MONKEY SERUM AMYLOID A (SAA) ELISA
Life Diagnostics, Inc., Catalog Number: SAA-3

INTRODUCTION
SAA is an acute phase serum protein that can be elevated up to 400-fold-fold in monkeys. As is the case in humans, measurement of SAA provides an excellent biomarker of inflammation and disease.

PRINCIPLE OF THE ASSAY
The assay uses two different peptide-specific monkey SAA antibodies; one for solid phase immobilization and the other, conjugated to horseradish peroxidase (HRP), for detection. Serum samples are first denatured by heating for 1 hour at 60°C. The denaturing step dissociates SAA from interfering factors. Subsequently, the denatured samples are diluted. Standards and denaturing step dissociates SAA from interfering factors.

Materials provided with the kit:
- SAA antibody coated 96-well plate (12 x 8-well strips)
- HRP Conjugate, 11 ml
- SAA stock (lyophilized)
- 20x Wash solution; TBS50-20, 50 ml
- Diluent; YD50-1, 50 ml
- TMB, TMB11-1, 11 ml
- Stop solution, SS11-1, 11 ml

Materials required but not provided:
- Pipettors and tips
- Distilled or deionized water
- Polypropylene or glass tubes
- Vortex mixer
- Absorbent paper or paper towels
- Plate incubator/shaker
- Plate washer
- Plate reader capable of measuring absorbance at 450 nm.
- Curve fitting software

STORAGE
The SAA stock should be stored at or below -20°C for optimum stability. The remainder of the kit should be stored at 2-8°C and the microtiter plate should be kept in a sealed bag with desiccant. Kits will remain stable for six months from the date of purchase.

GENERAL INSTRUCTIONS
1. All reagents should be allowed to reach room temperature before use.
2. Reliable and reproducible results will be obtained when the assay is carried out with a complete understanding of the instructions and with adherence to good laboratory practice.
3. The wash procedure is critical. Insufficient washing will result in poor precision and falsely elevated absorbance readings.
4. Laboratory temperature will influence absorbance readings. Our ELISA kits are calibrated using shaking incubators set at 150 rpm and 25°C. Performance of the assay at lower temperatures will result in lower absorbance values.

WASH SOLUTION PREPARATION
The wash solution is provided as a 20x stock. Prior to use, dilute the contents of the bottle (50 ml) with 950 ml of distilled or deionized water.

STANDARD PREPARATION
The stock vial contains lyophilized heat-treated SAA of known concentration (it must not be incubated at 60°C).

1. Reconstitute the stock with deionized or distilled water as described on the vial label. Mix gently several times over a period of 5 minutes.
2. Label 6 polypropylene tubes as 25, 12.5, 6.25, 3.125, 1.56 and 0.78 ng/ml.
3. Into the tube labeled 25 ng/ml, pipette the volume of diluent detailed on the stock vial label. Then add the indicated volume of stock and mix. This provides the 25 ng/ml standard.
4. Dispense 250 µl of diluent into the tubes labeled 12.5, 6.25, 3.125, 1.56 and 0.78 ng/ml.
5. Prepare the 12.5 ng/ml standard by mixing 250 µl of the 25 ng/ml SAA standard with 250 ml of diluent in the tube labeled 12.5 ng/ml.
6. Similarly prepare the remaining standards by serial dilution. Unused reconstituted stock should be stored frozen at or below -20°C if future use is intended (it is stable for at least one week at 2-8°C).

SAMPLE PREPARATION

Denaturation
1. Dispense 100 µl of each serum sample into a polypropylene microcentrifuge tube and tightly seal.
2. Incubate the samples at 60°C in a water bath for one hour.

Dilution
1. After denaturation, dilute 1.0 µl of denatured sample with 249 µl of diluent.

This procedure gives a 250-fold dilution of the original sample and presents SAA in a form that is recognizable by the antibodies used in the kit.

ASSAY PROCEDURE

1. Secure the desired number of 8-well strips in the holder. Unused strips should be stored in the re-sealed bag with desiccant at 2-8°C for future use.
2. Dispense 100 µl of standards and samples into the wells (we recommend that standards and samples be run in duplicate).
3. Add 100 µl of HRP-conjugate into each well.
4. Incubate on a plate shaker at 150 rpm and 25°C for one hour.
5. Empty and wash the microtiter wells 5x with 1x wash solution using a plate washer (400 µl/well).
6. Strike the wells sharply onto absorbent paper or paper towels to remove all residual droplets.
7. Dispense 100 µl of TMB into each well.
8. Incubate on an orbital micro-plate shaker at 150 rpm at 25°C for 20 minutes.
9. After 20-minutes, stop the reaction by adding 100 µl of Stop solution to each well.
10. Gently mix. It is important to make sure that all the blue color changes to yellow.
11. Read absorbance at 450 nm with a plate reader within 5 minutes.

**CALCULATION OF RESULTS**
1. Using curve fitting software, construct a standard curve by plotting absorbance values of the standards versus concentration.
2. Fit the standard curve to an appropriate model and derive the concentration of the samples (we recommend using a single site, total and nonspecific binding model).
3. Multiply the derived concentration by the dilution factor to determine the actual concentration in the serum sample.
4. If the A$_{450}$ values of samples fall outside the standard curve, samples should be diluted appropriately and re-tested.

**TYPICAL STANDARD CURVE**
A typical standard curve with absorbance at 450 nm on the Y-axis against SAA concentrations on the X-axis is shown below. This curve is for illustration only.

<table>
<thead>
<tr>
<th>SAA (ng/ml)</th>
<th>A$_{450}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>25</td>
<td>3.182</td>
</tr>
<tr>
<td>12.5</td>
<td>1.870</td>
</tr>
<tr>
<td>6.25</td>
<td>1.139</td>
</tr>
<tr>
<td>3.125</td>
<td>0.609</td>
</tr>
<tr>
<td>1.56</td>
<td>0.384</td>
</tr>
<tr>
<td>0.78</td>
<td>0.252</td>
</tr>
</tbody>
</table>

**REFERENCES**
2. Ossetrova NI, Sandgren DJ and Blakely WF. C-reactive protein and serum amyloid A as early-phase and prognostic indicators of acute radiation exposure in nonhuman primate total-body irradiation model. Radiation Measurements 46:1019-1024 (2011)