FL4 |
| Note: Visit bdbiosciences.com/cbasetup for setup protocols. | | |

- BD Falcon™ 12 × 75-mm sample acquisition tubes (Catalog No. 352008), or equivalent
- 15-mL conical, polypropylene tubes (BD Falcon, Catalog No. 352097), or equivalent
- FCAP Array software (Catalog No. 641488 [PC] or 645447 [Mac®])

**Materials
required for
plate loader-
equipped flow
cytometers**

- Millipore MultiScreen_{HTS}-BV 1.2- μ m Clear non-sterile 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)
-

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Assay preparation

This section covers the following topics:

- [Preparation of Human Th1/Th2 Cytokine Standards \(page 18\)](#)
- [Mixing Human Th1/Th2 Cytokine Capture Beads \(page 20\)](#)
- [Diluting samples \(page 22\)](#)

Preparation of Human Th1/Th2 Cytokine Standards

Purpose of this procedure The Human Th1/Th2 Cytokine Standards are lyophilized and must be reconstituted and serially diluted immediately before mixing with the Capture Beads and the PE Detection Reagent.

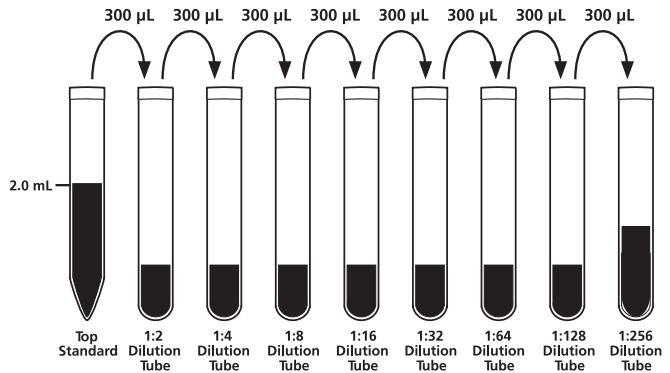
You must prepare fresh cytokine standards to run with each experiment. Do not store or reuse reconstituted or diluted standards.

Procedure

To reconstitute and serially dilute the standards:

1. Open one vial of lyophilized Human Th1/Th2 Standards. Transfer the standard spheres to a 15-mL conical, polypropylene tube. Label the tube “Top Standard.”
2. 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 reconstituted protein by pipet only. Do not vortex or mix vigorously.
3. Label eight 12 × 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 μ L of Assay Diluent to 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.
 - b. Continue making serial dilutions by transferring 300 μL 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.



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

Concentration of standards

See the table in [Performing the Human Th1/Th2 Cytokine II Assay \(page 24\)](#) for a listing of the concentrations (pg/mL) of all six recombinant proteins in each standard dilution.

Next step

Proceed to [Mixing Human Th1/Th2 Cytokine Capture Beads \(page 20\)](#).

Mixing Human Th1/Th2 Cytokine Capture Beads

Purpose of this procedure

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

Follow the procedure to mix the Capture Beads for all sample types. Then perform additional steps to incubate the beads in Serum Enhancement Buffer, if analyzing serum and/or plasma samples, to reduce the chances of false-positive results due to the effects of serum or plasma proteins.

Procedure

To mix the Capture Beads when testing any sample type:

1. Determine the number of assay tubes (including standards and controls) that are required for the experiment (eg, 8 unknowns, 9 cytokine standard dilutions, and 1 negative control = 18 assay tubes).
2. Vigorously vortex each Capture Bead suspension for a few seconds before mixing.
3. Add a 10- μ L aliquot of each Capture Bead, for each assay tube to be analyzed, into a single tube labeled “mixed Capture Beads” (eg, 10 μ L of IL-2 Capture Beads \times 18 assay tubes = 180 μ L of IL-2 Capture Beads required).
4. Vortex the bead mixture thoroughly.
5. If you are testing serum or plasma samples, you will need to incubate the Capture Beads in Serum Enhancement Buffer. Proceed to [Additional steps when testing serum and plasma samples \(page 21\)](#).

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 Human Th1/Th2 Cytokine II Assay \(page 24\)](#). If you need to dilute samples having high cytokine concentration, proceed to [Diluting samples \(page 22\)](#).

Additional steps when testing serum and plasma samples

Note: These steps can also be performed when testing cell culture supernatants.

When testing serum and/or plasma samples:

1. Perform the procedure to mix the beads in [Procedure \(page 20\)](#).
 2. Centrifuge the mixed Capture Beads at 200g for 5 minutes.
 3. Carefully aspirate and discard the supernatant.
 4. Resuspend the mixed Capture Beads pellet in Serum Enhancement Buffer (equal volume to amount removed in [step 3](#)) and vortex thoroughly.
 5. Incubate the mixed Capture Beads for 30 minutes at room temperature, protected from light.
-

Next step

To begin the assay, proceed to [Performing the Human Th1/Th2 Cytokine II Assay \(page 24\)](#). If you need to dilute samples having a high cytokine concentration, proceed to [Diluting samples \(page 22\)](#).

Diluting samples

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

Procedure To dilute samples with a known high cytokine 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.

Or, if you wish to begin staining your samples for the assay, proceed to [Performing the Human Th1/Th2 Cytokine II Assay \(page 24\)](#), and you can perform instrument setup during the 3-hour staining incubation.

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Assay procedure

- [Performing the Human Th1/Th2 Cytokine II Assay \(page 24\)](#)
- [Data analysis \(page 28\)](#)

Performing the Human Th1/Th2 Cytokine II Assay

- Before you begin**
- Prepare the standards as described in [Preparation of Human Th1/Th2 Cytokine Standards \(page 18\)](#).
 - Mix the Capture Beads as described in [Mixing Human Th1/Th2 Cytokine Capture Beads \(page 20\)](#).
 - If necessary, dilute the unknown samples. See [Diluting samples \(page 22\)](#).

Procedure for tubes

Follow these steps regardless of whether you are testing cell culture supernatants or serum/plasma samples. If testing serum and/or plasma samples, be sure to use the Capture Beads that you prepared specifically for the serum/plasma samples. See [Additional steps when testing serum and plasma samples \(page 21\)](#).

To perform the assay:

1. Vortex the mixed Capture Beads and add 50 μ L to all assay tubes.
2. Add 50 μ L of the Human Th1/Th2 Cytokine Standard dilutions to the control assay tubes as listed in the following table.

| Tube label | Concentration (pg/mL) | Cytokine 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 | 312.5 | 1:16 |
| 7 | 625 | 1:8 |

| Tube label | Concentration (pg/mL) | Cytokine Standard dilution |
|------------|-----------------------|----------------------------|
| 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 tubes.
4. Add 50 μL of the Human Th1/Th2 - II PE Detection Reagent to all assay tubes.
5. Incubate the assay tubes for 3 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 200g for 5 minutes.
7. Carefully aspirate and discard the supernatant from each assay tube.
8. Add 300 μL 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 μ L 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)
 - Human Th1/Th2 - II 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 3 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 μ L 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 \(page 28\)](#).

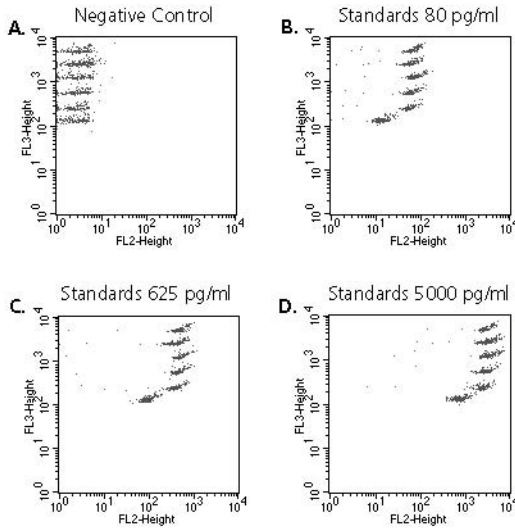
Data analysis

How to analyze

Analyze BD CBA Human Th1/Th2 Cytokine Kit II 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*.

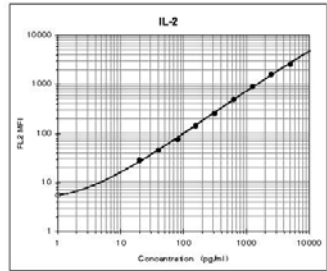
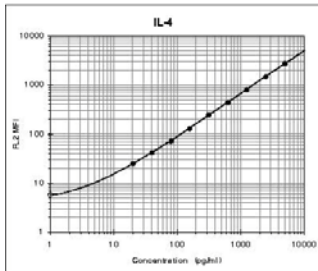
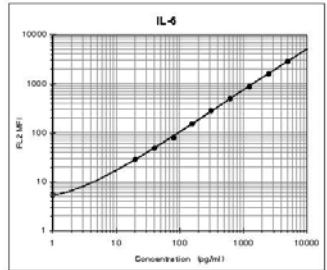
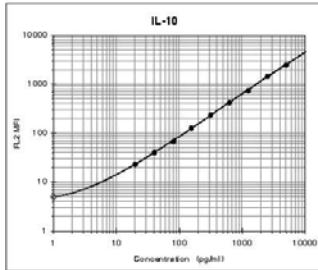
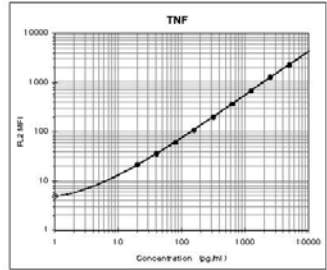
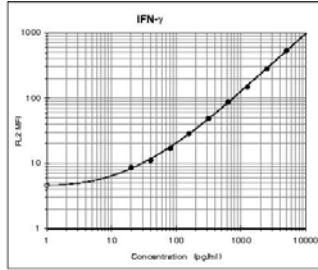
Typical Data

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



Standard curve examples

The following graphs represent standard curves from the BD CBA Human Th1/Th2 Cytokine Kit II.



5

Performance

- [Theoretical limit of detection \(page 32\)](#)
- [Recovery \(page 33\)](#)
- [Linearity \(page 35\)](#)
- [Specificity \(page 37\)](#)
- [Precision \(page 38\)](#)

Theoretical limit of detection

Experiment details

The individual standard curve range for a given cytokine defines the minimum and maximum quantifiable levels (20 pg/mL and 5,000 pg/mL) using the BD CBA Human Th1/Th2 Cytokine Kit II. 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 cytokine using the BD CBA Human Th1/Th2 Cytokine Kit II 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-2 | 3.3 | 0.2 | 2.6 |
| IL-4 | 2.3 | 0.2 | 2.6 |
| IL-6 | 2.6 | 0.2 | 3.0 |
| IL-10 | 2.4 | 0.2 | 2.8 |
| TNF | 2.0 | 0.2 | 2.8 |
| IFN- γ | 2.1 | 0.3 | 7.1 |

Recovery

Experiment details

Individual cytokine proteins were spiked into various matrices at three different levels within the assay range. The matrices used in these experiments were not diluted before addition of the cytokine protein. The plasma samples in these experiments were EDTA treated. Results are compared with the same concentrations of the cytokines spiked in the Standard Diluent, as follows.

Recovery data

| Cytokine | Matrix | Standard spike concentration (pg/mL) | Observed in given matrix (pg/mL) | % Recovery |
|----------|---------------------------|--------------------------------------|----------------------------------|------------|
| IL-2 | Pooled donor sera (n=5) | 2,500 | 2,028 | 81% |
| | | 625 | 439 | 70% |
| | | 80 | 67 | 84% |
| | Pooled donor plasma (n=5) | 2,500 | 1,870 | 87% |
| | | 625 | 439 | 70% |
| | | 80 | 50 | 62% |
| | Cell culture supernatant | 2,500 | 2,471 | 99% |
| | | 625 | 737 | 118% |
| | | 80 | 86 | 107% |
| IL-4 | Pooled donor sera (n=5) | 2,500 | 1,937 | 78% |
| | | 625 | 475 | 76% |
| | | 80 | 70 | 88% |
| | Pooled donor plasma (n=5) | 2,500 | 1,863 | 76% |
| | | 625 | 785 | 78% |
| | | 80 | 58 | 73% |
| | Cell culture supernatant | 2,500 | 2,367 | 95% |
| | | 625 | 684 | 110% |
| | | 80 | 84 | 104% |

| Cytokine | Matrix | Standard spike concentration (pg/mL) | Observed in given matrix (pg/mL) | % Recovery |
|----------|---------------------------|--------------------------------------|----------------------------------|------------|
| IL-6 | Pooled donor sera (n=5) | 2,500 | 2,187 | 88% |
| | | 625 | 533 | 85% |
| | | 80 | 73 | 91% |
| | Pooled donor plasma (n=5) | 2,500 | 1,959 | 84% |
| | | 625 | 491 | 79% |
| | | 80 | 61 | 76% |
| | Cell culture supernatant | 2,500 | 2,523 | 101% |
| | | 625 | 720 | 115% |
| | | 80 | 83 | 104% |
| IL-10 | Pooled donor sera (n=5) | 2,500 | 2,004 | 80% |
| | | 625 | 506 | 81% |
| | | 80 | 60 | 76% |
| | Pooled donor plasma (n=5) | 2,500 | 1,967 | 80% |
| | | 625 | 499 | 80% |
| | | 80 | 62 | 77% |
| | Cell culture supernatant | 2,500 | 2,530 | 101% |
| | | 625 | 719 | 115% |
| | | 80 | 83 | 104% |
| TNF | Pooled donor sera (n=5) | 2,500 | 1,907 | 76% |
| | | 625 | 487 | 78% |
| | | 80 | 60 | 75% |
| | Pooled donor plasma (n=5) | 2,500 | 1,773 | 70% |
| | | 625 | 477 | 76% |
| | | 80 | 58 | 72% |
| | Cell culture supernatant | 2,500 | 2,552 | 102% |
| | | 625 | 743 | 119% |
| | | 80 | 85 | 107% |

| Cytokine | Matrix | Standard spike concentration (pg/mL) | Observed in given matrix (pg/mL) | % Recovery |
|---------------|---------------------------|--------------------------------------|----------------------------------|------------|
| IFN- γ | Pooled donor sera (n=5) | 2,500 | 1,390 | 56% |
| | | 625 | 403 | 65% |
| | | 80 | 46 | 57% |
| | Pooled donor plasma (n=5) | 2,500 | 1,494 | 67% |
| | | 625 | 441 | 71% |
| | | 80 | 61 | 77% |
| | Cell culture supernatant | 2,500 | 2,184 | 87% |
| | | 625 | 711 | 114% |
| | | 80 | 79 | 99% |

Linearity

Experiment details

In two experiments, the following matrices were spiked with IL-2, IL-4, IL-6, IL-10, TNF, and IFN- γ and were then serially diluted with Assay Diluent.

Linearity data

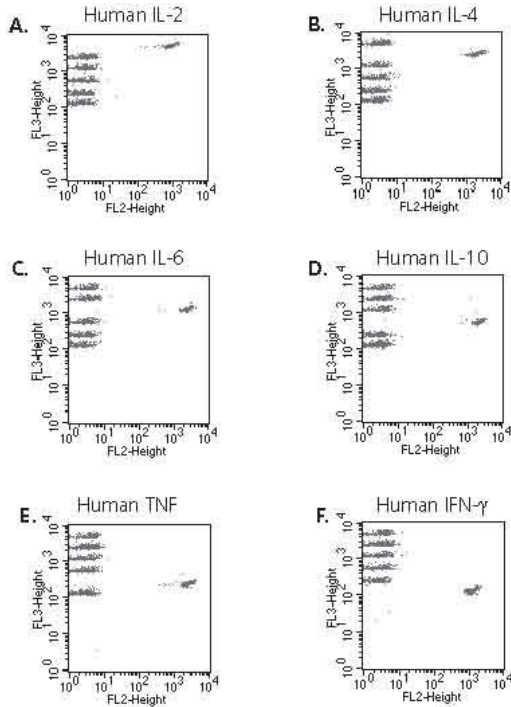
| Matrix | Dilution | Observed (pg/mL) | | | | | |
|---------------------------|----------|------------------|-------|-------|-------|-------|---------------|
| | | IL-2 | IL-4 | IL-6 | IL-10 | TNF | IFN- γ |
| Pooled donor sera (n=5) | Neat | 3,733 | 3,926 | 4,017 | 3,886 | 3,765 | 2,462 |
| | 1:2 | 2,020 | 2,147 | 2,175 | 2,169 | 1,920 | 1,407 |
| | 1:4 | 1,025 | 1,210 | 1,166 | 1,122 | 1,144 | 821 |
| | 1:8 | 497 | 611 | 600 | 582 | 630 | 450 |
| | 1:16 | 237 | 303 | 305 | 289 | 303 | 220 |
| | 1:32 | 120 | 153 | 141 | 147 | 155 | 112 |
| | 1:64 | 61 | 82 | 78 | 79 | 86 | 59 |
| | 1:128 | 28 | 40 | 40 | 39 | 43 | 42 |
| | 1:256 | 15 | 21 | 20 | 19 | 21 | 19 |
| | Slope | 1.09 | 1.00 | 1.00 | 0.99 | 0.95 | 1.19 |
| Pooled donor plasma (n=5) | Neat | 3,600 | 3,752 | 3,656 | 3,788 | 3,523 | 2,980 |
| | 1:2 | 1,981 | 2,225 | 2,092 | 2,166 | 2,088 | 1,630 |
| | 1:4 | 922 | 1,103 | 1,043 | 1,070 | 1,072 | 824 |
| | 1:8 | 448 | 596 | 547 | 558 | 600 | 435 |
| | 1:16 | 216 | 316 | 294 | 300 | 328 | 219 |
| | 1:32 | 100 | 163 | 159 | 154 | 174 | 113 |
| | 1:64 | 50 | 79 | 75 | 75 | 86 | 63 |
| | 1:128 | 23 | 40 | 38 | 36 | 45 | 36 |
| | 1:256 | 14 | 21 | 20 | 19 | 23 | 28 |
| | Slope | 1.12 | 0.99 | 0.97 | 0.99 | 0.93 | 1.17 |
| Cell culture medium | Neat | 4,376 | 4,468 | 4,706 | 4,561 | 4,554 | 4,350 |
| | 1:2 | 2,622 | 2,499 | 2,610 | 3,109 | 2,679 | 2,186 |
| | 1:4 | 1,277 | 1,212 | 1,289 | 1,288 | 1,345 | 1,028 |
| | 1:8 | 616 | 622 | 634 | 657 | 668 | 508 |
| | 1:16 | 286 | 307 | 316 | 331 | 335 | 251 |
| | 1:32 | 137 | 156 | 160 | 160 | 171 | 128 |
| | 1:64 | 69 | 83 | 78 | 83 | 87 | 66 |
| | 1:128 | 33 | 40 | 43 | 41 | 44 | 37 |
| | 1:256 | 15 | 19 | 21 | 20 | 21 | 27 |
| | Slope | 1.11 | 1.02 | 1.00 | 1.00 | 0.99 | 1.11 |

Specificity

Experiment details

The antibody pairs used in the BD CBA Human Th1/Th2 Cytokine Kit II assay have been screened for specific reactivity with their specific cytokines. Analysis of samples containing only a single recombinant cytokine protein found no cross-reactivity or background detection of cytokine in other Capture Bead populations using this assay. The following data shows the detection of individual cytokines.

Specificity data



Precision

Intra-assay precision

Ten replicates of each of three different levels of IL-2, IL-4, IL-6, IL-10, TNF, and IFN- γ were tested.

| Cytokine | Actual Mean Conc (pg/mL) | Standard deviation | %CV |
|---------------|--------------------------|--------------------|-----|
| IL-2 | 64 | 3 | 4 |
| | 514 | 25 | 5 |
| | 2,172 | 51 | 2 |
| IL-4 | 73 | 3 | 4 |
| | 590 | 21 | 3 |
| | 2,477 | 71 | 3 |
| IL-6 | 70 | 2 | 3 |
| | 571 | 14 | 2 |
| | 2,383 | 77 | 3 |
| IL-10 | 70 | 2 | 3 |
| | 577 | 10 | 2 |
| | 2,447 | 70 | 3 |
| TNF | 73 | 3 | 4 |
| | 581 | 12 | 2 |
| | 2,422 | 71 | 3 |
| IFN- γ | 64 | 3 | 4 |
| | 493 | 17 | 3 |
| | 1,916 | 85 | 4 |

Inter-assay

Three different levels of IL-2, IL-4, IL-6, IL-10, TNF, and IFN- γ (80, 625, and 2,500 pg/mL) were tested in four experiments conducted by different operators.

Note: The number of replicates refers to the total number of assay tubes tested at a given concentration of protein.

| Cytokine | Number of replicates | Actual Mean Conc (pg/mL) | Standard deviation | %CV |
|---------------|----------------------|--------------------------|--------------------|-----|
| IL-2 | 8 | 64 | 6 | 9 |
| | 8 | 538 | 34 | 6 |
| | 8 | 2,446 | 100 | 4 |
| IL-4 | 8 | 70 | 5 | 7 |
| | 8 | 576 | 17 | 3 |
| | 8 | 2,535 | 86 | 3 |
| IL-6 | 8 | 71 | 5 | 7 |
| | 8 | 581 | 17 | 3 |
| | 8 | 2,502 | 88 | 3 |
| IL-10 | 8 | 74 | 5 | 6 |
| | 8 | 586 | 22 | 4 |
| | 8 | 2,562 | 95 | 4 |
| TNF | 8 | 77 | 5 | 6 |
| | 8 | 613 | 21 | 3 |
| | 8 | 2,562 | 88 | 3 |
| IFN- γ | 8 | 56 | 6 | 11 |
| | 8 | 542 | 33 | 6 |
| | 8 | 2,260 | 121 | 5 |

6

Reference

- [Troubleshooting \(page 42\)](#)
- [References \(page 44\)](#)

Troubleshooting

Recommended actions

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

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

Note: The BD CBA Human Th1/Th2 Cytokine Kit II assay has been shown to detect IL-4, IL-6, TNF, and IFN- γ produced by the activation of cells from the non-human primate rhesus and cynomolgus models. Direct quantitation of cytokines from the rhesus and cynomolgus models has not been validated using this kit and results may vary.

| Problem | Suggested solution |
|---|---|
| Variation between duplicate samples. | Vortex Capture Beads before pipetting. Beads can aggregate. |
| Low bead number in samples. | Avoid aspiration of beads during the wash step. 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 Human Th1/Th2 - II 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 observed during analysis or distribution is unequal. | Ensure that equal volumes of beads were added to each assay tube. 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 | Suggested solution |
|--|--|
| Debris (FSC/SSC) during sample acquisition. Also for plasma samples. | Increase the FSC threshold or further dilute samples. 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 cytokine concentration. Ensure that instrument settings have been optimized using the Cytometer Setup Beads. |
| Standards assay tubes show low fluorescence or poor standard curve. | Check that all components are properly prepared and stored. Use a new vial of standard with each experiment and once reconstituted, do not use after 12 hours. Ensure that incubation times were of proper length. |
| All samples are positive or above the high standard mean 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 by the user. |

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