

AssayMax™ Human alpha-2-Macroglobulin ELISA Kit

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For any questions regarding troubleshooting or performing the assay, please contact our support team at support@assaypro.com.

Thank you for choosing Assaypro.

Assay Summary

Step 1. Add 50 μ l of Standard or Sample per well. Incubate 2 hours.

Step 2. Wash, then add 50 μ l of Biotinylated Antibody per well. Incubate 1 hour.

Step 3. Wash, then add 50 μ l of SP Conjugate per well. Incubate 30 minutes.

Step 4. Wash, then add 50 μ l of Chromogen Substrate per well. Incubate 20 minutes.

Step 5. Add 50 μ l of Stop Solution per well. Read at 450 nm immediately.

Symbol Key



Consult instructions for use.

Assay Template

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AssayMax™ Human alpha-2-Macroglobulin (A2M) ELISA Kit

Catalog No. EM1115-1
Sample insert for reference use only

Introduction

Alpha-2-macroglobulin (A2M, alpha-2-M) is a major serum protein with diverse functions, including inhibition of protease activity and binding of growth factors, cytokines, and disease factors (1). Increased serum alpha-2-macroglobulin has been suggested to be associated with multiple sclerosis (MS) [2], glomerular disease (3), and liver disease (4).

Principle of the Assay

The AssayMax™ Human alpha-2-Macroglobulin ELISA (Enzyme-Linked Immunosorbent Assay) Kit is designed for detection of A2M in human milk, saliva, CSF, and cell culture samples. This assay employs a quantitative sandwich enzyme immunoassay technique that measures human A2M in approximately 4 hours. A polyclonal antibody specific for human A2M has been pre-coated onto a 96-well microplate with removable strips. A2M in standards and samples is sandwiched by the immobilized antibody and a biotinylated polyclonal antibody specific for human A2M, 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.

Caution and Warning

- 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.

Reagents

- Human alpha-2-Macroglobulin Microplate: A 96-well polystyrene microplate (12 strips of 8 wells) coated with a polyclonal antibody against human A2M.
- Sealing Tapes: Each kit contains 3 precut, pressure sensitive sealing tapes that can be cut to fit the format of the individual assay.
- Human alpha-2-Macroglobulin Standard: Human A2M in a buffered protein base (1000 ng, lyophilized).
- Biotinylated Human alpha-2-Macroglobulin Antibody (50x): A 50-fold concentrated biotinylated polyclonal antibody against human A2M (120 μl).
- EIA Diluent Concentrate (10x): A 10-fold concentrated buffered protein base (30 ml).
- Wash Buffer Concentrate (20x): A 20-fold concentrated buffered surfactant (30 ml, 2 bottles).
- SP Conjugate (100x): A 100-fold concentrate (80 μl).
- Chromogen Substrate (1x): A stabilized peroxidase chromogen substrate tetramethylbenzidine (7 ml).
- **Stop Solution (1x):** A 0.5 N hydrochloric acid solution to stop the chromogen substrate reaction (11 ml).

Storage Condition

- Upon arrival, immediately store components of the kit at recommended temperatures up to the expiration date.
- 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.

Other Supplies Required

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

Sample Collection, Preparation, and Storage

- Milk: Collect milk using sample tube. Centrifuge samples at 800 x g for 10 minutes. A 40-fold sample dilution is suggested into EIA Diluent or within the range of 4x – 400x; 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.
- Saliva: Collect saliva using sample tube. Centrifuge samples at 800 x g for 10 minutes. A 4-fold sample dilution is suggested into EIA Diluent or within the range of 1x 40x; 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 40-fold sample dilution is suggested into EIA Diluent or within the range of 4x 400x; 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 Supernatant: Centrifuge cell culture media at 1500 rpm for 10 minutes at 4°C to remove debris and collect supernatant. If necessary, dilute samples into EIA 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) = 100-fold dilution	A) B)	4 μl sample : 396 μl buffer (100x) 4 μl of A : 396 μl buffer (100x) = 10000-fold dilution		
	Assuming the needed volume is less than or equal to 400 $\mu\text{l}.$		Assuming the needed volume is less than or equal to 400 µl.		
1000x			100000x		
A) B)	4 μl sample : 396 μl buffer (100x) 24 μl of A : 216 μl buffer (10x) = 1000-fold dilution	A) B) C)	4 μl sample : 396 μl buffer (100x) 4 μl of A : 396 μl buffer (100x) 24 μl of B : 216 μl buffer (10x) = 100000-fold dilution		
	Assuming the needed volume is less than or equal to 240 μ l.		= 100000-101α dilution Assuming the needed volume is less than or equal to 240 μl.		

Reagent Preparation

- Freshly dilute all reagents and bring all reagents to room temperature before use.
- **EIA Diluent Concentrate (10x):** Dilute the EIA 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 alpha-2-Macroglobulin Standard: Reconstitute the Human alpha-2-Macroglobulin Standard (1000 ng) with 2 ml of EIA Diluent to generate a 500 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 from the standard stock solution (500 ng/ml) 4-fold with EIA Diluent to produce 125, 31.25, 7.813, and 1.953 ng/ml solutions. EIA 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	[A2M] (ng/ml)
P1	1 part Standard (500 ng/ml)	500
P2	1 part P1 + 3 parts EIA Diluent	125
P3	1 part P2 + 3 parts EIA Diluent	31.25
P4	1 part P3 + 3 parts EIA Diluent	7.813
P5	1 part P4 + 3 parts EIA Diluent	1.953
P6	EIA Diluent	0.0

- Biotinylated Human alpha-2-Macroglobulin Antibody (50x): Spin down the antibody briefly and dilute the desired amount of the antibody 50fold with EIA Diluent to produce a 1x solution. The undiluted antibody should be stored at -20°C.
- Wash Buffer Concentrate (20x): Dilute the Wash Buffer Concentrate 20fold 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 EIA Diluent to produce a 1x solution. The undiluted conjugate should be stored at -20°C.

Assay Procedure

- 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 desiccants inside. Reseal the pouch
 securely to minimize exposure to water vapor and store in a vacuum
 desiccator.
- Add 50 µl of Human alpha-2-Macroglobulin 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.
- 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.
- Add 50 µl of Biotinylated Human alpha-2-Macroglobulin 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.
- Wash the microplate as described above.
- 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.
- 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 20 minutes or until the optimal blue color density develops.
- 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.
- 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.

Data Analysis

- 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

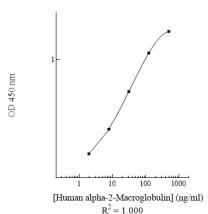
The typical data is provided for reference only. Individual laboratory
means may vary from the values listed. Variations between laboratories
may be caused by technique differences.

Standard Point	ng/ml	OD	Average OD	
P1	500	2.284	2.310	
ΓI	300	2.336	2.310	
P2	125	1.234	1.215	
ΓZ	123	1.196	1.213	
P3	31.25	0.395	0.388	
PS	31.25	0.381	0.566	
P4	7.813	0.132	0.127	
P4		0.122	0.127	
P5	1.953	0.058	0.061	
PO	1.955	0.064	0.061	
P6	0.0	0.033	0.033	0.034
FO		0.035	0.034	
Sample: Poo	lad Milk (40v)	0.409	0.418	
Sample: Pool	ieu ivilik (40X)	0.427	0.418	

Standard Curve

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

Human alpha-2-Macroglobulin Standard Curve



Performance Characteristics

 The minimum detectable dose of human A2M as calculated by 2SD from the mean of a zero standard was established to be 1.1 ng/ml.

- Intra-assay precision was determined by testing three reference control samples twenty times in one assay.
- Inter-assay precision was determined by testing three reference control samples in twenty assays.

	Intra-Assay Precision			Inter	-Assay Pred	ision
Sample	1	2	3	1	2	3
n	20	20	20	20	20	20
CV (%)	6.2%	4.4%	6.0%	11.4%	9.1%	10.5%
Average CV (%)	5.5%				10.3%	

Recovery

Standard Added Value	7.8 – 125 ng/ml	
Recovery %	89 – 112%	
Average Recovery %	97%	

Linearity

• Milk samples were serially diluted to test for linearity.

Average Percentage of Expected Value (%)			
Sample Dilution Milk			
20x	95%		
40x	99%		
80x	105%		

Cross-Reactivity

Species	Cross-Reactivity (%)
Canine	None
Bovine	None
Monkey	50%
Mouse	None
Rat	None
Swine	None
Rabbit	None

- No significant cross-reactivity observed with APP, complement C3, complement C4, complement C5, IL-1 beta, insulin, and TGF-beta-1.
- 10% FBS in culture media will not affect the assay.

Troubleshooting

Use of improper components • Check the expiration date listed before use. • Do not interchange components from differ • Check that the correct wash buffer is being in the correct wash of the correct wash. • Check the expiration date listed before use. • Do not interchange components from differ wash. • Check that the correct wash outfler is being in the correct wash of the correct wash of the correct wash of the correct wash outfler is being in the correct wash of the correct wash outfler is being in the correct wash outfler in the	
Check that the correct wash buffer is being i Check that all wells are empty after aspiration Improper wash step	
Check that all wells are empty after aspiration Improper wash step Check that the microplate washer is dispense	used.
Improper wash step • Check that the microplate washer is dispens	
	on.
 If washing by pipette, check for proper piper 	ing properly.
	tting
c technique.	
Splashing of reagents while loading wells Inconsistent volumes loaded into wells Pipette properly in a controlled and careful Check pipette calibration. Check pipette for proper performance.	manner.
• Pipette properly in a controlled and careful	manner.
• Check pipette calibration.	
Check pipette for proper performance.	
Insufficient mixing of Thoroughly agitate the lyophilized compone	ents after
reagent dilutions reconstitution.	
• Thoroughly mix dilutions.	
Check the microplate pouch for proper seali	-
• Check that the microplate pouch has no pur	
microplate Check that three desiccants are inside the m	nicroplate
pouch prior to sealing.	
Microplate was left • Each step of the procedure should be perfor	rmed
unattended between uninterrupted.	
Omission of stop	a list of stone
Omission of step Consult the provided procedure for complet Steps performed in Consult the provided procedure for the corr	
unattended between steps Omission of step Steps performed in incorrect order Insufficient amount of reagents added to wells Wash step was skipped Improper wash buffer Improper reagent preparation Insufficient or prolonged incubation Occupant the provided procedure for complete consult the provided procedure for the correct wash buffer occurately because the consult the provided procedure for all wash dilutions of all reagents. Unsufficient or prolonged incubation uninterrupted. Occupant the provided procedure for complete occurate the provided procedure for the dilutions of all reagents. Occupant the provided procedure for all wash dilutions of all reagents. Occupant the provided procedure for all wash dilutions of all reagents.	ect order.
Insufficient amount of • Check pipette calibration.	
reagents added to • Check pipette for proper performance.	
insufficient amount of reagents added to wells Wash step was skipped • Check pipette calibration. • Check pipette for proper performance. • Check pipette calibration.	
≥ ⊆ Wash step was skipped • Consult the provided procedure for all wash	steps.
Improper wash buffer • Check that the correct wash buffer is being	used.
Improper reagent • Consult reagent preparation section for the	correct
preparation dilutions of all reagents.	
Insufficient or • Consult the provided procedure for correct	incubation
' '	
periods	
Sandwich ELISA: If samples generate OD values to the state of the	
than the highest standard point (P1), dilute s	samples
further and repeat the assay. Non-optimal sample further and repeat the assay. Competitive ELISA: If samples generate OD v	alues leures
Non-optimal sample of dilution	
further and repeat the assay.	samples
User should determine the optimal dilution	factor for
samples.	140101 101
Non-optimal sample dilution Output Outpu	different
reagents samples or reagents during the assay proced	
Contents of wells • Verify that the sealing film is firmly in place	
evaporate the assay in the incubator or at room tempe	
Pipette properly in a controlled and careful	manner.
Improper pipetting • Check pipette calibration.	
Check pipette for proper performance.	

Insufficient mixing of reagent dilutions	Thoroughly agitate the lyophilized components after reconstitution. Thoroughly mix dilutions.
	more again, mix anations.

References

- (1) Pineda-Salgado L et al. (2005) Gene Expr Patterns. 6(1):3-10.
- (2) Jensen PE et al. (2004) Biochim Biophys Acta. 1690(3):203-7.
- (3) Yang AH et al. (1997) Nephrol Dial Transplant. 12(3):465-9.
- (4) Shiota G et al. (1995) J Med. 26(5-6):295-308.

Version 6.6