

- While plate is being blocked, prepare the appropriate sample dilutions (if necessary) and standards.
- Wash plate 4 times with Wash Buffer.
- Add 100 μ L/well of standards or samples to the appropriate wells. If dilution is required, samples should be diluted in 1X Assay Diluent A before adding to the wells.
- Seal plate and incubate at RT for 2 hours with shaking.
- Wash plate 4 times with Wash Buffer.
- Add 100 μ L of diluted Detection Antibody solution to each well, seal plate and incubate at RT for 1 hour with shaking.
- Wash plate 4 times with Wash Buffer.
- Add 100 μ L of diluted Avidin-HRP solution to each well, seal plate and incubate at RT for 30 minutes with shaking.
- Wash plate 5 times with Wash Buffer. For this final wash, soak wells in Wash Buffer for 30 seconds to 1 minute for each wash. This will help minimize background.
- Add 100 μ L of freshly mixed TMB Substrate Solution and incubate **in the dark** for 15 minutes*. Positive wells should turn blue in color. It is not necessary to seal the plate during this step.
- Stop reaction by adding 100 μ L of Stop Solution to each well. Positive wells should turn from blue to yellow.
- Read absorbance at 450 nm within 15 minutes. If the reader can read at 570 nm, the absorbance at 570 nm can be subtracted from the absorbance at 450 nm.

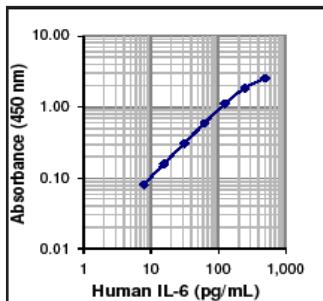
*Optimal substrate incubation time depends on laboratory conditions and the optical linear ranges of ELISA plate readers.

Calculation of Results

Plot the standard curve on log-log axis graph paper with analyte concentration on the x-axis and absorbance on the y-axis. Draw a best fit line through the standard points. To determine the unknown analyte concentrations in the samples, find the absorbance value of the unknown on the y-axis and draw a horizontal line to the standard curve. At the point of intersection, draw a vertical line to the x-axis and read the corresponding analyte concentration. If the samples were diluted, multiply by the appropriate dilution factor. The data is best calculated with computer-based curve-fitting software using a 5- or 4-parameter logistics curve-fitting algorithm. If a test sample's absorbance value falls outside the standard curve ranges, that test sample needs to be reanalyzed at a higher or lower dilution as appropriate.

Typical Data

Standard Curve: This standard curve was generated at BioLegend for demonstration purposes only. A standard curve must be run with each assay.



Assay Procedure Summary

- Coat plate with 100 μ L diluted Capture Antibody
incubate overnight, between 2°C and 8°C
- Wash 4 times
Add 200 μ L 1X Assay Diluent A
Incubate 1 hr, RT, shaking
- Wash 4 times
Add 100 μ L diluted standards and samples
Incubate 2 hrs, RT, shaking
- Wash 4 times
Add 100 μ L diluted Detection Antibody
Incubate 1 hr, RT, shaking
- Wash 4 times
Add 100 μ L Avidin-HRP
Incubate 30 min. RT, shaking
- Wash 5 times
Add 100 μ L TMB Substrate Solution
Incubate 15 min. RT, in the dark
- Add 100 μ L Stop Solution
- Read absorbance at 450nm and 570nm

Performance Characteristics

Sensitivity: The expected minimum detectable concentration of IL-6 for this set is 4 pg/ml.

Specificity: No cross reactivity was observed when this kit was used to analyze multiple human, mouse and rat recombinant proteins.

Troubleshooting

High Background:

- Background wells were contaminated.
- Matrix used had endogenous analyte.
- Plate was insufficiently washed.
- TMB Substrate Solution was contaminated.

No signal:

- Incorrect or no antibodies were added.
- Avidin-HRP was not added.
- Substrate solution was not added.
- Wash buffer contains sodium azide.

Low or poor signal for the standard curve:

- Standard was incompletely reconstituted or was stored improperly.
- Reagents were added to wells with incorrect concentrations.
- Plate was incubated with improper temperature, timing or agitation.

Signal too high, standard curves saturated:

- Standard was reconstituted with less volume than required.
- One or more reagent incubation steps were too long.
- Plate was incubated with inappropriate temperature, timing, or agitation.

Sample readings out of range:

- Samples contain no or below detectable levels of analyte.
- Samples contain analyte concentrations greater than highest standard point.

High variations in samples and/or standards:

- Pipetting errors may have occurred.
- Plate washing was inadequate or nonuniform.
- Samples were not homogenous.
- Samples or standard wells were contaminated.

References Using This Set

- Ishola Jr, D. A., et al., 2006. Kidney International 70: 724
- Kim, I., et al., 2006. Blood 108: 737
- Wimmer, A., et al., 2006. Blood 108:3722.

BioLegend, Inc.

BioLegend is ISO 9001:2008 and ISO 13485:2003 Certified

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