

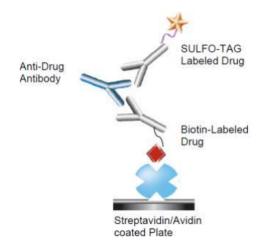
(Unwanted) Immunogenicity – the anti-drug antibody format



MSD Bridging assay (ECLIA) - commonly used in the assessment of unwanted immunogenicity







Meso Scale Discovery®, Bridging Immunogenicity Assays

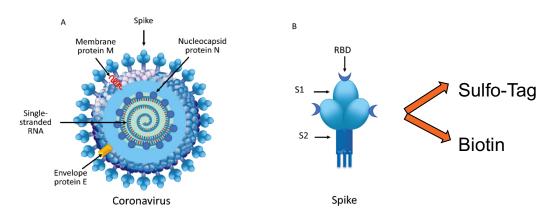
Why not an ELISA?

- Better sensitivity and analytical working range
- Homogenous solution phase incubation simplifies workflow
- Shorter assay times higher throughput
- Not species specific

Early challenges in method development



- Early prototype assays used S1 and RBD fragments to investigate bridging assay potential
- Full length Spike protein was not available until June/July 2020
- Full length assays required additional development to reduce background
 - > Buffer optimization
 - > Different challenge ratios
 - > Concentration in assay
- Positive controls not specific (only cross reactive from SARS)



The Virus Itself Rossi, et al. (2020) *Infection* volume 48, p.665–669

Final assay – format and precision

- Block Streptavidin MSD Plate
- Dilute sample 1 in 20
- Diluted sample is incubated for 1 hour with master mix containing equal concentrations of biotinylated- and sulfotagged full length spike protein
- Reaction mix is added to blocked plate for 1 hour
- Plate is washed an read on sector imager
- All liquid handling performed on an Integra ViaFlo

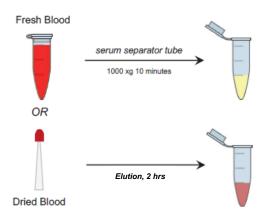


	left	middle	right	
NC	146	145	135	
CP-PC	1238	1299	1223	
PC	5524	5462	4915	
EPC	72557	73614	67643	
NC	154	158	150	
CP-PC	1326	1318	1213	
PC	5408	5392	5088	
EPC	73628	71590	70321	

	Mean (n=6)	%CV
VC	148	5.4
CP-PC	1270	4.0
PC	5298	4.5
EPC	71559	3.2

Sample types – serum and WB micro-sample





Neoteryx Mitra® VAMS and collection kits

Benefits

At home sampling, no need for a clinic visit or venipuncture

CE Marked, FDA Class 1 devices

Devices can be shipped directly to us

Barcoded, logged straight into our LIMS system for chain of custody and ease of reporting



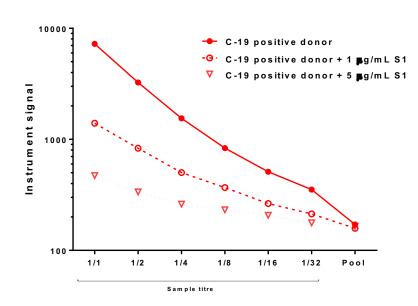




Questions over approach

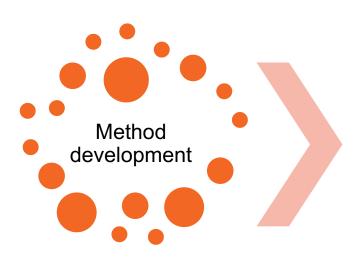


- Do we need a tiered approach? What about confirmatory analysis??
- What about how we normally validate an ADA assay?
 Selectivity etc..
- Analytical Sensitivity, PCs are not as good as real positive samples – can we justify not having the 100 ng/mL box ticked?
- Should we not be analyzing in duplicate?
- What regulations should we be working to?



The pathway to assay roll out*





□ Serum□ Mitra VAMS

Analytical Validation

- □ Serum
- Cut point
- Precision
- Stability

Labware LIMS

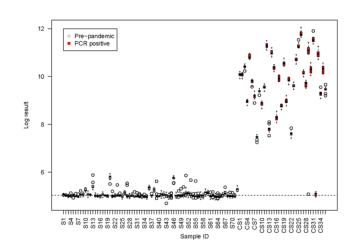
- Serum
- Verify CP with greater n
- Cross-reactivities
- Sensitivity/Specificity
- Equivalence with other assays
- Mitra VAMS
- Verify CP
- Concordance with serum

Clinical Verification

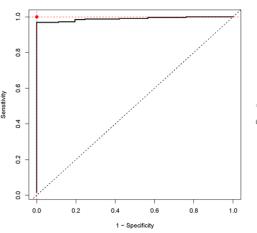
^{*} Verification and validation methodology and sample sets for evaluation of assays for SARS-CoV-2 (COVID 19), Royal College of Pathologists. document reference number: G222-3 (2020).

Assay Validation - Cut point establishment

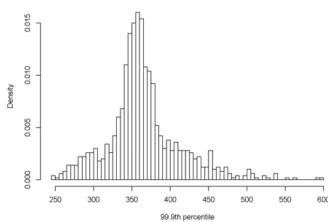




Given the purpose of the test, the threshold was set such that the specificity is as close to 100% as possible, while maximising the sensitivity. The point on the curve which is closest to the top left corner is at specificity 1 and sensitivity 0.9691, which is attained at thresholds between 385 and 1,400 RFU.



ROC Curve, the red dot represents a perfect test with 100% specificity and sensitivity



Bootstrapped values for the 99th percentile of the distribution of prepandemic samples

Assay Validation – Precision & Stability



 Precision – Intra/Inter-run and inter-analyst (3 analysts, 6 runs each of three plates)

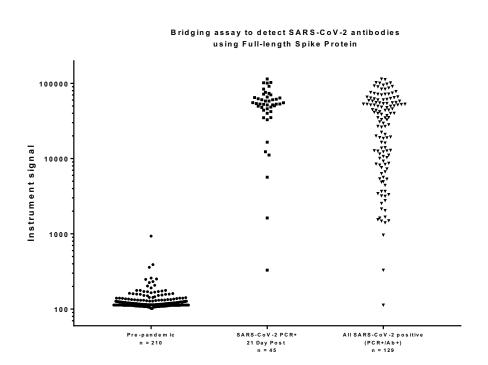
		NC	PC1	PC2	PC3
	Mean	115	973	4181	56106
Serum	Std Dev.	16.6	169	741	10886
	CV%	14.5	17.4	17.7	19.4
	n	156	156	156	138
		NO	D04	D00	DO2
		NC	PC1	PC2	PC3
Mitra	Mean	112	1057	4587	19982
	SD	18.5	181	817	4566
Eluate	CV%	16.4	17.1	17.8	22.9
	n	80	80	80	80

- PC1 and PC2 = Seracare Accurun controls.
 PC3 is high responding clinical sample
- All plate values are normalized to PC1, hence a positive sample is >1.

- Stability
 - o Serum
 - Benchtop 24 hrs, 3 x Freeze/Thaw, Long term frozen at -80°
 - Mitra sample
 - Dried tip stability at RT and 35°C for 7 days (covers postage period)
 - Eluate Benchtop 24 hrs, 3 x
 Freeze/Thaw, Long term frozen at -80°

Serum clinical verification





Days post PCR confirmation	N	Reactive	Non- Reactive	% Positive	Sensitivity % (95% CI)
11-20	19	19	0	100	
21-30	44	43	1*	98	
31-50	1	1	0	100	
From 21 days	45	44	1*	98.0	88.4 - 99.6

The assessment of sensitivity was performed on a cohort of COVID-19 patient samples where infection by SARS-CoV-2 had been confirmed by a PCR test 21 days prior to the sample being taken. In this case the assay demonstrated 98% Sensitivity.

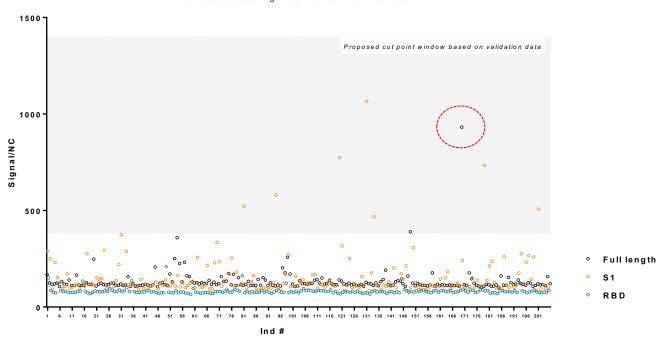
Positive Ab samples by Comparator assay	N	Positive by LGC assay	Negative by LGC assay
Roche Elecsys Anti-Sars-CoV-2 serology assay	47	47	0
Abbot SARS-CoV-2 IgG assay	43	42	1*
Siemens SARS-CoV-2 Total (COV2T) assay	67	67	0

^{*} Sample confirmed as Ab negative by both Roche Elecsys and Siemens assays









Serum clinical verification



 Assessment of serum samples for specificity used 377 prepandemic samples including the following disease state or interference assessments

Confounder samples

- 39 Coronavirus HKU Ab+
- 39 Coronavirus OC43 Ab+
- 40 Coronavirus 229E Ab+
- o 38 Coronavirus NL63 Ab+
- 4 Parainfluenza Ab+
- 4 Influenza A Ab+
- 4 Influenza B Ab+
- 4 Respiratory Syncytial Virus Ab+
- 2 Rheumatoid Factor
- 2 HIV+
- o 4 Enterovirus Ab+
- o 31 EBV Nuclear Antigen positive
- o 24 CMV Ab+
- o 16 HBs Ab+
- o 2 Immune thrombocytopenia (ITP)

Interference samples

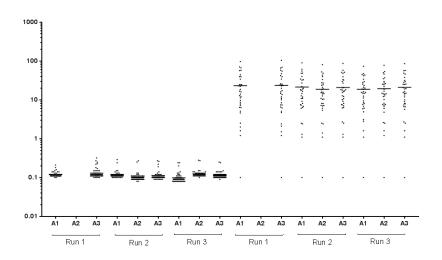
- o 2x hyperlipidaemia patient samples
- o 2x hyperlipidaemia (spiked to 4mg/mL)
- 2x hyperbilirubinaemia (spiked to 30 μg/mL)
- 2x haemolysed (3% equivalent to (>250 mg/dL of free haemoglobin)
- 2x Biotin (spiked to 1200 ng/mL)

Category	N	Reactive	Non-Reactive	Specificity (%)	95% CI
Negative samples: Pre-COVID era	301	0	301	100	
Interference samples	10	0	10	100	
Confounder samples	66	0	66	100	
Total	377	0	377	100	98.7 – 100%

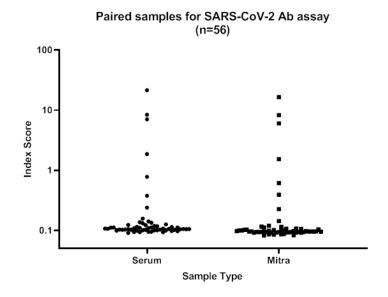
Additional Mitra Eluate clinical verification



Comparison between serum values and surrogate mitra samples (comprising of red blood cells from a healthy donor combined with serum from pre-pandemic or confirmed COVID-19 patients).



Paired venous draw serum and capillary "finger prick" Mitra samples, from volunteers at LGC, were assessed for concordance.



Labware LIMS





This is the plate review screen after data import.
On the right hand side we have the plate level data.

Top table shows the cut point control data.

Middle table is the Positive and Negative QC data.

At the bottom we find the unknown sample results.

This is the Covid Home screen, the workflow is as follows:

- Create new clients/sites as required
- Upload Mitra tip kit barcodes
- Linking a shipment with a client.
- Sample batching and QC checking
- After analysis run reviewed, accepted or rejected.
- Reporting

Science for a safer world Exit Log Out	Covid-1	9 Run Re	eview	Printer: FOR-LBL-01- 1XT	LGC	Suptamber 9 2020		
tun: COVID_I2	0-08-006 Date of A	nalysis: 26-A	UG-20 Analyst	: WALDA.KAYE			Accept Run	Plate Data
							Reject Run	CPC Mean: 749.5
								CPC SD: 6.6
			Cut Po					CPC CV%: 0.9
								0
OVID_QC_CPC-3886		3/No	750	Add	Reject			NC Mean: 104.8
COVID_QC_CPC-3886		6 No	760	Add	Reject			NC SD: 11.3
OVID_QC_CPC-3886		45 No	749	Add	Reject			NC CV%: 10.8
OVID_QC_CPC-3887		48 No	752	Add	Reject			
OVID_QC_CPC-3887		93 No 96 No	746 740	Add Add	Reject Reject			NC Range: 50-250
OVID_QC_CPC-3887	0	20 140	740	Add	Reject			PC Mean: 999.5
			oc	Samples				PC SD: 6.3
· ID	0 Type	0 a 10 d a		control of the contro		00		
OVID OC NC-38860		2 No	100 Yes	0.1 Negative	Add Comment 0	Reject	e 🕠 Rej. Keason	
COVID_QC_NC-38860		2 No 5 No	100 Yes	0.1 Negative 0.1 Negative	Add	Reject		PC Range: 750-1250
OVID_QC_NC-38866		44 No	99 Yes	0.1 Negative	Add	Reject		Warnings
OVID OC NC-38869		47 No	102 Yes	0.1 Negative	Add	Reject		
OVID OC NC-38872		92 No	93 Yes	0.1 Negative	Add	Reject		None
OVID_QC_NC-38875		95 No	125 Yes	0.2 Negative	Add	Reject		
OVID_QC_PC-38859		1 No	1000 Yes	1.3 Positive	Add	Reject		
OVID_QC_PC-38862		4 No	1010 Yes	1.3 Positive	Add	Reject		
OVID_QC_PC-38865		43 No	999 Yes	1.3 Positive	Add	Reject		
OVID_QC_PC-38868		46 No	1002 Yes	1.3 Positive	Add	Reject		
COVID_QC_PC-38871		91 No	993 Yes	1.3 Positive	Add Add	Reject		
COVID_QC_PC-38874	Covid 19 Positive Control	94 No	993 Yes	1.3 Positive	Add	Reject		
				Validation Samples				
'ID 0	Type	o Pos	0 Mod 0	Response 0 Cut Off In	ndex 0 Result		Comment	Add Comment
	f Primary Sample		7 No	100	0.1 Negative			Add
	d Primary Sample		8 No	350	0.5 Negative			Add
	d Primary Sample		9 No	650	0.9 Negative			Add
	d Primary Sample		10 No	75	0.1 Negative			Add
	d Primary Sample		11 No	125	0.2 Negative			Add
	d Primary Sample		12 No	1120	1.5 Positive			Add
	d Primary Sample d Primary Sample		13 No 14 No	100 110	0.1 Negative 0.1 Negative			Add Add
	d Primary Sample d Primary Sample		14 No	450	0.1 Negative 0.6 Negative			Add
	d Primary Sample		15 No	900	1.2 Positive			Add
	1 Primary Sample		17 No	250	0.3 Negative			Add
	d Primary Sample		18 No	240	0.3 Negative			Add
					0.000			

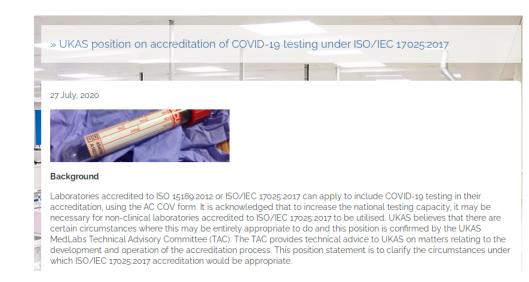
All data is fake data created for testing so it may be inconsistent

Whats next...



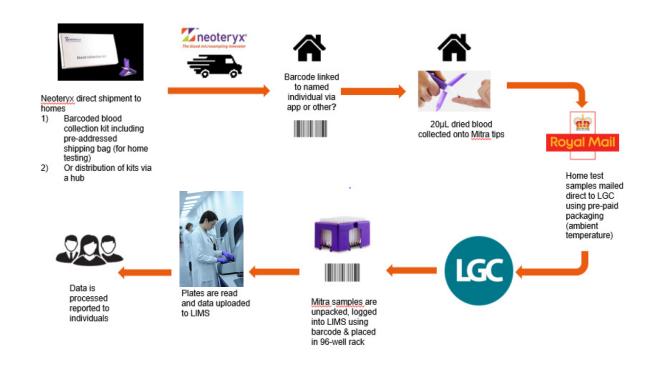
The MHRA are very clear that an assay such as this would be classed as a diagnostic medical device.

As such, it requires a CE mark (done) and performance under an ISO15189 or ISO17025 quality system (pending inspection).



An end—to—end solution (its not all about the assay…)





Thank you for listening

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