CHO Host Cell Protein ELISA Kit

Catalog #: ENZ-KIT128

Complete kit for the determination of Chinese Hamster Ovary (CHO) host cell protein contamination in bulk products expressed in CHO expression systems.

For the latest product information, including support documentation, visit us online:
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Please read entire booklet before proceeding with the assay.

Carefully note the handling and storage conditions of each kit component.

Please contact Enzo Life Sciences Technical Support if necessary.
DESCRIPTION

The CHO Host Cell Protein (HCP) ELISA kit is designed to quantitatively measure host cell protein contamination in bulk products expressed in Chinese Hamster Ovary (CHO) expression systems. Please read the complete kit insert before performing this assay. A series of CHO protein standards is provided to generate a standard curve for the assay and all unknown sample concentrations should be read off this standard curve. CHO standards or diluted unknown samples are pipetted into the provided 96-well plate which has been pre-coated with anti-CHO HCP antibodies to capture CHO proteins from biologics samples. Following an incubation to allow capture of the CHO protein by the antibodies on the plate, a second anti-CHO HCP antibody, conjugated with biotin, is added and incubated to allow it to bind to the captured CHO proteins. After 45 min incubation, the plate is washed and a Streptavidin-HRP (Horseradish Peroxidase) conjugate is added and incubated for 30 min. The Streptavidin-HRP conjugate will be captured by any biotin labeled antibody bound to the plate. Following a wash step to remove unbound conjugate, TMB substrate is added and is converted by the captured HRP to a colored product in proportion to the amount of HCP bound to the plate. After a short incubation to allow color development, the reaction is stopped and the intensity of the generated color is detected in a spectrophotometer plate reader capable of measuring 450nm wavelength. A standard curve will be generated from the CHO protein standards and used to calculate the concentration of CHO proteins in the unknown samples, after making suitable correction for the dilution of the sample.
SAFETY WARNINGS & PRECAUTIONS

FOR RESEARCH USE ONLY. NOT FOR USE IN DIAGNOSTIC PROCEDURES.

- As with all such products, this kit should only be used by qualified personnel who have had laboratory safety instruction. The complete insert should be read and understood before attempting to use the product.

- This kit utilizes a peroxidase-based readout system. Buffers, including other manufacturers Wash Buffers, containing sodium azide will inhibit color production from the enzyme. Make sure all buffers used for samples are azide free. Ensure that any plate washing system is rinsed well with deionized water prior to using the supplied Wash Buffer as prepared on Page 4.

- The Stop Solution is acid. The solution should not come in contact with skin or eyes. Take appropriate precautions when handling this reagent.
MATERIALS SUPPLIED

1. **Coated Clear 96 Well Plates**
   A clear plastic microtiter plate(s) with break apart strips coated with Rabbit anti-CHO HCP IgG. (1 plate)

2. **CHO Protein Standards**
   600 µL of CHO proteins at a concentration of 2430 ng/mL, sufficient for generating a standard curve from 3.3 ng/mL to 810 ng/mL. (1 tube with HCP standard)

3. **5x Dilution Buffer**
   Buffer used for dilution of antibodies and Streptavidin-HRP conjugate. The 15 mL of concentrate should be diluted to 75 mL with 60 mL of deionized or distilled water (15 mL)

4. **10x PBS-T**
   After dilution, this is used for wash solution (PBS with 0.1% Tween-20). The 30 mL of concentrate should be diluted to 300 mL with 270 mL of deionized or distilled water. (30 mL)

5. **Reporting antibody (90 µL)**
   A biotin labeled Rabbit polyclonal antibody specific for CHO cell proteins. Immediately prior to the assay, dilute the reporting antibody by adding 75 µL to 15 mL of 1x Dilution Buffer.

6. **Streptavidin-HRP Conjugate**
   A Streptavidin – Horse Radish Peroxidase conjugate in a special stabilizing solution. Immediately prior to the assay, dilute the entire 375 µL into 15 mL of 1x Dilution buffer to give a 0.1 μg/mL working stock. (4μg/ml, 420 µL/tube)

7. **TMB Substrate**
   Use directly without dilution. (15 mL)

8. **Stop Solution**
   A 1M solution of sulfuric acid. CAUSTIC. Use directly without dilution. (15 mL)

9. **Plate Sealer**
   (one)
OTHER MATERIALS NEEDED

1. Distilled or deionized water
2. Single- and multi-channel micro-pipettes with disposable tips to accurately dispense volumes 5-250 µL.
3. Plastic tubes (i.e. 1.5 mL) for sample and standard dilution.
4. Reagent reservoirs for sample addition.
5. Colorimetric 96 well microplate reader capable of reading optical density at 450nm.
6. Software for converting raw relative optical density readings from the plate reader and carrying out four parameter logistic curve (4-parameter) fitting. Contact your plate reader manufacturer for details.
PROCEDURAL NOTES

1. Allow diluted reagents and buffers to reach room temperature (18-25°C) prior to starting the assay. Once the assay has been started, all steps should be completed in sequence and without interruption. You do not want the plate to dry out in between steps as this can cause high backgrounds or erroneous results. Make sure that required reagents and buffers are ready when needed. Prior to adding to the plate, reagents should be mixed gently (not vortexed) by swirling.

2. Avoid contamination of reagents, pipette tips and wells. Use new disposable tips and reservoirs, do not return unused reagent to the stock bottles / vials and do not mix caps of stock solutions.

3. Incubation time can affect results. All wells should be handled in the same order for each step.

4. Microplate washing is important and can affect results by giving erroneous results or high backgrounds. We recommend a multichannel pipette to add 250 µL of buffer to each well across the plate, followed by a dumping out of contents (to a sink or other receptacle) with a rapid wrist motion. The plate should then be tapped firmly on a paper towel to shake out any remaining liquid. Avoid prolonged incubation is wash buffer when performing wash steps.

5. When making additions to the plate, be careful to avoid damaging the coating, for example by scratching the bottoms or the sides of the wells. One technique to avoid this is to make additions (for a right-handed person) from left to right across the plate, supporting the pipette tips on the right edge of the well with each addition and thus avoiding contact with the bottom or sides of the wells.

6. During the incubation times, the plate should be covered to minimize evaporation from the wells. This can be done with the adhesive covers provided or by stacking an empty plate on top.

7. After the last wash step and prior to adding the TMB substrate, wipe the bottom of the plate with a clean paper towel to ensure that moisture or fingerprints do not interfere with the OD reading.
8. Once the TMB substrate is added it will be converted by the captured HRP to a blue colored product. Generally we find that a 10 to 15 min incubation is sufficient for enough color development to discern differences between the standards and the reaction should be stopped at this point. Bear in mind that, given sufficient time, even a small amount HRP is capable of converting all the TMB to product and if this happens it will be difficult to discern differences between differing concentrations of HCP. Keeping OD450 values well below 2.0 will result in greatest accuracy as at high absorbance values very little light is reaching the detector and measurements are error prone. (Remember that at an OD of 1.0 only 10% of the light is being detected and at an OD of 2.0 only 1% of the light is reaching the detector).

SAMPLE PREPARATION SUGGESTIONS

For drug substance, it is recommended to use 4mg/mL or less for the HCP quantification; method qualification and ICH validation should also use the same recommended protein concentration. For in-process samples running for the first time, pre-rProtein A column fraction needs to be serially diluted 4 times using a 1:10 dilution, one or two of the dilutions will have HCP levels that fall onto the HCP standard curve. After the first test, it is sufficient to test at one dilution where the HCP readings fell in the range of 10-270ng/mL in the standard curve. For the post-rProtein A fraction, we recommend you serially dilute twice with 1:10 dilution. For any downstream samples (cation exchange and anion exchange column fraction), the original sample and a 1:10 dilution sample should be tested for the first test. For spike recovery testing in drug substance, it is recommended to use 20-30 ng/mL HCP whereas for in-process samples a 100 ng/mL spike is recommended. We also point out that the HCP ELISA may not achieve linear dilution for drug substances because of the complexity of the drug composition whereas for in-process samples, linear dilution can be expected.
ASSAY PROCEDURE

1. Use the plate layout sheet on the back page to plan sample layout on plate and also aid in proper sample and standard identification after the assay. We recommend that assays are carried out in duplicate or (preferably) triplicate in order to minimize spurious results.

2. Dilute the 10x PBS-T and 5x Dilution buffer with water to 1x-strength. Check both concentrate bottles for precipitates before proceeding and if found warm slightly in a water bath to dissolve before proceeding. The 30 mL of 10xPBS-T should be diluted to 300 mL with 270 mL water and the 15 mL of 5x Dilution Buffer should be diluted to 75 mL with 60 mL water.

3. Prepare the HCP standards by numbering seven 1.5 mL tubes and adding 1 mL of Dilution buffer to each. Cap the seventh tube….this will be the blank (0 ng/mL HCP). To tube one add 500µl of the provided 2430 ng/mL HCP stock and mix well…this will be the 810 ng/mL standard. Then serially dilute 500 µL of tube one across tubes two through six to generate the remainder of the standards. Pipette 100 µL of each standard into the plate according to the plate map.

4. Dilute your sample in 1x dilution buffer; for best accuracy perform serial dilutions over a wide range such that multiple dilutions will span the range of 10 – 800 ng/mL. For drug substance the typical testing protein concentration is 2 mg/ml; it is not recommended to test drug substance at concentrations higher than 4 mg/ml. Pipette 100 µL of samples or standards into wells in the plate. Cover plate and incubate 1.5 hr at room temperature.

5. During the above incubation, dilute the reporting antibody by adding 75 µL to 15 mL of 1x Dilution Buffer.

6. Wash plate by emptying contents and adding 250 µL of wash buffer to each well. Empty wells again and tap the plate firmly upside down on a paper towel to fully empty well. Repeat twice.

7. Pipette 100 µL of Reporting Antibody into each well. Cover plate and incubate plate 45 min at room temperature.

8. During the above incubation, dilute the 4 µg/mL Streptavidin-HRP conjugate to 0.1 µg/mL by adding the entire 375 µL to 15 mL of 1x Dilution Buffer.
9. Wash plate by emptying contents and adding 250 µL of wash buffer to each well. Empty wells again and tap the plate firmly upside down on a paper towel to fully empty well. Repeat twice.

10. Pipette 100 µL of Streptavidin-HRP conjugate into wells. Cover plate and incubate plate 30 min. at room temperature.

11. Wash plate by emptying contents and adding 250 µL of wash buffer to each well. Empty wells again and tap the plate firmly upside down on a paper towel to fully empty well. Repeat twice.

12. Add 100 µL of TMB substrate to each well. Monitor color development and stop reaction by adding 100 µL of Stop Solution to each well when color development within standards is sufficient. Generally 15-20 min time will be sufficient. Longer incubation times may increase background.

13. Read the optical density generated from each well in a plate reader capable of reading at 450nm, use three wells without sample as blank such as H1-H3.

14. Either graph the results on log graph paper or use the plate reader’s built-in 4-parameter fit software capabilities to calculate HCP concentration for each sample.

**CALCULATION OF RESULTS**

Average the triplicate OD readings for each standard, sample and background wells to give a mean OD reading. Subtract the averaged background values from the mean OD values to give a net OD value and create a standard curve using either log graph paper of 4-parameter fit software. Match OD values for the unknowns to [HCP] using the standard curve, remembering to correct for dilution:
TYPICAL RESULTS

The results shown below are for illustration only and should not be used to calculate results.

<table>
<thead>
<tr>
<th>Sample</th>
<th>Mean OD</th>
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</thead>
<tbody>
<tr>
<td>NSB (0ng/mL HCP)</td>
<td>0.243</td>
</tr>
<tr>
<td>3.3 ng/mL HCP</td>
<td>0.296</td>
</tr>
<tr>
<td>10 ng/mL HCP</td>
<td>0.368</td>
</tr>
<tr>
<td>30 ng/mL HCP</td>
<td>0.501</td>
</tr>
<tr>
<td>90 ng/mL HCP</td>
<td>0.762</td>
</tr>
<tr>
<td>270ng/mL HCP</td>
<td>1.332</td>
</tr>
<tr>
<td>810ng/mL HCP</td>
<td>2.278</td>
</tr>
</tbody>
</table>
VALIDATION SUMMARY

Validation assays were performed on the CHO HCP ELISA kit. Precision was quantified by calculating the coefficient of variation (%CV). %CV is calculated by dividing the standard deviation by the mean. Validation assays were performed using samples that fall within the high, medium, and low range of the standard curve. An intra-assay experiment was performed to assess well-to-well variation in the assay. Negligible well-to-well variation was observed in the assay. Lot-to-lot variation was also assessed. Negligible lot-to-lot variation was observed.

Intra-Assay

<table>
<thead>
<tr>
<th>Number of Tests</th>
<th>Mean (ng/mL)</th>
<th>Standard Deviation</th>
<th>%CV</th>
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<tbody>
<tr>
<td>24</td>
<td>90.68</td>
<td>5.96</td>
<td>6.6%</td>
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<tr>
<td>24</td>
<td>48.93</td>
<td>1.78</td>
<td>3.6%</td>
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<tr>
<td>24</td>
<td>30.4</td>
<td>1.21</td>
<td>4.0%</td>
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<tr>
<td>24</td>
<td>12.26</td>
<td>0.54</td>
<td>4.0%</td>
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Inter-Assay

<table>
<thead>
<tr>
<th>Standard (ng/mL)</th>
<th>Mean OD Values (Lot 1)</th>
<th>Mean OD Values (Lot 2)</th>
<th>Mean OD Values (Lot 3)</th>
<th>Overall Mean</th>
<th>Standard Deviation</th>
<th>%CV</th>
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<tbody>
<tr>
<td>90</td>
<td>2.22</td>
<td>2.22</td>
<td>2.44</td>
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<tr>
<td>30</td>
<td>1.10</td>
<td>1.03</td>
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<td>10</td>
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### Plate Template

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