Human GIP (active)

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MSD Customer Service Phone: 1-240-314-2795 Fax: 1-301-990-2776 Email: CustomerService@ mesoscale.com

Scientific Support

Phone: 1-240-314-2798 Email: ScientificSupport@ mesoscale.com

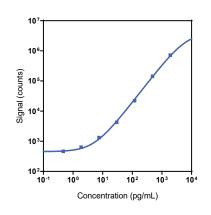
Company Address

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

	Product Options	Catalog Number	Description				
Ŋ®	Multiplex	K151ACM, K251ACM	U-PLEX Metabolic Group 1 (human)				
		K1516NK-1/-2/-4	U-PLEX Human GIP (active) Assay with SECTOR [™] plates				
	Singleplex	K1516NK-21/-22/-24	U-PLEX Human GIP (active) Assay with QuickPlex® plates				
		K2516NK-2/-4	U-PLEX Human GIP (active) Assay with 384-well plates				
	Antibody Set	B216N-2/-3	U-PLEX Human GIP (active) Antibody Set				
	Protocol	U-PLEX Product Inserts are available at <u>www.mesoscale.com</u>					

The U-PLEX[®] 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 Human GIP (active) 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 on 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)		
GIP (active)	1.3	1.0-1.3		

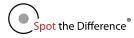
The Calibrator curve was fitted with a 4-parameter logistic model with a $1/Y^2$ weighting. The lower limit of detection (LLOD) is a calculated concentration corresponding to 2.5 standard deviations above the background (zero Calibrator).

Precision

Control Average Conc. (pg/mL)		Average Intra-run Conc. (%CV)	Inter-run Conc. (%CV)		
High	757	2.9	100		
Mid	212	2.6	13.3		
Low	76	4.2	17.0		

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.

For Research Use Only. Not for use in diagnostic procedures.





MSD® U-PLEX Human GIP (active)

Tested Samples

Sample Type	Serum (N=12)	EDTA Plasma (N=12)	P800 Plasma (N=8)
Median (pg/mL)	ND	2.5	32
Range (pg/mL)	NA	ND-2.5	7.7-93
% Detected	0	8	100

Normal serum, EDTA plasma, and P800 plasma samples were diluted 4-fold prior to the assay. ND = non-detectable (<LLOD); NA = not applicable due to 0% detected

Dilution Linearity

	Serum		EDTA Plasma P800 Plasma Cell Culture Media			A Plasma P800 Plasma			ledia		
Fold Dilution	Average % Recovery	% Recovery Range	Fold Dilution	Average % Recovery	% Recovery Range	Fold Dilution	Average % Recovery	% Recovery Range	Fold Dilution	Average % Recovery	% Recovery Range
2	79	57-101	2	91	72-102	2	89	80-96	2	83	69-96
8	113	102-139	8	102	99-108	8	102	98-106	8	112	95-138
16	115	85-163	16	103	99-114	16	107	98-114	16	123	98-156

Normal human serum, EDTA plasma, P800 plasma, and cell culture media were spiked with Calibrator and tested at different dilutions. Percent recovery at each dilution level was normalized to the dilution-adjusted, 4-fold concentration. Samples may benefit from additional dilution with assay diluent to reduce matrix effects.

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

Spike Recovery

	Serum		EDTA Plasma		P800 I	Plasma	Cell Culture Media		
Spike Level	Average % Recovery	% Recovery Range	Average % % Recovery Recovery Range		Average % Recovery	% Recovery Range	Average % Recovery	% Recovery Range	
High	81	47-99	98	87-109	88	76-94	86	76-106	
Mid	80	49-96	99	87-106	90	86-94	88	78-104	
Low	85	52-99	101	89-110	93	89-95	90	80-105	

Normal serum, EDTA plasma, P800 plasma, and cell culture media were spiked with Calibrator at 3 levels. Spiked samples were diluted 4-fold 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 GIP (active) Antibody Set was tested individually against a larger panel of analytes for nonspecific binding (BAFF, BDNF, C-Peptide, CTACK, Desghrelin, ENA-78, Eotaxin, Eotaxin-2, Eotaxin-3, EPO, FGF-21, FGF-23, FLT3L, Fractalkine, FSH, G-CSF, Ghrelin (Ser3-octanoylated), GIP (1–42), GIP (3-42), GLP-1 (7–36), GLP-1 (9–36), GM-CSF, GRO- α , I-309, IFN- α 2a, IFN- β , IFN- γ , IL-1 α , IL-1 β , IL-1RA, IL-2, IL-2R α , IL-3, 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-17C, IL-17D, IL-17E/IL-25, IL-17F, IL-18, IL-21, IL-22, IL-23, IL-27, IL-29/IFN- λ 1, IL-33, Insulin, IP-10, I-TAC, Leptin, LH, MCP-1, MCP-2, MCP-4, M-CSF, MDC, MIF, MIP-1 α , MIP-1 β , MIP-5, PIGF, PP, Proinsulin, PYY (3-36), SDF-1 α , TNF- α , TNF- β , TPO, TRAIL, TSLP, VEGF-A, YKL-40, and β -NGF). Nonspecific binding was less than 2.0%.

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

The GIP (inactive) and GIP (total) assays will cross-react with the GIP (active) assay. We do not recommend multiplexing the GIP (active) assay with the GIP (inactive) or GIP (total) assays on the same plate.

Diluent Compatibility

The data included in this document were collected with Assay Diluent 13 (supplemented with 1,000 KIU/mL Aprotinin [provided] and 100 µM diprotin A [not provided]) and Antibody Diluent 11. MSD offers a range of assay and antibody diluents for separate purchase. Depending on your assay needs, other diluents may be tested. Diprotin A should be purchased separately.

Assay Components

Calibrator: GIP (active) is included in Human GIP (active) Calibrator. The human GIP (active) Calibrator is a synthetic peptide. Antibodies: The U-PLEX Human GIP (active) Assay uses a mouse monoclonal antibody for capture and a mouse monoclonal antibody for detection. Assay generation: A

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

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