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# AlphaLISA<sup>®</sup> Interleukin 17A (porcine) Detection Kit

Product number: AL572HV/C/F

Research Use Only. Not for use in diagnostic procedures.

# **Product Information**

Application:	This kit is designed for the quantitative determination of porcine Interleukin 17A (pIL-17A) in serum and cell culture media using a homogeneous AlphaLISA assay (no wash steps).
Sensitivity:	Lower Detection Limit (LDL): 3.8 pg/mL Lower Limit of Quantification (LLOQ): 11.8 pg/mL
	EC <sub>50</sub> : 14.5 ng/mL
Dynamic range:	3.8 – 100 000 pg/mL

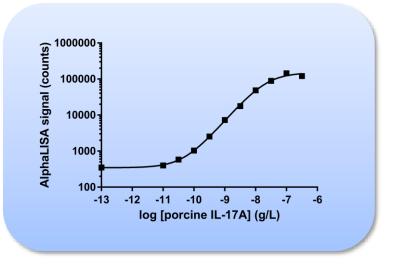


Figure 1. Typical sensitivity curve in AlphaLISA HiBlock Buffer. The data was generated using a white Optiplate<sup>™</sup>-384 microplate and the EnVision<sup>®</sup> Multilabel Plate Reader 2103 with Alpha option.

**Storage:** Store kit in the dark at +4°C. For reconstituted analyte aliquot and store at -20 °C. Avoid freeze-thaw cycles.

**Stability:** This kit is stable for at least 12 months from the manufacturing date when stored in its original packaging and the recommended storage conditions.

### **Analyte of Interest**

Interleukin 17A (IL-17A) is a 155 aa protein with a unique structure only shared with interleukin 17F. IL-17A acts as an early response component in inflammation. Upon detection of an abnormal element, macrophages will secrete interleukin 17A. This cytokine plays two major roles. The first is as a potent chemoattractant for neutrophils and NK killer cells. Its second role is to stimulate the secretion of inflammatory cytokines such as TGF $\beta$ , TNF $\alpha$ , IL2 and IL6 amongst others. These roles make IL17A a clear marker of inflammation in animals and also serve to measure early response to infection and/or vaccination. This kit has been designed for the detection of porcine IL17A in porcine serum

### **Description of the AlphaLISA Assay**

AlphaLISA technology allows the detection of molecules of interest in buffer, cell culture media, serum and plasma in a highly sensitive, quantitative, reproducible and user-friendly mode. In this AlphaLISA assay, a Biotinylated Anti-pIL-17A Antibody binds to the Streptavidin-coated Alpha Donor beads, while another Anti-pIL-17A Antibody is conjugated to AlphaLISA Acceptor beads. In the presence of the pIL-17A, the beads come into close proximity. The excitation of the Donor beads provokes the release of singlet oxygen molecules that triggers a cascade of energy transfer in the Acceptor beads, resulting in a sharp peak of light emission at 615 nm (Figure 2).

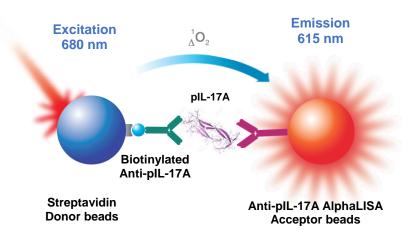


Figure 2. AlphaLISA Assay Principle.

### **Precautions**

- The Alpha Donor beads are light-sensitive. All the other assay reagents can be used under normal light conditions. All Alpha assays using the Donor beads should be performed under subdued laboratory lighting (< 100 lux). Green filters (LEE 090 filters (preferred) or Roscolux filters #389 from Rosco) can be applied to light fixtures.</li>
- Take precautionary measures to avoid contamination of the reagent solutions.
- The Biotinylated Anti-Analyte Antibody contains sodium azide. Contact with skin or inhalation should be avoided.

# **Kit Content: Reagents and Materials**

Kit components	AL572HV (100 assay points***)	AL572C (500 assay points***)	AL572F (5000 assay points***)
AlphaLISA Anti-pIL-17A Acceptor beads stored in PBS, 0.05% Kathon, pH 7.2	20 μL @ 5 mg/mL (1 brown tube, <u>white</u> cap)	50 μL @ 5 mg/mL (1 brown tube, <u>white</u> cap)	500 µL @ 5 mg/mL (1 brown tube, <u>white</u> cap)
Streptavidin (SA)-coated Donor beads stored in 25 mM HEPES, 100 mM NaCl, 0.05% Kathon, pH 7.4	80 μL @ 5 mg/mL (1 brown tube, <u>black</u> cap)	200 µL @ 5 mg/mL (1 brown tube, <u>black</u> cap)	2 X 1 mL @ 5 mg/mL (2 brown tubes, <u>black</u> caps)
Biotinylated Anti-pIL-17A Antibody stored in PBS, 0.1% Tween-20, 0.05% NaN <sub>3</sub> , pH 7.4	20 µL @ 500 nM (1 tube, <u>black</u> cap)	50 μL @ 500 nM (1 tube, <u>black</u> cap)	500 μL @ 500 nM (1 tube, <u>black</u> cap)
Lyophilized Recombinant Porcine IL-17A*	1 μg (1 tube, <u>clear</u> cap)	1 μg (1 tube, <u>clear</u> cap)	1 μg (1 tube, <u>clear</u> cap)
AlphaLISA HiBlock Buffer (10X)** (1)	2 mL, 1 small bottle	10 mL, 1 medium bottle	100 mL, 1 large bottle

- \* Reconstitute lyophilized analyte in 100 µL Milli-Q<sup>®</sup> grade H<sub>2</sub>O. The reconstituted analyte should be used within 60 minutes or aliquoted into screw-capped polypropylene vials and stored at -20°C for future experiments. Avoid freeze-thaw cycles. One vial contains an amount of analyte sufficient for performing 10 standard curves. Additional vials can be ordered separately (cat # AL572S).
- \*\* Extra buffer can be ordered separately (cat # AL004C: 10 mL, cat # AL004F: 100 mL).
- (1) HiBlock Buffer may appear cloudy, especially after storage at cold temperature. Bring to room temperature prior to use and dilution.
- \*\*\* The number of assay points is based on an assay volume of 100 μL in 96-well plates or 50 μL in 96- or 384-well assay plates using the kit components at the recommended concentrations.

Sodium azide should **not** be added to the stock reagents. High concentrations of sodium azide (> 0.001 % final in the assay) might decrease the AlphaLISA signal. Note that sodium azide from the Biotinylated Antibody stock solution will not interfere with the AlphaLISA signal (0.0001% final in the assay).

#### Specific additional required reagents and materials:

The following materials are recommended:

Item	Suggested source	Catalog #
TopSeal <sup>™</sup> -A Plus Adhesive Sealing Film	Revvity Inc.	6050185
EnVision <sup>®</sup> -Alpha Reader	Revvity Inc.	-

### Recommendations

#### IMPORTANT: PLEASE READ THE RECOMMENDATIONS BELOW BEFORE USE

- The volume indicated on each tube is guaranteed for single pipetting. Multiple pipetting of the reagents may reduce the theoretical amount left in the tube. To minimize loss when pipetting beads, it is preferable not to pre-wet the tip.
- Centrifuge all tubes (including lyophilized analyte) before use to improve recovery of content (2000g, 10-15 sec). Re-suspend all reagents by vortexing before use.
- Use Milli-Q<sup>®</sup> grade H<sub>2</sub>O (18 MΩ•cm) to dilute 10X AlphaLISA HiBlock Buffer and to reconstitute the lyophilized analyte.
- When diluting the standard or samples, <u>change tips</u> between each standard or sample dilution. When loading
  reagents in the assay microplate, <u>change tips</u> between each standard or sample addition and after each set
  of reagents.
- When reagents are added to the microplate, make sure the liquids are at the bottom of the well.
- Small volumes may be prone to evaporation. It is recommended to cover microplates with TopSeal-A Adhesive Sealing Films to reduce evaporation during incubation. Microplates can be read with the TopSeal-A Film in place.
- The AlphaLISA signal is detected with an EnVision Multilabel Plate Reader equipped with the Alpha option using the AlphaScreen standard settings (e.g. Total Measurement Time: 550 ms, Laser 680 nm Excitation Time: 180 ms, Mirror: D640as, Emission Filter: M570w, Center Wavelength 570 nm, Bandwidth 100 nm, Transmittance 75%).
- AlphaLISA signal will vary with temperature and incubation time. For consistent results, identical incubation times and temperature should be used for each plate.
- The standard curves shown in this technical data sheet are provided for information only. A standard curve must be generated for each experiment.

### Assay Procedure

- The manual described below is an example for generating one standard curve in a 50 µL final assay volume (48 wells, triplicate determinations). The manuals also include testing samples in 452 wells. If different amount of samples are tested, <u>the volumes of all reagents have to be adjusted accordingly</u>, as shown in the table <u>below</u>. These calculations do not include excess reagent to account for losses during transfer of solutions or dead volumes.
- The standard dilution manual is provided for information only. As needed, the number of replicates or the range of concentrations covered can be modified.
- Use of four background points in triplicate (12 wells) is recommended when LDL/LLOQ is calculated. One background point in triplicate (3 wells) can be used when LDL/LLOQ is not calculated.

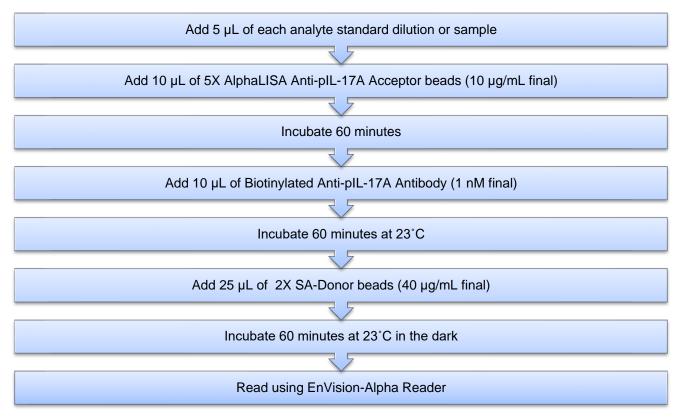
			Volume				
Format	# of data points	Final	Sample	AlphaLISA Acceptor Beads	Biotinylated Antibody	SA- Donor beads	Plate recommendation
AL572HV	100	100 µL	10 µL	20 µL	20 µL	50 µL	White OptiPlate-96 (cat # 6005290) White ½ AreaPlate-96 (cat # 6005560)
	250	100 µL	10 µL	20 µL	20 µL	50 µL	White OptiPlate-96 (cat # 6005290) White ½ AreaPlate-96 (cat # 6005560)
AL572C	500	50 µL	5 µL	10 µL	10 µL	25 µL	White ½ AreaPlate-96 (cat # 6005560) White OptiPlate-384 (cat # 6007290) Light gray AlphaPlate <sup>™</sup> -384 (cat # 6005350)
	1 250	20 µL	2 µL	4 µL	4 µL	10 µL	Light gray AlphaPlate-384 (cat # 6005350) ProxiPlate <sup>™</sup> -384 Plus (cat # 6008280) White OptiPlate-384 (cat # 6007290)
	2 500	10 µL	1 µL	2 µL	2 µL	5 µL	Light gray AlphaPlate-1536 (cat # 6004350)
	5 000	50 µL	5 µL	10 µL	10 µL	25 µL	White ½ AreaPlate-96 (cat # 6005560) White OptiPlate-384 (cat # 6007290) Light gray AlphaPlate-384 (cat # 6005350)
AL572F	12 500	20 µL	2 µL	4 µL	4 µL	10 µL	Light gray AlphaPlate-384 (cat # 6005350) ProxiPlate-384 Plus (cat # 6008280) White OptiPlate-384 (cat # 6007290)
	25 000	10 µL	1 µL	2 µL	2 µL	5 µL	Light gray AlphaPlate-1536 (cat # 6004350)

3 Step Manual described below is for 500 assay points including one standard curve (48 wells) and samples (452 wells). If different amount of samples are tested, the volumes of all reagents have to be adjusted accordingly.

- 1) <u>Preparation of 1X AlphaLISA HiBlock Buffer</u>: Add 5 mL of 10X AlphaLISA HiBlock Buffer to 45 mL Milli-Q<sup>®</sup> grade H<sub>2</sub>O.
- 2) <u>Preparation of plL-17A analyte standard dilutions</u>:
  - a. Reconstitute lyophilized pIL-17A (1  $\mu$ g) in 100  $\mu$ L Milli-Q<sup>®</sup> grade H<sub>2</sub>O.
  - b. Prepare standard dilutions as follows in 1X AlphaLISA HiBlock Buffer (change tip between each standard dilution):

Take	Vol. of	Vol. of	[pIL-17A] in standard curve	
Tube	pIL-17Α (μL)	diluent (µL) *	(g/mL in 5 µL)	(pg/mL in 5 µL)
A	10 µL of reconstituted pIL-17A	90	1.00E-06	1 000 000
В	60 µL of tube A	140	3.00E-07	300 000
С	60 µL of tube B	120	1.00E-08	100 000
D	60 μL of tube C	140	3.00E-08	30 000
E	60 μL of tube D	120	1.00E-08	10 000
F	60 μL of tube E	140	3.00E-09	3 000
G	60 μL of tube F	120	1.00E-09	1 000
Н	60 μL of tube G	140	3.00E-10	300
I	60 μL of tube H	120	1.00E-10	100
J	60 μL of tube I	140	3.00E-11	30
K	60 μL of tube J	120	1.00E-11	10
L	60 μL of tube K	140	3.00E-12	3
M ** (background)	0	100	0	0
N ** (background)	0	100	0	0
O ** (background)	0	100	0	0
P ** (background)	0	100	0	0

- Dilute standards in diluent (e.g. 1X AlphaLISA HiBlock Buffer).
   At low concentrations of analyte, a significant amount of analyte can bind to the vial. Therefore, load the analyte standard dilutions in the assay microplate within 60 minutes of preparation.
- \*\* Four background points in triplicate (12 wells) are used when LDL is calculated. If LDL does not need to be calculated, one background point in triplicate can be used (3 wells).
- 3) Preparation of 5X Anti-pIL-17A AlphaLISA Acceptor beads (50 µg/mL):
  - a. Prepare just before use.
  - b. Add 50 µL Anti-pIL-17A Acceptor beads to 4950 µl of 1X AlphaLISA HiBlock Buffer.
- 4) Preparation of 5X biotinylated Anti-pIL-17A antibody (5 nM):
  - a. Prepare just before use.
  - b. Add 50 µL 500 nM Biotinylated Anti-pIL-17A Antibody to 4950 µl of 1X AlphaLISA HiBlock Buffer.
- 5) Preparation of 2X Streptavidin (SA) Donor beads (80 µg/mL):
  - a. Prepare just before use.
  - b. Keep the beads under subdued laboratory lighting.
  - c. Add 200  $\mu L$  of 5 mg/mL SA-Donor beads to 12 300  $\mu L$  of 1X AlphaLISA HiBlock Buffer.



# **Data Analysis**

- Calculate the average count value for the background wells.
- Generate a standard curve by plotting the AlphaLISA counts versus the concentration of analyte. A log scale can be used for either or both axes. No additional data transformation is required.
- Analyze data according to a nonlinear regression using the 4-parameter logistic equation (sigmoidal dose-response curve with variable slope) and a 1/Y<sup>2</sup> data weighting (the values at maximal concentrations of analyte after the hook point should be removed for correct analysis).
- The LDL is calculated by interpolating the average background counts (12 wells without analyte) + 3 x standard deviation value (average background counts + (3xSD)) on the standard curve.
- The LLOQ as measured here is calculated by interpolating the average background counts (12 wells without analyte) + 10 x standard deviation value (average background counts + (10xSD)) on the standard curve. Alternatively, the true LLOQ can be determined by spiking known concentrations of analyte in the matrix and measuring the percent recovery, and then determining the minimal amount of spiked analyte that can be quantified within a given limit (usually +/- 20% or 30% of the real concentration).
- Read from the standard curve the concentration of analyte contained in the samples.
- If samples have been diluted, the concentration read from the standard curve must be multiplied by the dilution factor.

# **Assay Performance Characteristics**

AlphaLISA assay performance described below was determined using the 3 step manual using AlphaLISA HiBlock Buffer as assay buffer. The analytes (standards) were prepared in HiBlock, DMEM, RPMI, HiBlock Buffer + 10% FBS ,or 100% FBS and all other components were prepared in HiBlock.

#### Assay Sensitivity:

The LDL was calculated as described above. The values correspond to the lowest concentration of analyte that can be detected in a volume of 5  $\mu$ L sample using the recommended assay conditions.

LDL (pg/mL)*	(Analyte diluent)	# of experiments
3.8	HiBlock	6
12.3	DMEM	6
74.2	RPMI **	6
6.7	HiBlock Buffer + 10% FBS	6
7.8	100% FBS	6

- Note that LDL can be decreased (i.e. sensitivity increased) by increasing the volume of analyte in the assay (e.g. use 10 µL of analyte in a final assay volume of 50 µL).
- \*\* RPMI has a large impact on signal and sensitivity and is not recommended for this assay.
  - Assay Precision:

The following assay precision data were calculated from the three independent assays using two different kit lots. In each lot, the analytes were prepared in HiBlock, DMEM, HiBlock Buffer + 10% FBS or 100% FBS. All other components were prepared in HiBlock. Each assay consisted of one standard curve comprising 12 data points (each in triplicate) and 12 background wells (no analytes). The assays were performed in 384-well format.

• Intra-assay precision:

The intra-assay precision was determined using a total of 16 independent determinations in triplicate with 30 ng/mL samples. Shown as CV%.

plL-17A	HiBlock	DMEM	HiBlock Buffer + 10% FBS	100% FBS
CV (%)	5	2	3	1

• Inter-assay precision:

The inter-assay precision was determined using a total of 3 independent determinations with 9 measurements for 30 ng/mL sample. Shown as CV%.

plL-17A	HiBlock	DMEM	HiBlock Buffer + 10% FBS	100% FBS
CV (%)	7	11	7	15

#### • Spike Recovery:

Three known concentrations of analyte were spiked into HiBlock, DMEM, HiBlock Buffer + 10% FBS, or 100% FBS. All samples, including non-spiked diluents were measured in the assay. Note that the analytes for the respective standard curves were prepared in HiBlock, DMEM, HiBlock Buffer + 10% FBS, or 100% FBS. All other assay components were diluted in HiBlock.

Spiked	% Recovery       HiBlock     DMEM       HiBlock FBS     100%			
pIL-17A (ng/mL)				
10	95	101	89	112
3	108	97	94	106
1	88	93	79	97

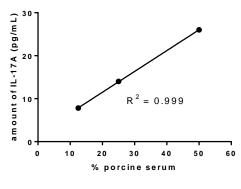
### <u>Specificity:</u>

Cross-reactivity of the pIL-17A AlphaLISA Kit was tested using the following proteins at 100 ng/mL in HiBlock.

Tested Proteins	% Cross Reactivity
Porcine IL-17F	0
Bovine IL-17A	1.3
Human IL-17A	0.4

### • Porcine Serum Experiments

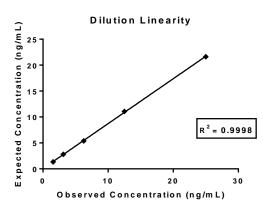
Detection of IL-17A in porcine serum



o Dilution Linearity

A pIL-17A-spiked (50 ng/mL) Normal Porcine Serum sample was diluted with 100% FBS and the assay was performed along with a standard curve using the analyte prepared in 100% FBS. Concentrations of pIL-17A in diluted porcine serum were determined by interpolating to the standard curve. Excellent dilution linearity ( $R^2 > 0.9998$ ) was achieved in the pIL-17A-spiked porcine serum samples that were diluted  $\ge 2$  fold. Endogenous levels were subtracted from observed results. The results are shown in table and figure below.

Dilution Factor (x)	Expected plL-17A (ng/mL)	Observed pIL- 17A (ng/mL)
1	50	44
2	25.0	26
4	12.5	14.8
8	6.25	7.9
16	3.12	4
32	1.56	1.8
64	0.8	0.88
128	0.4	0.41
256	0.2	0.19
512	0.1	0.07
0	0	0



### o Spike Recovery

Three known amounts of pIL-17A were spiked into Normal Porcine Serum (10, 3, and 1 ng/mL pIL-17A in spiked samples) and then the samples were diluted 2-fold into 100% FBS. The standard was prepared in 100% FBS and all other reagents were prepared in HiBlock. The spike recoveries of pIL-17A were determined and the results are shown in table below.

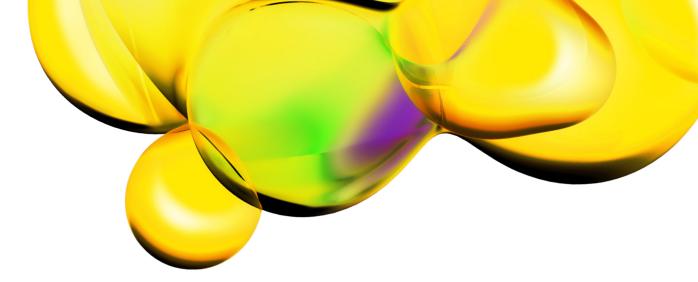
	Diluent: 100% FBS			
	Spiked sample (Normal Porcine Seru			
Spike (ng/mL)	Concentration (ng/mL)	Recovery (%)		
No spike	0.025	N/A		
30	34	113		
10	8.8	88		
3	2.5	82		

Excellent recovery was achieved for all three spikes tested.

## **Troubleshooting Guide**

You will find detailed recommendations for common situations you might encounter with your AlphaLISA Assay kit at: <u>www.revvity.com</u>

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Revvity, Inc. 940 Winter Street Waltham, MA 02451 USA www.revvity.com

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