# Sensitive Detection and Quantification of Low Abundant Cytokines Using Discovery's Multiplex Immunoassay Services 

## Introduction

The proteomic service laboratories at Discovery Life Sciences offer services based on the Lumine ${ }^{\circledR} \times \mathrm{MAP}$ technology for multiplexed immunoassays including assay customization, execution, and data analysis for high-sensitivity protein quantitation. Herein we demonstrate the success of our multiplexed immunoassay services using the ProcartaPlexTM Human High Sensitivity panel and the Luminex xMAP® INTELLIFLEX system to profile and compare plasma cytokine levels in male self-reported tobacco users to those who reported no tobacco use.

## Methods

This study analyzed a cohort of male tobacco users (both cigarettes and smokeless tobacco) compared to those who reported no tobacco use (Figure 1). Plasma samples were obtained from Discovery's biospecimen repository from consented donors and categorized according to their self-reported tobacco use: never used, previous use, current use of cigarettes, and current use of smokeless tobacco.

Cytokine levels in the plasma samples were measured using the High Sensitivity 9-Plex Human ProcartaPlexTM. The panel analyzes 9 protein targets in a single well using Luminex


Figure 1. Cohort metrics. Average ages of men who used smokeless tobacco $(n=6)$, cigarettes $(n=8)$ or were nonsmokers ( $n=10$ ). xMAP® ${ }^{\circledR}$ technology for highly sensitive detection of analytes. The assay was run according to the ProcartaPlexTM Multiplex Immunoassays user's guide1 with an overnight incubation (total time $=20$ hours). Each sample was analyzed in duplicate and the average concentration of each cytokine in the panel is shown in Table 2.

## Results:

- Absolute quantification of each target analyte was achieved by using a standard curve, which was generated by incorporating standard analytes into the assay. (Figure 2).
- Most analytes ( $88 \%$ ) were measured at values below LLOQ (lower limit of quantification) and above ULOQ (Upper limit of quantification) with Discovery's established protocol showing that our protocol outperforms the stated kit sensitivity and dynamic range.
- Differences in cytokine levels between tobacco users and non-users were observed (Table 2). In this small cohort study, statistically significant (Mann-Whitney rank test; $p<0.05$ ) differences for a subset of cytokines between reported cigarette users and non-users were noted (Figure 3).

Table 2: Cytokine measurement results.

| Analyte | LLOQ\|UPLOQ <br> Reported <br> COA | LLOQ\|UPLOQ <br> Experimental | Average <br> (pg/mL) |  |  |  |
| :--- | :---: | :---: | :---: | :---: | :---: | :---: |
|  | $(\mathbf{p g / m L})$ |  | Smokeless <br> Tobacco | Cigarettes | All Tobacco <br> [combined] | Non-Users |
| IFN Gamma | $1.26 \mid 1295$ | $0.32 \mid 5180$ | 0.9 | 1.1 | 1.0 | 0.6 |
| IL-1 beta | $0.28 \mid 1130$ | $0.07 \mid 1130$ | 1.6 | 2.0 | 1.8 | 0.8 |
| IL-10 | $0.17 \mid 690$ | $0.67 \mid 690$ | ND | ND | ND | ND |
| IL-12p70 | $0.77 \mid 3170$ | $0.2 \mid 3170$ | 6.2 | 7.2 | 6.7 | 3.3 |
| IL-17A (CTLA-8) | $0.30 \mid 1220$ | $0.07 \mid 1220$ | 1.3 | 1.5 | 1.4 | 0.6 |
| IL-2 | $0.88 \mid 3600$ | $0.22 \mid 3600$ | 2.6 | 3.5 | 3.1 | 1.8 |
| IL-4 | $1.29 \mid 5270$ | $0.32 \mid 5270$ | 1.3 | 1.8 | 1.6 | 0.7 |
| IL-6 | $1.29 \mid 1320$ | $0.32 \mid 5280$ | 2.5 | 6.8 | 4.8 | 1.7 |
| TNF alpha | $0.62 \mid 2520$ | $0.15 \mid 2520$ | 2.9 | 3.7 | 3.3 | 1.5 |



Figure 2. Example calibration curve for IL-12p70.


Figure 3. Statistically significant differences among groups for a subset of measure cytokines.

## Summary

The results from this small cohort study demonstrate Discovery's ability to profile cytokine levels in plasma samples with highly sensitive detection and qualification using Invitrogen ProcartaPlexTM cytokine assay on the Luminex xMAP® INTELLIFLEX system. The high sensitivity of this assay can accurately measure subtle changes in protein levels, and when combined with multiplexing capabilities, it allows for rapid profiling of cytokine levels in complex biological samples.

## Reference

1 ProcartaPlex Multiplex Immunoassay User Guide, Revision A. 0 (31.) April 07, 2020.
Publication Number MAN0019074

## Discovery offers Luminex®-Based Multiplex Immunoassay Services

Discovery has a full suite of multiplex immunoassays and optimized workflows for proteomic profiling services. Leverage Discovery's extensive proteomics expertise to deliver accurate and reliable results.

The Luminex xMAP® ${ }^{\circledR}$ INTELLIFLEX System is a compact, high throughput flow-based multiplex platform that performs simultaneous measurements on up to 80 protein or RNA targets from a small amount of sample with sensitivity to detect low-level analytes. Each target is compared to a unique standard curve to ensure accurate quantitation that covers 5.5 logs of dynamic range.

The ProcartaPlexTM multiplexed immunoassays make use of the Luminex xMAP® Technology to enable simultaneous analysis of multiple proteins and RNA in a single sample. Select from over 90 preconfigured multiplex panels (2- to 80 -plex) in 96 well plate formats or mix and match targets to create custom assays tailored to your needs.

| Luminex Description | Sample Types | Deliverables | Applications | Standard Sample Numbers | Standard Sample Volume | Variables for TAT and Price |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| - Immunoassay <br> - Bead-based ELISA <br> - Absolute quantification <br> - High sensitivity <br> - High throughput <br> - Preconfigured E customizable | - Plasma <br> - Serum <br> - CSF <br> - Tissue <br> - Cells <br> - Conditioned media | Absolute quantification of selected protein targets | - Proteomic profiling <br> - Biomarker profiling | Up to 80 per plate | $25 \mu \mathrm{~L}$ | Number of panels Number of samples |

## About Discovery Life Sciences

Discovery combines the world's largest commercial biospecimen inventory and procurement network with preeminent multi-omic biomarker services covering genomics, pathology, proteomics, cell biology, and ADME Tox to accelerate new therapies supported by biomarker and companion diagnostic programs for cancer, infectious disease, and other rare and complex conditions. HudsonAlpha Discovery, a division of Discovery, offers a wide spectrum of tools and techniques required for identification and validation of clinical biomarkers, including RNA, whole genome, whole exome, targeted panel, hi-fidelity long-read, single cell, and epigenomic next generation sequencing. Working with Discovery as a single strategic partner helps accelerate biomarker strategies with actionable data and enables the ability to customize study designs as required to take a more targeted and nuanced approach to biomarker discovery and validation.

