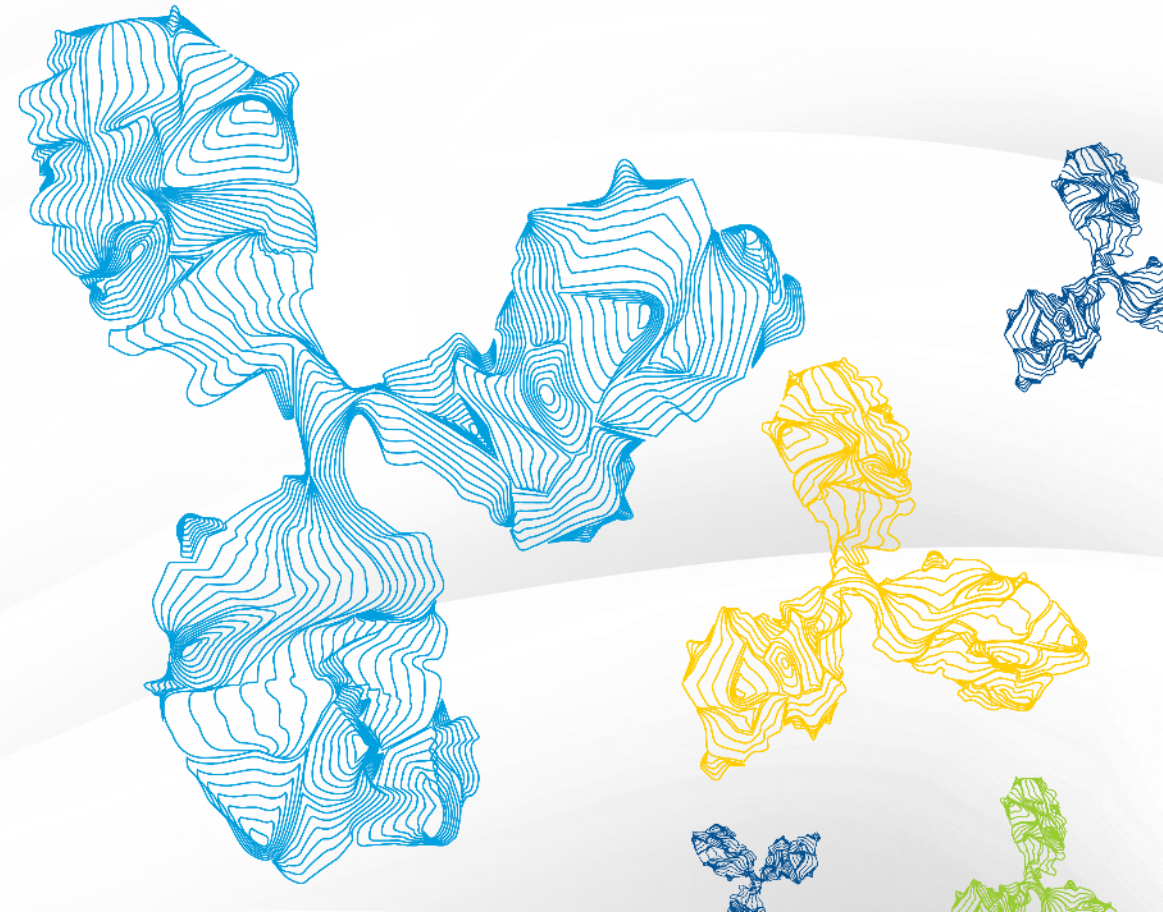


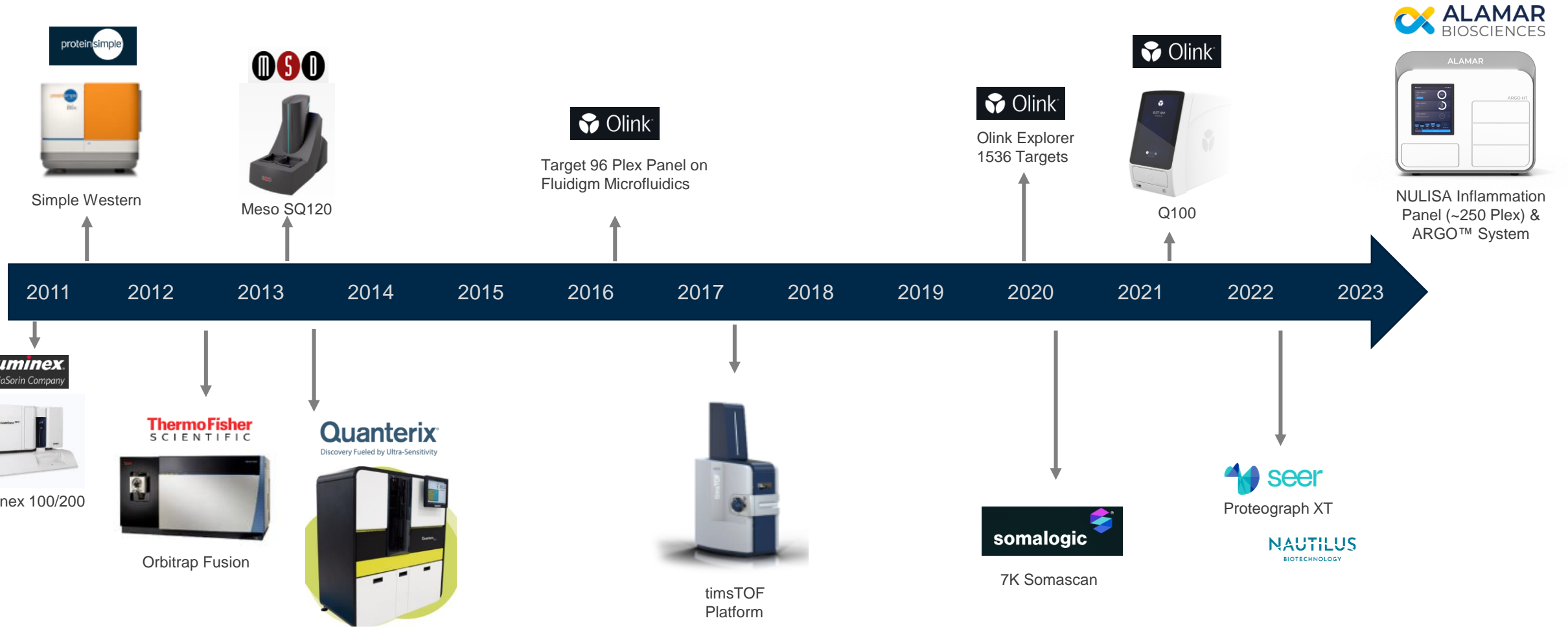
NULISA: Fully automated ultra-sensitive immunoassay platform for profiling fluid-based protein biomarkers

Doug Hinerfeld, Ph.D. Senior Director
of Application Support

WACD
October 13, 2023



Proteomics: The Next Frontier in Life Science Tools and Diagnostics



Alamar Biosciences



Mission: Power precision proteomics to enable the earliest detection of disease



Fast Facts

Founded
2018

Team
70+ Full-Time Employees

Based
San Francisco Bay Area

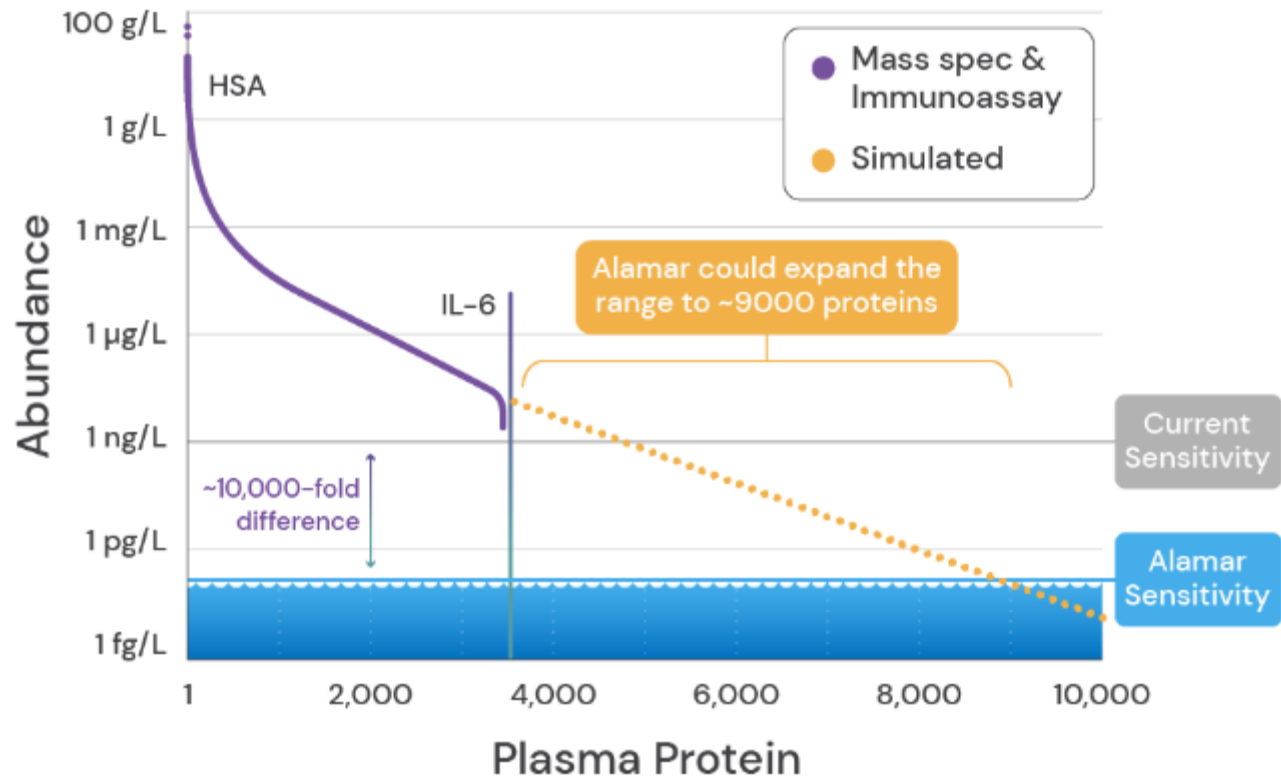
Stage
Development

Funding
\$120M Series A/B+

- **NULISA™** - ultra-sensitive Attomolar (10^{-18} M) NGS enabled digital immunoassay platform for protein analysis from 1 – 1000's of plex
- **Attobody™**: bi-paratopic antibodies with picomolar affinity
- **ARGO™** - Fully automated solutions to perform assay across a range of multiplex levels and applications from discovery to the clinic

Automated solutions for **ultra-sensitive** protein analysis across a range of **multiplex** level to detect **critical biomarkers** from discovery to the clinic.

NULISA™ platform enables greater sensitivity and precision in high-plex proteomics



- Best-in-class Attomolar (fg/ml) sensitivity increases the depth of measurable proteins
- Enabling detection of known biomarkers at lower levels
- Enabling discovery of novel biomarkers that are currently undetectable
- Enabling clearer definition of health baseline biomarker levels

NULISA™: the most sensitive immunoassay liquid biopsy platform



Unprecedented sensitivity

- Attomolar (fg/mL) level of detection
- Quantification of low abundance proteins
- Enables discovery of critical biomarkers

Widest dynamic range

- >12 logs dynamic range
- Quantification across biological range of protein levels
- In one reaction w/o dilution

Flexible, multiplex chemistry

- Single target or multiplex
- 100s to 1000s of targets in a single experiment
- Broad sample compatibility

Fully automated

- Ease of use
- Minimal hands-on time
- High reproducibility

Precision proteomics at the push of a button

Full solution of instrument, reagent kits, and software



ARGO™ HT System



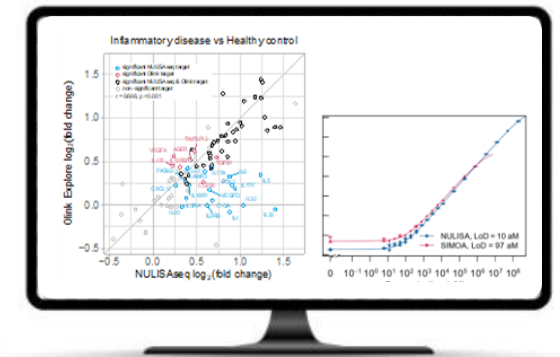
Consumables

Simple, robust, reproducible



Instrument

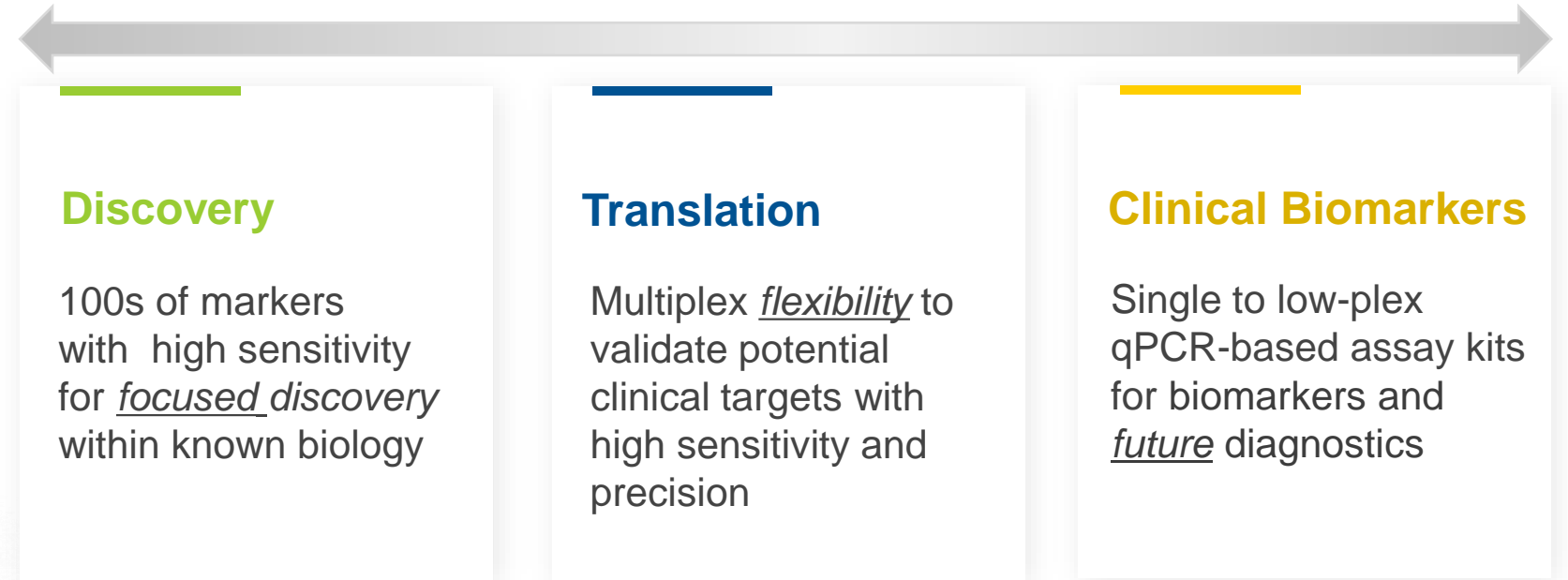
ARGO all-in-one instrument
w/ onboard qPCR



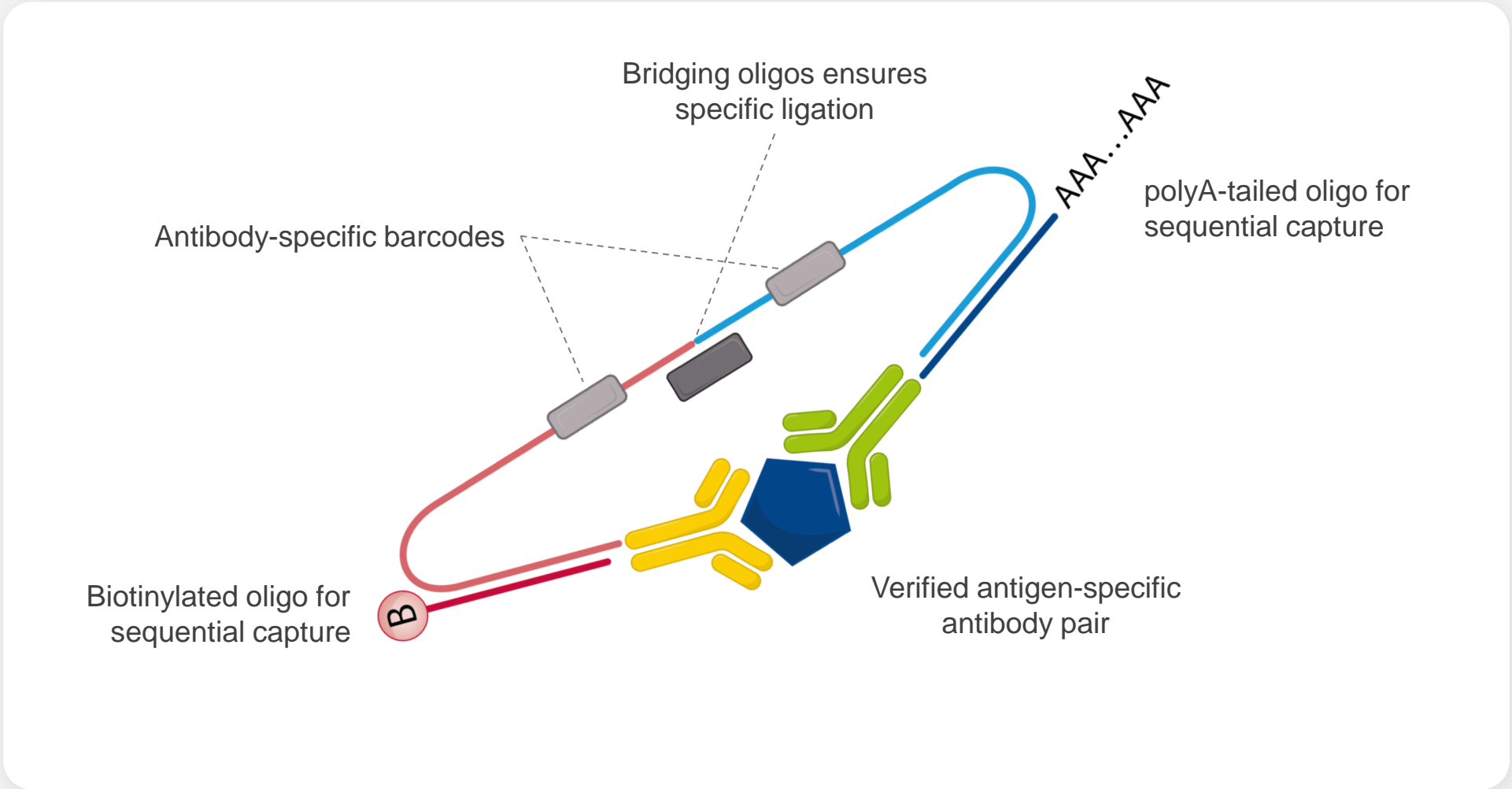
Software

From samples to insights

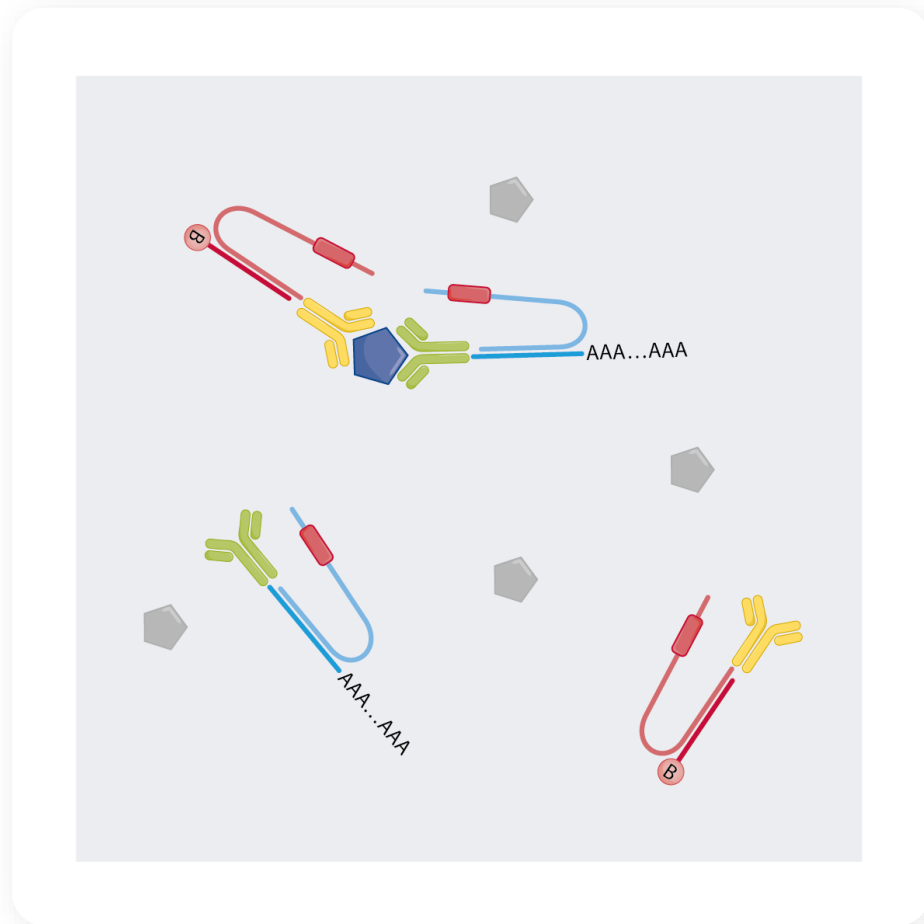
Scalability and Flexibility All-in-one instrument



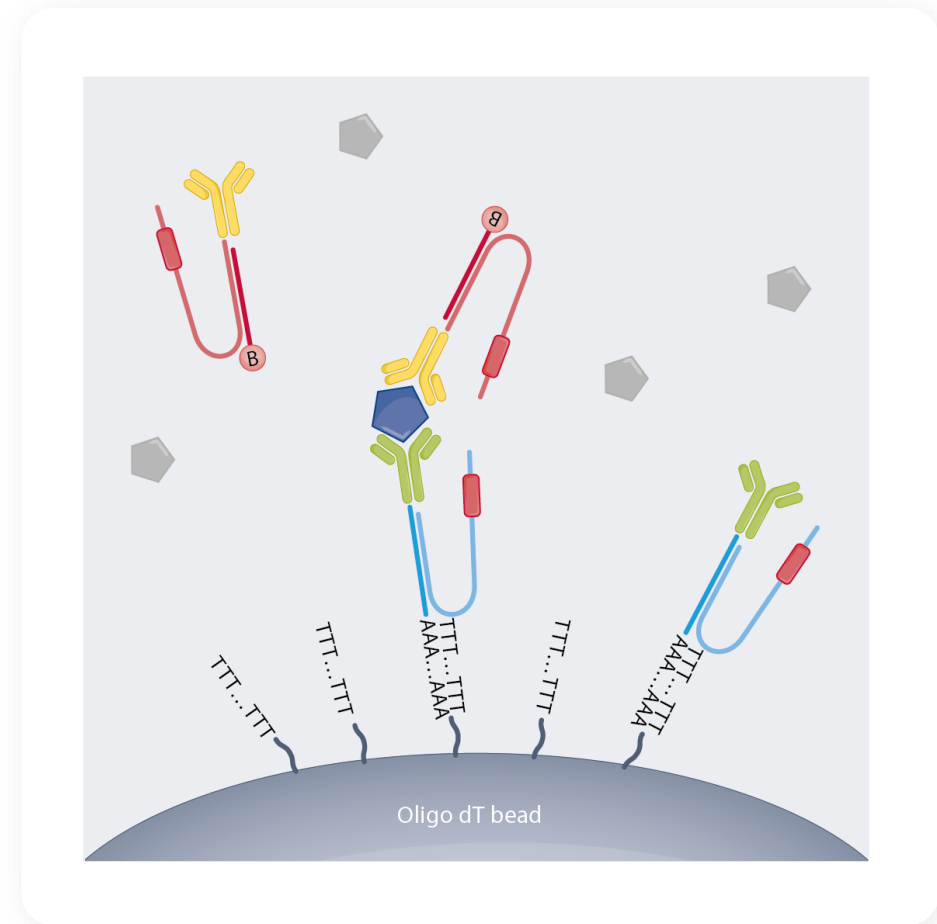
NULISA Technology and how it works?



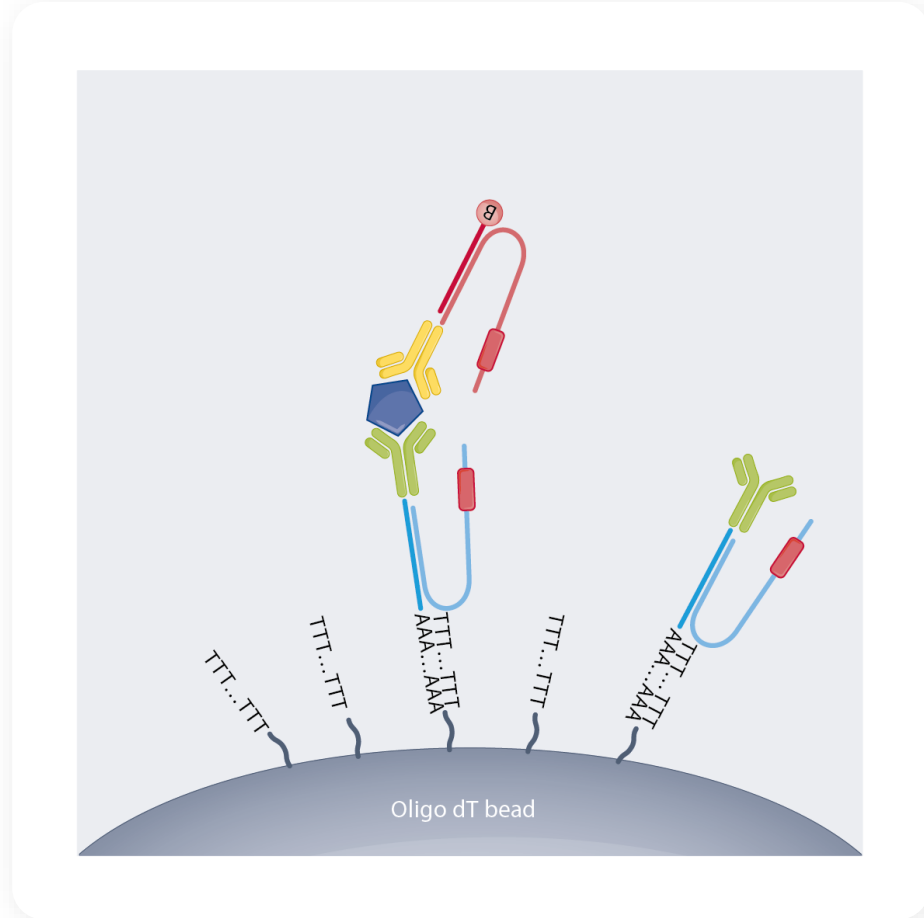
- **Step 1:** in solution antibody – antigen immune complex formation



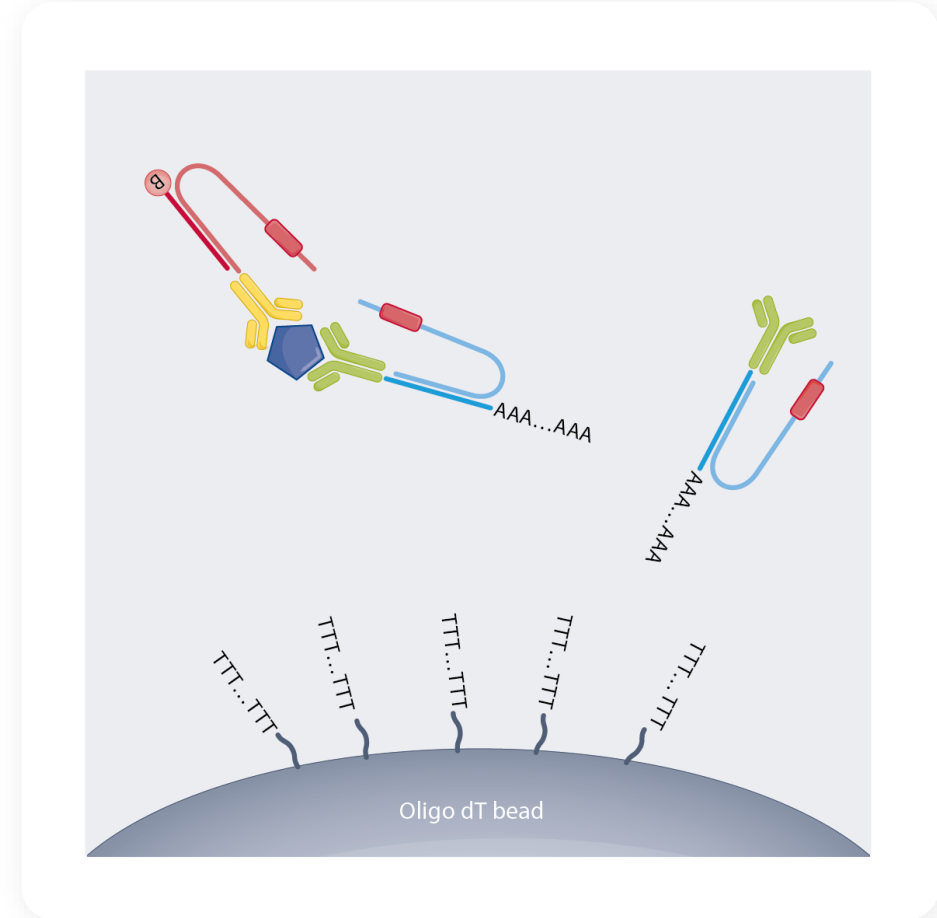
- **Step 2:** 1st bead-based capture – polyA-linked capture antibody on poly-dT beads



- **Step 3:** wash – removal of all free detection antibodies, analytes and matrix; retention of specific analyte bound by capture antibody



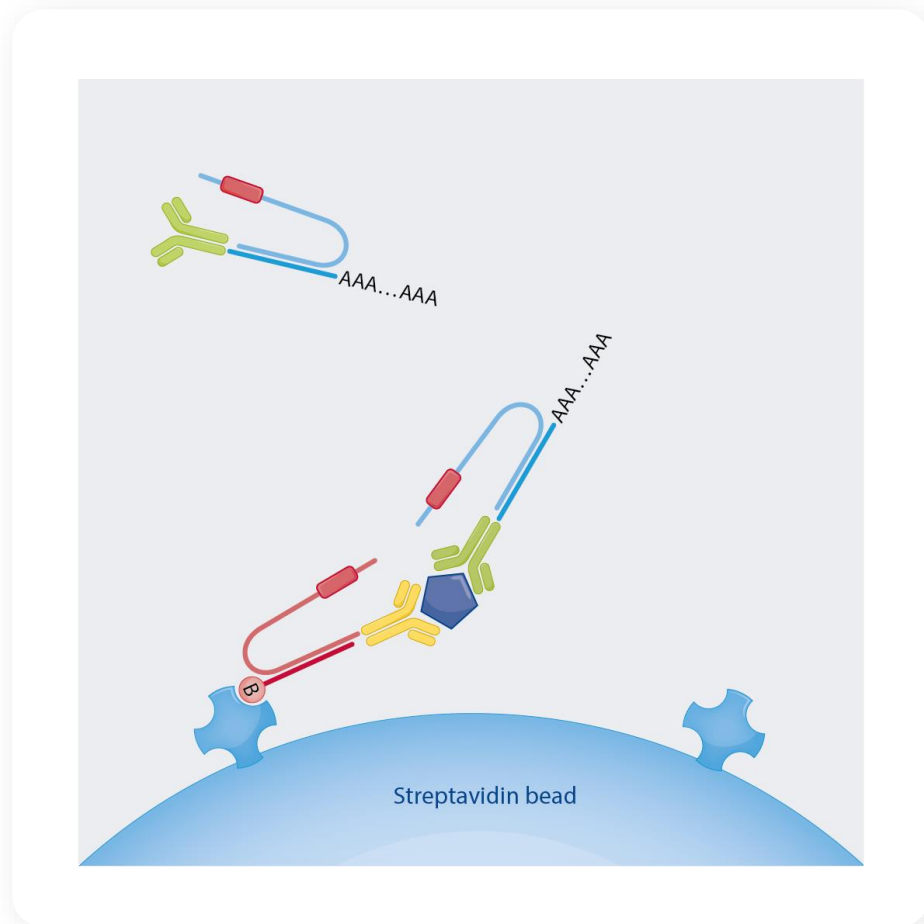
- **Step 4:** release of capture antibody/analyte complex



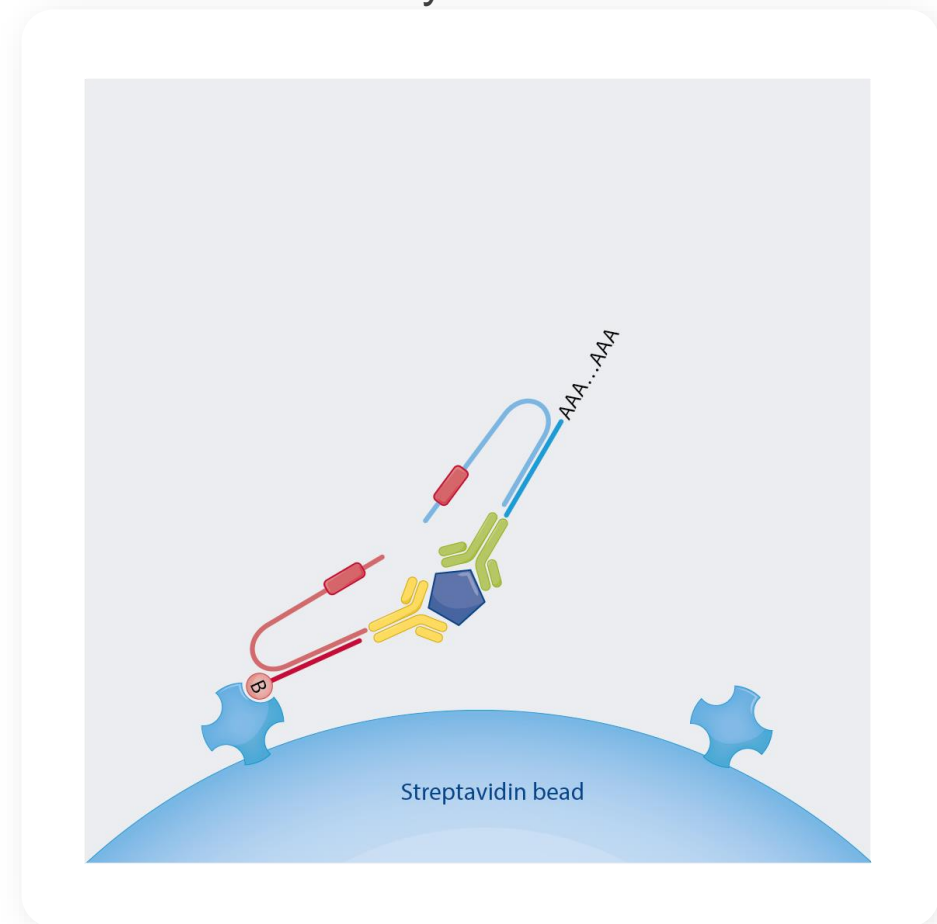
NULISA Workflow



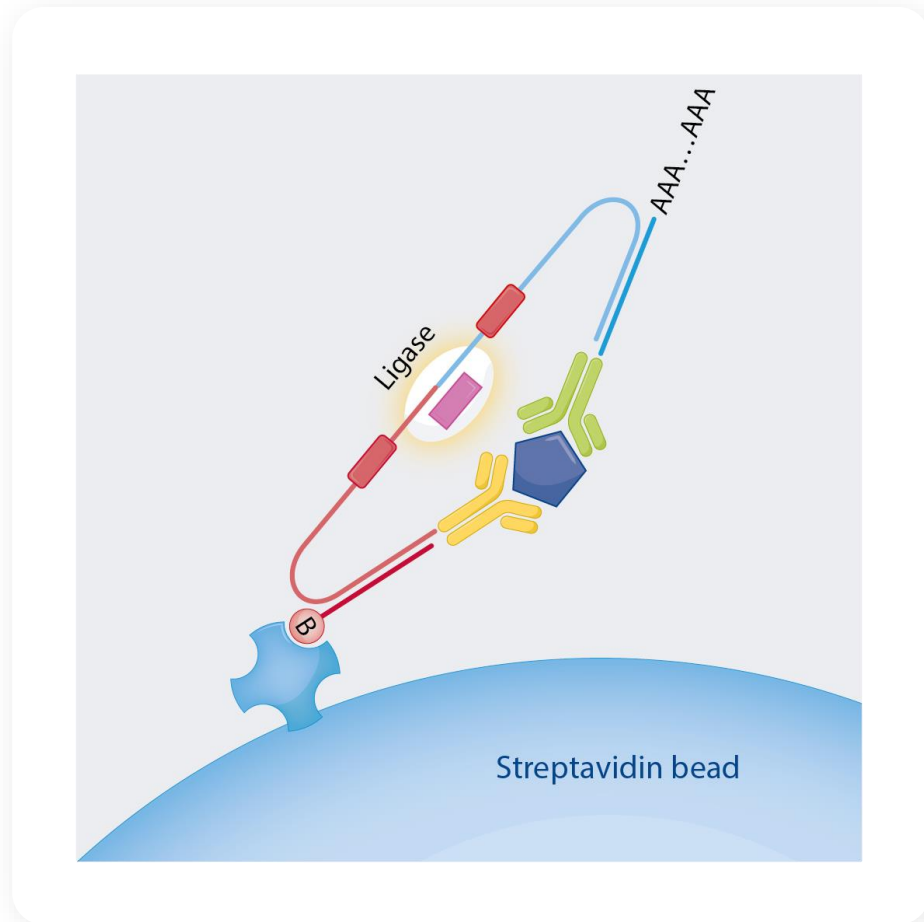
- **Step 5:** 2nd bead-based capture via biotinylated oligo binding to streptavidin



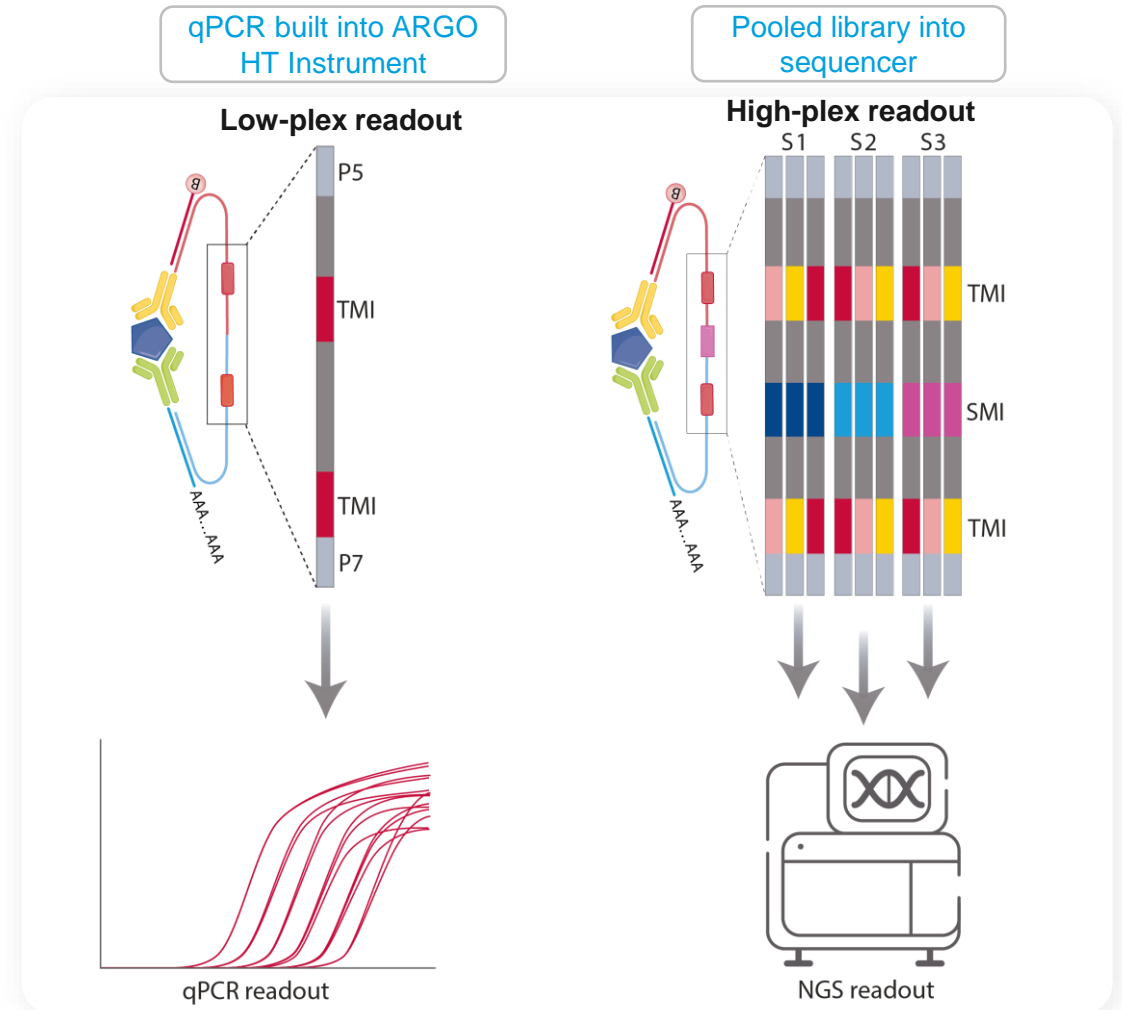
- **Step 6:** wash – removal of all antibodies and residual matrix not complexed with analyte and 2nd detection antibody



- **Step 7:** ligation of specific bar-coded oligos



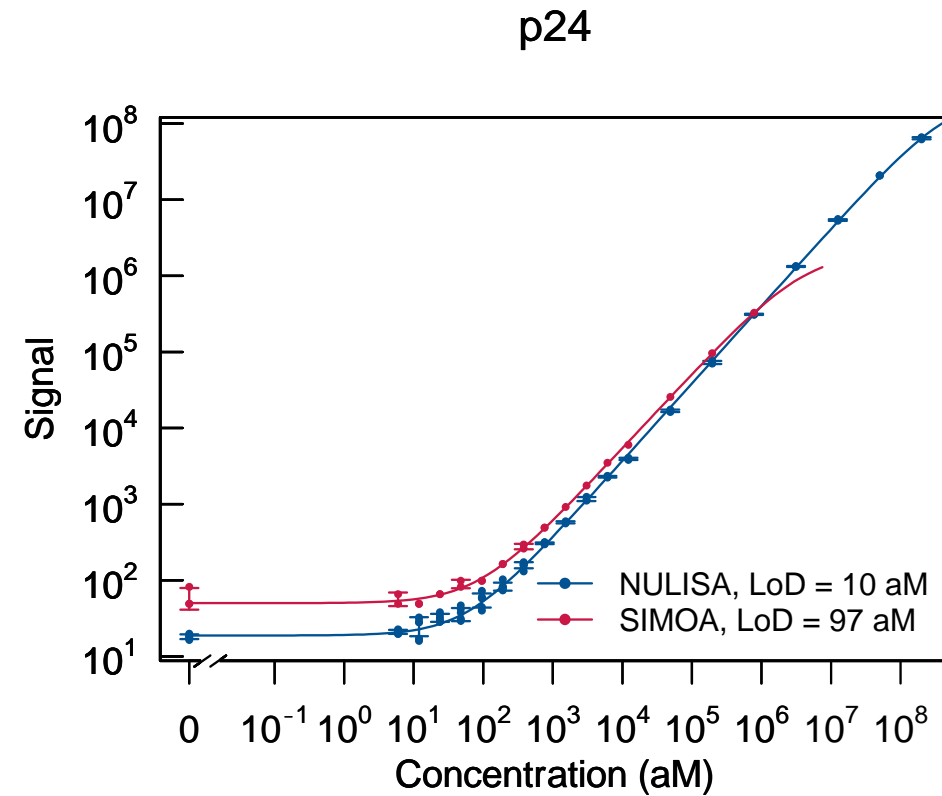
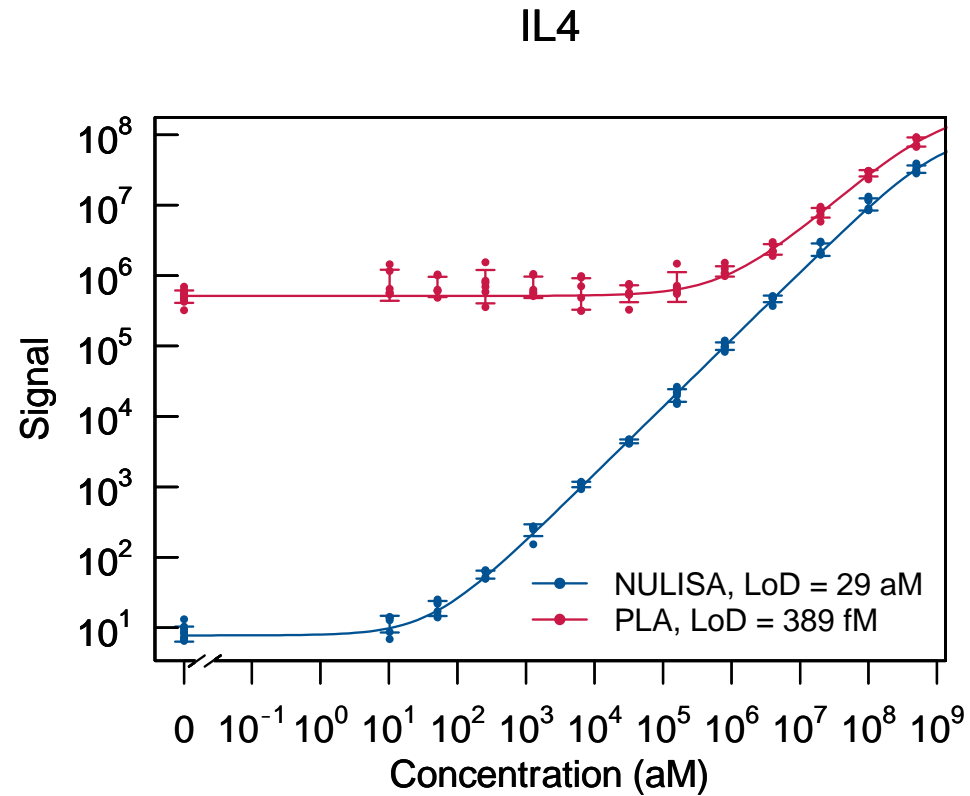
- **Step 8:** wash, then qPCR or NGS



NULISA achieves the highest sensitivity with the same antibodies using proprietary assay background suppression



IL-4 and HIV p24 NULISA using the same antibodies: NULISA vs PLA and SIMOA



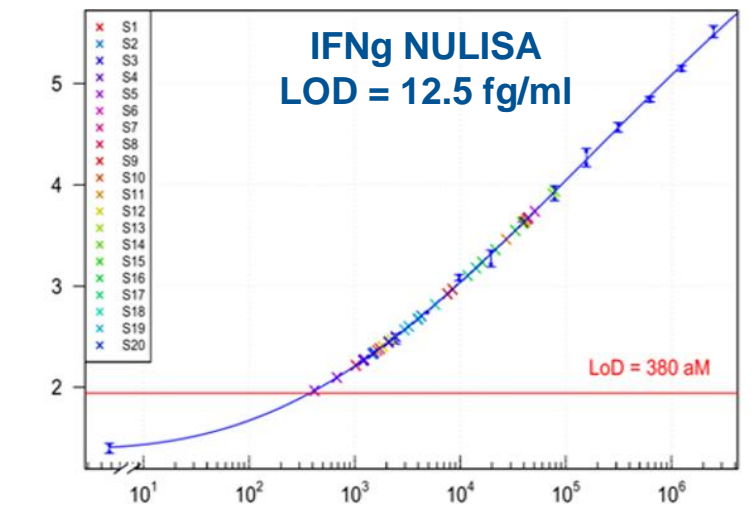
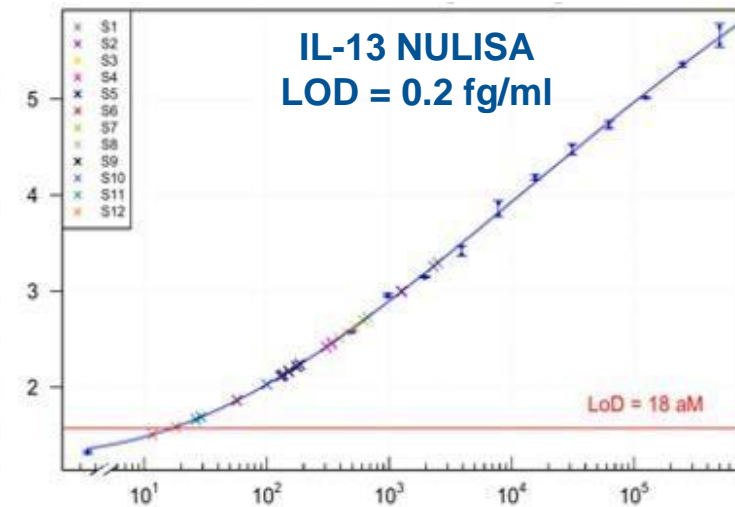
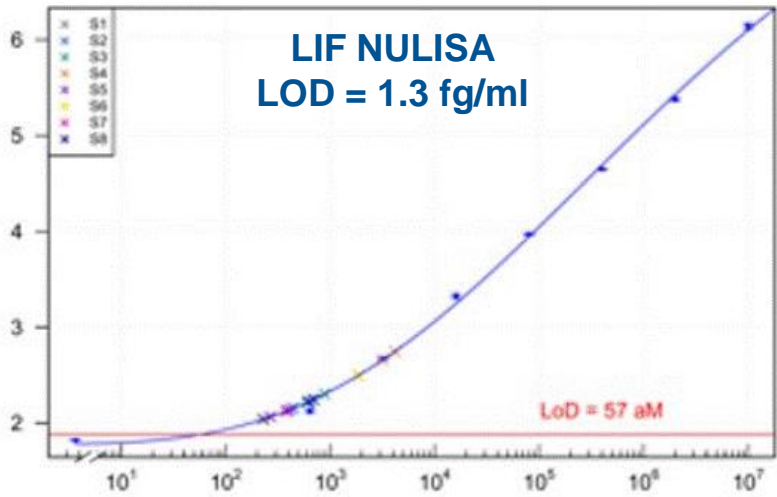
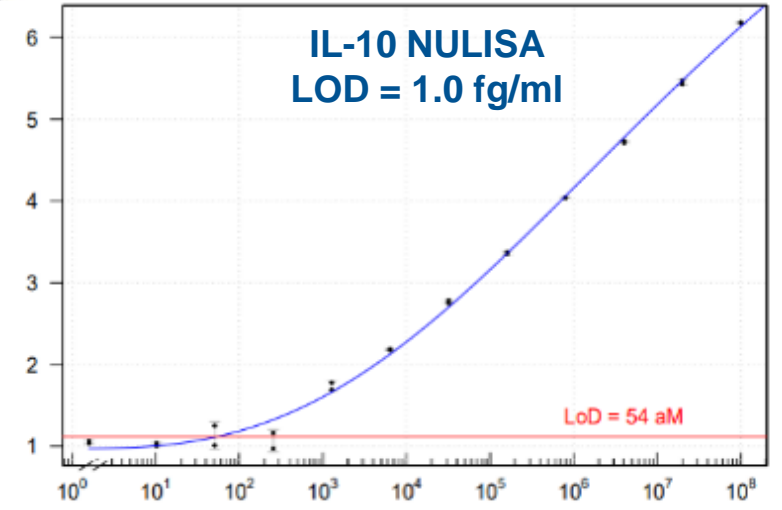
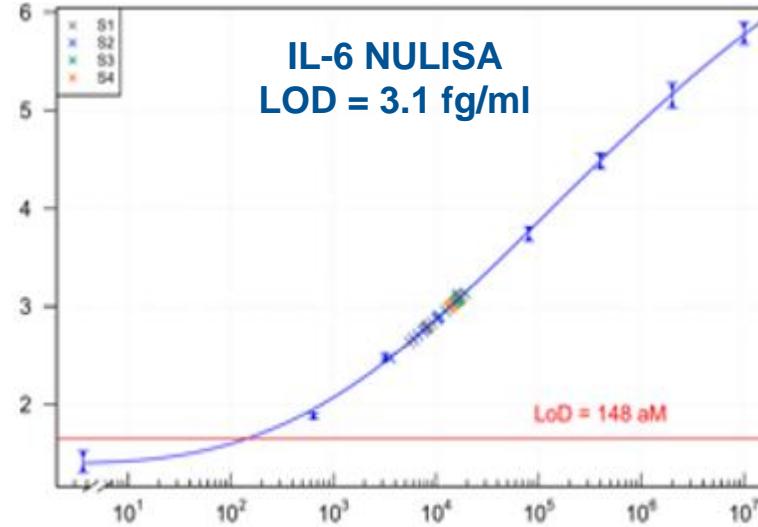
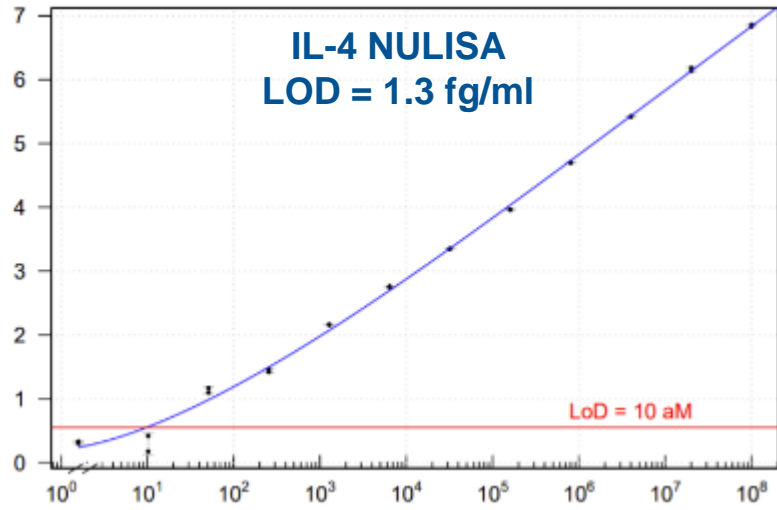
NULISA lowers the limit of detection and increases the quantifiable dynamic range with less sample

IL-4 assay using 100 pM capture/detection antibodies and 500 pM IL-4 at 5-fold dilution.

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Source: <https://www.biorxiv.org/content/10.1101/2023.04.09.536130v1>

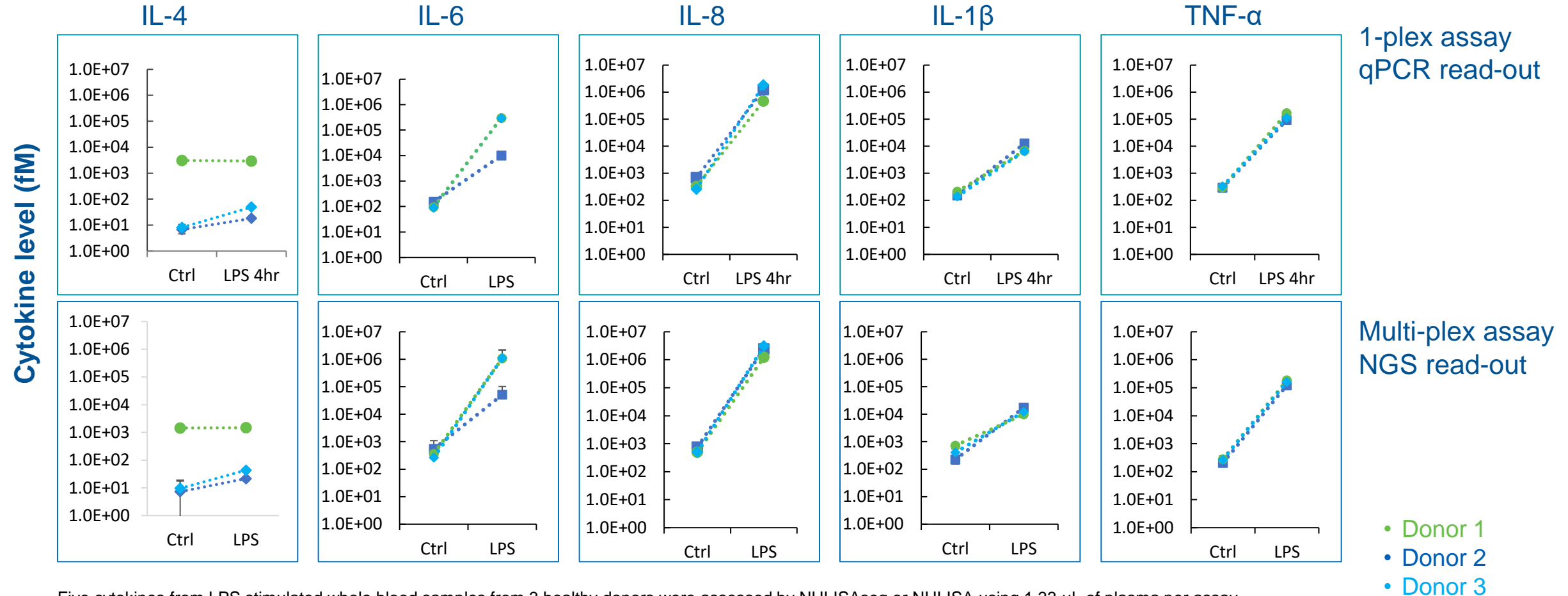
Routine and reliable attomolar (fg/mL) LOD for single-plex NULISA



NULISA “sample-in-data-out” platform enables high precision analysis of single and multiplexed analytes



Concordance, high sensitivity and wide dynamic range for single-plex qPCR and multi-plex NGS NULISA readouts



Five cytokines from LPS stimulated whole blood samples from 3 healthy donors were assessed by NULISAseq or NULISA using 1.33 μ L of plasma per assay

NULISAseq™ Inflammation Panel



Most complete coverage of cytokines and chemokines on the market

250+
biomarkers in 10µL

Attomolar sensitivity
(10 fg/mL)

~12 logs
dynamic range
without dilution

Highly reproducible
CV <10%

Customizable
Subset panels and panel+

TARGETS

AGER	CCL23	CD83	CXCL13	GFAP	IL-13	IL-22	IL-9	MMP9	TAFA5	TNFSF10
AGRP	CCL24	CD93	CXCL14	GRN	IL-13RA2	IL-23A EBI3	IRAK4	MPO	TEK	TNFSF11
ANGPT1	CCL25	CEACAM5	CXCL16	GZMA	IL-15	IL-23A IL-12B	KDR	MUC16	TGFB1	TNFSF12
ANGPT2	CCL26	CHI3L1	CXCL2	GZMB	IL-15RA	IL-24	KITLG	NAMPT	TGFB3	TNFSF13
ANXA1	CCL27	CLEC4A	CXCL3	HAVCR1	IL-16	IL-27 EBI3	KLRK1	NCR1	THBS2	TNFSF13B
AREG	CCL28	CNTF	CXCL5	HGF	IL-17A	IL-2RA	KNG1	NGF	THPO	TNFSF14
BDNF	CCL3	CRP	CXCL6	HLA-DRA	IL-17A IL-17F	IL-2RB	LAG3	NTF3	TIMP1	TNFSF15
BMP7	CCL4	CSF1	CXCL8	ICAM1	IL-17B	IL-32	LAMP3	OSM	TIMP2	TNFSF18
BST2	CCL5	CSF1R	CXCL9	ICOSLG	IL-17C	IL-33	LCN2	PDCD1	TLR3	TNFSF4
C1QA	CCL7	CSF2	EGF	IFNA1; IFNA13	IL-17F	IL-34	LGALS9	PDCD1LG2	TNF	TNFSF8
CALCA	CCL8	CSF2RB	EPO	IFNA2	IL-17RA	IL-36A	LIF	PDGFA	TNFRSF11A	TNFSF9
CCL1	CD200	CSF3	FASLG	IFNB1	IL-17RB	IL-36G	LIL-RB2	PDGFB	TNFRSF11B	TREM1
CCL11	CD200R1	CSF3R	FGF19	IFNG	IL-18	IL-37	LTA	PGF	TNFRSF13B	TREM2
CCL13	CD27	CST7	FGF2	IFNL1	IL-18BP	IL-3RA	LTA LTB	PTX3	TNFRSF13C	VCAM1
CCL14	CD274	CTF1	FGF21	IFNL2;IFNL3	IL-18R1	IL-4	MDK	S100A12	TNFRSF14	VEGFA
CCL15	CD276	CTLA4	FGF23	IFNW1	IL-19	IL-4R	MERTK	S100A9	TNFRSF17	VEGFC
CCL16	CD3E	CTSS	FLT1	IKBKKG	IL-1B	IL-5	MICA	SCG2	TNFRSF18	VEGFD
CCL17	CD4	CX3CL1	FLT3LG	IL-10	IL-1R1	IL-5RA	MICB	SDC1	TNFRSF1A	VSNL1
CCL19	CD40	CXADR	FLT4	IL-10RB	IL-1R2	IL-6	MIF	SELE	TNFRSF1B	VSTM1
CCL2	CD40LG	CXCL1	FTH1	IL-11	IL-1RL1	IL-6R	MMP1	SELP	TNFRSF21	WNT16
CCL20	CD46	CXCL10	FURIN	IL-12A IL-12B	IL-1RN	IL-6ST	MMP12	SIRPA	TNFRSF4	WNT7A
CCL21	CD70	CXCL11	GDF15	IL-12B	IL-2	IL-7	MMP3	SLAMF1	TNFRSF8	
CCL22	CD80	CXCL12	GDF2	IL-12RB1	IL-20	IL-7R	MMP8	SPP1	TNFRSF9	

The screenshot displays the bioRxiv preprint interface. At the top, the bioRxiv logo and the text 'THE PREPRINT SERVER FOR BIOLOGY' are visible. A yellow banner at the top left contains the text: 'bioRxiv does not certify COVID-19-related research. A reminder that they have not been formally peer-reviewed and should not guide health-related behavior or be reported in the mass as conclusive.' The main title of the preprint is 'NULISA: a novel proteomic liquid biopsy platform with attomolar sensitivity and high multiplexing'. Below the title, the authors are listed: Wei Fan, Jennie Bao, Qiyu Mei, Ishara S Aravasa, Anoma Sahaan, Andrei Kamarch, Kark Cho, Pliang Hsu, Xiaohu Qi, Xiaomei Xu, Binshu Zhang, Yu Tianming, Rahn Niazali, Wang Ping, Li Kang, Eyal Liraz Zhan, Wan-Jia Chen, Junjie Lee, Chen-Hsiang Park, Chao-Peng Kwan, Zhen-Jia Chen, Gregorio Gonzalez, Susanna V. Schmidt, Akashdeep Ojha, Jesse S. Bahn, COVIMUNE Consortium, Kazuo Buehler, Dwight Kuo, Li-Ping Binshu Zhang, Jovita Chen, Erika Trivan, Yixi Yuhua Luo, Xiaodun He. The date is April 10, 2023. The abstract begins with 'The blood proteome holds great promise for precision medicine but poses daunting challenges due to the low abundance of the majority of plasma proteins and the vast dynamic range across the proteome. We report the development and validation of a novel proteomic analysis technology - Nucleic acid Linked Immuno-Sandwich Assay (NULISA™) - that incorporates a dual capture and release mechanism to suppress the assay background to the minimum, thus drastically improving the signal-to-noise ratio. NULISA improves the sensitivity of the proximity ligation assay by over 10,000-fold to the attomolar level, which is enabled by antibody-conjugated DNA sequences that mediate the purification of immunocomplexes and contain targeted analysis specific barcodes for next-generation sequencing-based, highly multiplexed analysis. To demonstrate its performance and utility, we developed a 200-plex NULISA targeting 124 cytokines and chemokines and 80 other immune response-related proteins that demonstrated superior sensitivity for detecting low-abundance proteins and high concordance with other immunoprecipitation assays. The ultra-high sensitivity enabled the detection of previously difficult-to-detect but biologically important, low-abundance biomarkers in patients with autoimmune diseases and COVID-19. Fully automated NULISA uniquely addresses longstanding challenges in the proteomic analysis of liquid biopsy samples and makes broad and in-depth proteomic analysis accessible to the general research community and future diagnostic applications.' The page also includes a 'Subject Areas' list, a 'Competing Interest Statement', and a 'Copyright' notice. The bottom right corner features the Chan Zuckerberg Initiative logo.

NULISA: a novel proteomic liquid biopsy platform with attomolar sensitivity and high multiplexing

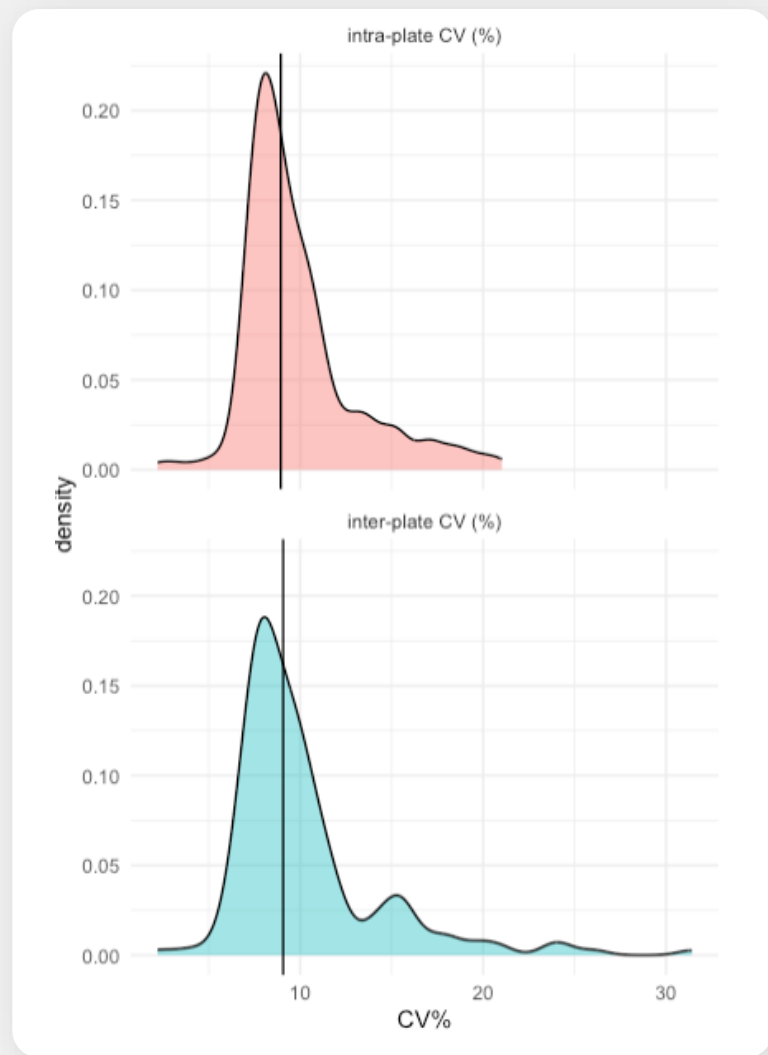


<https://www.biorxiv.org/content/10.1101/2023.04.09.536130v2>

NULISA has tighter CV% versus Olink

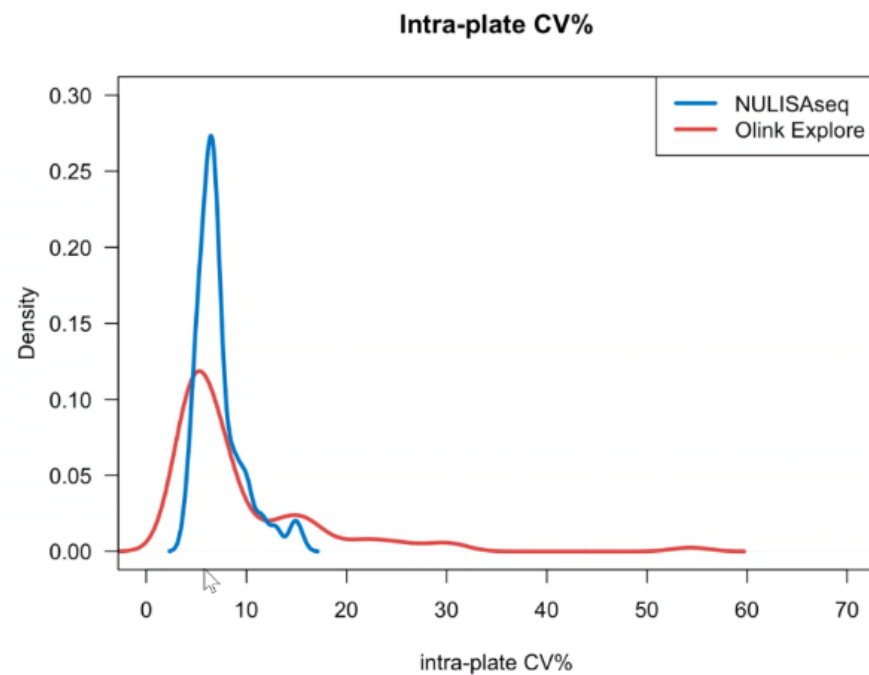


200-plex NULISA
Intra- and inter-plate median CV within 10%



Intra-plate CV for NULISA and Olink for 92 shared targets in 159 plasma samples

	CV% Plate 1 & 2 Average
NULISAseq	7.3 (2.4) / 6.6 [4.1, 15.3]
Olink Explore	9.4 (8.0) / 6.5 [2.6, 54.4]

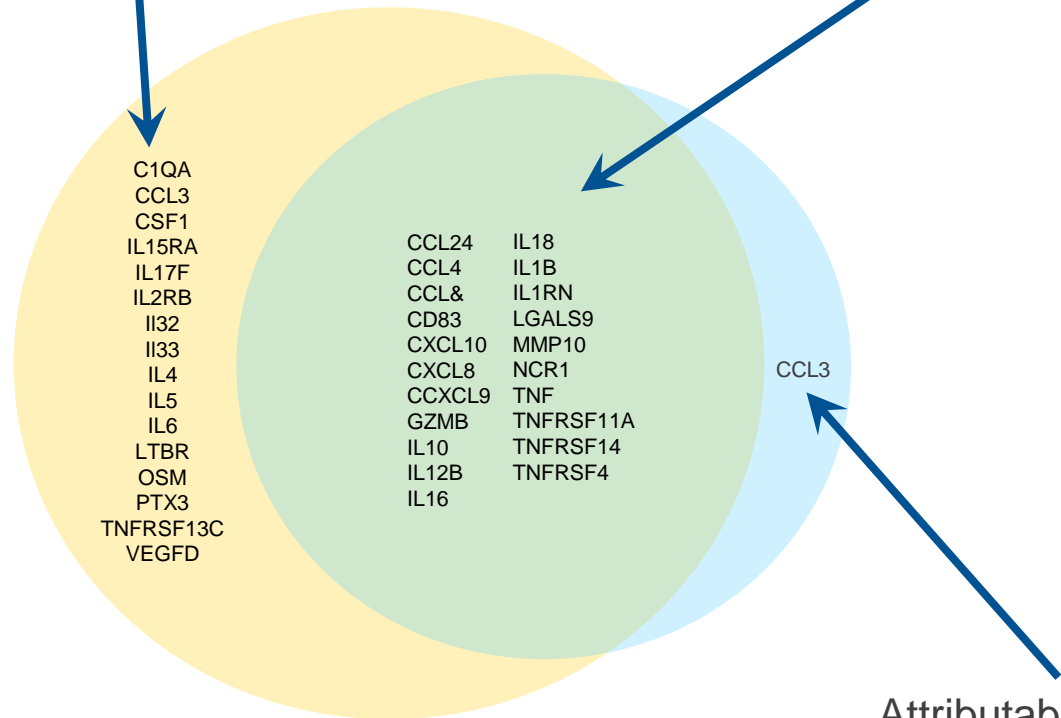


NULISA shows greater detection of differential expression of key inflammatory markers versus Olink



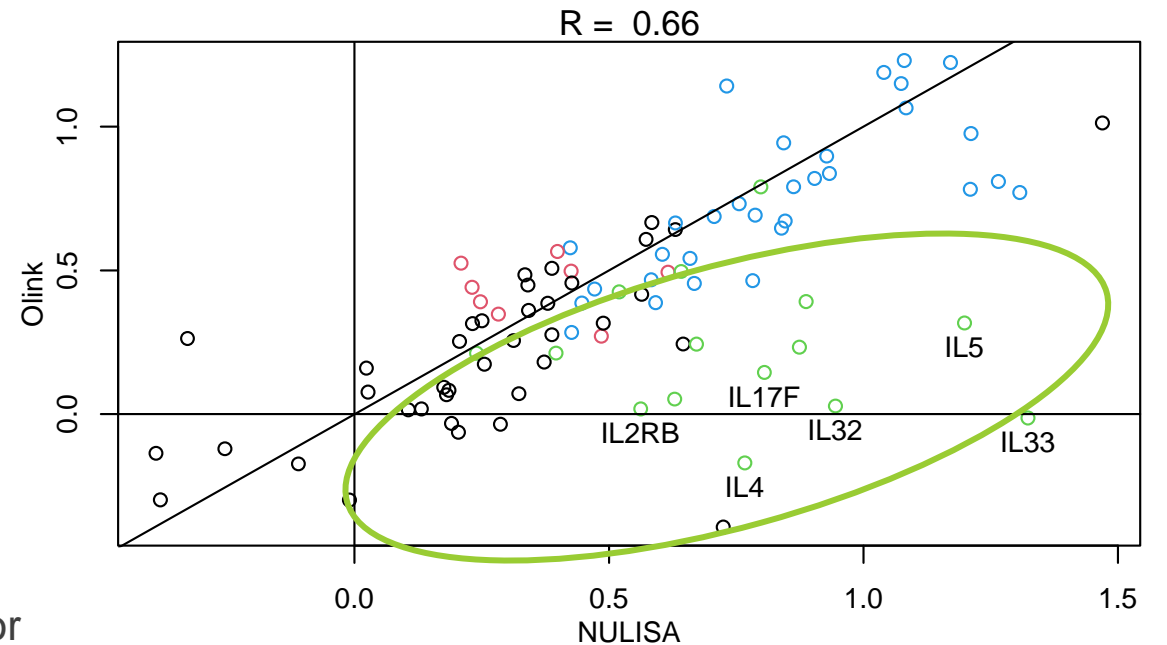
NULISA identified significant differences

NULISA & Olink identify significant differences



Attributable to poor antibody performance

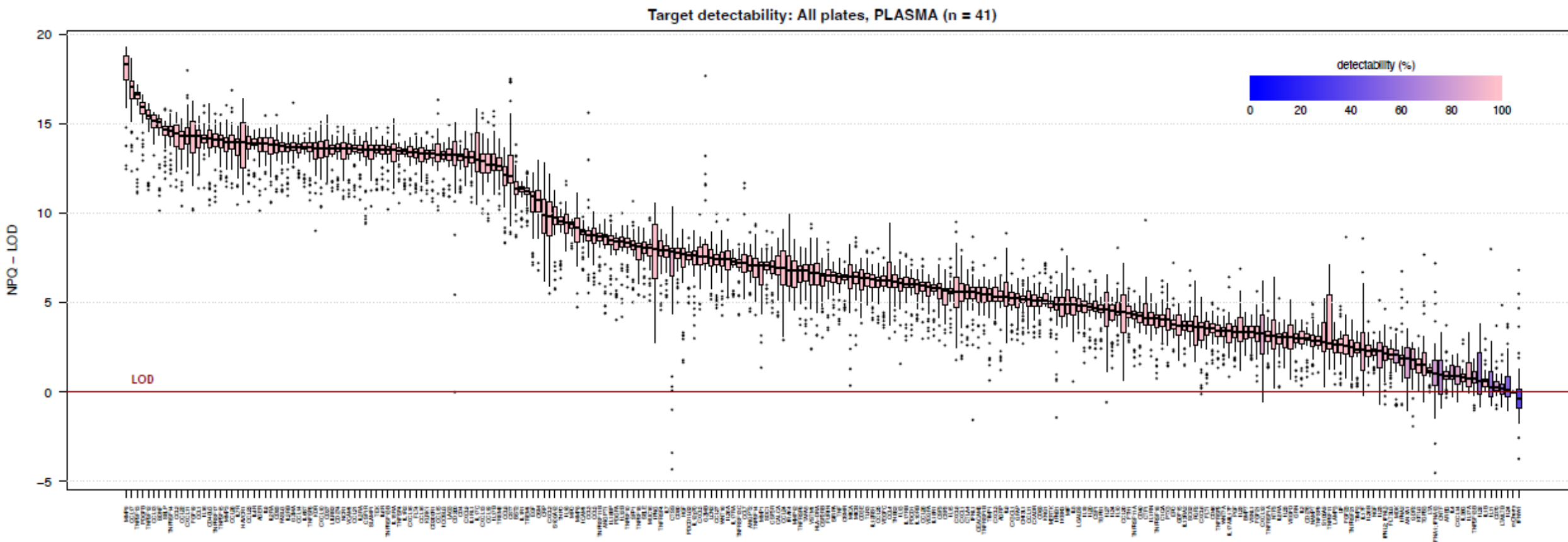
NULISA vs Olink fold-differences: Inflammatory disease and healthy donor plasma
Inflammation vs. Healthy



Detectability of NULISA Inflammation Panel in plasma



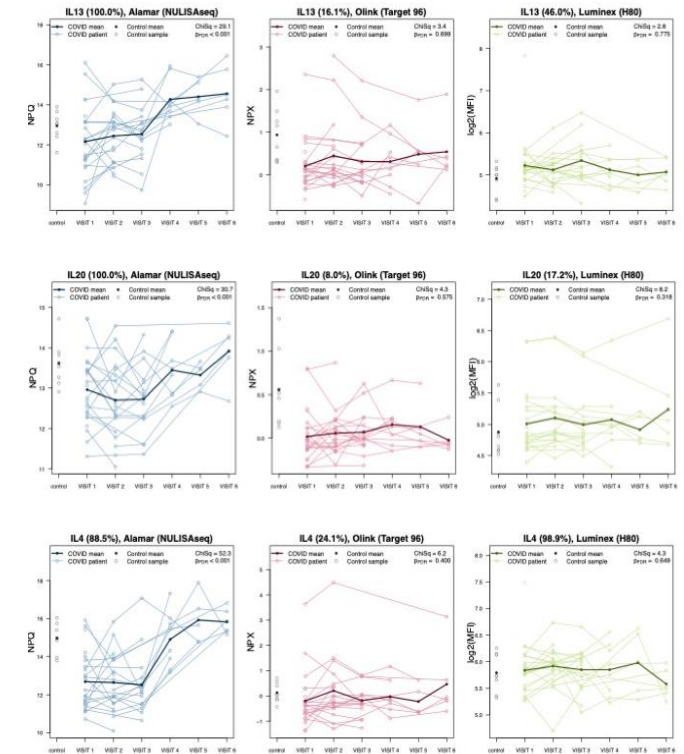
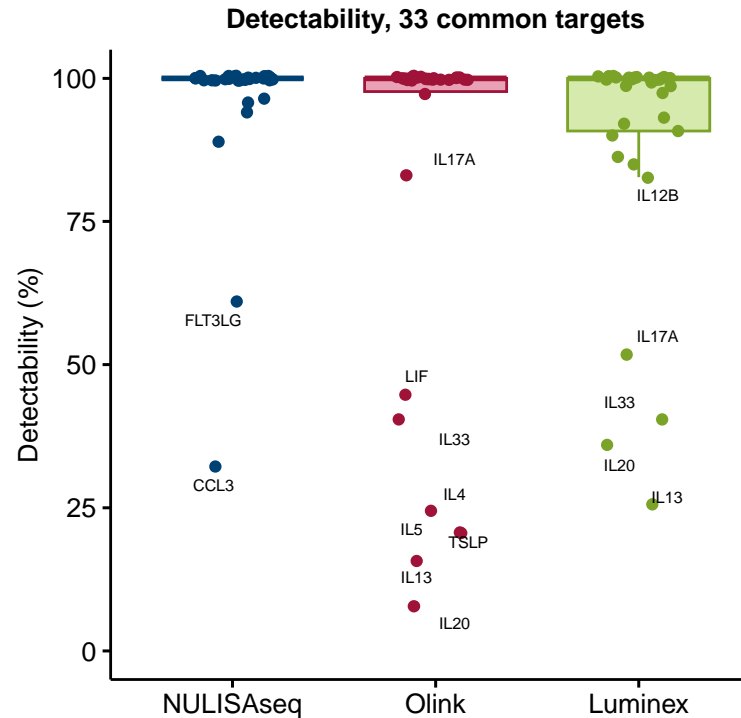
99.6% of targets were detectable in at least 50% of samples



Stanford Human Immune Monitoring Core Adopts NULISA and ARGO on performance and workflow automation



- NIH IMPACC Study of COVID-19 patients
- Three-way comparison between NULISAseq Inflammation Panel, Olink and Luminex.
- NULISAseq showed the highest detectability, which translated to biological insights not obtained from the other two platforms
- Feedback from this customer: **impressed with high sensitivity and fully automated workflow**
- Signed up as first beta customer.



"NULISA platform showed a higher sensitivity for a fair number of analytes and beyond the system's sensitivity, the fact that it comes packaged as an automated platform is attractive, especially given that PEA has historically been a complicated process."



Targets (120 Plex)

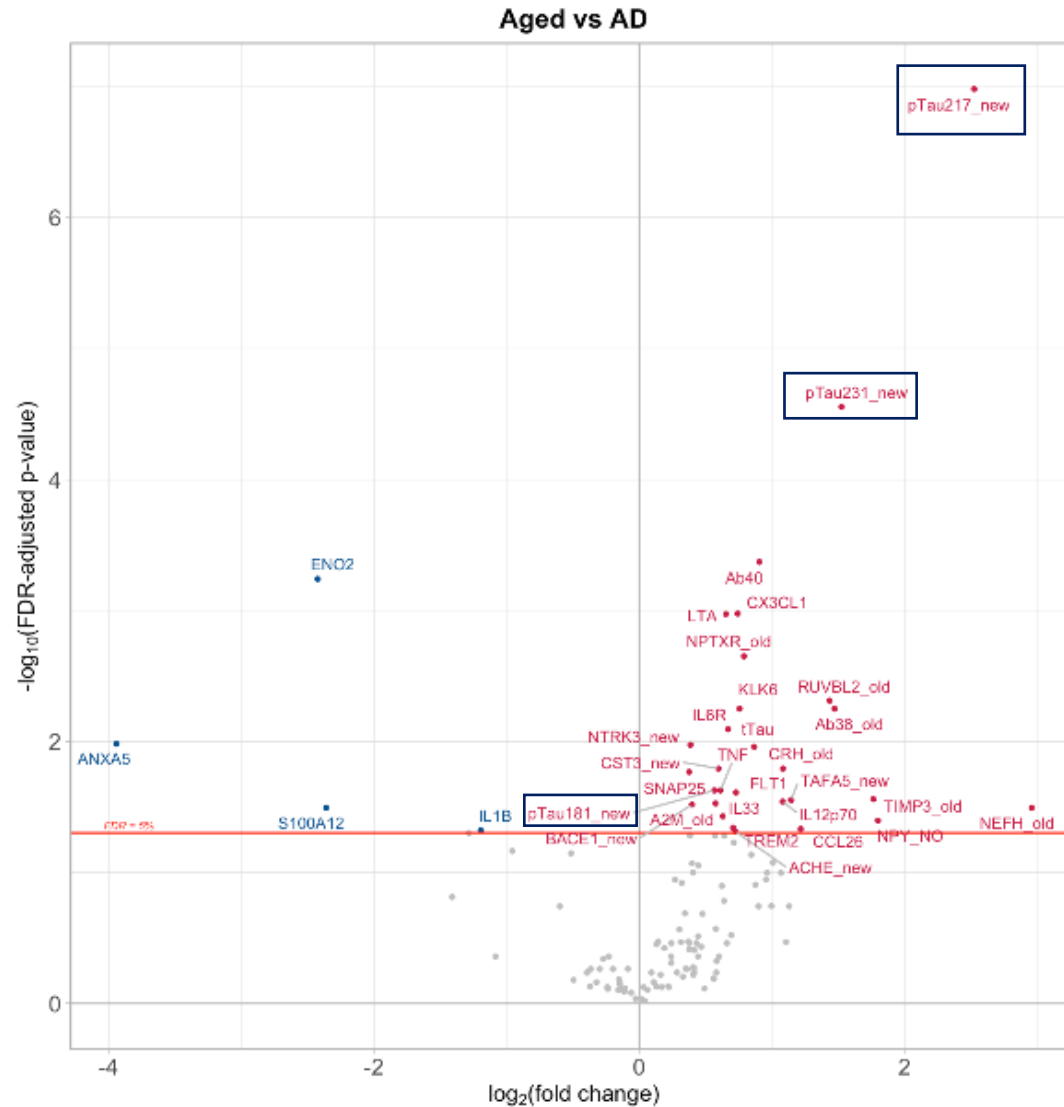
Amyloid Beta 38	GDI1	MCP1	PTN
Amyloid Beta 40	GDNF	MCP4	REST
Amyloid Beta 42	GFAP	MDC	RUVBL2
ACHE	GM-CSF	MDH1	S100A12
AGRN	GOT1	MIP1a/CCL3	S100B
ANXA5	Hemoglobin subunit Alpha	Mip1b/CCL4	SAA1
ARSA	ICAM1	MME	SFRP1
BACE1	IFN-gamma	Mesothelin	SFTPD
BASP1	IGF1	NCAM1	SLIT2
BDNF	IGFBP7	NEFH	SNAP25
CALB2	IL-1 beta	NF-L/NEFL	SNCAagg
CD40L/TNFSF5	IL-10	NGF	SNCAmono
CD63	IL-12p70	NPTX1	SOD1
CHIT1	IL-13	NPTX2	SQSTM1
CNTN2	IL-15	NPTXR	sTREM1
CRH	IL16	NPY	TAFA5
CRP	IL-17A	NRGN	TARC/CCL17
CST3	IL18	PARK7	TDP43
CX3CL1/Fractalkine	IL-2	PDGFRB	Tie-2/TEK
CXCL1/GROa	IL-33	PDLIM5	TIMP3
ENO2	IL-4	PGK1	TNF-a
Eotaxin	IL5	PLGF	TREM2
Eotaxin-3	IL-6	POSTN	tTau
FABP3	IL-6R a	PRDX6	UCHL1
FCN2	IL-7	PSEN1	VCAM1/CD106
FGF basic	IL8	p-TARDBP-409	VEGF R1
FOLR1	IL9	pTau181	VEGF R2
GAP43	IP-10	pTau217	VEGF-A
GDF15	KLK6	pTau231	VEGF-D

**Designed with input from
pharma and academic
KOLs**

**Enabled by NULISA's
combination of ultra-
sensitivity and high-plex**

**Early-stage disease
markers (pTau217, NFL,
GFAP) and neuro-
inflammation markers in
single panel**

CNS Disease Panel: Detecting key differences in plasma between age-matched and Alzheimer's Disease samples



Significantly up-regulated markers include:

- Phospho-Tau181
- Phospho-Tau217
- Phospho-Tau231

Significantly down-regulated markers include:

- S100A12
- ENO2

ARGO™ HT Instrument

Precision Proteomics at the Push of a Button



Simple, automated workflow
<30 min hands-on time

Integrated data analysis
Cloud-based analytics

Minimal sample input
10 – 20 μ L

Single plex results <8 hrs
On-board qPCR analysis

Multiplex workflow
Outputs NGS-ready libraries

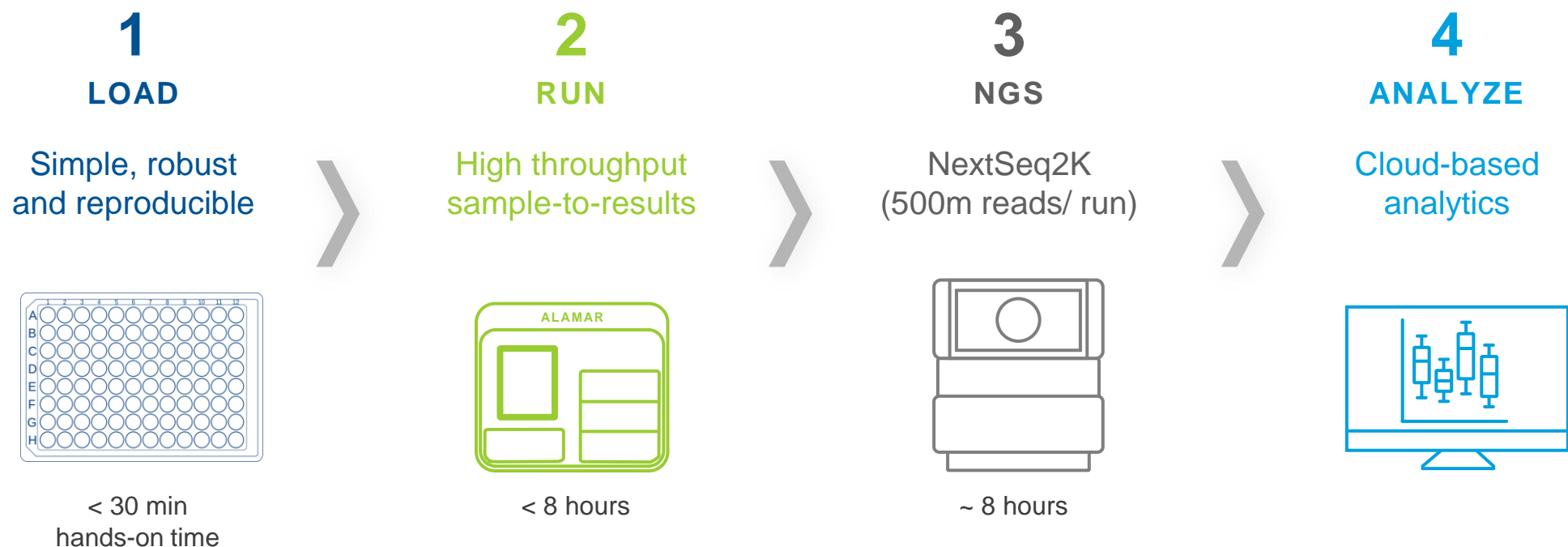
High throughput
Up to 288 samples in 3 x 96-well plates per day

Proteomics at the push of a button



Enabling global access of proteomic analysis from sample to data

Fully Automated Workflow with **ARGO™ HT**



ARGO™ System offers 3x faster and 20x simpler workflow solution



7
Instruments

>200 Step
Workflow

3 Days
Sample to Result

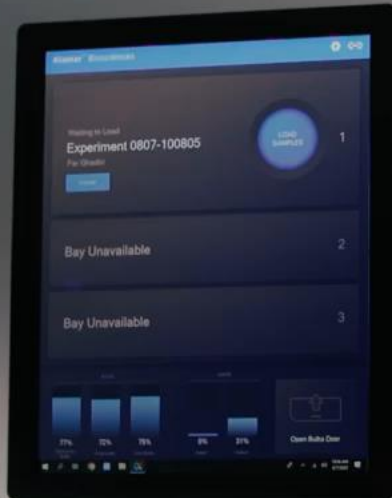


2
Instruments

10 Step
Workflow

1 Day
Sample to Result

ALAMAR



Thank you for your attention

Contact:

Douglas Hinerfeld, PhD
dhinerfeld@alamarbio.com



For more information
Visit www.alamarbio.com

READ the pre-print publication

