

Human c-Met/HGFR ELISA Kit (hcMet-ELISA)

Cat. No. EK0744

96 Tests in 8 x 12 divisible strips

Background

c-Met (MET or MNNG HOS Transforming gene) is a proto-oncogene that encodes a protein known as hepatocyte growth factor receptor (HGFR). MET proto-oncogene has a total length of 125,982 bp, and is located on the 7q31 locus of chromosome 7. MET is a membrane receptor that is essential for embryonic development and wound healing. Activation of MET triggers mitogenesis and morphogenesis.

ScienCell's human c-Met ELISA kit is based on standard sandwich enzyme-linked immune-sorbent assay technology. Human c-Met-specific polyclonal antibodies are pre-coated onto 8 x 12 divisible strips. The human-specific detection polyclonal antibodies are biotinylated. The test samples and biotinylated detection antibodies are subsequently added to the wells and then washed with PBS or TBS buffer. Avidin-Biotin-Peroxidase Complex is added and unbound conjugates are washed away with PBS or TBS buffer. HRP substrate TMB is used to visualize HRP enzymatic reaction. TMB is catalyzed by HRP to produce a blue color product that changes to yellow after adding acidic stop solution. The intensity of yellow is proportional to the amount of human c-Met in the sample that is captured on the strips.

Size 96 Tests in 8 x 12 divisible strips

Assay type Sandwich ELISA

Range 62.5 pg/ml-4000 pg/ml

Sensitivity < 5 pg/ml

Specificity No detectable cross-reactivity with any other cytokine.

Storage Store at 4°C for frequent use, at -20°C for infrequent use.

Avoid multiple freeze-thaw cycles.

Shipping Shipped on gel ice.

Expiration Four months at 4° C and eight months at -20° C.

Application For quantitative detection of human c-Met in cell culture supernatants, serum and plasma (heparin,

EDTA).

Kit components 1. Lyophilized recombinant human c-Met standard: 10ng/tube×2.

2. 8 x 12 divisible strips pre-coated with anti- human c-Met antibody.

3. Sample diluent buffer: 30 ml

- 4. Biotinylated anti-human c-Met antibody: 130µl, dilution 1:100.
- 5. Antibody diluent buffer: 12ml.
- 6. Avidin-Biotin-Peroxidase Complex (ABC): 130µl, dilution 1:100.
- 7. TMB color developing agent: 10ml.
- 8. TMB stop solution: 10ml.

Materials

1. Microplate reader.

Required But

2. Automated plate washer.

Not Provided

- 3. Adjustable pipettes and pipette tips. Multichannel pipettes are recommended for large amount of samples.
- 4. Clean tubes and Eppendorf tubes.
- 5. Washing buffer (neutral PBS or TBS).

Preparation of 0.01M TBS: Add 1.2g Tris, 8.5g NaCl; 450µl of purified acetic acid or 700µl of concentrated hydrochloric acid to 1000ml H2O and adjust pH to 7.2-7.6. Finally, adjust the total volume to 1L.

Preparation of 0.01 M PBS: Add 8.5g NaCl, 1.4g Na2HPO4 and 0.2g NaH2PO4 to 1000ml distilled water and adjust pH to 7.2-7.6. Finally, adjust the total volume to 1L.

Usage

This product is for research use only. It is not approved for use in humans, animals, or *in vitro* diagnostic procedures.

Reference

- 1. Bottaro DP, Rubin JS, Faletto DL, Chan AM, Kmiecik TE, Vande Woude GF, Aaronson SA (February 1991). "Identification of the hepatocyte growth factor receptor as the met proto-oncogene product". Science 251 (4995): 802–4.
- 2. Galland F, Stefanova M, Lafage M, Birnbaum D (1992). "Localization of the 5' end of the MCF2 oncogene to human chromosome 15q15----q23". Cytogenet. Cell Genet. 60 (2): 114–6.
- 3. Gentile A, Trusolino L, Comoglio PM (March 2008). "The Met tyrosine kinase receptor in development and cancer". Cancer Metastasis Rev. 27 (1): 85–94.

Protocol for Human c-Met ELISA (96-well format)

Notes before you begin

- 1. To inspect the validity of experiment operation and the appropriateness of sample dilution proportion, a pilot experiment using standards and a small number of samples is recommended.
- 2. The TMB Color developing agent should be colorless and transparent before using.
- 3. Before using the kit, spin tubes and bring down all components to the bottom of tubes.
- 4. A duplicate well assay is recommended for both standard and samples.
- 5. Do not let wells dry, as this will inactivate active components in wells.
- 6. Do not reuse tips and tubes to avoid cross contamination.
- 7. Avoid using reagents from different batches.
- 8. In order to avoid marginal effect of plate incubation due to temperature difference (reaction may be stronger in the marginal wells), it is suggested that the diluted ABC and TMB solution be pre-warmed in 37°C for 30 minutes before use.

Preparation

Sample Preparation and Storage

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 supernatants**: Remove particulates by centrifugation, assay immediately or aliquot and store 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, 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.

Sample Dilution Guideline

The user needs to estimate the concentration of the target protein in the sample and select a proper dilution factor so that the diluted target protein concentration falls near the middle of the linear regime in the standard curve. Dilute the sample using the provided diluent buffer. The following is a guideline for sample dilution. Several trials may be necessary in practice. **The sample must be mixed well with the diluent buffer**.

- **High target protein concentration (40-400 ng/ml)**. The working dilution is 1:100. i.e. Add 3 μl sample into 297 μl sample diluent buffer.
- **Medium target protein concentration (4-40 ng/ml)**. The working dilution is 1:10. i.e. Add 25 μl sample into 225 μl sample diluent buffer.
- Low target protein concentration (62.5-4000 pg/ml). The working dilution is 1:2. i.e. Add 120 μl sample to 120 μl sample diluent buffer.
- Very Low target protein concentration (\leq 62.5 pg/ml). No dilution necessary, or the working dilution is 1:2.

Reagent Preparation and Storage

- A. Reconstitution of the human C-MET standard: C-MET standard solution should be prepared no more than 2 hours prior to the experiment. Two tubes of C-MET standard (10 ng per tube) are included in each kit. Use one tube for each experiment.
 - 10,000 pg/ml of human C-MET standard solution: Add 1 ml sample diluent buffer into one tube, keep the tube at room temperature for 10 minutes and mix thoroughly.
 - 4000pg/ml of human C-MET standard solution: Add 0.4 ml of the above 10ng/ml C-MET standard solution into 0.6 ml sample diluent buffer and mix thoroughly.
 - 2000pg/ml→62.5pg/ml of human C-MET standard solutions: Label 6 Eppendorf tubes with 2000pg/ml, 1000pg/ml, 500pg/ml, 250pg/ml, 125pg/ml, 62.5pg/ml respectively. Aliquot 0.3ml of the sample diluent buffer into each tube. Add 0.3ml of the above 4000pg/ml C-MET standard solution into 1st tube and mix. Transfer 0.3 ml from 1st tube to 2nd tube and mix. Transfer 0.3ml from 2nd tube to 3rd tube and mix, and so on.

Note: The standard solutions are best used within 2 hours. The 10 ng/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.

- B. Preparation of biotinylated anti-human C-MET antibody working solution: The solution should be prepared no more than 2 hours prior to the experiment.
 - The total volume should be: 0.1ml/well x (the number of wells). (Allowing 0.1-0.2 ml more than total volume).
 - Biotinylated anti-human C-MET antibody should be diluted in 1:100 with the antibody diluent buffer and mixed thoroughly.
- C. Preparation of Avidin-Biotin-Peroxidase Complex (ABC) working solution: The solution should be prepared no more than 1 hour prior to the experiment.

- The total volume should be: 0.1ml/well x (the number of wells). (Allowing 0.1-0.2 ml more than total volume).
- Avidin-Biotin-Peroxidase Complex (ABC) should be diluted in 1:100 with the ABC dilution buffer and mixed thoroughly.

Assay Procedure

The ABC working solution and TMB color developing agent 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 C-MET detection curve should be prepared for each experiment. The user will decide sample dilution fold by crude estimation of C-MET amount in samples.

- 1. Aliquot 0.1ml per well of the 4000pg/ml, 2000pg/ml, 1000pg/ml, 500pg/ml, 250pg/ml, 125pg/ml, 62.5pg/ml human C-MET standard solutions into the pre-coated 8 x 12 divisible strips. Add 0.1ml of the sample diluent buffer into the control well (**blank well**). Add 0.1ml of each properly diluted sample of human cell culture supernatants, serum or plasma (heparin, EDTA) to each empty well. See "**Sample Dilution Guideline**" above for details. We recommend that each human C-MET standard solution and each sample is measured in duplicate.
- 2. Seal the strips with the cover and incubate at 37°C for 90 minutes.
- 3. Remove the cover, discard the strips' contents, and blot the strips onto paper towels or other absorbent material. **Do NOT** let the wells completely dry at any time.
- 4. Add 0.1ml of biotinylated anti-human C-MET antibody working solution into each well and incubate the strips at 37°C for 60 minutes.
- 5. Wash the strips 3 times with 0.01M TBS or 0.01M PBS, and each time let washing buffer stay in the wells for 1 minute. Discard the washing buffer and blot the strips onto paper towels or other absorbent material. (**Strips Washing Method**: Discard the solution in the wells without touching the side walls. Blot the strips onto paper towels or other absorbent material. Soak each well with at least 0.3 ml PBS or TBS buffer for 1~2 minutes. Repeat this process two additional times for a total of THREE washes. Note: For automated washing, aspirate all wells and wash THREE times with PBS or TBS buffer, overfilling wells with PBS or TBS buffer. Blot the strips onto paper towels or other absorbent material).
- 6. Add 0.1ml of prepared ABC working solution into each well and incubate the strips at 37°C for 30 minutes.
- 7. Wash the strips 5 times with 0.01M TBS or 0.01M PBS, and each time let washing buffer stay in the wells for 1-2 minutes. Discard the washing buffer and blot the strips onto paper towels or other absorbent material. (See Step 5 for strip washing method).
- 8. Add 90 μl of prepared TMB color developing agent into each well and incubate the strips at 37°C in dark for 25-30 minutes (**Note**: For reference only, the optimal incubation time should be determined by end user. And the shades of blue can be seen in the wells with the four most concentrated human C-MET standard solutions; the other wells show no obvious color).
- 9. Add 0.1ml of prepared TMB stop solution into each well. The color changes to yellow immediately.
- 10. Read the O.D. absorbance at 450 nm 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 blank 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 human C-MET 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.

Summary

- 1. Add samples and standards and incubate the strips at 37°C for 90 minutes. Do not wash.
- 2. Add biotinylated antibodies and incubate the strips at 37°C for 60 minutes. Wash strips 3 times with 0.01M TBS.
- 3. Add ABC working solution and incubate the strips at 37°C for 30 minutes. Wash strips 5 times with 0.01M TBS.
- 4. Add TMB color developing agent and incubate the strips at 37°C in dark for 25-30 minutes.
- 5. Add TMB stop solution and read.

Typical Data Obtained from Human c-Met/HGFR

(TMB reaction incubate at 37°C for 26 minutes)

Concentration	0.0	62.5	125	250	500	1000	2000	4000
(pg/ml)								
Absorbance	0.053	0.112	0.147	0.227	0.374	0.813	1.470	2.373
(450 nm)								

Typical Human c-Met/HGFR ELISA Kit Standard Curve

This standard curve was generated for demonstration purpose only. A standard curve must be run with each assay.

