

APCS (Human) ELISA Kit

Catalog Number KA1841

96 assays

Version: 12

Intended for research use only



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Introduction

Background

Serum amyloid P component (SAP, serum AP, APCS), a 25 kDa pentameric protein, is a normal plasma protein and a universal non-fibrillar constituent of amyloid deposits (1). SAP is a pentraxin similar to C-reactive protein (CRP). SAP and CRP are suggested to be associated with cardiovascular disease (2-3). SAP is also suggested to be linked with Alzheimer's disease (4) and liver disease (5).

Principle of the Assay

The APCS (Human) ELISA Kit is designed for detection of SAP in human plasma, serum, urine, CSF and cell culture samples. This assay employs a quantitative sandwich enzyme immunoassay technique that measures human SAP in approximately 4 hours. A polyclonal antibody specific for human SAP has been pre-coated onto a 96-well microplate with removable strips. SAP in standards and samples is sandwiched by the immobilized antibody and a biotinylated polyclonal antibody specific for human SAP, which is recognized by a streptavidin-peroxidase (SP) conjugate. All unbound material is washed away and a peroxidase enzyme substrate is added. The color development is stopped and the intensity of the color is measured.



General Information

Materials Supplied

List of component

Component	Amount	
Human Serum Amyloid P Microplate: A 96-well polystyrene microplate (12 strips of 8	00 (0v42) welle	
wells) coated with a polyclonal antibody against human SAP.	96 (8x12) wells	
Sealing Tapes: Pressure sensitive sealing tapes that can be cut to fit the format of the	0 "1	
individual assay.	3 slides	
Human Serum Amyloid P Standard: Human SAP in a buffered protein base	20.55	
(lyophilized).	20 ng	
Biotinylated Human Serum Amyloid P Antibody (50x): A 50-fold concentrated	120 µL	
biotinylated polyclonal antibody against human SAP.		
MIX Diluent Concentrate (10x): A 10-fold concentrated buffered protein base.	30 mL	
Wash Buffer Concentrate (20x): A 20-fold concentrated buffered surfactant.	30 mL x 2	
SP Conjugate (100x): A 100-fold concentrate.	80 μL	
Chromogen Substrate (1x): A stabilized peroxidase chromogen substrate	7 ml	
tetramethylbenzidine.	7 mL	
Stop Solution: A 0.5 N hydrochloric acid solution to stop the chromogen substrate	11 mL	
reaction.	I I IIIL	

Storage Instruction

- ✓ Upon arrival, immediately store components of the kit at recommended temperatures up to the expiration
- ✓ Store SP Conjugate and Biotinylated Antibody at -20°C.
- ✓ Store Microplate, Diluent Concentrate (10x), Wash Buffer, Stop Solution, and Chromogen Substrate at 2-8°C.
- ✓ Unused microplate wells may be returned to the foil pouch with the desiccant packs and resealed. May be stored for up to 30 days in a vacuum desiccator.
- ✓ Store Standard at 2-8°C before reconstituting with Diluent and at -20°C after reconstituting with Diluent.

Materials Required but Not Supplied

- ✓ Microplate reader capable of measuring absorbance at 450 nm.
- ✓ Pipettes (1-20 μ L, 20-200 μ L, 200-1000 μ L and multiple channel).
- ✓ Deionized or distilled reagent grade water.



Precautions for Use

- ✓ This product is for Research Use Only and is not intended for use in diagnostic procedures.
- ✓ Prepare all reagents (diluent buffer, wash buffer, standard, biotinylated antibody, and SP conjugate) as instructed, prior to running the assay.
- ✓ Prepare all samples prior to running the assay. The dilution factors for the samples are suggested in this insert. However, the user should determine the optimal dilution factor.
- ✓ Spin down the SP conjugate vial and the biotinylated antibody vial before opening and using contents.
- ✓ The Stop Solution is an acidic solution.
- ✓ The kit should not be used beyond the expiration date.



Assay Protocol

Reagent Preparation

- ✓ Freshly dilute all reagents and bring all reagents to room temperature before use.
- ✓ MIX Diluent Concentrate (10x): Dilute the MIX Diluent Concentrate 10-fold with reagent grade water to produce a 1x solution. When diluting the concentrate, make sure to rinse the bottle thoroughly to extract any precipitates left in the bottle. Mix the 1x solution gently until the crystals have completely dissolved. Store for up to 30 days at 2-8°C.
- ✓ Human Serum Amyloid P Standard: Reconstitute the Human Serum Amyloid P Standard (20 ng) with 2 mL of MIX Diluent to generate a 10 ng/mL standard stock solution. Allow the vial to sit for 10 minutes with gentle agitation prior to making dilutions. Prepare duplicate or triplicate standard points by serially diluting the standard stock solution (10 ng/mL) 2-fold with equal volume of MIX Diluent to produce 5, 2.5, 1.25, 0.625, 0.313, 0.156, and 0.078 ng/mL solutions. MIX Diluent serves as the zero standard (0 ng/mL). Any remaining stock solution should be stored at -20°C and used within 30 days. Avoid repeated freeze-thaw cycles.

Standard Point	Dilution	[SAP] (ng/mL)
P1	1 part Standard (10 ng/mL) +1 part MIX Diluent	5.00
P2	1 part P1 + 1 part MIX Diluent	2.500
P3	1 part P2 + 1 part MIX Diluent	1.250
P4	1 part P3 + 1 part MIX Diluent	0.625
P5	1 part P4 + 1 part MIX Diluent	0.313
P6	1 part P5 + 1 part MIX Diluent	0.156
P7	1 part P6 + 1 part MIX Diluent 0.0	
P8	MIX Diluent	0.000

- Biotinylated Human Serum Amyloid P Antibody (50x): Spin down the antibody briefly and dilute the desired amount of the antibody 50-fold with MIX Diluent to produce a 1x solution. The undiluted antibody should be stored at -20°C.
- Wash Buffer Concentrate (20x): Dilute the Wash Buffer Concentrate 20-fold with reagent grade water to produce a 1x solution. When diluting the concentrate, make sure to rinse the bottle thoroughly to extract any precipitates left in the bottle. Mix the 1x solution gently until the crystals have completely dissolved.
- SP Conjugate (100x): Spin down the SP Conjugate briefly and dilute the desired amount of the conjugate 100-fold with MIX Diluent to produce a 1x solution. The undiluted conjugate should be stored at -20°C.

Sample Preparation

✓ 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 and collect plasma. A 40000-fold sample dilution is suggested into



MIX Diluent; however, user should determine optimal dilution factor depending on application needs. The undiluted samples can be stored at -20°C or below for up to 3 months. Avoid repeated freeze-thaw cycles (EDTA or Heparin can also be used as an anticoagulant).

- ✓ 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. A 40000-fold sample dilution is suggested into MIX Diluent; however, user should determine optimal dilution factor depending on application needs. The undiluted samples can be stored at -20°C or below for up to 3 months. Avoid repeated freeze-thaw cycles.
- ✓ Urine: Collect urine using sample pot. Centrifuge samples at 800 x g for 10 minutes. A 2-fold sample dilution is suggested into MIX Diluent or within the range of 1x-20x; however, user should determine optimal dilution factor depending on application needs. The undiluted samples can be stored at -20°C or below for up to 3 months. Avoid repeated freeze-thaw cycles.
- ✓ CSF: Collect cerebrospinal fluid (CSF) using sample pot. Centrifuge samples at 3000 x g for 10 minutes. A 20-fold sample dilution is suggested into MIX Diluent; however, user should determine optimal dilution factor depending on application needs. The undiluted samples can be stored at -80°C for up to 3 months. Avoid repeated freeze-thaw cycles.
- ✓ Cell Culture Supernatants: Centrifuge cell culture media at 1500 rpm for 10 minutes at 4°C to remove debris and collect supernatants. If necessary, dilute samples into MIX Diluent; user should determine optimal dilution factor depending on application needs. The undiluted samples can be stored at -80°C. Avoid repeated freeze-thaw cycles.

Applicable samples may also include biofluids, cell culture, and tissue homogenates. If necessary, user should determine optimal dilution factor depending on application needs.

✓ Refer to Dilution Guidelines for further instruction.

Guidelines for Dilutions of 100-fold or Greater		
(for reference only; please follow the insert for specific dilution suggested)		
100x	10000x	
A) 4 μL Sample: 396 μL buffer (100x)	A) 4 μL Sample: 396 μL buffer (100x)	
= 100-fold dilution	B) 4 μL of A: 396 μL buffer (100x)	
	= 10000-fold dilution	
Assuming the needed volume is less than or Assuming the needed volume is less that		
equal to 400 μL. equal to 400 μL.		
1000x	100000x	
A) 4 μL Sample: 396 μL buffer (100x)	A) 4 μL sample: 396 μL buffer (100x)	
B) 24 μL of A: 216 μL buffer (10x)	B) 4 μL of A: 396 μL buffer (100x)	
= 1000-fold dilution	C) 24 µL of B: 216 µL buffer (10x)	
	= 100000-fold dilution	
Assuming the needed volume is less than or	Assuming the needed volume is less than or	
equal to 240 μL.	equal to 240 μL.	



Assay Procedure

- 1. Prepare all reagents, standard solutions, and samples as instructed. Bring all reagents to room temperature before use. The assay is performed at room temperature (20-25°C).
- 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.
- 3. Add 50 µL of Human Serum Amyloid P Standard or sample to each well. Gently tap plate to thoroughly coat the wells. Break any bubbles that may have formed. Cover wells with a sealing tape and incubate for 2 hours. Start the timer after the last addition.
- 4. Wash the microplate manually or automatically using a microplate washer. Invert the plate and decant the contents; hit 4-5 times on absorbent material to completely remove the liquid. If washing manually, wash five times with 200 μL of Wash Buffer per well. Invert the plate each time and decant the contents; hit 4-5 times on absorbent material to completely remove the liquid. If using a microplate washer, wash six times with 300 μL of Wash Buffer per well; invert the plate and hit 4-5 times on absorbent material to completely remove the liquid.
- 5. Add 50 μL of Biotinylated Human Serum Amyloid P Antibody to each well. Gently tap plate to thoroughly coat the wells. Break any bubbles that may have formed. Cover wells with a sealing tape and incubate for 1 hour.
- 6. Wash the microplate as described above.
- 7. Add 50 μL of SP Conjugate to each well. Gently tap plate to thoroughly coat the wells. Break any bubbles that may have formed. Cover wells with a sealing tape and incubate for 30 minutes. Turn on the microplate reader and set up the program in advance.
- 8. Wash the microplate as described above.
- Add 50 μL of Chromogen Substrate to each well. Gently tap plate to thoroughly coat the wells. Break any bubbles that may have formed. Incubate in ambient light for 12 minutes or until the optimal blue color density develops.
- 10. Add 50 μL of Stop Solution to each well. The color will change from blue to yellow. Gently tap plate to ensure thorough mixing. Break any bubbles that may have formed.
- 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.



- ✓ Assay Summary
- 1. Add 50 µL of Standard or Sample per well. Incubate 2 hours.
- 2. Wash, then add 50 μL of Biotinylated Antibody per well. Incubate 1 hour.
- 3. Wash, then add 50 μ L of SP Conjugate per well. Incubate 30 minutes.
- 4. Wash, then add 50 μL of Chromogen Substrate per well. Incubate 12 minutes.
- 5. Add 50 μ L of Stop Solution per well. Read at 450 nm immediately.



Data Analysis

Calculation of Results

- ✓ Calculate the mean value of the duplicate or 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 (OD) 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
- ✓ Typical Data

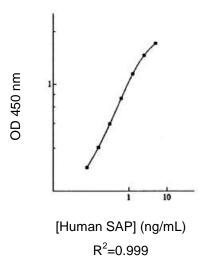
The typical data is provided for reference only. Individual laboratory means may vary from the values listed. Variations between laboratories may be caused by technique differences.

		<u> </u>	A OD
Standard Point	ng/mL	OD	Average OD
P1	5.000	2.196	2.232
1 1	3.000	2.268	2.232
P2	2.500	1.727	1.761
FZ	2.300	1.795	1.701
P3	1.250	1.246	1.222
P3	1.250	1.198	1.222
P4	0.605	0.770	0.755
P4	0.625	0.740	0.755
DE	P5 0.313	0.443	0.450
Po		0.469	0.456
P6	0.450	0.280	0.207
Po	0.156	0.294	0.287
P7	0.078	0.188	0.192
F /	0.078	0.196	0.192
P8	0.0	0.065	0.064
P0	0.0	0.063	0.064
Sample: Pooled Normal		0.993	0.070
Sodium Citrate F	Sodium Citrate Plasma (40000x)		0.979
Sample: Po	oled Normal	1.007	4.040
Serum (40000x)		1.029	1.018



✓ Standard Curve

The curve is provided for illustration only. A standard curve should be generated each time the assay is performed.



✓ Reference Value

Normal human SAP plasma and serum levels range from 29 – 48 $\mu g/mL$.

Plasma and serum samples from healthy adults were tested (n=40). On average, human SAP level was 35.7 μg/mL.

Sample	n	Average Value (µg/mL)
Pooled Normal Plasma	10	36.0
Normal Plasma	20	31.9
Pooled Normal Serum	10	39.3

Performance Characteristics

- ✓ The minimum detectable dose of human SAP as calculated by 2SD from the mean of a zero standard was established to be 26 pg/mL.
- ✓ Intra-assay precision was determined by testing three plasma samples twenty times in one assay.
- ✓ Inter-assay precision was determined by testing three plasma samples in twenty assays.

	Intra-Assay Precision			Inte	r-Assay Precis	sion
Sample	1	2	3	1	2	3
n	20	20	20	20	20	20
CV (%)	4.9%	4.4%	4.2%	10.2%	9.8%	9.1%
Average CV (%)	4.5%				9.7%	



Spiking Recovery

Recovery was determined by spiking two plasma samples with different SAP concentrations.

	Unspiked	Spiking			
Sample	Sample	Value	Expected	Observed	Recovery (%)
	(ng/mL)	(ng/mL)			
		1.5	2.8	2.9	104%
1	1.3	3.0	4.3	4.4	102%
		7.5	8.8	7.6	86%
		1.5	5.0	4.8	96%
2	3.5	3.0	6.5	6.7	103%
		7.5	11.0	10.4	95%
	98%				

Linearity

Plasma and serum samples were serially diluted to test for linearity.

Average Percentage of Expected Value (%)			
Sample Dilution	ution Plasma Serum		
20000x 109% 104%		104%	
40000x 88%		97%	
80000x	96%	103%	

Cross-Reactivity

Species	Cross Reactivity (%)
Canine	None
Bovine	None
Monkey	<5%
Mouse	None
Rat	None
Swine	None
Rabbit	None

10% FBS in culture media will not affect the assay.



Resources

Troubleshooting

Issue	Causes	Course of Action		
	Han of improvements	Check the expiration date listed before use.		
	Use of improper components	Do not interchange components from different lots.		
		Check that the correct wash buffer is being used.		
		Check that all wells are empty after aspiration.		
		Check that the microplate washer is dispensing		
	Improper wash step	properly.		
		If washing by pipette, check for proper pipetting		
		technique.		
io	Splashing of reagents while loading	Pipette properly in a controlled and careful manner.		
ecis	wells			
Low Precision		Pipette properly in a controlled and careful manner.		
Lo	Inconsistent volumes loaded into wells	Check pipette calibration.		
		Check pipette for proper performance.		
		Thoroughly agitate the lyophilized components after		
	Insufficient mixing of reagent dilutions	reconstitution.		
		Thoroughly mix dilutions.		
	Improperly sealed microplate	Check the microplate pouch for proper sealing.		
		Check that the microplate pouch has no punctures.		
		Check that three desiccants are inside the		
		microplate pouch prior to sealing.		
	Microplate was left unattended between	Each step of the procedure should be performed		
	steps	uninterrupted.		
ensity	Omission of step	Consult the provided procedure for complete list of		
Inte	Offission of step	steps.		
gnal	Steps performed in incorrect order	Consult the provided procedure for the correct		
h Się	Steps performed in incorrect order	order.		
Hig	Insufficient amount of reagents added to	Check pipette calibration.		
w or	wells	Check pipette for proper performance.		
/ Lo	Wash step was skipped	Consult the provided procedure for all wash steps.		
Unexpectedly Low or High Signal Int	Improper wash buffer	Check that the correct wash buffer is being used.		
cbec	Improper reagent properation	Consult reagent preparation section for the correct		
Jne	Improper reagent preparation	dilutions of all reagents.		
	Insufficient or prolonged incubation	Consult the provided procedure for correct		
	periods	incubation time.		



		Sandwich ELISA: If samples generate OD values
		higher than the highest standard point (P1), dilute
		samples further and repeat the assay.
	Non entimal comple dilution	Competitive ELISA: If samples generate OD values
	Non-optimal sample dilution	lower than the highest standard point (P1), dilute
		samples further and repeat the assay.
		User should determine the optimal dilution factor
ve F		for samples.
Deficient Standard Curve Fit	Contamination of reagents	A new tip must be used for each addition of
dard		different samples or reagents during the assay
Stan		procedure.
ent 8	Contents of wells evaporate	Verify that the sealing film is firmly in place before
efici		placing the assay in the incubator or at room
۵		temperature.
		Pipette properly in a controlled and careful manner.
	Improper pipetting	Check pipette calibration.
		Check pipette for proper performance.
		Thoroughly agitate the lyophilized components
	Insufficient mixing of reagent dilutions	after reconstitution.
		Thoroughly mix dilutions.

References

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Plate Layout

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