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Email: info@biomatik.com https://www.biomatik.com

User Manual

Catalog Number: EKC31046

Product Name: Cynomologus Monkey C-Peptide ELISA Kit

Detection Range: 0.5 ng/mL-8 ng/mL

Intended Use: For quantitative determination of Cynomologus Monkey C-

Peptide concentrations in serum, plasma, tissue homogenates.

Precautions: For research use only. Not for use in diagnostic procedures.

Manual Version: 202301V1

Storage:

Unopened kit	6 months when stored at 2 - 8°C.
Opened Kit	May be stored up to 1 month at 2 - 8°C. Keep it in sealed aluminum foil bag and avoid moisture.

The product manual may be updated as a result of continuous improvements.

Always refer to the hard copy manual included in the kit for your experiment.



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Kit Components

Reagents	Quantity	
Assay plate (12 x 8 coated Microwells)	1(96 wells)	
Standard	6 x 0.5 ml	
HRP-conjugate	1 x 6 ml	
Wash Buffer (20 x concentrate)	1 x 15 ml	
Substrate A	1 x 7 ml	
Substrate B	1 x 7 ml	
Stop Solution	1 x 7 ml	
Adhesive Strip (For 96 wells)	4	
Instruction manual	1	

Working Principle

This assay adopts quantitative sandwich enzyme immunoassay technique. Antibody specific to C-Peptide has been pre-coated onto a microplate. Standards and samples are pipetted into the wells with a Horseradish Peroxidase (HRP) conjugated antibody specific to C-Peptide. Following a wash to remove any unbound reagent, a substrate solution is added to the wells and color develops in proportion to the amount of C-Peptide bound in the initial step. At last, measure the intensity of color after stopping color development.

Sensitivity

The minimum detectable dose of mouse C-Peptide is less than 0.25 ng/mL. The sensitivity of this assay, or Lower Limit of Detection (LLD) was defined as the lowest cynomologus monkey C-Peptide concentration

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that could be differentiated from zero. It was determined by the mean O.D

value of 20 replicates of the zero standard added by their three standard

deviations

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Specificity

This assay has high sensitivity and excellent specificity. No significant

cross-reactivity or interference between Cynomologus Monkey C-Peptide

and analogues was observed. Limited by current skills and knowledge, it

is impossible for us to complete the cross-reactivity detection between

Cynomologus Monkey C-Peptide and all the analogues, therefore, cross

reaction may still exist.

Precision

Intra-assay Precision (Precision within an assay): CV%<15%

Three samples of known concentration were tested twenty times on one

plate to assess.

Inter-assay Precision (Precision between assays): CV%<15%

Three samples of known concentration were tested in twenty assays to

assess.

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Other Supplies Required

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Microplate reader capable of measuring absorbance at 450 nm,

with the correction wavelength set at 600 nm or 630 nm.

An incubator which can provide stable incubation conditions up to

37°C±0.5°C.

• Squirt bottle, manifold dispenser, or automated microplate washer.

Absorbent paper for blotting the microtiter plate.

100ml and 500ml graduated cylinders.

Deionized or distilled water.

Pipettes and pipette tips.

Test tubes for dilution.

Sample Collection & Storage

• Serum Use a serum separator tube (SST) and allow samples to

clot for two hours at room temperature or overnight at 4°C before

centrifugation at 1000 ×g for 15 mins. Remove serum and assay

immediately or aliquot and store samples at -20°C or -80°C. Avoid

repeated freeze-thaw cycles.





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Plasma Collect plasma using EDTA or heparin as an anticoagulant.

Within 30 mins after collecting samples, centrifuge samples at 1000 x g, 2

- 8°C, for 15 mins. Assay immediately or aliquot and store samples at -

20°C or -80°C. Avoid repeated freeze-thaw cycles.

Tissue Homogenates Rinse 100mg tissue with 1X PBS,

homogenize it in 1 ml of 1X PBS and store it overnight at -20°C. The cell

membranes break after two freeze-thaw cycles, then centrifuge

homogenates at 5000 x g, 2 - 8°C for 5 mins. Take the supernate for

assay. Alternatively, aliquot and store the supernate at -20°C or -80°C.

Centrifuge samples again after thawing. Avoid repeated freeze-thaw

cycles.

Notes:

1. Biomatik is only responsible for the kit itself, not for the samples

consumed during the assay. The user need to calculate the possible

amount of the samples to be used in the whole test. Please reserve

sufficient samples in advance.

2. Samples to be used within 5 days may be stored at 2-8°C, otherwise,

samples must be stored at -20°C (≤1month) or -80°C (≤2month) to avoid

contamination and loss of bioactivity.

3. Grossly hemolyzed samples are not suitable to use.

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4. It would be necessary to run a preliminary experiment for validation, if the samples are not indicated in the manual.

5. Please predict the concentration before assaving. If results were not within the range of the standard curve, users would need to estimate the

optimal sample dilutions for their particular experiments.

6. Tissue or cell extraction samples prepared by chemical lysis buffer may cause unexpected ELISA results due to the impacts of certain

chemicals.

7. Considering the possibility of mismatch between antigen from other resources and antibody in our kits (e.g., antibody targets conformational epitope rather than linear epitope), some native or recombinant proteins from other manufacturers may not be recognized by our products.

8. Due to factors including cell viability, cell number and sampling time.

samples from cell culture supernatant may not be detected by the kit.

9. Recommend to use fresh samples for the test. If you store samples for long time, protein degradation and denaturalization may occur in those

samples and finally lead to wrong results.

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Reagents Preparation

- Kindly use graduated containers to prepare the reagent.
 Please don't prepare the reagent directly in the Diluent vials in the kit.
- Bring all reagents to room temperature (18-25°C) before use for 30 mins.
- Distilled Water is recommended. Contaminated water or container for reagents preparation will affect the test result.

Wash Buffer(1x)- If crystals have formed in the concentrate, warm up to room temperature and mix gently until the crystals have completely dissolved. Dilute 15 ml of Wash Buffer Concentrate (20 x) into deionized or distilled water to prepare 300 ml of Wash Buffer (1 x)

Standard Concentration

Standard	S0	S1	S2	S3	S4	S5
Concentration	0	0.5	1	2	4	8
(ng/ml)						

Key Notes

 Do not mix or substitute reagents with those from other lots or sources.

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If samples generate higher values than the highest standard, dilute

the samples with Sample Diluent and repeat the assay.

Any variation in Sample Diluent, operator, pipetting technique,

washing technique, incubation time or temperature, and kit age can

cause variation in binding.

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This assay is designed to eliminate interference by soluble

receptors, binding proteins, and other factors present in biological

samples. Until all factors have been tested in the Immunoassay, the

possibility of interference cannot be excluded.

Precautions

The Stop Solution provided with this kit is an acid solution. Wear eve. hand, face.

and clothing protection when using this material.

Assay Procedures

Bring all reagents and samples to room temperature before use.

Centrifuge the sample again after thawing before the assay. It is

recommended to assay all samples and standards in duplicate.

1. Prepare all reagents, working standards, and samples as directed in

the previous sections.

2. Refer to the Assay Layout Sheet to determine the number of wells to be

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used and put any remaining wells and the desiccant back into the pouch and

3. Set a blank well without any solution.

seal the ziploc, store unused wells at 4°C.

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4. Add 50µl of standard or sample per well. Standard test in duplicate.

5. Add 50ul of HRP-conjugate to each well (not to Blank well). Mix well and

then incubate at 37°C for one hour.

6. Aspirate each well and wash, repeating the process two times for a total of

three washes. Wash by filling each well with Wash Buffer (200ul) using a

squirt bottle, multi-channel pipette, manifold dispenser, or autowasher, and

let it stand for 10 seconds, complete removal of liquid at each step is

essential to good performance. After the last wash, remove any remaining

wash Buffer by aspirating or decanting. Invert the plate and blot it against

clean paper towels.

7. Add 50ul of **Substrate A** and 50ul of **Substrate B** to each well, mix well.

Incubate at 37°C for 15 minutes. Keeping the plate away from drafts and

other temperature fluctuations in the dark.

8.Add 50ul of Stop Solution to each well, gently tap the plate to ensure

thorough mixing.

9. Determine the optical density of each well within 10 mins, using a

microplate reader set to 450 nm.

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Notes:

The final experimental results will be closely related to validity of 1

the products, operation skills of the end users and the experimental

environments.

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2. Samples or reagents addition: Please use the freshly prepared

Standard. Please carefully add samples to wells and mix gently to avoid

foaming. Do not touch the well wall as possible. For each step in the

procedure, total dispensing time for addition of reagents or samples to

the assay plate should not exceed 10 mins. This will ensure equal

elapsed time for each pipetting step, without interruption. Duplication of

all standards and specimens, although not required, is recommended. To

avoid cross-contamination, change pipette tips between additions of each standard level, between sample additions, and between reagent additions.

Also, use separate reservoirs for each reagent.

3. Incubation: To ensure accurate results, proper adhesion of plate

sealers during incubation steps is necessary. Do not allow wells to sit

uncovered for extended periods between incubation steps. Once

reagents have been added to the well strips, DO NOT let the strips DRY

at any time during the assay. Incubation time and temperature must be

observed.





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- 4. Washing: The wash procedure is critical. Complete removal of liquid at each step is essential to good performance. After the last wash, remove any remaining Wash Solution by aspirating or decanting and remove any drop of water and fingerprint on the bottom of the plate. Insufficient washing will result in poor precision and falsely elevated absorbance reading. When using an automated plate washer, adding a 30 second soak period following the addition of wash buffer, and/or rotating the plate 180 degrees between wash steps may improve assay precision.
- 5. Controlling of reaction time: Observe the change of color after adding TMB Substrate (e.g. observation once every 10 mins), TMB Substrate should change from colorless or light blue to gradations of blue. If the color is too deep, add Stop Solution in advance to avoid excessively strong reaction which will result in inaccurate absorbance reading.
- TMB Substrate is easily contaminated. TMB Substrate should remain colorless or light blue until added to the plate. Please protect it from light.
- 7. Stop Solution should be added to the plate in the same order as the TMB Substrate. The color developed in the wells will turn from blue to yellow upon addition of the Stop Solution. Wells that are green in color indicate that Stop Solution has not mixed thoroughly with TMB Substrate.

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Calculation of Results

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Using the professional soft "Curve Expert" to make a standard

curve is recommended, which can be downloaded from our web.

Average the duplicate readings for each standard and sample and

subtract the average zero standard optical density.

Create a standard curve by reducing the data using computer software

capable of generating a four parameter logistic (4-PL) curve-fit. As an

alternative, construct a standard curve by plotting the mean absorbance

for each standard on the x-axis against the concentration on the y-axis

and draw a best fit curve through the points on the graph. The data may

be linearized by plotting the log of the C-Peptide concentrations versus

the log of the O.D. and the best fit line can be determined by regression

analysis. This procedure will produce an adequate but less precise fit of

the data

If samples have been diluted, the concentration read from the standard

curve must be multiplied by the dilution factor.