

# The Challenges of Developing and Validating Biomarker assays

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# Introduction



## Focus of presentation

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- Difference between endogenous and recombinant analyte
  - Stability
  - Drug interference
  
- Determination of Acceptance criteria
  - Context of Use
  - Project timelines

# Endogenous vs recombinant analyte

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## Use of recombinant analyte:

- Required for early stage antibody screening
  - Removes the effect of matrix
  - Low endogenous analyte levels
- Required for calibration curves and QC's
  - Analytical range

## Difference between endogenous and recombinant analyte:

- Recombinant analyte raised *in vitro*
- SNPs
- Post translational modifications (Glycosylation, oxidation, acetylation)

# Endogenous Vs recombinant analyte- Stability



## Case study one- Analyte X

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Assay information:

- PD Biomarker assay on the MSD platform for a FTIH study
- Assay range 69714-34 fg/mL
- To cover the analytical range VC's were a combination of pooled human serum and spiked pooled human serum

Validation control	Nominal Concentration (fg/mL)*	Make up
VC1	94	Endogenous analyte
VC2	234	Endogenous analyte
VC3	2022	Approx. 1500 fg/mL recombinant spike
VC4	17011	Approx. 27000 fg/mL recombinant spike
VC5	46484	Approx. 60000 fg/mL recombinant spike

\*Nominal concentration determined from pre-validation P&A runs

# Endogenous Vs recombinant analyte (Stability)



## Case study one- Analyte X

### Freeze thaw stability

3 Freeze/Thaw cycles

VC	Nominal concentration (fg/mL)	BCC (fg/mL)	% Recovery
VC1	94	85	90.4
VC2	234	233	99.6
VC3	2022	1951	96.5
<b>VC4</b>	<b>17011</b>	<b>35784</b>	<b>210.5</b>

### Conclusions & learnings:

- Recombinant analyte X behaves differently to endogenous analyte X
- Not always feasible to perform stability assessments in validation which cover the analytical range.
- Is it more appropriate to monitor stability using study samples?

### Bench top stability

2 hour RT stability

VC	Nominal concentration (fg/mL)	BCC (fg/mL)	% Recovery
VC1	94	74.5	79.3
VC2	234	196	83.8
VC3	2022	1997	98.8
<b>VC4</b>	<b>17011</b>	<b>27596</b>	<b>162.2</b>

24 hour RT stability

VC	Nominal concentration (fg/mL)	BCC (fg/mL)	% Recovery
VC1	94	87	92.6
VC2	234	246	105.1
VC3	2022	1676	82.9
<b>VC4</b>	<b>17011</b>	<b>41952</b>	<b>246.6</b>

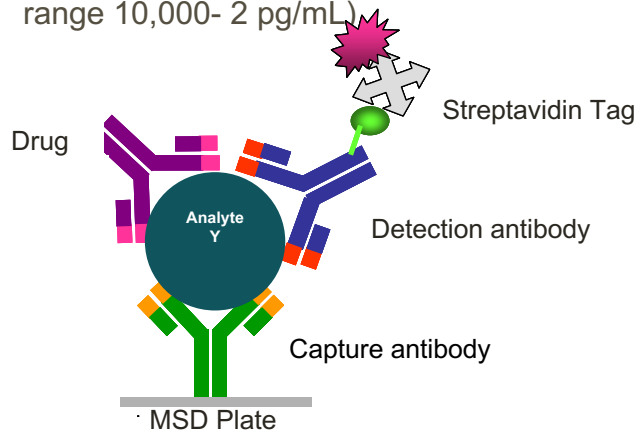
# Endogenous Vs recombinant analyte (drug interference)



## Case study two- Analyte Y

### Assay background:

- Total target engagement biomarker assay on the MSD platform for a FTIH single escalation dose study
- Assay was required to be highly sensitive (Assay range 10,000- 2 pg/mL)



### Initial Method development results:

- Commercial antibodies were screened to find a pair which could detect total analyte Y
  - Surrogate matrix was spiked with recombinant analyte Y and incubated with and without drug

ECL Units with <u>No</u> drug present	ECL units with drug present	%Recovery
1488	1545	103%

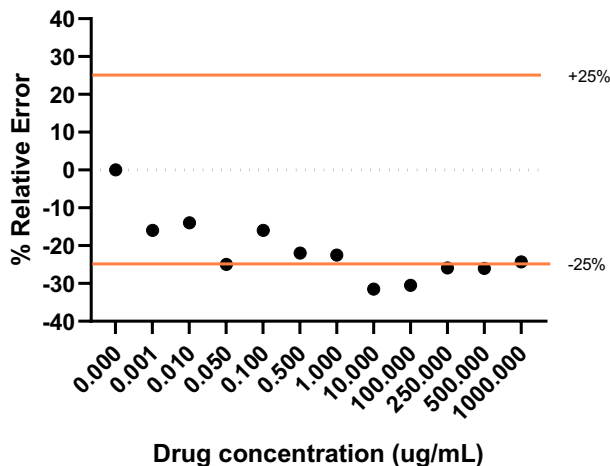
- Method development and optimization continued using the above antibody pair

# Endogenous Vs recombinant analyte (drug interference)



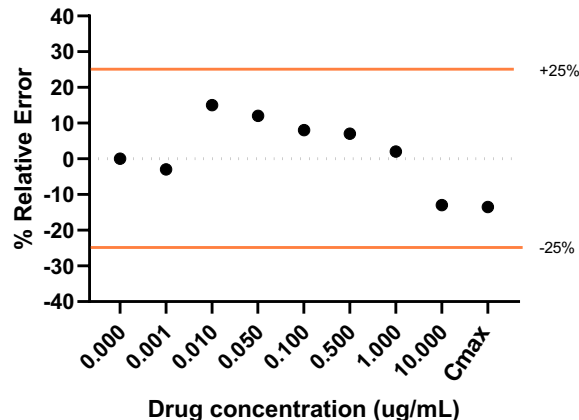
## Case study two- Analyte Y

Drug interference in endogenous samples



Drug saturation

drug interference in endogenous samples with drug saturation



How could this unacceptable drug interference be solved?

- Use of different buffers
- Use of different MSD plate types
- Different capture and detection antibody

**Conclusion:** Saturating the assay with drug reduced drug interference

**Learnings:** Importance of introducing endogenous samples as early on as possible in method development

# Endogenous Vs recombinant analyte

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## Key learnings:

- Importance of introducing endogenous analyte as soon possible during method development
- Impact of reagent availability
- Impact of study time lines



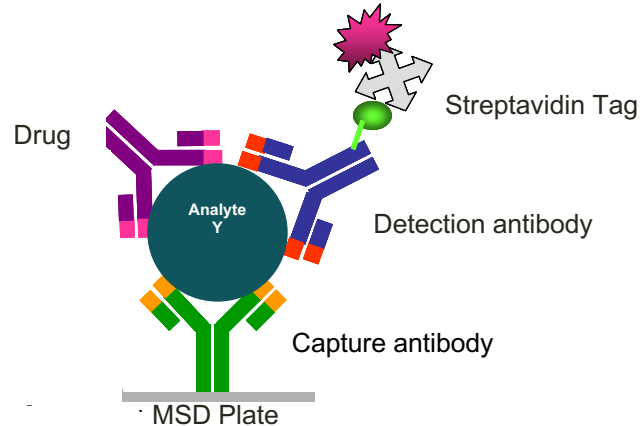
# Acceptance criteria



## Case study 2- Analyte Y

### Assay background:

- Total target engagement biomarker assay on the MSD platform for a FTIH single escalation dose study
- Assay was required to be highly sensitive (assay range 10,000- 2 pg/mL)
- Calibration curve was prepared using recombinant analyte in surrogate matrix
- To cover the analytical range VC's were combination of pooled human serum and spiked pooled human serum



# Acceptance criteria



## Case study two- Analyte Y

### Initial Method development:

- Echo 525 acoustic dispenser was used prepare calibration curve
- Addition of reagents and samples were added to the assay plate manually.
- assay precision and accuracy were acceptable



### Validation A:

- Precision and accuracy VC's was unacceptable

VC	%RE	%CV
1 (LLOQ)	1.72	28.01
2	1.82	15.91
3	-5.19	17.88
4	-7.55	17.56
5	-13.01	22.82
6 (ULOQ)	-14.16	31.45

- The precision and accuracy was causing other validation parameters to fail

### Further method development

- Method was fully automated to try and improve the precision and accuracy.
- Micropro 300 was used to stamp reagents and samples onto the assay plate



# Acceptance criteria



## Case study two- Analyte Y

### Validation B:

- Automation had resulted in the calibrators having very tight %CV and %RE across the assay range

Calibrator	1 (ULOQ)	2	3	4	5	6	7	8	9 (ULOQ)
%CV	3.91	3.36	4.11	0.23	6.03	0.69	2.67	6.50	4.99
%RE	1.00	-1.67	-3.60	3.00	6.00	3.33	-7.20	-2.00	6.50

- Validation showed higher level of variability in the QC's than our target acceptance criteria (<25% %CV, %RE)

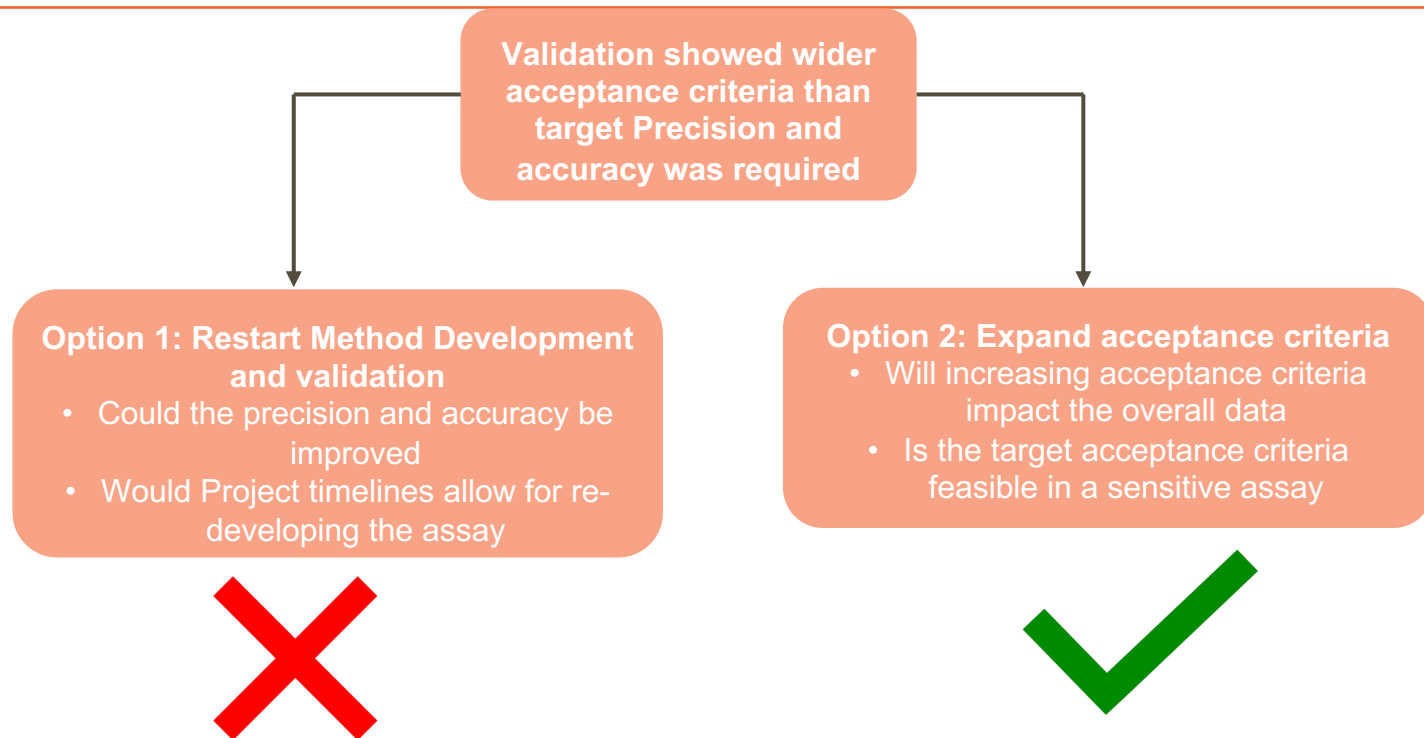
QC	1 (rep 1)	1 (rep2)	2 (rep 1)	2 (rep 2)	3 (rep 1)	3 (rep 2)	4 (rep 2)	4 (rep 1)
%CV	10.55	2.51	2.90	6.73	2.27	0.94	8.03	2.65
%RE	<b>38.35</b>	-4.13	18.90	2.44	<b>27.73</b>	3.18	<b>27.04</b>	2.97

- Could dilute samples to remove matrix effect however this would increase the LLOQ

# Acceptance criteria



## Case study two- Analyte Y



# Acknowledgements

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**Questions**