## ADVANCES IN ADA, PK, AND BIOMARKER IMMUNOASSAYS TO MEET DEMANDS FOR ASSAY SPEED AND PERFORMANCE

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#### AGENDA

- Singlet analysis for ADA detection
- Development of a generic ADA assay for preclinical studies
- Extending sensitivity for PK and biomarker assays



#### Research Article

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Bioanalysis

# SINGLET VS DUPLICATE GYROLAB PEMBROLIZUMAB IMMUNOGENICITY ASSAY COVANCE

Goals of microfluidic Gyrolab ADA assay:

Automate assay using Mixing CD 96

Evaluate singlet vs duplicate analysis

Evaluate drug tolerance of assay

Comparing singlet and duplicate immunogenicity assay in human plasma for pembrolizumab using Gyrolab<sup>®</sup>

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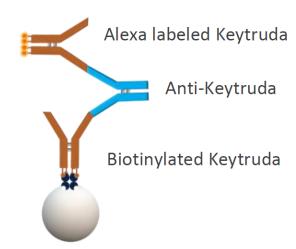
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#### PEMBROLIZUMAB METHOD — BRIDGING ASSAY

- Capture: Biotinylated pembrolizumab
- Detect: Alexa Fluor 647-labeled pembrolizumab
- Positive control:
   Hu IgG1 anti-pembrolizumab Ab (BioRad)
- Mixing CD 96
- Gyrolab xP workstation



· Method sequence: analyte, acidic buffer, master mix/neutralization







#### Precision of screening assay — no improvement with duplicates

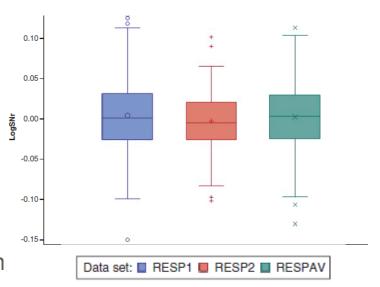
### Between run precision over six runs, analyzed by three analysts.

Precision	Positive control	Run 1	Run 2	Ave
Whole CD precision	LPC	7.1%	3.4%	4.0%
	HPC	6.0%	7.7%	6.3%
Inter-run precision	LPC	8.0%	8.4%	7.9%
	HPC	5.5%	6.1%	5.6%

CD: Compact disc; HPC: High positive control; LPC: Low positive con

LPC: 50 ng/mL; HPC: 10,000 ng/mL

## Box plots of screening data (outliers excluded)



Singlet and duplicate (ave. of 2 runs) values showed no significant improvements to the data quality by using the average result over a single measurement



#### DRUG TOLERANCE

- Assay drug tolerance calculated to be 1796 μg/mL pembrolizumab at 50 ng/mL PC
  - Tested at 50, 100, 200, 500 and 1000 μg/mL
  - Positive control at 50, 100 and 250 μg/mL
  - All response levels above cut point

- Higher drug tolerance than Gyrolab assay protocol
  - Gyrolab assay protocol drug tolerance of 640 μg/mL at 100 ng/mL PC

	Calculated drug tolerance for each cutpoint (μg/mL)				
PC level (ng/mL)	Measurement 1 (± difference to ave)	Measurement 2 (± difference to ave)	Ave		
50	1779 (-17)	1832 (+36)	1796		
100	2642 (-14)	2685 (+29)	2656		
250	2616 (-5)	2632 (+11)	2621		

#### SUMMARY

- Gyrolab ADA assay using mixing CD automates acid dissociation
- High precision (<15%) for both Gyrolab assay protocol and Covance developed assays
- Drug tolerance (640 and >1000 µg/mL for Gyrolab and Covance assays, respectively) appropriate for clinical use
- Automation and use of singlet analysis increases productivity for bioanalytical laboratories
- Significant time savings

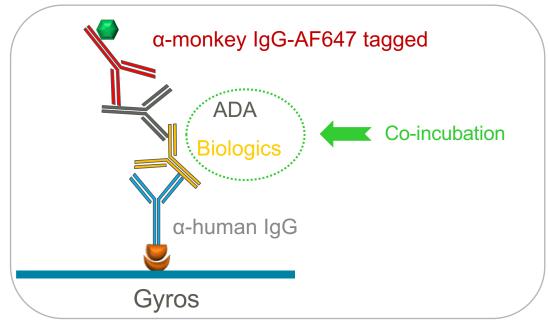


#### GENERIC ADA ASSAY

Developed by: Romain Gauchet and Franck Levasseur, DMPK Department – Biokinetics Servier



#### ADA GENERIC METHOD: FORMAT IN MONKEY



<u>Benefits</u>: « Plug and Play method » → no need to tag each new Biologics with biotin, just need to coat biotinylated antihuman IgG to quantify all ADAs targeting the Biologics under development

<u>Limitations</u>: Method only works for IgG-based biologics; not working with other therapeutic protein structures (e.g hormone)

The challenge to validate the format of this generic method has been to prepare a generic positive control and to identify a suitable and reliable cut-point strategy





#### GENERIC METHOD VS SPECIFIC METHOD

#### **Tested Biologics:**

- Compound A (monoclonal Ab)
- Compound B (IgG-based compound)
- Compound C (IgG-based compound)

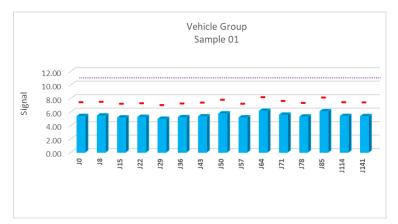
#### Comparison of ADA results between validated/qualified method and generic method

Compound A	Monoclonal antibody	lgG4	Bridging	
Compound B	IgG-based compound	lgG1	Sandwich	Generic method
Compound C	IgG-based compound	lgG1	Sandwich	

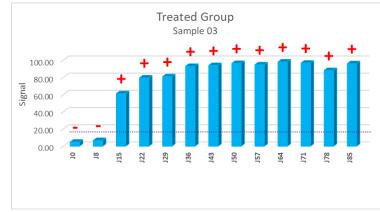


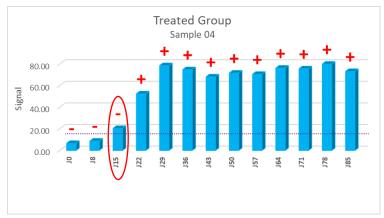


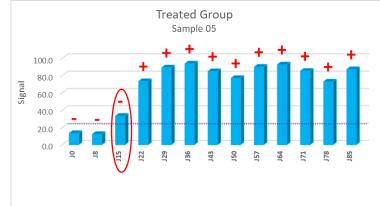
#### GENERIC METHOD VS SPECIFIC METHOD: COMPOUND A











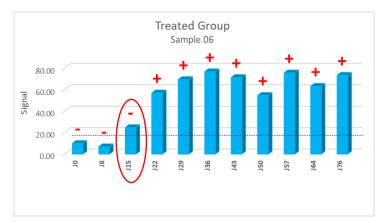
Calculated Preliminary Cut-Point = 1.25\* Mean NC

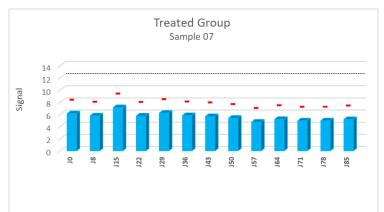
-/+: Specific Method Results

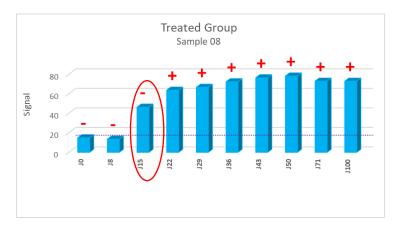


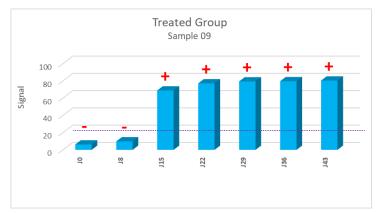


#### GENERIC METHOD VS SPECIFIC METHOD: COMPOUND A









..... Calculated Preliminary Cut-Point = 1.25\*

Mean NC

-/+: Specific Method Results

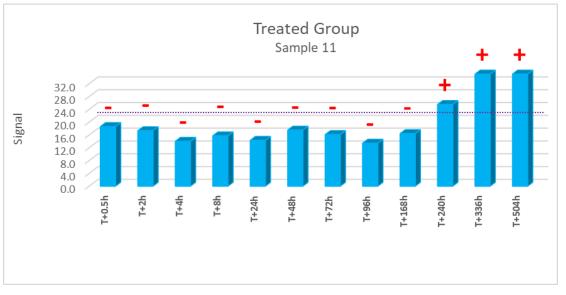
Good correlation between Specific and Generic Method Format works well and trends of signal correlate between both methods





#### GENERIC METHOD VS SPECIFIC METHOD: COMPOUND B





..... Calculated Preliminary Cut-Point = 1.25\* Mean NC

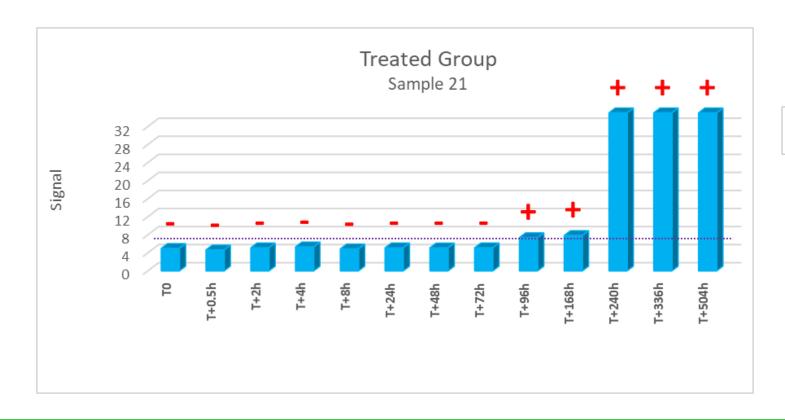
-/+: Specific Method Results

Good correlation between Specific and Generic Method
Format works well and trends of signal correlate between both methods





#### GENERIC METHOD VS SPECIFIC METHOD: COMPOUND C



... Calculated Preliminary Cut-Point = 1.25\* Mean NC

-/+: Specific Method Results

Good correlation between specific and generic methods:

Identical ADA status and same trend for signal level across samples

→ Generic format performs well





#### PC PRODUCTION

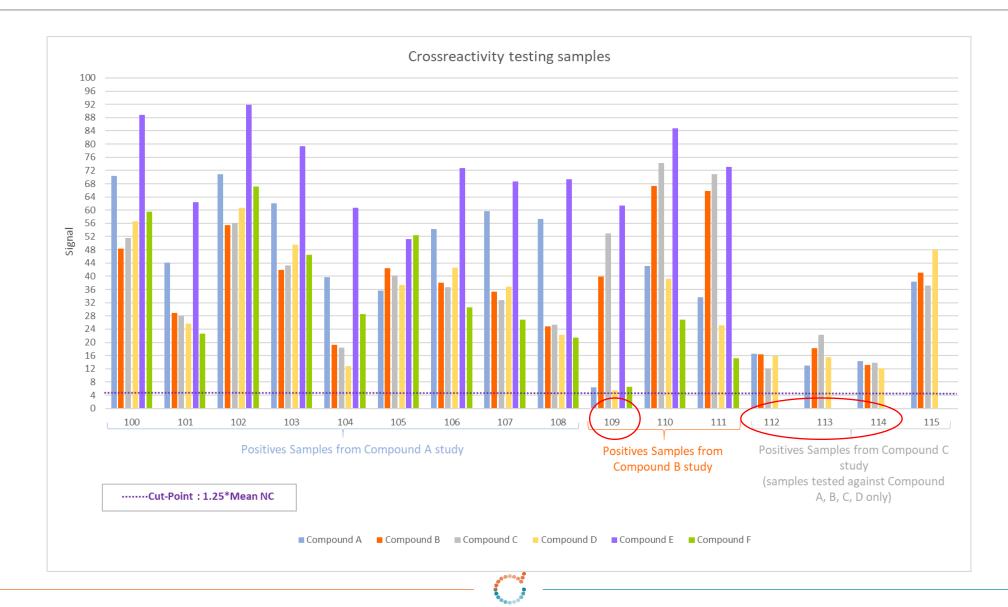
- Selection of ADA+ samples from 3 monkey studies and cross-reactivity testing against 5 Biologics
- Selection of crossreacting ADA samples to constitute a high polyspecific positive control pool:

	Biologics tested for ADA cross-reaction						
	Compound A mAb IgG4						
Positive Samples from     Compound A study		9 🗸	9 🗸	9 🗸	9 🗸	9 🗸	
3 Positive Samples from Compound B study	2 🗸		3 🗸	2 🗸	3 🗸	2 🗸	
4 Positive Samples from Compound C study	4 🗸	4 🗸		4 🗸	Not Tested	Not Tested	





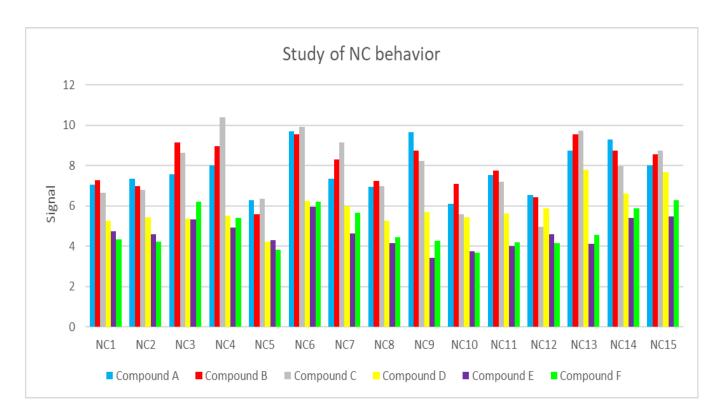
#### PC PRODUCTION





#### **CUT-POINT STRATEGY**

Naïve monkey samples (ADA negative) behavior against 6 different Biologics:



	Mean	SD	CV%
Compound A	7.75	1.16	15.0
Compound B	8.00	1.19	14.9
Compound C	7.83	1.62	20.8
Compound D	5.87	0.93	15.8
Compound E	4.63	0.70	15.0
Compound F	4.89	0.94	19.2
Compound A to F	6.49	1.80	27.7





#### **CUT-POINT STRATEGY**

#### **Conclusion:**

- 6 Tested Biologics → structural homology
- CV > 27% → Conservative approach applying a dynamic cut-point to reduce the risk of high variability against other Biologics
- Dynamic cut-point = mean negative control (NC) of 15 ADA negative samples + 3SD



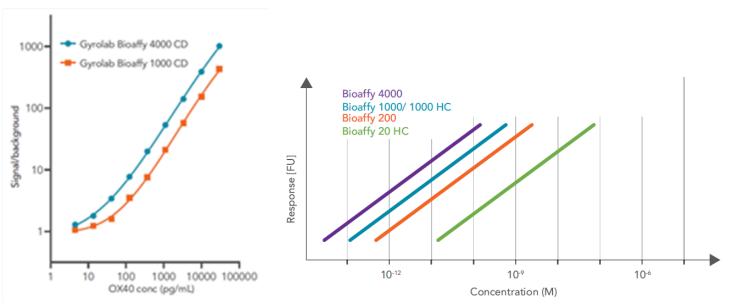


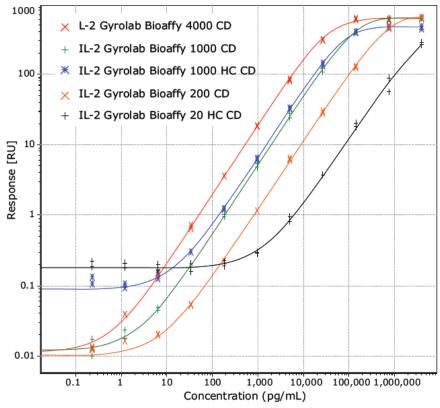
#### BIOMARKER APPLICATIONS



#### EXTENDING THE ANALYTICAL RANGE

- Combined analytical range of 6 logs
- Sample volume determines assay sensitivity
- Seamless assay transfer between CDs



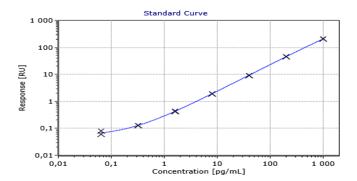


Note: The higher binding capacity of CDs using the high capacity (HC) streptavidin particle increases the upper limit of detection but may also increase background binding.



#### CYTOKINE ASSAYS WITH EXTENDED SENSITIVITY





Assay range	LOD (pg/mL)	LLOQ (pg/mL)	ULOQ (pg/mL)
On plate	~ 0.1	~ 0.3	~ 800
In neat serum	~ 0.2	~ 0.6	~ 1 600



anti-human/cyno IFN-γ mAb

rh IFN-γ

anti-human/cyno IFN-γ mAb

IFN	l- gam	ıma

100		Standard	Curve		
Response [RU]	*	Standard	X	*	*
0,001	0,1		10	100	1 000
0,01	5,1	Concent	ration [pg/mL]		1 000

Assay range	LOD (pg/mL)	LLOQ (pg/mL)	ULOQ (pg/mL)
On plate	~ 0.2	~ 0.3	~ 800
In neat serum	~ 0.4	~ 0.6	~ 1 600



anti-human/cyno IL-4 mAb

rh IL-4

anti-human/cyno IL-4 mAb

	1,000		Standard Cur	/e		
	100				×	
Response [RU]	10			×		
Respor	1		×			
	0.1	***************************************				
	0.1	1	10 Concentrati	100 on [pg/mL]	1,000	10,000

Assay range	LOD (pg/mL)	LLOQ (pg/mL)	ULOQ (pg/mL)
On plate	~ 0.6	~ 1	~ 1 200
In neat serum	~ 1.2	~ 2	~ 2 400

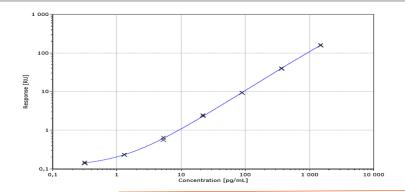
TNF-Alpha



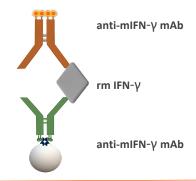
anti-human/cyno TNF-α mAb

rh TNF-α

anti-human/cyno TNF-α mAb



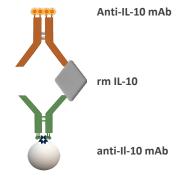
Assay range	LOD (pg/mL)	LLOQ (pg/mL)	ULOQ (pg/mL)
On plate	~0.5	~1	~1 000
In neat matrix	~1	~2	~2 000





1 000 -		Standard Cur	ve		
100				×	
[] 10			×		
Response [RU]		×			
0,1	×				
0,01	10	100 Concentrati	1 000 ion [pg/mL]	10 000	100 000

Assay range	LOD (pg/mL)	LLOQ (pg/mL)	ULOQ (pg/mL)
On plate	~0.3	~4	~15 000
In neat matrix	~0.6	~8	~30 000



	100-				×
_					
Response [RU	10	 		×	
Res			×		
	1-	 ×			

Assay range	LOD (pg/mL)	LLOQ (pg/mL)	ULOQ (pg/mL)
On plate	~0.5	~1	~4 000
In neat matrix	~1	~2	~8 000

anti-mTNF-α mAb
rm TNF-α
anti-mTNF-α mAb

TNF-Alpha

IL-10

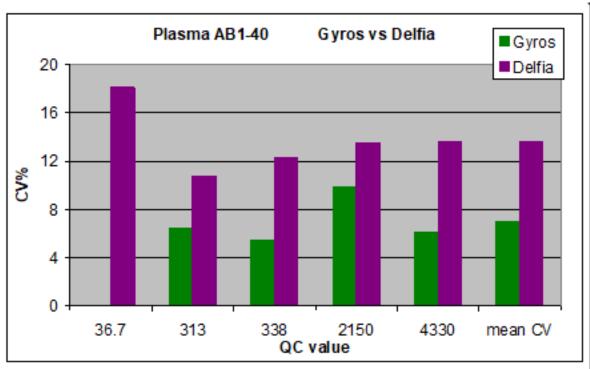


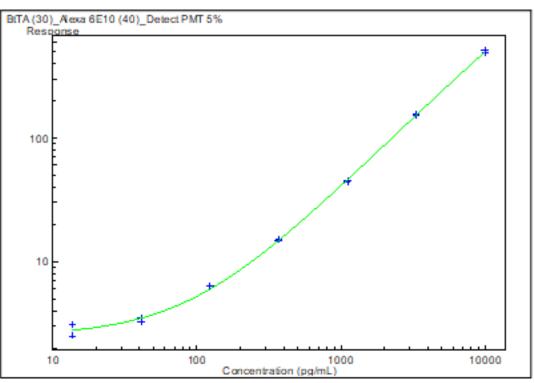
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HISTORICAL ABETA AMYLOID 1-40



#### PREVIOUS ABETA AMYLOID 1-40 DATA





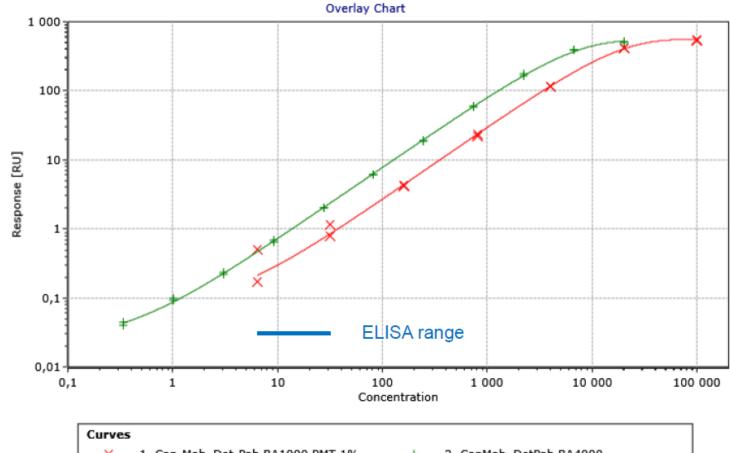
Courtesy of John Allinson, ICON PLC





#### BA 1000 vs BA 4000





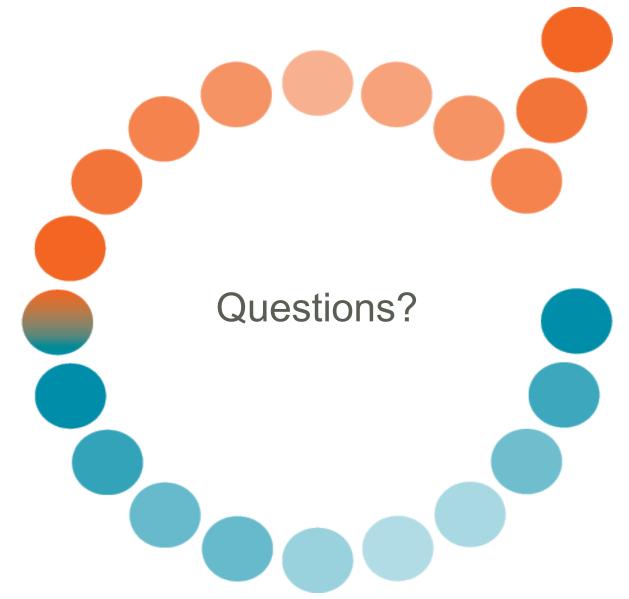
1. Cap-Mab\_Det-Pab BA1000 PMT 1% 2. CapMab\_DetPab BA4000

**BA4000 CD IMPROVED SENSITIVITY** 

#### SUMMARY

- Using the mixing CD with a Gyrolab ADA assay automates acid dissociation for significant time savings in immunogenicity assays
- High precision (<15%) for Covance pembrolizumab immunogenicity assay assays allowed singlet analysis
  - Drug tolerance 640 and >1000 μg/mL for Gyrolab (data not shown) and Covance assay, respectively appropriate for clinical use
  - Automation and use of singlet analysis increases productivity for bioanalytical laboratories with significant time and reagent savings
- Extended sensitivity for cytokine assays and high precision biomarker analysis is demonstrated by the Bioaffy 4000





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