

# Human CD120b ELISA Kit

# **User Manual**

Catalog # CEK1059

(Version 1.2B)

Sandwich Enzyme-Linked Immunosorbent Assay for Quantitative Detection of Human CD120b Concentrations in Cell Culture Supernatants, Serum, Plasma, Urine.

For research use only. Not for diagnostic or therapeutic procedures.



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# I. INTRODUCTION

Tumor necrosis factor receptor p75 (TNF-R p75, or TNF-R2) is a 75-kDa type I transmembrane protein expressed predominantly on cells of hematopoietic lineage. TNF-R p75 belongs to the TNF receptor superfamily characterized by cysteine-rich extracellular regions composed of three to six disulfide-linked domains. The tumor necrosis factor receptor II (TNFRII) gene localizes to 1p36. 2, a genomic region characteristically deleted in neuroblastomas and other malignancies. In addition, TNFRII is the principal mediator of the effects of TNF on cellular immunity, and it may cooperate with TNFRI in the killing of nonlymphoid cells.



#### **II. ASSAY PRINCIPLES**

The Cohesion Bioscience Human CD120b ELISA (Enzyme-Linked Immunosorbent Assay) kit is an in vitro enzyme-linked immunosorbent assay for the quantitative measurement of Human CD120b in Cell Culture Supernatants, Serum, Plasma, Urine. This assay employs an antibody specific for Human CD120b coated on a 96-well plate. Standards and samples are pipetted into the wells and CD120b present in a sample is bound to the wells by the immobilized antibody. The wells are washed and biotinylated anti-Human CD120b antibody is added. After washing away unbound biotinylated antibody, HRP-conjugated streptavidin is pipetted to the wells. The wells are again washed, a TMB substrate solution is added to the wells and color develops in proportion to the amount of CD120b bound. The Stop Solution changes the color from blue to yellow, and the intensity of the color is measured at 450 nm.



#### **III. KIT COMPONENTS**

Component	Volume
96-well Plate Coated with Anti-Human CD120b Antibody	8 wells x 12 Strips
Human CD120b Standard	10 ng x 2
Biotin-Labeled Detection Antibody (100X)	120 μl
Streptavidin-HRP (100X)	120 μl
Standard/Sample Diluent	30 ml
Detection Antibody Diluent	12 ml
Streptavidin-HRP Diluent	12 ml
Wash Buffer (20X)	30 ml
TMB Substrate Solution	12 ml
Stop Solution	12 ml
Plate Adhesive Strips	3 Strips
Technical Manual	1 Manual

# IV. STORAGE AND STABILITY

All kit components are stable at 2 to 8 °C. Standard (recombinant protein) should be stored at -20 °C or -80 °C (recommended at -80 °C) after reconstitution. Opened Microplate Wells or reagents may be store for up to 1 month at 2 to 8 °C. Return unused wells to the pouch containing desiccant pack, reseal along entire edge. Note: the kit can be used within one year if the whole kit is stored at -20 °C. Avoid repeated freeze-thaw cycles.



# V. MATERIALS REQUIRED BUT NOT PROVIDED

- 1. Microplate reader capable of measuring absorbance at 450 nm.
- 2. Adjustable pipettes and pipette tips to deliver 2  $\mu$ l to 1 ml volumes.
- 3. Adjustable 1-25 ml pipettes for reagent preparation.
- 4. 100 ml and 1 liter graduated cylinders.
- 5. Absorbent paper.
- 6. Distilled or deionized water.
- 7. Computer and software for ELISA data analysis.
- 8. Tubes to prepare standard or sample dilutions.

#### VI. HEALTH AND SAFETY PRECAUTIONS

1. Reagents provided in this kit may be harmful if ingested, inhaled or absorbed through the skin. Please carefully review the MSDS for each reagent before conducting the experiment.

Stop Solution contains 2 N Sulfuric Acid (H<sub>2</sub>SO<sub>4</sub>) and is an extremely corrosive agent. Please wear proper eye, hand and face protection when handling this material. When the experiment is finished, be sure to rinse the plate with copious amounts of running water to dilute the Stop Solution prior to disposing the plate.



#### VII. REAGENT PREPARATION

#### 1. Sample Preparation

Store samples to be assayed within 24 hours at 2-8°C. For long-term storage, aliquot and freeze samples at -20°C. Avoid repeated freeze-thaw cycles.

**Cell culture supernates**: Remove particulates by centrifugation, assay immediately or aliquot and store samples at -20°C.

**Serum**: Allow the serum to clot in a serum separator tube (about 4 hours) at room temperature. Centrifuge at approximately 1000 X g for 15 minutes. Analyze the serum immediately or aliquot and store samples at -20°C.

**Plasma**: Collect plasma using heparin or EDTA as an anticoagulant. Centrifuge for 15 minutes at 1500 X g within 30 minutes of collection. Assay immediately or aliquot and store samples at -20°C.

**Cell Lysates:** Collect cells and rinse cells with PBS. Homogenize and lyse cells throughly in lysate solution. Centrifuge cell lysates at approximately 10000 X g for 5 minutes to remove debris. Aliquots of the cell lysates were removed and assayed. **Bone Tissue:** Extract demineralized bone samples in 4 M Guanidine-HCl and protease inhibitors. Dissolve the final sample in 2 M Guanidine-HCl.

**Tissue Homogenates:** The preparation of tissue homogenates will vary depending upon tissue type. Rinse tissue with 1X PBS to remove excess blood, homogenized in 20 mL of 1X PBS and stored overnight at  $\leq -20$  °C. After two freeze-thaw cycles were performed to break the cell membranes, the homogenates were centrifuged for 5 minutes at 5000 x g. The supernate was removed immediately and assayed. Alternatively, aliquot and store samples at  $\leq -20$  °C.

**Note:** Some lysis buffer, such as RIPA can not be used. Some components will affect the binding.

**Urine**: Urinary samples should be cleared by centrifugation and then can be used directly without dilution. Storage at -20°C.

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# 2. Human CD120b Standard Preparation

Reconstitute the lyophilized Human CD120b Standard by adding 1 ml of Standard/Sample Diluent to make the 10,000 pg/ml standard stock solution. Allow solution to sit at room temperature for 5 minutes, then gently vortex to mix completely. Use within one hour of reconstituting. Two tubes of the standard (10 ng per tube) are included in each kit. Use one tube for each experiment. Perform 2-fold serial dilutions of the top standards to make the standard curve within the range of this assay (7.8 pg/ml - 500 pg/ml) as below. Standard/Sample Dilution Buffer serves as the zero standard (0 pg/ml).

Standard	Add	Into
500 pg/ml	50 $\mu l$ of the Standard (10,000 pg/ml)	950 $\mu l$ of the Standard/Sample Diluent
250 pg/ml	500 $\mu l$ of the Standard (500 pg/ml)	500 $\mu l$ of the Standard/Sample Diluent
125 pg/ml	500 $\mu l$ of the Standard (250 pg/ml)	500 $\mu l$ of the Standard/Sample Diluent
62.5 pg/ml	500 $\mu l$ of the Standard (125 pg/ml)	500 $\mu l$ of the Standard/Sample Diluent
31.25 pg/ml	500 $\mu l$ of the Standard (62.5 pg/ml)	500 $\mu l$ of the Standard/Sample Diluent
15.625 pg/ml	500 $\mu l$ of the Standard (31.25 pg/ml)	500 $\mu l$ of the Standard/Sample Diluent
7.8125 pg/ml	500 $\mu l$ of the Standard (15.625 pg/ml)	500 $\mu l$ of the Standard/Sample Diluent
0 pg/ml	1 ml of the Standard/Sample Diluent	

**Note:** The standard solutions are best used within 2 hours. The 10,000 pg/ml standard solution should be stored at 4°C for up to 12 hours, or at -20°C for up to 48 hours. Avoid repeated freeze-thaw cycles.

Biotin-Labeled Detection Antibody Working Solution Preparation
 The Biotin-Labeled Detection Antibody should be diluted in 1:100 with the Detection
 Antibody Diluent and mixed thoroughly. The solution should be prepared no more
 than 2 hours prior to the experiment.

4. Streptavidin-HRP Working Solution Preparation



The Streptavidin-HRP should be diluted in 1:100 with the Streptavidin-HRP Diluent and mixed thoroughly. The solution should be prepared no more than 1 hour prior to the experiment.

5. Wash Buffer Working Solution Preparation

Pour entire contents (30 ml) of the Wash Buffer Concentrate into a clean 1,000 ml graduated cylinder. Bring final volume to 600 ml with glass-distilled or deionized water (1:20).



#### VIII. ASSAY PROCEDURE

The Streptavidin-HRP Working Solution and TMB Substrate Solution must be kept warm at 37°C for 30 minutes before use. When diluting samples and reagents, they must be mixed completely and evenly. Standard detection curve should be prepared for each experiment. The user will decide sample dilution fold by crude estimation of protein amount in samples.

1. Add 100  $\mu l$  of each standard and sample into appropriate wells.

2. Cover well and incubate for 90 minutes at room temperature or over night at 4°C with gentle shaking.

3. Remove the cover, discard the solution and wash plate 3 times with Wash Buffer Working Solution, and each time let Wash Buffer Working Solution stay in the wells for 1 - 2 minutes. Blot the plate onto paper towels or other absorbent material. Do NOT let the wells completely dry at any time.

4. Add 100  $\mu$ l of Biotin-Labeled Detection Antibody Working Solution into each well and incubate the plate at 37°C for 60 minutes.

5. Wash plate 3 times with Wash Buffer Working Solution, and each time let Wash Buffer Working Solution stay in the wells for 1 - 2 minutes. Discard the Wash Buffer Working Solution and blot the plate onto paper towels or other absorbent material.
6. Add 100 μl of Streptavidin-HRP Working Solution into each well and incubate the plate at 37°C for 45 minutes.

7. Wash plate 5 times with Wash Buffer Working Solution, and each time let wash buffer stay in the wells for 1 - 2 minutes. Discard the wash buffer and blot the plate onto paper towels or other absorbent material.

8. Add 100  $\mu l$  of TMB Substrate Solution into each well and incubate plate at 37°C in dark for 10-20 minutes.

9. Add 100  $\mu l$  of Stop Solution into each well. The color changes into yellow immediately.



10. Read the O.D. absorbance at 450nm in a microplate reader within 30 minutes after adding the Stop Solution.

For calculation, (the relative O.D.450) = (the O.D.450 of each well) - (the O.D.450 of Zero well). The standard curve can be plotted as the relative O.D.450 of each standard solution (Y) vs. the respective concentration of the standard solution (X). The concentration of the samples can be interpolated from the standard curve.

**Note:** If the samples measured were diluted, multiply the dilution factor to the concentrations from interpolation to obtain the concentration before dilution.



# IX. ASSAY PROCEDURE SUMMARY



- Add 100 μl Standard or Sample
- Wash plate 3 times with Wash Buffer Working Solution
- Add 100 µl Biotin-Labeled Detection Antibody Working Solution
- Wash plate 3 times with Wash Buffer Working Solution
- Add 100 μl Streptavidin-HRP Working Solution
- Wash plate 5 times with Wash Buffer Working Solution
- Add 100 μl TMB Substrate Solution
- Add 100 µl Stop Solution
- Read the plate at 450nm



# X. TYPICAL DATA

each assay. y = -9E-06x<sup>2</sup> + 0.0092x + 0.0753 2.5  $R^2 = 0.9976$ 2 Absorbance (450nm) 1.5 1 0.5 0 100 200 300 400 500 600 0 Human CD120b Concentration (pg/ml)

The standard curve is for demonstration only. A standard curve must be run with

#### XI. SENSITIVITY

The minimum detectable dose of Human CD120b is typically less than 4 pg/ml.

#### XII. SPECIFICITY

The Human CD120b ELISA Kit allows for the detection and quantification of endogenous levels of natural and/or recombinant Human CD120b proteins within the range of 7.8 pg/ml - 500 pg/ml.



# XIII. CROSS REACTIVITY

No detectable cross-reactivity with other relevant proteins.

#### XIV. REFERENCES

 Santee, S. M.; Owen-Schaub, L. B. Human tumor necrosis factor receptor p75/80 (CD120b) gene structure and promoter characterization. J. Biol. Chem. 271: 21151-21159, 1996.

2. Beltinger, C. P.; White, P. S.; Maris, J. M.; Sulman, E. P.; Jensen, S. J.; LePaslier, D.; Stallard, B. J.; Goeddel, D. V.; de Sauvage, F. J.; Brodeur, G. M. Physical mapping and genomic structure of the human TNFR2 gene. Genomics 35: 94-100, 1996.



# XV. TROUBLESHOOTING GUIDE

Problem	Possible Cause	Solution
High signal and background in	<ul> <li>Insufficient washing</li> </ul>	<ul> <li>Increase number of washes</li> </ul>
all wells		<ul> <li>Increase time of soaking</li> </ul>
		between in wash
	<ul> <li>Too much Streptavidin-HRP</li> </ul>	Check dilution, titration
	<ul> <li>Incubation time too long</li> </ul>	<ul> <li>Reduce incubation time</li> </ul>
	Development time too long	<ul> <li>Decrease the incubation</li> </ul>
		time before the stop solution
		is added
No signal	<ul> <li>Reagent added in incorrect</li> </ul>	<ul> <li>Review protocol</li> </ul>
	order, or incorrectly prepared	
	<ul> <li>Standard has gone bad (If</li> </ul>	<ul> <li>Check the condition of</li> </ul>
	there is a signal in the sample	stored standard
	wells)	
	<ul> <li>Assay was conducted from an</li> </ul>	<ul> <li>Reagents allows to come to</li> </ul>
	incorrect starting point	20 - 30 °C before performing
		assay
Too much signal-whole plate	<ul> <li>Insufficient washing-unbound</li> </ul>	<ul> <li>Increase number of washes</li> </ul>
turned uniformly blue	Streptavidin-HRP remaining	Carefully
	<ul> <li>Too much Streptavidin-HRP</li> </ul>	Check dilution
	<ul> <li>Plate sealer or reservoir</li> </ul>	<ul> <li>Use fresh plate sealer and</li> </ul>
	reused, resulting in presence of	reagent reservoir for each
	residual Streptavidin-HRP	step
Standard curve achieved but	<ul> <li>Plate not developed long</li> </ul>	<ul> <li>Increase substrate solution</li> </ul>
poor discrimination between	enough	incubation time
point	<ul> <li>Improper calculation of</li> </ul>	<ul> <li>Check dilution, make new</li> </ul>
	standard curve dilution	standard curve
No signal when a signal is	<ul> <li>Sample matrix is masking</li> </ul>	<ul> <li>More diluted sample</li> </ul>
expected, but standard curve	detection	Recommended
looks fine		
Samples are reading too high,	<ul> <li>Samples contain protein levels</li> </ul>	<ul> <li>Dilute samples and run</li> </ul>
but standard curve is fine	above assay range	Again
Edge effect	<ul> <li>Uneven temperature around</li> </ul>	<ul> <li>Avoid incubating plate in</li> </ul>
	work surface	areas where environmental
		conditions vary
		<ul> <li>Use plate sealer</li> </ul>



# XVI. TECHNICAL SUPPORT

For troubleshooting, information or assistance, please go online to www.cohesionbio.com or contact us at techsupport@cohesionbio.com

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XVII. NOTES