ab108803 – Apolipoprotein AI (APOA1) Human ELISA Kit

Instructions for Use

For the quantitative measurement of Human Apolipoprotein AI (APOA1) in plasma, serum, urine, saliva, milk, CSF and cell culture samples.

This product is for research use only and is not intended for diagnostic use.
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INTRODUCTION

1. BACKGROUND

Abcam’s Apolipoprotein AI (APOA1) Human in vitro ELISA (Enzyme-Linked Immunosorbent Assay) kit is designed for the quantitative measurement of apolipoprotein AI concentrations in plasma, serum, urine, saliva, milk, CSF and cell culture samples.

An Apolipoprotein AI specific antibody has been precoated onto 96-well plates and blocked. Standards or test samples are added to the wells and subsequently an Apolipoprotein AI specific biotinylated detection antibody is added and then followed by washing with wash buffer. Streptavidin-Peroxidase Complex is added and unbound conjugates are washed away with wash buffer. TMB is then used to visualize Streptavidin-Peroxidase enzymatic reaction. TMB is catalyzed by Streptavidin-Peroxidase to produce a blue color product that changes into yellow after adding acidic stop solution. The density of yellow coloration is directly proportional to the amount of Apolipoprotein AI captured in plate.

Human apolipoprotein AI comprises about 70% of the high-density lipoproteins (HDL) protein mass and apolipoprotein AII another 15–20%. APOA1, a 243-amino acid molecule that contains a series of highly homologous amphipathic alpha-helices, is a 28-kDa single polypeptide that lacks glycosylation or disulfide linkages. About 5–10% of Human plasma apolipoprotein AI exists in a lipoprotein-unassociated state. Apolipoprotein AI appears to have effects on reverse cholesterol transport, and anti-inflammation. Oxidation of specific amino acid residues in apolipoprotein AI may impair cholesterol efflux from macrophages. A majority of HDL functionality is derived from the ability of apolipoprotein AI to sequester phospholipids and cholesterol and interact with plasma enzymes and cellular receptors. During reverse cholesterol transport, HDL interacts with lecithin: cholesteryl acyltransferase (LCAT) and cellular receptors, including ATP-binding cassette transporter protein AI (ABCA1) and the scavenger receptor class B type I in an ordered fashion that is reflected by HDL particle lipid composition. A high-affinity HDL receptor for apolipoprotein AI is
INTRODUCTION

beta-chain of ATP synthase on the surface of hepatocytes. The plasma concentration of apolipoprotein AI is one of the best indicators of susceptibility to cardiovascular disease.
2. ASSAY SUMMARY

Primary capture antibody
Prepare all reagents, samples and standards as instructed.

Sample
Add standard or sample to each well used. Incubate at room temperature.

Primary detector antibody
Wash and add prepared biotin antibody to each well. Incubate at room temperature.

Streptavidin Label
Wash and add prepared Streptavidin-Peroxidase Conjugate. Incubate at room temperature.

Colored product
Add Chromogen Substrate to each well. Incubate at room temperature. Add Stop Solution to each well. Read immediately.
3. **PRECAUTIONS**

Please read these instructions carefully prior to beginning the assay.

Modifications to the kit components or procedures may result in loss of performance.

4. **STORAGE AND STABILITY**

Store kit at 4°C immediately upon receipt, apart from the SP Conjugate & Biotinylated Antibody, which should be stored at -20°C.

Refer to list of materials supplied for storage conditions of individual components. Observe the storage conditions for individual prepared components in sections 9 & 10.

5. **MATERIALS SUPPLIED**

<table>
<thead>
<tr>
<th>Item</th>
<th>Amount</th>
<th>Storage Condition (Before Preparation)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Apolipoprotein AI Microplate (12 x 8 well strips)</td>
<td>96 wells</td>
<td>4°C</td>
</tr>
<tr>
<td>Apolipoprotein AI Standard</td>
<td>1 vial</td>
<td>4°C</td>
</tr>
<tr>
<td>10X Diluent N Concentrate</td>
<td>30 mL</td>
<td>4°C</td>
</tr>
<tr>
<td>Biotinylated Human Apolipoprotein AI Antibody</td>
<td>1 vial</td>
<td>-20°C</td>
</tr>
<tr>
<td>100X Streptavidin-Peroxidase Conjugate (SP Conjugate)</td>
<td>80 µL</td>
<td>-20°C</td>
</tr>
<tr>
<td>Chromogen Substrate</td>
<td>8 mL</td>
<td>4°C</td>
</tr>
<tr>
<td>Stop Solution</td>
<td>12 mL</td>
<td>4°C</td>
</tr>
<tr>
<td>20X Wash Buffer Concentrate</td>
<td>2 x 30 mL</td>
<td>4°C</td>
</tr>
<tr>
<td>Sealing Tapes</td>
<td>3</td>
<td>N/A</td>
</tr>
</tbody>
</table>
6. MATERIALS REQUIRED, NOT SUPPLIED

These materials are not included in the kit, but will be required to successfully utilize this assay:

- 1 Microplate reader capable of measuring absorbance at 450 nm.
- Precision pipettes to deliver 1 µL to 1 mL volumes.
- Adjustable 1-25 mL pipettes for reagent preparation.
- 100 mL and 1 liter graduated cylinders.
- Absorbent paper.
- Distilled or deionized water.
- Log-log graph paper or computer and software for ELISA data analysis.
- 9 tubes to prepare standard or sample dilutions.

7. LIMITATIONS

- Do not mix or substitute reagents or materials from other kit lots or vendors.
8. TECHNICAL HINTS

- Samples generating values higher than the highest standard should be further diluted in the appropriate sample dilution buffers.
- Avoid foaming or bubbles when mixing or reconstituting components.
- Avoid cross contamination of samples or reagents by changing tips between sample, standard and reagent additions.
- Ensure plates are properly sealed or covered during incubation steps.
- Complete removal of all solutions and buffers during wash steps.
- This kit is sold based on number of tests. A ‘test’ simply refers to a single assay well. The number of wells that contain sample, control or standard will vary by product. Review the protocol completely to confirm this kit meets your requirements. Please contact our Technical Support staff with any questions.
9. REAGENT PREPARATION

Equilibrate all reagents to room temperature (18-25°C) prior to use. Prepare fresh reagents immediately prior to use. If crystals have formed in the concentrate, mix gently until the crystals have completely dissolved.

9.1 1X Diluent N

Dilute the 10X Diluent N Concentrate 1:10 with reagent grade water. Mix gently and thoroughly. Store for up to 1 month at 4°C.

9.2 1X Wash Buffer

Dilute the 20X Wash Buffer Concentrate 1:20 with reagent grade water. Mix gently and thoroughly.

9.3 1X Biotinylated Apolipoprotein Al Detector Antibody

9.3.1 The stock Biotinylated Apolipoprotein Al Antibody must be diluted with 1X Diluent N according to the label concentration to prepare 1X Biotinylated Apolipoprotein Al Antibody for use in the assay procedure. Observe the label for the “X” concentration on the vial of Biotinylated Apolipoprotein Al Antibody.

9.3.2 Calculate the necessary amount of 1X Diluent N to dilute the Biotinylated Apolipoprotein Al Antibody to prepare a 1X Biotinylated Apolipoprotein Al Antibody solution for use in the assay procedure according to how many wells you wish to use and the following calculation:

<table>
<thead>
<tr>
<th>Number of Wells Strips</th>
<th>Number of Wells</th>
<th>((V_T) ) Total Volume of 1X Biotinylated Antibody (µL)</th>
</tr>
</thead>
<tbody>
<tr>
<td>4</td>
<td>32</td>
<td>1,760</td>
</tr>
<tr>
<td>6</td>
<td>48</td>
<td>2,640</td>
</tr>
<tr>
<td>8</td>
<td>64</td>
<td>3,520</td>
</tr>
<tr>
<td>10</td>
<td>80</td>
<td>4,400</td>
</tr>
<tr>
<td>12</td>
<td>96</td>
<td>5,280</td>
</tr>
</tbody>
</table>

Any remaining solution should be frozen at -20°C.
ASSAY PREPARATION

Where:

\( C_S \) = Starting concentration \((X)\) of stock Biotinylated Apolipoprotein Al Antibody (variable)

\( C_F \) = Final concentration (always = 1X) of 1X Biotinylated Apolipoprotein Al Antibody solution for the assay procedure

\( V_T \) = Total required volume of 1X Biotinylated Apolipoprotein Al Antibody solution for the assay procedure

\( V_A \) = Total volume of \((X)\) stock Biotinylated Apolipoprotein Al Antibody

\( V_D \) = Total volume of 1X Diluent N required to dilute \((X)\) stock Biotinylated Apolipoprotein Al Antibody to prepare 1X Biotinylated Apolipoprotein Al solution for assay procedures

Calculate the volume of \((X)\) stock Biotinylated Antibody required for the given number of desired wells:

\[
\frac{C_F}{C_S} \times V_T = V_A
\]

Calculate the final volume of 1X Diluent N required to prepare the 1X Biotinylated Apolipoprotein Al Antibody:

\[
V_T - V_A = V_D
\]

Example:

NOTE: This example is for demonstration purposes only. Please remember to check your antibody vial for the actual concentration of antibody provided.

\( C_S \) = 50X Biotinylated Apolipoprotein Al Antibody stock

\( C_F \) = 1X Biotinylated Apolipoprotein Al Antibody solution for use in the assay procedure

\( V_T \) = 3,520 µL (8 well strips or 64 wells)

\[
(1X/50X) \times 3,520 \mu L = 70.4 \mu L
\]

\[
3,520 \mu L - 70.4 \mu L = 3,449.6 \mu L
\]

\( V_A \) = 70.4 µL total volume of \((X)\) stock Biotinylated Apolipoprotein Al Antibody required

\( V_D \) = 3,449.6 µL total volume of 1X Diluent N required to dilute the 50X stock Biotinylated Antibody to prepare 1X Biotinylated Apolipoprotein Al Antibody solution for assay procedures

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9.3.3 First spin the Biotinylated Apolipoprotein AI Antibody vial to collect the contents at the bottom.

9.3.4 Add calculated amount $V_A$ of stock Biotinylated Apolipoprotein AI Antibody to the calculated amount $V_D$ of 1X Diluent N. Mix gently and thoroughly.

9.4 **1X SP Conjugate**

Spin down the 100X Streptavidin-Peroxidase Conjugate (SP Conjugate) briefly and dilute the desired amount of the conjugate 1:100 with 1X Diluent N.

*Any remaining solution should be frozen at -20°C.*
10. STANDARD PREPARATIONS

- Prepare serially diluted standards immediately prior to use. Always prepare a fresh set of standards for every use.
- Any remaining standard should be stored at -20°C after reconstitution and used within 30 days.
- This procedure prepares sufficient standard dilutions for duplicate wells.

10.1 Reconstitution of the Apolipoprotein AI Standard vial to prepare a 100 ng/mL Apolipoprotein AI Stock Standard:

10.1.1 First consult the Apolipoprotein AI Standard vial to determine the mass of protein in the vial.

10.1.2 Calculate the appropriate volume of 1X Diluent N to add when resuspending the Apolipoprotein AI Standard vial to produce a 100 ng/mL Apolipoprotein AI Stock Standard by using the following equation:

\[
C_S = \text{Starting mass of Apolipoprotein AI Standard (see vial label) (µg)}
\]

\[
C_F = 100 \text{ ng/mL Apolipoprotein AI Stock Standard final required concentration}
\]

\[
V_D = \text{Required volume of 1X Diluent N for reconstitution (µL)}
\]

Calculate total required volume 1X Diluent N for resuspension:

\[
(C_S / C_F) \times 1,000 = V_D
\]
Example:

NOTE: This example is for demonstration purposes only. Please remember to check your standard vial for the actual amount of standard provided.

\[ C_S = 3,200 \text{ ng of Apolipoprotein AI Standard in vial} \]
\[ C_F = 400 \text{ ng/mL Apolipoprotein AI Stock Standard final concentration} \]
\[ V_D = \text{Required volume of 1X Diluent N for reconstitution} \]
\[(3,200 \text{ ng} / 400 \text{ ng/mL}) \times 1,000 = 8,000 \mu L\]

10.1.3 First briefly spin the Apolipoprotein AI Standard Vial to collect the contents on the bottom of the tube.

10.1.4 Reconstitute the Apolipoprotein AI Standard vial by adding the appropriate calculated amount \( V_D \) of 1X Diluent N to the vial to generate the 400 ng/mL Apolipoprotein AI Stock Standard. Mix gently and thoroughly.

10.2 Allow the reconstituted 100 ng/mL Apolipoprotein AI Stock Standard to sit for 10 minutes with gentle agitation prior to making subsequent dilutions.

10.3 Label eight tubes #1-8.

10.4 Prepare the 100 ng/mL Standard #1 by adding 500 µL of the reconstituted 100 ng/mL Apolipoprotein AI Stock Standard to tube #1.

10.5 Add 120 µL of 1X Diluent N to tube #2 – 8.

10.6 To prepare Standard #2, add 120 µL of the Standard #1 into tube #2 and mix gently.

10.7 To prepare Standard #3, add 120 µL of the Standard #2 into tube #3 and mix gently.
10.8 Using the table below as a guide, prepare subsequent serial dilutions.

10.9 1X Diluent N serves as the zero standard, 0 ng/mL (tube #8).

**Standard Dilution Preparation Table**

<table>
<thead>
<tr>
<th>Standard #</th>
<th>Volume to Dilute (µL)</th>
<th>Volume Diluent N (µL)</th>
<th>Total Volume (µL)</th>
<th>Starting Conc. (ng/mL)</th>
<th>Final Conc. (ng/mL)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Step 10.4</td>
<td></td>
<td></td>
<td>100.0</td>
<td></td>
</tr>
<tr>
<td>2</td>
<td>120</td>
<td>120</td>
<td>240</td>
<td>100.0</td>
<td>50.00</td>
</tr>
<tr>
<td>3</td>
<td>120</td>
<td>120</td>
<td>240</td>
<td>50.00</td>
<td>25.00</td>
</tr>
<tr>
<td>4</td>
<td>120</td>
<td>120</td>
<td>240</td>
<td>25.00</td>
<td>12.50</td>
</tr>
<tr>
<td>5</td>
<td>120</td>
<td>120</td>
<td>240</td>
<td>12.50</td>
<td>6.250</td>
</tr>
<tr>
<td>6</td>
<td>120</td>
<td>120</td>
<td>240</td>
<td>6.250</td>
<td>3.125</td>
</tr>
<tr>
<td>7</td>
<td>120</td>
<td>120</td>
<td>240</td>
<td>3.125</td>
<td>1.563</td>
</tr>
<tr>
<td>8</td>
<td>-</td>
<td>120</td>
<td>120</td>
<td>-</td>
<td>0</td>
</tr>
</tbody>
</table>
11. SAMPLE PREPARATION

11.1 Plasma
Collect plasma using one tenth volume of 0.1 M sodium citrate as an anticoagulant. Centrifuge samples at 3000 x g for 10 minutes. Dilute samples 100,000-fold into 1X Diluent N or within the range of 20,000x – 200,000x, and assay. Depending on application needs, user should determine proper dilutions. The undiluted samples can be stored at -20°C or below for up to 3 months. Avoid repeated freeze-thaw cycles (EDTA and Heparin can also be used as an anticoagulant).

11.2 Serum
Samples should be collected into a serum separator tube. After clot formation, centrifuge samples at 3000 x g for 10 minutes and remove serum. Dilute samples 100,000-fold into 1X Diluent N or within the range of 20,000x – 200,000x, and assay. Depending on application needs, user should determine proper dilutions. The undiluted samples can be stored at -20°C or below for up to 3 months. Avoid repeated freeze-thaw cycles.

11.3 Urine
Collect urine using sample pot. Centrifuge samples at 800 x g for 10 minutes. Dilute samples at 1:2 into 1X Diluent N or within the range of 1x – 10x, and assay. Depending on application needs, user should determine proper dilutions. The undiluted samples can be stored at -20°C or below for up to 3 months. Avoid repeated freeze-thaw cycles.

11.4 Cell culture supernatants
Centrifuge cell culture media at 3,000 x g for 10 minutes to remove debris. Collect supernatants and assay. Store samples at -20°C or below. Avoid repeated freeze-thaw cycles.
11.5 **Cell Lysate**

Rinse cell with cold PBS and then scrape the cell into a tube with 5 ml cold PBS with 0.5 M EDTA. Centrifuge suspension at 1,500 rpm for 10 minutes at 4°C and aspirate supernatant. Re-suspend the pellet in ice-cold Lysis Buffer (10 mM Tris, pH 8.0, 130 mM NaCl, 1% Triton X-100, protease inhibitor cocktail). For every 1 x 10^6 cells add approximately 100 μL of ice-cold Lysis Buffer. Incubate on ice for 60 min. Centrifuge at 13,000 rpm for 30 minutes at 4°C and collect supernatant for assay.

11.6 **Saliva**

Collect saliva using sample tube. Centrifuge samples at 800 x g for 10 minutes. Dilute samples at 1:32 into 1X Diluent N and assay. The undiluted samples can be stored at -20°C or below for up to 3 months. Avoid repeated freeze-thaw cycles.

11.7 **CSF**

Collect cerebrospinal fluid (CSF) using sample pot. Centrifuge samples at 3000 x g for 10 minutes. Dilute samples at 1:40 into 1X Diluent N or within the range of 5x – 200x and assay. The undiluted samples can be stored at -80°C for up to 3 months. Avoid repeated freeze-thaw cycles.

11.8 **Milk**

Collect milk using sample tube. Centrifuge samples at 800 x g for 10 minutes. Use undiluted samples or dilute into 1X Diluent N 32x and assay. Store samples at -20°C or below for up to 3 months. Avoid repeated freeze-thaw cycles.
12. PLATE PREPARATION

- The 96 well plate strips included with this kit are supplied ready to use. It is not necessary to rinse the plate prior to adding reagents.
- Unused well plate strips should be returned to the plate packet and stored at 4°C.
- For statistical reasons, we recommend each sample should be assayed with a minimum of two replicates (duplicates).
- Well effects have not been observed with this assay. Contents of each well can be recorded on the template sheet included in the Resources section.
13. ASSAY PROCEDURE

- Equilibrate all materials and prepared reagents to room temperature (18 - 25°C) prior to use.
- It is recommended to assay all standards, controls and samples in duplicate.

13.1 Prepare all reagents, working standards and samples as instructed. Equilibrate reagents to room temperature before use. The assay is performed at room temperature (18-25°C).

13.2 Remove excess microplate strips from the plate frame and return them immediately to the foil pouch with desiccant inside. Reseal the pouch securely to minimize exposure to water vapor and store in a vacuum desiccator.

13.3 Add 50 μL of Apolipoprotein A1 Standard or sample per well. Cover wells with a sealing tape and incubate for two hours. Start the timer after the last sample addition.

13.4 Wash five times with 200 μL of 1X Wash Buffer manually. Invert the plate each time and decant the contents; tap it 4-5 times on absorbent paper towel to completely remove the liquid. If using a machine, wash six times with 300 μL of 1X Wash Buffer and then invert the plate, decant the contents; tap it 4-5 times on absorbent paper towel to completely remove the liquid.

13.5 Add 50 μL of 1X Biotinylated Apolipoprotein A1 Antibody to each well and incubate for one hour.

13.6 Wash microplate as described above.

13.7 Add 50 μL of 1X SP Conjugate to each well and incubate for 30 minutes. Turn on the microplate reader and set up the program in advance.

13.8 Wash microplate as described above.

13.9 Add 50 μL of Chromogen Substrate per well and incubate for about 15 minutes or till the optimal blue colour density
develops. Gently tap plate to ensure thorough mixing and break the bubbles in the well with pipette tip.

13.10 Add 50 μL of Stop Solution to each well. The color will change from blue to yellow.

13.11 Read the absorbance on a microplate reader at a wavelength of 450 nm immediately. If wavelength correction is available, subtract readings at 570 nm from those at 450 nm to correct optical imperfections. Otherwise, read the plate at 450 nm only. Please note that some unstable black particles may be generated at high concentration points after stopping the reaction for about 10 minutes, which will reduce the readings.
14. **CALCULATIONS**
Calculate the mean value of the triplicate readings for each standard and sample. To generate a Standard Curve, plot the graph using the standard concentrations on the x-axis and the corresponding mean 450 nm absorbance on the y-axis. The best-fit line can be determined by regression analysis using log-log or four-parameter logistic curve-fit. Determine the unknown sample concentration from the Standard Curve and multiply the value by the dilution factor.

15. **TYPICAL DATA**
**TYPICAL STANDARD CURVE** – Data provided for demonstration purposes only. A new standard curve must be generated for each assay performed.
16. TYPICAL SAMPLE VALUES

SENSITIVITY –
The minimum detectable dose of Apolipoprotein AI is typically 3 ng/mL.

RECOVERY –
Standard Added Value: 12.5 – 100 ng/mL
Recovery %: 91 – 108.
Average Recovery %: 98

LINEARITY OF DILUTION –

<table>
<thead>
<tr>
<th>Plasma Dilution</th>
<th>Average % Expected Value</th>
</tr>
</thead>
<tbody>
<tr>
<td>25000X</td>
<td>105</td>
</tr>
<tr>
<td>50000X</td>
<td>101</td>
</tr>
<tr>
<td>100000X</td>
<td>98</td>
</tr>
</tbody>
</table>

PRECISION –

<table>
<thead>
<tr>
<th></th>
<th>Intra-Assay</th>
<th>Inter-Assay</th>
</tr>
</thead>
<tbody>
<tr>
<td>% CV</td>
<td>4.0</td>
<td>10.0</td>
</tr>
</tbody>
</table>
17. ASSAY SPECIFICITY

<table>
<thead>
<tr>
<th>Species</th>
<th>% Cross Reactivity</th>
</tr>
</thead>
<tbody>
<tr>
<td>Canine</td>
<td>None</td>
</tr>
<tr>
<td>Bovine</td>
<td>None</td>
</tr>
<tr>
<td>Monkey</td>
<td>&lt;5</td>
</tr>
<tr>
<td>Mouse</td>
<td>None</td>
</tr>
<tr>
<td>Rat</td>
<td>None</td>
</tr>
<tr>
<td>Swine</td>
<td>None</td>
</tr>
<tr>
<td>Rabbit</td>
<td>None</td>
</tr>
</tbody>
</table>

No significant cross reactivity observed with Apo-AI, Apo-B, Apo-CII, Apo-CIII, and Apo-E.
# Troubleshooting

<table>
<thead>
<tr>
<th>Problem</th>
<th>Cause</th>
<th>Solution</th>
</tr>
</thead>
<tbody>
<tr>
<td>Poor standard curve</td>
<td>Improper standard dilution</td>
<td>Confirm dilutions made correctly</td>
</tr>
<tr>
<td></td>
<td>Standard improperly reconstituted (if applicable)</td>
<td>Briefly spin vial before opening; thoroughly resuspend powder (if applicable)</td>
</tr>
<tr>
<td></td>
<td>Standard degraded</td>
<td>Store sample as recommended</td>
</tr>
<tr>
<td></td>
<td>Curve doesn't fit scale</td>
<td>Try plotting using different scale</td>
</tr>
<tr>
<td>Low signal</td>
<td>Incubation time too short</td>
<td>Try overnight incubation at 4°C</td>
</tr>
<tr>
<td></td>
<td>Target present below detection limits of assay</td>
<td>Decrease dilution factor; concentrate samples</td>
</tr>
<tr>
<td></td>
<td>Precipitate can form in wells upon substrate addition when concentration of target is too high</td>
<td>Increase dilution factor of sample</td>
</tr>
<tr>
<td></td>
<td>Using incompatible sample type (e.g. serum vs. cell extract)</td>
<td>Detection may be reduced or absent in untested sample types</td>
</tr>
<tr>
<td></td>
<td>Sample prepared incorrectly</td>
<td>Ensure proper sample preparation/dilution</td>
</tr>
<tr>
<td>Large CV</td>
<td>Bubbles in wells</td>
<td>Ensure no bubbles present prior to reading plate</td>
</tr>
<tr>
<td></td>
<td>All wells not washed equally/thoroughly</td>
<td>Check that all ports of plate washer are unobstructed; wash wells as recommended</td>
</tr>
<tr>
<td></td>
<td>Incomplete reagent mixing</td>
<td>Ensure all reagents/master mixes are mixed thoroughly</td>
</tr>
<tr>
<td></td>
<td>Inconsistent pipetting</td>
<td>Use calibrated pipettes and ensure accurate pipetting</td>
</tr>
<tr>
<td></td>
<td>Inconsistent sample preparation or storage</td>
<td>Ensure consistent sample preparation and optimal sample storage conditions (e.g. minimize freeze/thaw cycles)</td>
</tr>
<tr>
<td>Problem</td>
<td>Cause</td>
<td>Solution</td>
</tr>
<tr>
<td>------------------------------</td>
<td>------------------------------------</td>
<td>--------------------------------------------------------------------------</td>
</tr>
<tr>
<td>Wells are insufficiently</td>
<td>Wash wells as per protocol</td>
<td></td>
</tr>
<tr>
<td>washed</td>
<td>recommendations</td>
<td></td>
</tr>
<tr>
<td>Contaminated wash buffer</td>
<td>Make fresh wash buffer</td>
<td></td>
</tr>
<tr>
<td>Waiting too long to read</td>
<td>Read plate immediately</td>
<td></td>
</tr>
<tr>
<td>plate after adding STOP</td>
<td>after adding STOP solution</td>
<td></td>
</tr>
<tr>
<td>solution</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Improper storage of</td>
<td>Store all reagents as recommended</td>
<td></td>
</tr>
<tr>
<td>ELISA kit</td>
<td>Please note all reagents may not</td>
<td></td>
</tr>
<tr>
<td></td>
<td>have identical storage requirements.</td>
<td></td>
</tr>
<tr>
<td>Using incompatible sample</td>
<td>Detection may be reduced or absent</td>
<td></td>
</tr>
<tr>
<td>type (e.g. Serum vs. cell</td>
<td>in untested sample types</td>
<td></td>
</tr>
<tr>
<td>extract)</td>
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</table>
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