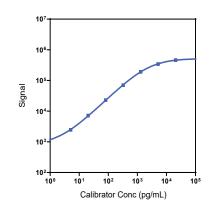
# **U**-PLEX<sup>®</sup> NHP MIP-3α

	Product Options	Catalog Number	Description				
om®	Multiplex	K15068M, K25068M	U-PLEX Biomarker Group 1 (NHP)				
		K156UZK-1/-2/-4	U-PLEX NHP MIP-3 $\alpha$ Assay with SECTOR <sup>TM</sup> plates				
	Singleplex	K156UZK-21/-22/-24	U-PLEX NHP MIP-3 $\alpha$ Assay with QuickPlex® plates				
		K256UZK-2/-4	U-PLEX NHP MIP-3 $\alpha$ Assay with 384-well plates				
	Antibody Set	B26UZ-2/-3	U-PLEX NHP MIP-3 $\alpha$ Antibody Set				
	Assay Protocol	tocol U-PLEX Product Inserts are available at <u>www.mesoscale.com</u>					

The U-PLEX<sup>®</sup> platform was designed to provide ultimate flexibility for detection of biomarkers in a wide variety of sample types. This datasheet provides the representative performance of the U-PLEX NHP MIP-3 $\alpha$  Assay tested on U-PLEX 96-well SECTOR plates run as a multiplex. The data do not represent the product specifications. Under your experimental conditions, the assay may perform differently from the representative data. U-PLEX assays are offered in either singleplex or multiplex; both are available in 96- or 384-well plates. See a U-PLEX product insert for instrument compatibility.

### Representative Calibration Curve and Sensitivity



Assay	Median LLOD (pg/mL)	LLOD Range (pg/mL)		
MIP-3a	0.27	0.24-0.41		

The Calibrator curve was fitted with a 4-parameter logistic model with a  $1/Y^2$  weighting. The lower limit of detection (LLOD) is a calculated concentration corresponding to 2.5X the standard deviations above the background (zero Calibrator).

### Precision

	Control	Average Conc. (pg/mL)	Average Intra-run Conc. %CV	Inter-run Conc. %CV
	High	NA	NA	NA
MIP-3α	Mid	957	5.7	14.2
	Low	95	5.3	14.3

For Research Use Only. Not for use in diagnostic procedures. Controls were made by spiking Calibrator into assay diluent at 3 levels within the quantitative range of the assay. Average intra-run concentration %CV is the average %CV of the control replicates within an individual run. Inter-run concentration %CV is the variability of controls across multiple runs. NA = not applicable due to 0% detected

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## Ordering Information

WW

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#### Scientific Support

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#### **Company Address**

MESO SCALE DISCOVERY® A division of Meso Scale Diagnostics, LLC. 1601 Research Boulevard Rockville, MD 20850-3173 USA

#### Spike Recovery

		Serum (N=5)		Plasma	a (N=5)	Cell Culture Media (N=5)		
	Spike Level	Average % Recovery	% Recovery Range	Average % Recovery	% Recovery Range	Average % Recovery	% Recovery Range	
Ormanialaura	High	59	39-83	60	55-69	88	75-104	
Cynomolgus Monkey	Mid	53	33-74	52	38-63	72	49-92	
WORKEy	Low	45	26-63	45	26-62	87	83-96	
Dheeue	High	52	47-63	73	50-87	88	75-104	
Rhesus Monkey	Mid	49	45-62	73	49-86	72	49-92	
wonkey	Low	45	39-51	67	44-81	87	83-96	

Normal serum, EDTA plasma, and cell culture media were spiked with Calibrator at 3 levels. Undiluted samples were tested to determine the expected concentration of the analyte. Samples may benefit from additional dilution with assay diluent to reduce matrix effects.

% Recovery = (measured concentration / expected concentration) x 100

## **Tested Samples**

	Sample Type	Serum (N=11)	Plasma (N=11)	Cell Culture Media (N=10)
	Median (pg/mL)	12.6	7.9	4.0
Cynomolgus Monkey	Range (pg/mL)	1.6-24	4.0-59	0.8-14
WORKCy	% Detected	100	100	100
	Median (pg/mL)	9.9	9.6	9.6
Rhesus Monkey	Range (pg/mL)	1.4-17	2.9-14	0.8-23
Wonkey	% Detected	100	100	100

Normal serum, EDTA plasma, and cell culture media were diluted 2-fold prior to the assay.

### **Dilution Linearity**

	Serum (N=5)			Plasma (N=5)			Cell Culture Media (N=5)		
	Fold Dilution	Average % Recovery	% Recovery Range	Fold Dilution	Average % Recovery	% Recovery Range	Fold Dilution	Average % Recovery	% Recovery Range
Oursemalause	2	124	110-138	2	124	107-135	2	136	126-147
Cynomolgus Monkey	4	152	122-188	4	148	137-168	4	140	128-148
WORKCy	8	160	119-216	8	154	141-188	8	153	140-160
Dhaava	2	124	103-137	2	124	108-160	2	136	126-147
Rhesus Monkey	4	133	107-163	4	133	112-176	4	140	128-148
Wonkey	8	141	112-175	8	137	106-176	8	153	140-160

Normal serum, EDTA plasma, and cell culture media were spiked with Calibrator and tested at different dilutions. Undiluted samples were tested to determine the expected concentration of the analyte. Samples may benefit from additional dilution with assay diluent to reduce matrix effects.

% Recovery = (measured concentration / expected concentration) x 100





## Specificity

To assess specificity, the MIP-3 $\alpha$  Antibody Set was tested individually against a larger panel of recombinant human analytes for nonspecific binding (CTACK, Eotaxin, Eotaxin-2, Eotaxin-3, ENA-78, FLT3L, Fractalkine, G-CSF, GM-CSF, GRO- $\alpha$ , I-309, IFN- $\alpha$ 2a, IFN- $\gamma$ , IL-1 $\alpha$ , IL-1 $\beta$ , IL-1RA, IL-2, IL-4, IL-5, IL-6, IL-7, IL-8, IL-9, IL-10, IL-12/IL-23p40, IL-12p70, IL-13, IL-15, IL-16, IL-17A, IL-17A/F, IL-17B, IL-17D, IL-17D, IL-17F, IL-18, IL-22, IL-23, IP-10, I-TAC, MCP-1, MCP-2, MCP-3, MCP-4, M-CSF, MDC, MIF, MIP-1 $\alpha$ , MIP-1 $\beta$ , MIP-3 $\alpha$ , MIP-3 $\beta$ , MIP-5, SDF-1 $\alpha$ , TARC, TNF- $\alpha$ , TNF- $\beta$ , TPO, TRAIL, VEGF-A, and YKL-40). Nonspecific binding was less than 0.5%.

MIP-3 $\alpha$  detection antibody interacts with capture antibodies for Eotaxin-2, Eotaxin-3, and IL-1 $\alpha$ , causing elevated background. Background at Eotaxin-3 exceeds 12,000 counts.

% Nonspecificity = (nonspecific signal / specific signal) x 100

## **Diluent Compatibility**

Diluents 57 and 3 are provided with this assay. MSD offers a range of assay and antibody diluents for separate purchase. Depending on your assay needs, other diluents may be tested.

## Assay Components

**Calibrator:** MIP- $3\alpha$  is included in Calibrator 4. The full-length recombinant protein is expressed in *E. coli*. **Antibodies:** The U-PLEX NHP MIP- $3\alpha$  Assay uses a rabbit polyclonal antibody for capture and a goat polyclonal antibody for detection. **Assay generation:** A

Note: This datasheet contains representative assay performance data. In custom multiplex formats, the assay may perform differently than the representative data shown.

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