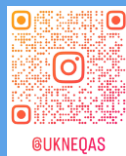




# UK NEQAS H&I

## Annual Participant's Meeting 2024-25





# Meet The Team!

Director: Deborah Pritchard

Manager: Amy De'Ath

Deputy Manager: Melanie Bartley

Healthcare Scientist Practitioner: Geraint Clarke

QA Technical Officer: Jack Jefferies

MLA: Sue Davies

[UKNEQASHandl@Wales.NHS.UK](mailto:UKNEQASHandl@Wales.NHS.UK)



# UK NEQAS for H&I Steering Committee 2024-25



Helena Lee (Chair)

Arthi Anand

Katy Derbyshire

Sylvia McConnell

Katherine Mounsey

Anthony Calvert

Sunil Daga (Clinical Representative)

Elizabeth Wroe (BSHI Representative to UK NQAAP)



Rhys Goodhead (Expert Advisor Scheme 5B)

Barbara McNamara (Expert Advisor Scheme 5B)

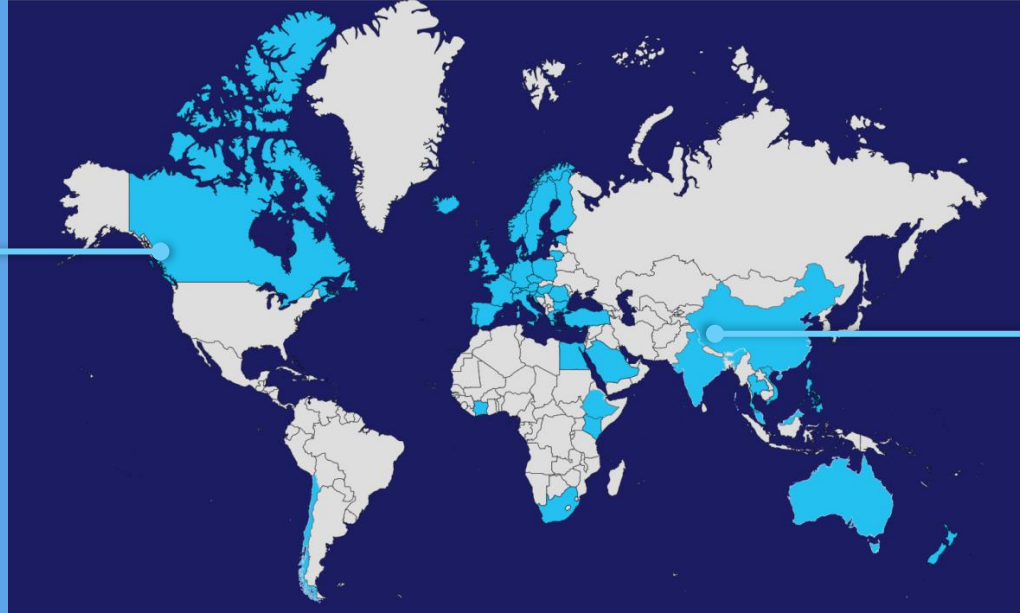
Tim Clench (Expert Advisor Scheme 5B)



# UK NEQAS for H&I: An Overview



>320 participants



>50 countries



# UK NEQAS for H&I is 50!



1975

- National Tissue Typing and Reference Laboratory (now OTDT) in Bristol initiated a quality control scheme for HLA typing (1A) and crossmatching (2A)
- Exercises included technical comparisons e.g. comparing batches of complement and the sensitivity of different techniques

1988

- Introduction of HLA antibody specification (Scheme 3)

1989

- First Annual Participant Meeting
- Idea for a UK professional society for tissue typers: BSHI formed

1990

- HLA disease association scheme – B27

1992

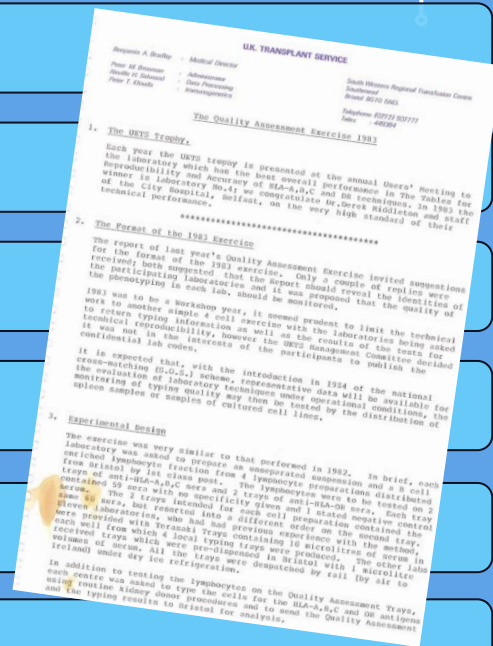
- HLA genotyping scheme introduced: Class II only (Class I not introduced until 1999)

1994

- First international participants joined

1998

- Founding member of the EFI EPT Committee



# UK NEQAS for H&I is 50!



2000

- UK NEQAS for H&I relocated to the Welsh Blood Service
- New HFE (5) and antibody detection (6) schemes introduced
- First educational schemes for HLA typing

2008

- Drug hypersensitivities introduced for Abacavir (7)

2011

- 2<sup>nd</sup> Field HLA Genotyping (4A2) developed
- HLA disease association (8) and KIR genotyping (9) introduced

2015

- ISO 17043 accreditation
- HPA genotyping pilot (10)

2016

- HPA Antibody pilot (11)

2018

- 3<sup>rd</sup> Field HLA Genotyping

2025

- HNA genotyping (12) and HNA antibody detection (13)



# Things To Note...



## Presentation Focus...

Performance, key trends,  
discussion points and 2025  
changes



## Further Details...

The presentation will be  
available to view on our  
website.



## Lab Locations...

Generally:  
1-100 = UK & Ireland.  
101+ = Rest of the world

# Scheme Assessments



- Most Schemes assessed on a consensus basis using a 75% consensus level i.e. 75% of reports must agree on a result for it to be assessed.
- Reference typing results are used for typing/disease schemes if consensus not reached plus educational schemes where required:
  - ▶ *e.g. Scheme 8: HLA Genotyping for Coeliac and Other HLA Associated Diseases*
- Equivocal result only accepted for Scheme 2B.
- All Not Tested (NT) results excluded from assessment.
- Labs that fail to return results or do not provide a valid reason for NT are assessed as unacceptable.



# Unsatisfactory Performance (UP)



- Each scheme has minimum annual performance criteria:

- ▶ *HLA Typing schemes 90%*
- ▶ *Crossmatching 85%*
- ▶ *Disease Association Schemes 100%*
- ▶ *Antibody Specificity 75%*
- ▶ *Antibody Detection 80%*



- Participants that do not meet the minimum criteria are classed as **unsatisfactory performers**.
- Must complete a root cause analysis and CAPA form.



# Changes for 2025-26



## Pilot Schemes

Scheme 12 – HNA Genotyping  
Scheme 13 – HNA Antibody Detection

## Sustainability

Paperless: no distribution slips  
Packaging: reduce plastic



## Courier

Reduction in costs  
Used for domestic and international



## Scheme Changes

Scheme 7 – HLA-related Pharmacogenetics  
Scheme 8 – now only HLA Associated Diseases



# Pilot Schemes



12

## HNA Genotyping



Assess participants ability to correctly determine HNA polymorphisms



4 blood samples in two distributions



Pilot Schemes are free of charge and not formally assessed

13

## HNA Antibody Detection/Specification



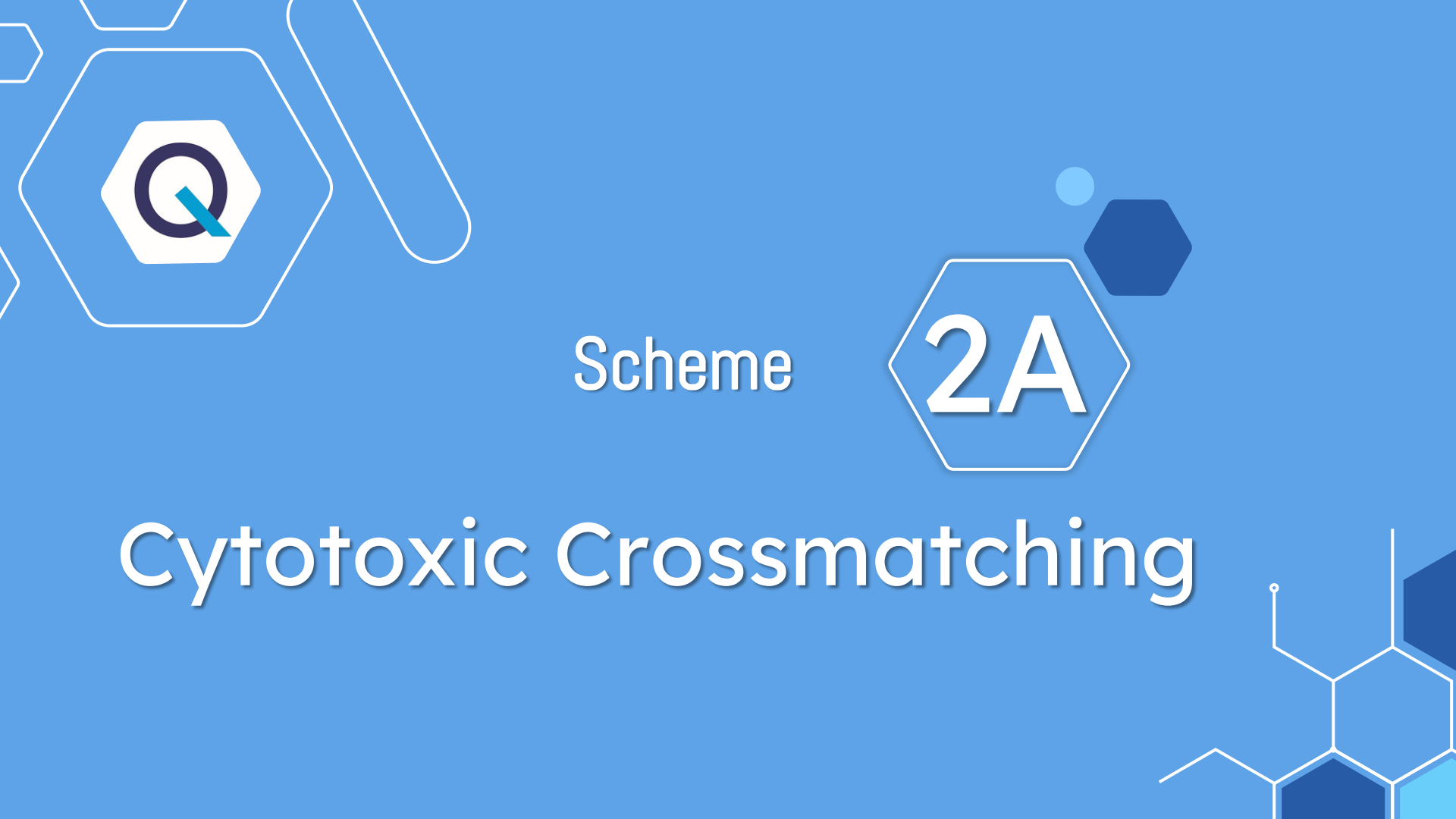
Assess participants ability to correctly detect the presence and specificity of HNA antibodies



4 serum/plasma samples in two distributions



Pilot Schemes are free of charge and not formally assessed

The background is a solid blue color. In the top left, there is a large white hexagon containing a dark blue 'Q' with a teal diagonal line. To its right is a white test tube shape. In the top right, there is a small light blue circle and a dark blue hexagon. In the bottom right, there is a complex molecular structure made of white lines and blue hexagons.

Scheme

2A

# Cytotoxic Crossmatching

# Scheme 2A – Cytotoxic Crossmatch



## Purpose

Assess participants ability to determine cell/serum cytotoxicity crossmatch status

## Satisfactory Performance

85% of reports agree with consensus in distribution year for each cell/DTT type



## Consensus

At least 75% agreement on pos/neg result

*10 blood samples, 40 serum samples over 5 distributions*



# Scheme 2A: Performance

| All cells with and without DTT                                | 2018             | 2019           | 2020         | 2021        | 2022           | 2023          | 2024          |
|---------------------------------------------------------------|------------------|----------------|--------------|-------------|----------------|---------------|---------------|
| Number of Participants (UK&I)                                 | 71<br>(18)       | 71<br>(22)     | 66<br>(16)   | 63<br>(15)  | 59<br>(15)     | 47<br>(10)    | 47<br>(8)     |
| Number with Unsatisfactory Performance<br>( $< 85\%$ ) (UK&I) | 16 (7)           | 5 (1)          | 7 (0)        | 4 (0)       | 6 (3)          | 2 (2)         | 5 (0)         |
| % Unsatisfactory Performance<br>(UK&I)                        | 22.5%<br>(38.8%) | 7.0%<br>(4.5%) | 10.6%<br>(0) | 6.3%<br>(0) | 10.2%<br>(20%) | 4.2%<br>(20%) | 10.8%<br>(0%) |

2024: 5 Unsatisfactory Performers (0 UK & Ireland)





# Scheme 2A: Performance by category

|                                        | PBL        | PBL +DTT   | T Cell     | T Cell +DTT | B Cell       | B Cell +DTT |
|----------------------------------------|------------|------------|------------|-------------|--------------|-------------|
| Crossmatches assessed (n=40)<br>(UK&I) | 31<br>(31) | 33<br>(33) | 36<br>(39) | 34<br>(36)  | 25<br>(33) ↓ | 30<br>(36)  |
| NT – Assessed samples only             | 17.1%      | 14.2%      | 37.8%      | 33.5%       | 56.9% ↑      | 53.6%       |
| % incorrect assignments                | 4.1%       | 3.0%       | 9.3%       | 7.1%        | 20.4% ↑      | 11.3%       |
| False Positive                         | 3.2%       | 1.5%       | 5.5%       | 5.3%        | 11.2% ↑      | 5.0%        |
| False Negative                         | 0.9%       | 1.5%       | 3.8%       | 1.8%        | 9.2% ↑       | 6.3%        |



# Scheme 2A: Unacceptable Performers 2024



| Lab ID | PBL -DTT | T -DTT | B -DTT | PBL + DTT | T + DTT | B + DTT | Lab Identified Error                  |
|--------|----------|--------|--------|-----------|---------|---------|---------------------------------------|
| 133    |          | 80%    |        |           |         |         | No response                           |
| 189    | 81.3%    |        |        | 82.4%     |         |         | Procedural error                      |
| 226    |          |        | 84.4%  |           |         |         | Interpretation criteria (pos cut off) |
| 232    |          |        |        |           |         | 83.3%   | Sample quality (delivery delays)      |
| 1412   | 75%      | 74.4%  |        |           |         |         | Poor cell prep / reagent issues       |





# Scheme 2A: Discussion

- Not all Scheme 2A results will reach consensus (that's ok!)
- B-cells are difficult (transport, non-specific binding)
- Only partially emulates clinical practice
- 2A is a technical assessment of cytotoxic crossmatching and should not be 'interpreted'
- Lab's need to ensure that all test parameters and acceptance criteria are met prior to reporting NEQAS samples
  - CDC assays are not quantitative so reliant on subjective assessment





Scheme



2B

Crossmatching by Flow Cytometry



# Scheme 2B: Crossmatching by Flow Cytometry



## Purpose

Assess participants ability to determine cell/serum flow crossmatch status

## Satisfactory Performance

85% reports agree with consensus in distribution year for each cell type



## Consensus

At least 75% agreement on pos/neg or equivocal result

*10 blood samples, 40 serum samples over 5 distributions*



# Scheme 2B: Performance

|                                                               | 2018            | 2019            | 2020         | 2021        | 2022            | 2023       | 2024       |
|---------------------------------------------------------------|-----------------|-----------------|--------------|-------------|-----------------|------------|------------|
| Number of Participants (UK&I)                                 | 83<br>(22)      | 84<br>(23)      | 80<br>(21)   | 80<br>(22)  | 84<br>(19)      | 83<br>(21) | 79<br>(20) |
| Number with Unsatisfactory Performance<br>( $< 85\%$ ) (UK&I) | 15<br>(2)       | 12<br>(1)       | 11<br>(0)    | 5<br>(0)    | 6<br>(2)        | 10<br>(0)  | 4<br>(0)   |
| % Unsatisfactory Performance (UK&I)                           | 18.1%<br>(9.1%) | 14.2%<br>(4.3%) | 13.8%<br>(0) | 6.3%<br>(0) | 7.1%<br>(10.5%) | 12%<br>(0) | 5%<br>(0)  |

2024: 4 Unsatisfactory Performers (0 UK & Ireland)



# Scheme 2B: Performance by Category



|                                           | T Cells          |                  |                | B Cells        |                |                |
|-------------------------------------------|------------------|------------------|----------------|----------------|----------------|----------------|
|                                           | UK&I             | RoW<br>PC        | RoW<br>WB      | UK&I           | RoW<br>PC      | RoW<br>WB      |
| Number of participants                    | 20               | 28               | 26             | 20             | 28             | 24             |
| Number of XM assessed<br>(>75% consensus) | 35/40<br>(87.5%) | 31/40<br>(77.5%) | 36/40<br>(90%) | 36/40<br>(90%) | 36/40<br>(90%) | 36/40<br>(90%) |
| Number of Positive XM                     | 24 (69%)         | 20 (65%)         | 22 (61%)       | 32 (89%)       | 29 (81%)       | 28 (78%)       |
| Number of Negative XM                     | 11 (31%)         | 11 (35%)         | 14 (39%)       | 4 (11%)        | 7 (19%)        | 8 (22%)        |
| Number of incorrect assignments           | 20               | 37               | 21             | 18             | 41             | 36             |
| Number of False Pos                       | 10               | 12               | 13             | 6              | 14             | 12             |
| Number of False Neg                       | 10               | 25               | 8              | 12             | 27             | 24             |
| Number of equivocal assignments           | 0                | 6                | 0              | 1              | 9              | 1              |
| Number of samples NT                      | 25               | 162              | 58             | 56             | 111            | 64             |

UK&I and RoW receive different blood samples



## Scheme 2B: Unacceptable Performers 2024

| Lab | T Cell | No. of results submitted | B Cell | No. of results submitted | Issue                                              |
|-----|--------|--------------------------|--------|--------------------------|----------------------------------------------------|
| 118 | 90%    | 24/40                    | 77%    | 24/40                    | Sample quality (delivery delay)                    |
| 139 | 87%    | 26/40                    | 78%    | 20/40                    | No response                                        |
| 189 | 42%    | 32/40                    | 41%    | 32/40                    | Technical issue (cytometer)                        |
| 191 | 97%    | 40/40                    | 83%    | 40/40                    | Sensitivity issue (delivery delays/poor cell prep) |



4 labs with UP (<85%)



Scheme

6

# HLA Antibody Detection



# Scheme 6: HLA Antibody Detection

## Purpose

Assess participants ability to determine **presence or absence** of HLA antibodies

## Satisfactory Performance

80% reports agree with consensus in distribution year



## Consensus

At least 75% agreement on presence/absence of HLA antibodies

*12 serum samples over 3 distributions*





# Scheme 6: Performance

3 Unsatisfactory Performers (1 UK&I)

|                                                       | 2018         | 2019         | 2020         | 2021       | 2022       | 2023        | 2024           |
|-------------------------------------------------------|--------------|--------------|--------------|------------|------------|-------------|----------------|
| Number of Participants (UK&I)                         | 88<br>(25)   | 82<br>(25)   | 74<br>(25)   | 71<br>(23) | 68<br>(23) | 68<br>(23)  | 67<br>(24)     |
| Number with Unsatisfactory Performance (< 80%) (UK&I) | 5 (0)        | 8 (0)        | 2 (0)        | 0 (0)      | 4 (0)      | 3 (0)       | 3 (1)          |
| % Unsatisfactory Performance                          | 5.7%<br>(0%) | 9.7%<br>(0%) | 2.7%<br>(0%) | 0%<br>(0%) | 5%<br>(0%) | 4.4%<br>(0) | 4.4%<br>(4.1%) |

38% negative

58% positive

4% samples not assessed





## Scheme 6: Unacceptable Performers 2024

| Lab  | Performance (<80%) | Issue                        |
|------|--------------------|------------------------------|
| 11   | 75.0%              | Interpretation (pos cut off) |
| 128  | 75.0%              | Interpretation / kit issue   |
| 1418 | 58.3%              | Interpretation / kit issue   |

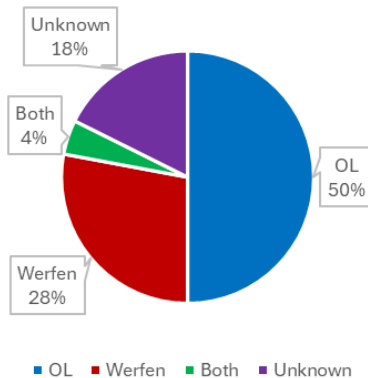
3 labs with UP (<85%)



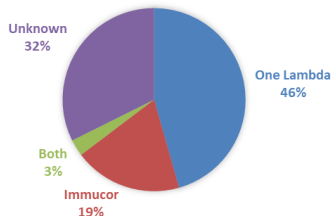


# Scheme 6: Kit Use and Performance

Manufacturer of Kit Used for Antibody Detection  
(n=68)



MANUFACTURER OF KIT USED FOR  
ANTIBODY DETECTION (N=68)



| 2024-25 | Class I           |     |               |     | Class II          |     |               |     |
|---------|-------------------|-----|---------------|-----|-------------------|-----|---------------|-----|
|         | One Lambda (n=34) | %   | Werfen (n=19) | %   | One Lambda (n=34) | %   | Werfen (n=19) | %   |
| 601     | Positive          | 100 | Positive      | 100 | Positive          | 100 | Positive      | 100 |
| 602     | Positive          | 100 | Positive      | 100 | Negative          | 84  | Negative      | 100 |
| 603     | Positive          | 100 | Positive      | 100 | Positive          | 100 | Positive      | 100 |
| 604     | Positive          | 67  | Negative      | 80  | Negative          | 100 | Negative      | 87  |
| 605     | Positive          | 100 | Positive      | 100 | Positive          | 100 | Positive      | 100 |
| 606     | Positive          | 100 | Positive      | 94  | Positive          | 100 | Positive      | 100 |
| 607     | Negative          | 94  | Negative      | 100 | Negative          | 94  | Negative      | 94  |
| 608     | Negative          | 84  | Negative      | 100 | Negative          | 77  | Negative      | 94  |
| 609     | Positive          | 100 | Positive      | 94  | Positive          | 100 | Positive      | 100 |
| 610     | Negative          | 66  | Negative      | 100 | Positive          | 100 | Positive      | 100 |
| 611     | Positive          | 100 | Positive      | 100 | Positive          | 97  | Positive      | 100 |
| 612     | Negative          | 97  | Negative      | 94  | Negative          | 97  | Negative      | 100 |



Scheme

3

# HLA Antibody Specificity Analysis



# Scheme 3: HLA Antibody Specificity Analysis



## Purpose

Assess participants ability to determine **specificity** of HLA antibodies



## Consensus

At least 75% agreement on presence of HLA antibodies, 95% agreement on absense.

## Satisfactory Performance

75% reports agree with consensus in distribution year



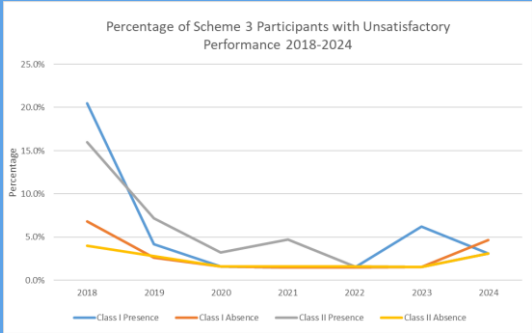
*10 serum samples over 3 distributions*



# Scheme 3: Performance



| Class I                                       |          | 2018       | 2019       | 2020       | 2021       | 2022       | 2023       | 2024       |
|-----------------------------------------------|----------|------------|------------|------------|------------|------------|------------|------------|
| Number of Participants (UK&I)                 |          | 73<br>(25) | 70<br>(25) | 64<br>(24) | 65<br>(24) | 65<br>(24) | 64<br>(24) | 63<br>(23) |
| Number with Unsatisfactory Performance (UK&I) | Presence | 15 (1)     | 3 (0)      | 1 (0)      | 1 (0)      | 1 (0)      | 4 (0)      | 2 (0)      |
|                                               | Absence  | 5 (0)      | 2 (0)      | 1 (0)      | 1 (0)      | 1 (0)      | 1 (0)      | 3 (0)      |
| % Unsatisfactory Performance                  | Presence | 20.5%      | 4.2%       | 1.6%       | 1.5%       | 1.5%       | 6.3%       | 3.2%       |
|                                               | Absence  | 6.8%       | 2.6%       | 1.6%       | 1.5%       | 1.5%       | 1.5%       | 4.8%       |



Overall 3  
labs with UP  
(0 UK&I)

| Class II                                      |          | 2018       | 2019       | 2020       | 2021       | 2022       | 2023       | 2024       |
|-----------------------------------------------|----------|------------|------------|------------|------------|------------|------------|------------|
| Number of Participants (UK&I)                 |          | 75<br>(25) | 69<br>(25) | 63<br>(24) | 64<br>(24) | 64<br>(24) | 64<br>(24) | 63<br>(23) |
| Number with Unsatisfactory Performance (UK&I) | Presence | 12 (0)     | 5 (0)      | 2 (0)      | 3 (0)      | 1 (0)      | 1 (0)      | 2 (0)      |
|                                               | Absence  | 3 (0)      | 2 (0)      | 1 (0)      | 1 (0)      | 1 (0)      | 1 (0)      | 2 (0)      |
| % Unsatisfactory Performance                  | Presence | 16.0%      | 7.2%       | 3.2%       | 4.7%       | 1.6%       | 1.6%       | 3.2%       |
|                                               | Absence  | 4.0 %      | 2.8%       | 1.6%       | 1.6%       | 1.6%       | 1.6%       | 3.2%       |



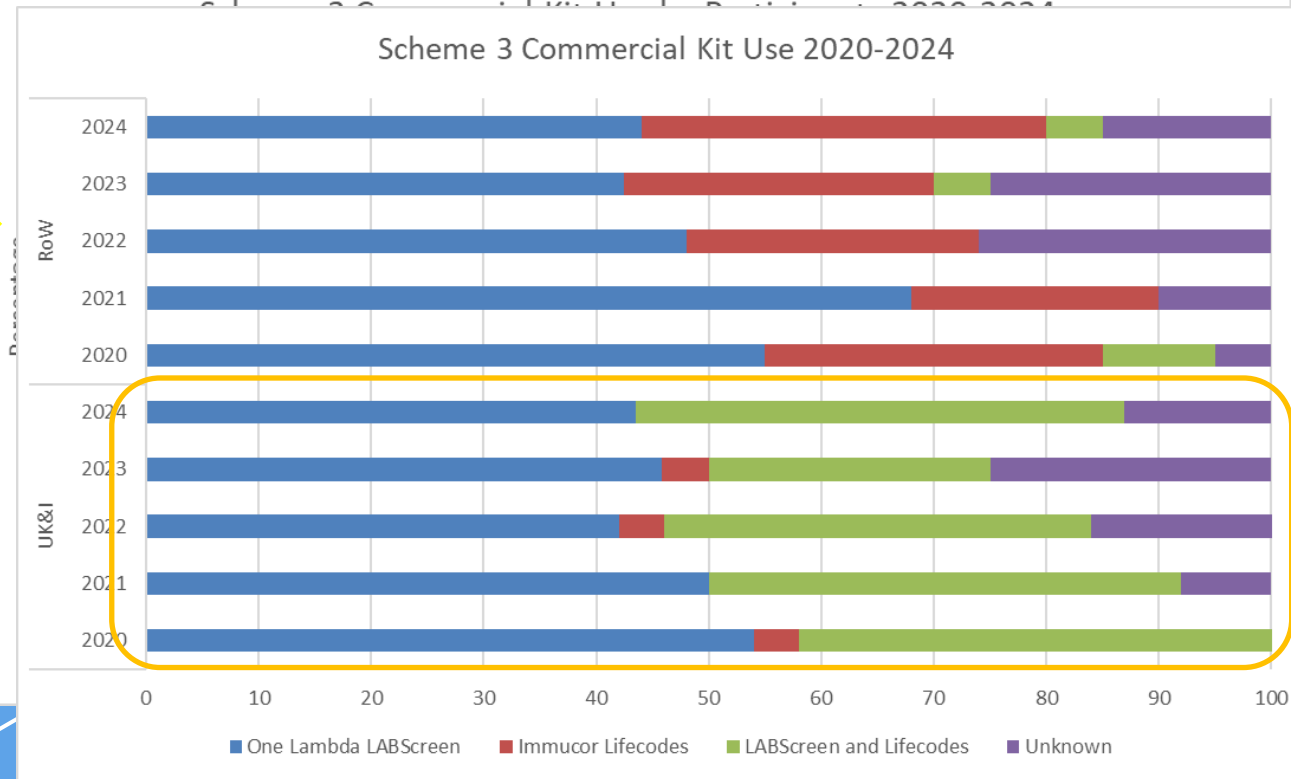
# Scheme 3: Unacceptable Performers 2024

3 labs (0 UK&I) with UP (<75%)

| Lab  | Class I  |         | Class II |         | CAPA                                  | Kit        |
|------|----------|---------|----------|---------|---------------------------------------|------------|
|      | Presence | Absence | Presence | Absence |                                       |            |
| 302  | 62%      | 53%     | 57%      | 72%     | No reply                              | Werfen     |
| 1312 | 74%      | 55%     | 48%      | 52%     | No reply                              | One Lambda |
| 1412 |          | 58%     |          |         | Interpretation issue<br>(pos cut off) | Werfen     |



# Scheme 3: Kit Use 2020-2024



Overall OL kits are the most widely used

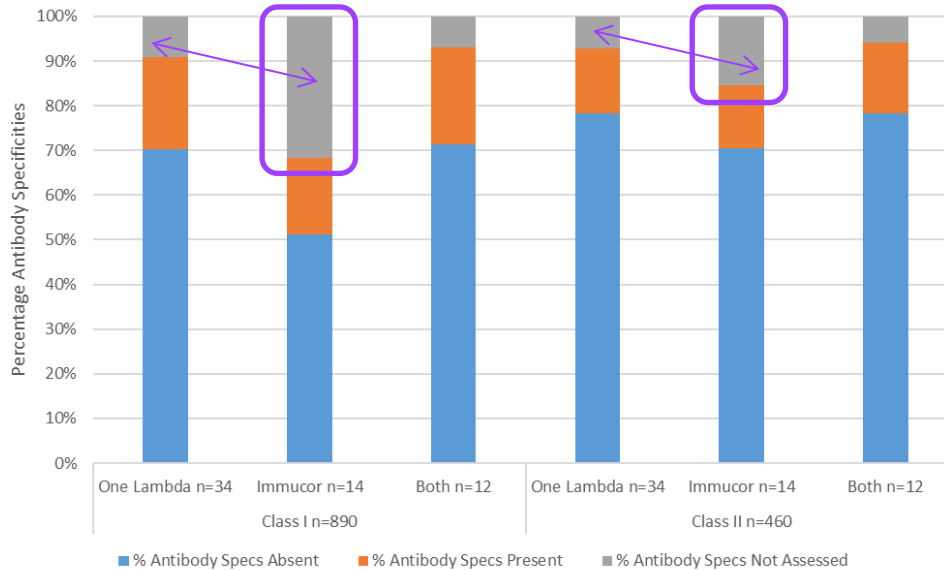
UK&I labs are more likely to use a combination of kits

Werfen only kit use more prevalent in RoW labs

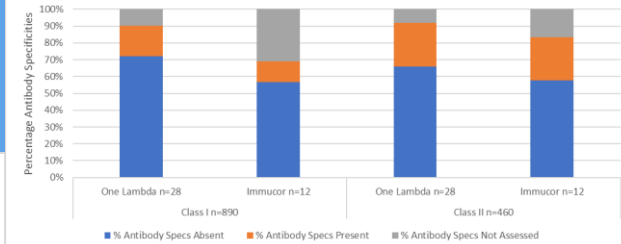
# Scheme 3: Results by Kit Use



Scheme 3 2024 - Comparison of HLA Antibody Specificity Results Between Kits



Scheme 3 2023 - Comparison of HLA Antibody Specificity Results Between Kits



Similar percentage of antibodies reach consensus present (orange) in both kits

Less concordance in 'absent' antibodies

Greater percentage of Class I antibodies classed as not assessed in Werfen group

# Scheme 3: Kit Use and Performance



Average overall satisfactory performance for detecting the 'presence' and 'absence' of antibodies was marginally higher for users of both kits.

2024-25

| Average Performance | Class I              |                  |                | Class II             |                  |                |
|---------------------|----------------------|------------------|----------------|----------------------|------------------|----------------|
|                     | One Lambda<br>(n=34) | Werfen<br>(n=14) | Both<br>(n=12) | One Lambda<br>(n=34) | Werfen<br>(n=14) | Both<br>(n=12) |
| Presence            | 94.1%                | 86.5%            | 97.2% ↑        | 94.5%                | 89.7%            | 97.8% ↑        |
| Absence             | 97.5%                | 94.2%            | 99.5% ↑        | 97.4%                | 95.3%            | 99.5% ↑        |

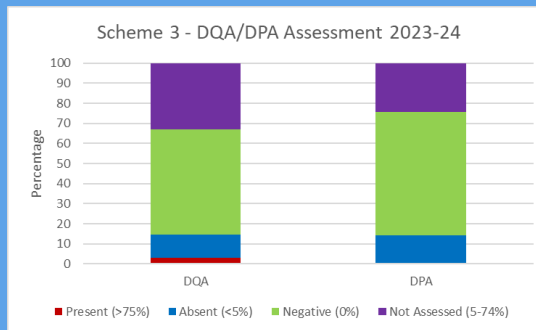
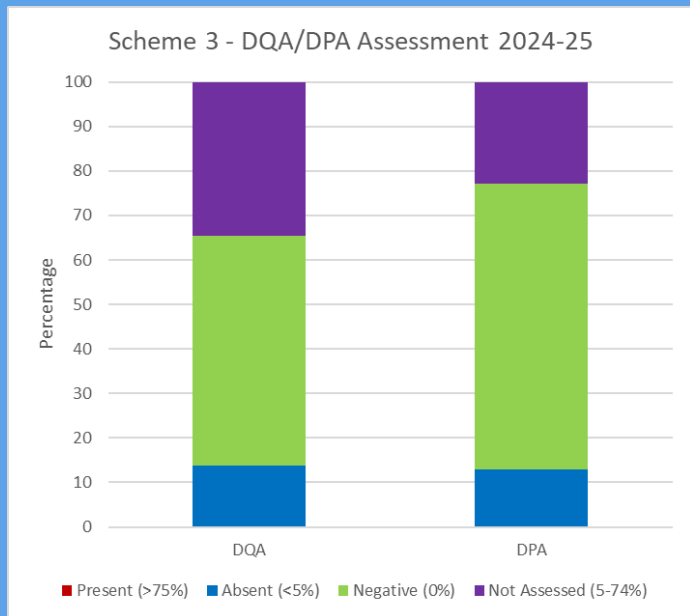
# Scheme 3: DQA/DPA Antibody Reporting



Reporting of antibodies to HLA-DQA and –DPA is optional and not assessed.

Overall 42/63 (67%) report DQA, 38/63 (60%) report DPA

Previously 58% and 50%



An analysis of the data submitted for DQA and DPA antibodies in 2024-25 and 2023-24 was performed.

Large proportion of samples are negative or consensus absent.

No antibodies deemed positives.

Approx 25-30% not assessed.



Scheme

11

HPA Antibody Detection/Specification



# Scheme 11: HPA Antibody Detection/Specification

## Purpose

Assess participants ability to correctly determine presence and specificity of HPA antibodies.

## Satisfactory Performance

At least 75% of specificities in agreement with the consensus result in a distribution year.



## Consensus

Presence of specificity determined by at least 75% agreement and absence determined by at least 95% agreement.

*8 serum/plasma samples over 2 distributions*





# Scheme 11: Performance

- 1 Unsatisfactory Performers (0 UK&I)

|                                                                  | 2018      | 2019      | 2020      | 2021      | 2022      | 2023      | 2024      |
|------------------------------------------------------------------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|
| Number of Participants<br>(UK&I)                                 | 35<br>(4) | 39<br>(5) | 42<br>(4) | 43<br>(4) | 43<br>(4) | 43<br>(4) | 44<br>(4) |
| Number with Unsatisfactory<br>Performance<br>( $< 75\%$ ) (UK&I) | 1<br>(0)  | 1<br>(0)  | 3<br>(0)  | 6<br>(0)  | 2<br>(0)  | 9<br>(0)  | 1<br>(0)  |
| % Unsatisfactory<br>Performance                                  | 2.9%      | 2.6%      | 7.1%      | 13.9%     | 4.5%      | 20.9%     | 2.3%      |





# Scheme 11: HPA Antibody Detection/Specification



| 2024 Sample | HPA Detection | HLA Detection | Expected Result     | HPA Antibody Consensus |                                             |
|-------------|---------------|---------------|---------------------|------------------------|---------------------------------------------|
|             |               |               |                     | Presence               | Absence                                     |
| 1           | 97.6% Pos     | 86.5% Pos     | HPA-5b              | HPA 5b 97.6%           | HPA GP1a/11a 97.2%%                         |
| 2           | 100% Neg      | 94.4% Neg     | HPA neg,<br>HLA neg | N/A                    | N/A                                         |
| 3           | 100% Pos      | 100% Neg      | HPA-5b,<br>HLA neg  | HPA 5b 100%            | HPA GP1a/11a 97.2%                          |
| 4           | 97.5% Pos     | 91.4% Neg     | HPA-1a              | HPA-1a 97.5%           | HPA-3a 92.3%, 4b 90.9%,<br>GPIIb/IIIa 91.7% |
| 5           | 100% Neg      | 94.3% Neg     | HPA neg,<br>HLA neg | N/A                    | N/A                                         |
| 6           | 60% Neg       | 100% Pos      | HPA-15b             | N/A                    | HPA 15b 60% GP11b/111a<br>97.3%             |
| 7           | 97.2% Neg     | 94.3% Neg     | HPA neg,<br>HLA neg | N/A                    | HPA 2b 97.2%                                |
| 8           | 97.1% Neg     | 94.3% Neg     | HPA neg,<br>HLA neg | N/A                    | HPA GP1b 97.1%                              |

# Scheme 11: Unacceptable Performers 2024

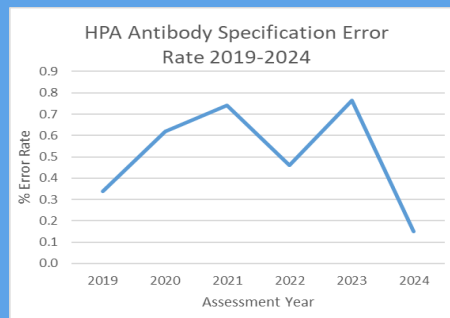


| Lab  | HPA Presence | HPA Absence | Samples reported | Method                  | Error       |
|------|--------------|-------------|------------------|-------------------------|-------------|
| 1378 | 50%          | 100%        | 8/8              | Immucor/Werfen<br>PakLx | No response |

1 lab with UP (<75%)

# Scheme 11: Analysis of Errors 2024

- Error rate extremely low (overall 0.15%) but errors often at clinically relevant polymorphisms.
- Errors found at HPA-1a (n=1, error rate 0.3%), 2b (n=1, error rate 0.3%), 5b (n=1, error rate 0.3%) and some glycoproteins.



| Errors 2024  | HPA-1a | HPA-1b | HPA-2a | HPA-2b | HPA-3a | HPA-3b | HPA-4a | HPA-4b | HPA-5a | HPA-5b | HPA-6a | HPA-6b | HPA-15a | HPA-15b | GP IIb/IIIa | GP Ia/IIa | GP Ib | GP IV | CD109 | Total |
|--------------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|---------|---------|-------------|-----------|-------|-------|-------|-------|
| False Pos    | 0      | 0      | 0      | 1      | 0      | 0      | 0      | 0      | 0      | 0      | 0      | 0      | 0       | 0       | 1           | 2         | 1     | 0     | 0     | 5     |
| False Neg    | 1      | 0      | 0      | 0      | 0      | 0      | 0      | 0      | 0      | 1      | 0      | 0      | 0       | 0       | 0           | 0         | 0     | 0     | 0     | 2     |
| Total Errors | 1      | 0      | 0      | 1      | 0      | 0      | 0      | 0      | 0      | 1      | 0      | 0      | 0       | 0       | 1           | 2         | 1     | 0     | 0     | 7     |
| % Error Rate | 0.3    | 0.0    | 0.0    | 0.3    | 0.0    | 0.0    | 0.0    | 0.0    | 0.0    | 0.3    | 0.0    | 0.0    | 0.0     | 0.0     | 0.3         | 0.7       | 0.4   | 0.0   | 0.0   | 0.15  |
| Total Tested | 320    | 304    | 288    | 288    | 312    | 312    | 284    | 268    | 320    | 328    | 108    | 108    | 120     | 120     | 292         | 292       | 280   | 264   | 120   | 4728  |

- False positive (n=5) more common than false negative (n=2) errors.

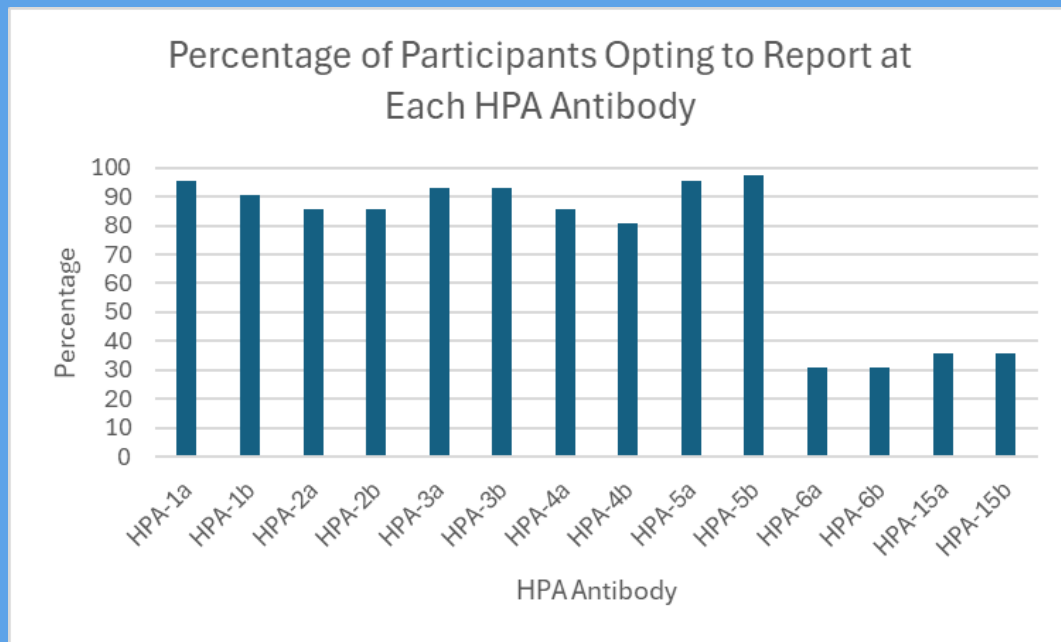
- Most labs had only 1 or 2 errors

| Number of Errors | Number of Labs |
|------------------|----------------|
| 1                | 3              |
| 2                | 2              |



# Scheme 11: Selection of HPA Antibodies for Assessment

- Introduced in 2024-25
- Labs can select any/all HPA antibodies for assessment based on their clinical strategy





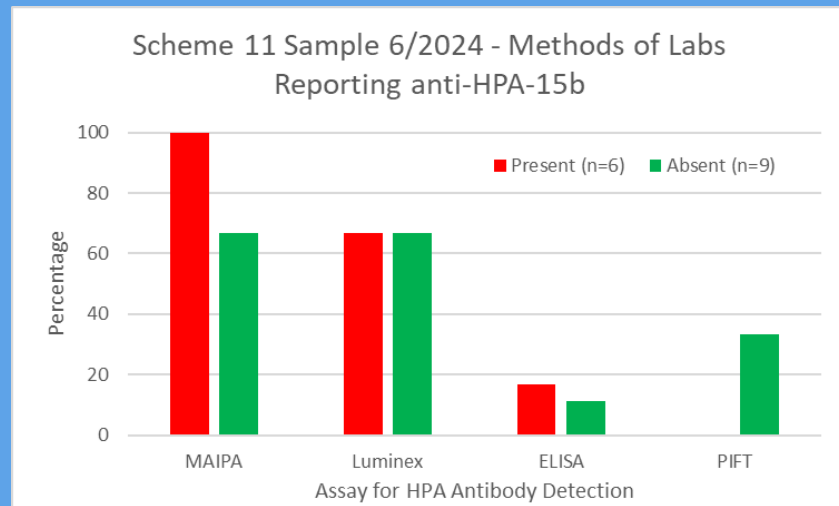
# Scheme 11: HPA-15 Detection

- 36% (15/42) of labs selected to be assessed for HPA-15 antibody detection
- We sent NIBSC Standard HPA-15b
- Sample 6/2024:

60% (9/15) reported HPA-15b **absent**

40% (6/15) reported HPA-15b **present**

100% reported HLA ab present



Key Data from the Schemes  
**Deborah Pritchard**  
UK NEQAS for H&I Director





Scheme



1A

HLA Phenotyping



# Scheme 1A: HLA Phenotyping

## Purpose

Assess participants ability to use serological and supplementary methods to correctly identify HLA phenotype

## Satisfactory Performance

9 or more complete HLA phenotypes in agreement with consensus per distribution year.



## Consensus

At least 75% agreement on each specificity.

*10 blood samples over 5 distributions*





# Scheme 1A: Performance

- 0 labs with unsatisfactory performance

|                                                       | 2018   | 2019   | 2020   | 2021   | 2022   | 2023   | 2024   |
|-------------------------------------------------------|--------|--------|--------|--------|--------|--------|--------|
| Number of Participants (UK&I)                         | 38 (6) | 38 (5) | 34 (4) | 33 (2) | 28 (1) | 23 (0) | 19 (0) |
| Number with Unsatisfactory Performance (< 90%) (UK&I) | 6 (1)  | 8 (1)  | 3 (1)  | 2 (0)  | 2 (0)  | 2 (0)  | 0 (0)  |
| % Unsatisfactory Performance                          | 15.8%  | 21.1%  | 8.8%   | 6.1%   | 7.1%   | 8.7%   | 0%     |





# Scheme 1A: 2024 Incorrect Assignments

3/190 (1.6%) incorrect HLA types in 2024 reported by 3 labs:

2 reports that contained broad not split specificity (e.g. B40 v B60)

1 clerical/typo error

0 labs with  
unsatisfactory  
performance

# Scheme 1A: 2024 Incorrect Assignments (not resulting in UPs)



| Sample | ID        | Consensus         | Report            |
|--------|-----------|-------------------|-------------------|
| 1A 03  | 268       | A1, A26; B38, B57 | A1, A26; B68, B57 |
| 1A 05  | 147 + 159 | A3, A23; B49, B65 | A3, A23; B49, B14 |





Scheme



4A1

HLA Typing at 1<sup>st</sup> Field Resolution



# Scheme 4A1: HLA Typing at 1<sup>st</sup> Field Resolution

## Purpose

Assess participants ability to correctly determine HLA genotypes at the 1<sup>st</sup> field resolution.

## Satisfactory Performance

9 or more full HLA types in agreement with consensus/reference result in a distribution year.



## Consensus

At least 75% agreement on each allele. When consensus is not met, a reference result is used. Reference result is always used for DPB1 assessment

*10 blood samples over 3 distributions*





# Scheme 4A1: Performance

- 6 labs with unsatisfactory performance (2 UK&I)

|                                                          | 2018            | 2019         | 2020         | 2021         | 2022         | 2023          | 2024           |
|----------------------------------------------------------|-----------------|--------------|--------------|--------------|--------------|---------------|----------------|
| Number of Participants<br>(UK&I)                         | 105<br>(28)     | 100<br>(28)  | 88<br>(26)   | 82<br>(25)   | 81<br>(25)   | 81<br>(25)    | 84<br>(24)     |
| Number with Unsatisfactory<br>Performance (< 90%) (UK&I) | 15 (1)          | 4 (1)        | 8 (0)        | 6 (1)        | 7 (0)        | 11 (1)        | 6 (2)          |
| % Unsatisfactory<br>Performance                          | 14.3%<br>(3.6%) | 4%<br>(3.6%) | 9.1%<br>(0%) | 7.3%<br>(4%) | 8.4%<br>(0%) | 13.6%<br>(4%) | 7.1%<br>(8.3%) |



# Scheme 4A1: 2024-25 Incorrect Assignments

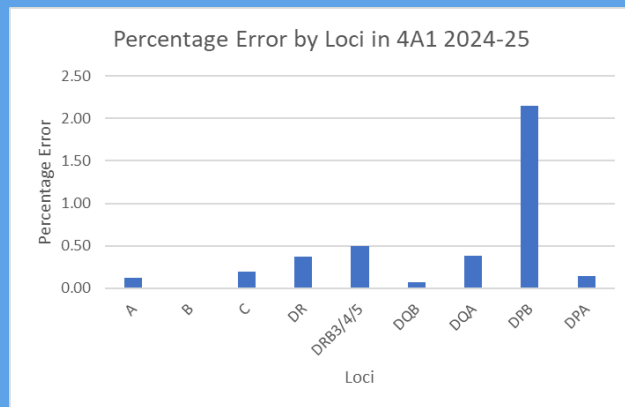
39/11493 (0.34%) errors reported by 20 different labs (3 UK&I) – last year 0.26%

29 samples contained an error:

- 21 samples with incorrect assignments *e.g. C\*13 rather than C\*07*
- 3 samples with nomenclature issue
- 3 missed assignment (reported homozygous when heterozygous)
- 2 samples with DRB3/4/5 presence/absence reported incorrectly

24 (83%) HLA types with one error  
5 (17%) HLA types with multiple errors

14 (70%) labs made 1 error  
3 (15%) labs made 2 errors  
3 (15%) lab made 3 errors





# Scheme 4A1: Unacceptable Performers 2024

| Lab  | Sample   | Error                                        | CAPA Response                              |
|------|----------|----------------------------------------------|--------------------------------------------|
| 41   | 01+02+03 | Multiple issues                              | Transcription error / interpretation error |
| 6    | 03+04+07 | Incorrect DPB1* assignments                  | Limitations of kit                         |
| 172  | 03+06+08 | A* and DQB1* missed / assignment errors      | Procedural error (new method)              |
| 1412 | 05+10    | DQA1* and DPA1* and DPB1* missed assignments | Procedural error (reagent issue)           |
| 1418 | 07+09    | Reporting errors                             | Transcription errors (staffing levels)     |
| 1443 | 09+10    | Multiple reporting errors                    | EQA specific result entry errors           |





Scheme



4A1i

Interpretive HLA Genotype



# Scheme 4A1: Interpretive HLA Genotype

## Purpose

Assess participants ability to correctly interpret their 4A1 genotype result to the 'split' specificity level.

## Satisfactory Performance

9 or more full HLA types in agreement with consensus/reference result in a distribution year.



## Consensus

At least 75% agreement on each specificity. When consensus is not met, a reference result is used.

*10 HLA genotypes from Scheme 4A1*





# Scheme 4A1i: Performance

- 5 lab with unsatisfactory performance (0 UK&I)

|                                                                | 2018       | 2019       | 2020       | 2021       | 2022       | 2023       | 2024       |
|----------------------------------------------------------------|------------|------------|------------|------------|------------|------------|------------|
| Number of Participants<br>(UK&I)                               | 40<br>(21) | 44<br>(22) | 44<br>(22) | 42<br>(21) | 40<br>(21) | 40<br>(21) | 41<br>(20) |
| Number with<br>Unsatisfactory<br>Performance (< 90%)<br>(UK&I) | 6 (0)      | 8 (1)      | 6 (2)      | 5 (1)      | 2 (0)      | 1 (0)      | 5 (0)      |
| % Unsatisfactory<br>Performance                                | 15.0%      | 18.1%      | 13.6%      | 11.9%      | 5.0%       | 2.5%       | 12.2%      |

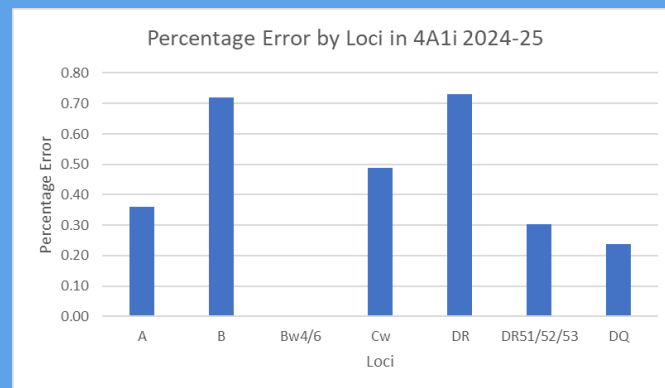
# Scheme 4A1i: 2024-25 Incorrect Assignments



- 24/5778 (0.42%) incorrect results reported by 7 different labs (0 UK&I) – last year 0.35%
- 17 samples contained an error:
  - 5 reporting at broad not split specificity level
  - 3 samples with incorrect assignments
  - 3 samples with missing assignment (reported homozygous when heterozygous)
  - 3 sample with incorrect uses of nomenclature
  - 3 samples with errors at presence/absence of DR51/52/53

2 (29%) labs made 1 error  
5 (71%) labs made 2-4 errors

12 (71%) HLA types with single errors  
5 (29%) HLA type with multiple errors





# Scheme 4A1i: Unacceptable Performers 2024

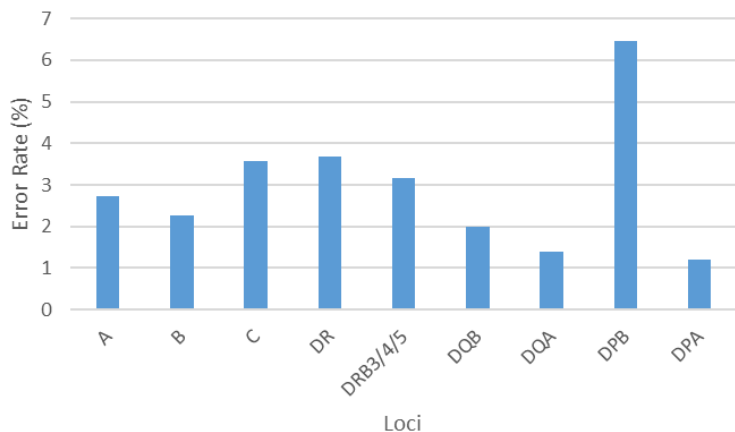
| Lab  | Sample     | Error                                              | CAPA Response                                      |
|------|------------|----------------------------------------------------|----------------------------------------------------|
| 1418 | 01-03 + 07 | Broads instead of splits/multiple reporting errors | Staff training / interpretation                    |
| 111  | 02+03+05   | Incorrect nomenclature                             | EQA specific reporting errors<br>(new participant) |
| 1412 | 02+05      | Multiple errors                                    | Transcription errors                               |
| 1372 | 02+06      | Multiple errors                                    | No response                                        |
| 1433 | 05+07      | Multiple errors                                    | Staff training / interpretation                    |



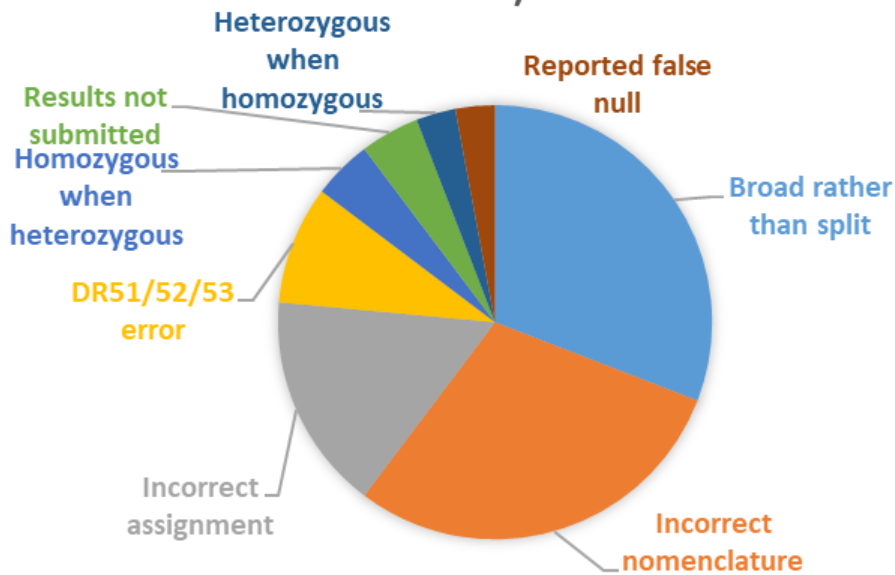


# Scheme 4A1: Types of Errors Over 5 Years

Scheme 4A1 - Error Rate by Loci (2020-2024)



SCHEME 4A1 - TYPES OF ERRORS (2020-2024)



# Scheme 4A1i: Serological Equivalents



- 4A1i Interpretative HLA Genotyping, which allows participants to translate genotypes to phenotypes
- Participants are expected to report to the 'split' specificity level using serological nomenclature, e.g. HLA-DQB1\*03:01 should be reported as DQ7 (DQ3)
- Knowledge and exposure of phenotyping and converting between genotypes and phenotypes may no longer be so commonplace

Reliance on LIMS/analysis software

No longer perform phenotyping

- Errors common due to reporting issues (broad rather split or using incorrect nomenclature)
- CAPA repeatedly cite issues due to staff training and knowledge
- Encourage utilisation of 4A1i for competency assessment

UK NEQAS  
Education | Quality | Global

Histocompatibility  
& Immunogenetics

**Scheme 4A1i – Competency Assessment Template**

**Task:** Convert HLA genotypes reported in Scheme 4A1 to 'split specificity' level phenotypes (minimising computer aided assistance).

**Learning Objective:** Accurately demonstrate knowledge of serological equivalents.

Name \_\_\_\_\_ Date \_\_\_\_\_

*Note resources used e.g. The HLA dictionary tool (if applicable):* \_\_\_\_\_

**Example:**

| Genotype               | A*      | A*      | B*      | B*      |         | C*      | C*      | DR*     | DR*     | DRB3/4/5* | DRB3/4/5*   | DQB1*       | DQB1*   | Score      |
|------------------------|---------|---------|---------|---------|---------|---------|---------|---------|---------|-----------|-------------|-------------|---------|------------|
| Sample ID              | 01      | 11      | 55      | 57      |         | 03:03   | 06      | 03:01   | 07      | DRB3*01   | DRB4*01N    | DQ          | 03:03   | Acceptable |
| Phenotype              | A       | A       | B       | B       | Bw4*    | Bw6*    | Cw      | Cw      | DR      | DR        | DRB1/52/53* | DRB1/52/53* | DQ      |            |
| Serological Equivalent | 1       | 11      | 55      | 57      | Present | Present | 9       | 6       | 17      | 7         | DR52        | N/A         | 2       | 9          |
| Assessment             | Correct | Correct | Correct | Correct | Correct | Correct | Correct | Correct | Correct | Correct   | Correct     | Correct     | Correct |            |

\*Indicate Presence or Absence  
\*\*Each full phenotype (compared to consensus result on UK NEQAS Portal) deemed acceptable

| Genotype               | A* | A* | B* | B* |      | C*   | C* | DR* | DR* | DRB3/4/5* | DRB3/4/5*   | DQB1*       | DQB1* | Score |
|------------------------|----|----|----|----|------|------|----|-----|-----|-----------|-------------|-------------|-------|-------|
| Sample ID              |    |    |    |    |      |      |    |     |     |           |             |             |       |       |
| Phenotype              |    |    |    |    | Bw4* | Bw6* | Cw | Cw  | DR  | DR        | DRB1/52/53* | DRB1/52/53* | DQ    | DQ    |
| Serological Equivalent |    |    |    |    |      |      |    |     |     |           |             |             |       |       |
| Assessment             |    |    |    |    |      |      |    |     |     |           |             |             |       |       |

\*Indicate Presence or Absence  
\*\*Each full phenotype (compared to consensus result on UK NEQAS Portal) deemed acceptable



UK NEQAS have developed a template:

<https://ukneqashandi.org.uk/app/uploads/2025/03/Scheme-4A1i-Template.docx>



Scheme



4A2

HLA Typing to 2<sup>nd</sup> or 3<sup>rd</sup> Field Resolution



# Scheme 4A2: HLA Typing to 2<sup>nd</sup> or 3<sup>rd</sup> Field Resolution



## Purpose

Assess participants ability to correctly determine HLA type to 2<sup>nd</sup> or 3<sup>rd</sup> field.

## Satisfactory Performance

9 or more full HLA types in agreement with consensus/reference genotype in a distribution year.



## Consensus

At least 75% agreement on each allele. If consensus is not met, a reference result is used.

*10 blood samples over 3 distributions*

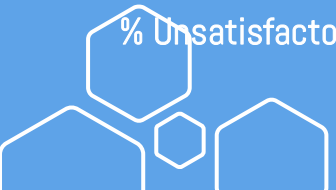


# Scheme 4A2: Performance

- 46/67 participants registered for 2<sup>nd</sup> field
- 22/67 participants registered for 3<sup>rd</sup> field
  
- 8 labs with unsatisfactory performance (1 UK&I)

7/8 labs with unsatisfactory performance completed CAPA

|                                                       | 2018       | 2019       | 2020       | 2021       | 2022       | 2023            | 2024            |
|-------------------------------------------------------|------------|------------|------------|------------|------------|-----------------|-----------------|
| Number of Participants (UK&I)                         | 63<br>(20) | 62<br>(20) | 64<br>(20) | 63<br>(22) | 61<br>(23) | 65<br>(23)      | 67<br>(23)      |
| Number with Unsatisfactory Performance (< 90%) (UK&I) | 9 (2)      | 9 (1)      | 7 (0)      | 6 (0)      | 4 (0)      | 9 (2)           | 8 (1)           |
| % Unsatisfactory Performance                          | 14.3%      | 14.5%      | 11.0%      | 11.1%      | 6.5%       | 13.8%<br>(8.7%) | 11.9%<br>(4.3%) |





# Scheme 4A2: Incorrect Assignments: 2<sup>nd</sup> Field

33/8846 (0.37%) incorrect HLA alleles reported by 12 labs (0 UK&I) – last year (0.55%)

- 14 reports of errors at the 2nd field

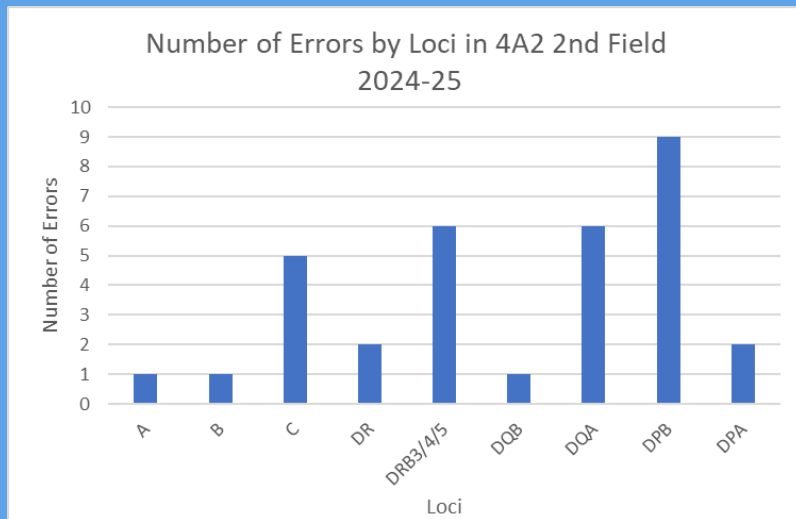
*e.g. DQA1\*03:02 rather than DQA1\*03:03*

- 3 samples with alleles in a string that should have been resolved
- 3 reports of the wrong HLA type
- 3 reports of incorrect nomenclature

*e.g. the use of P / G groups*

7 (58%) labs made 1 error  
5 (42%) labs made 3-4 errors

17 (74%) HLA types with a single error  
6 (26%) HLA types with multiple errors





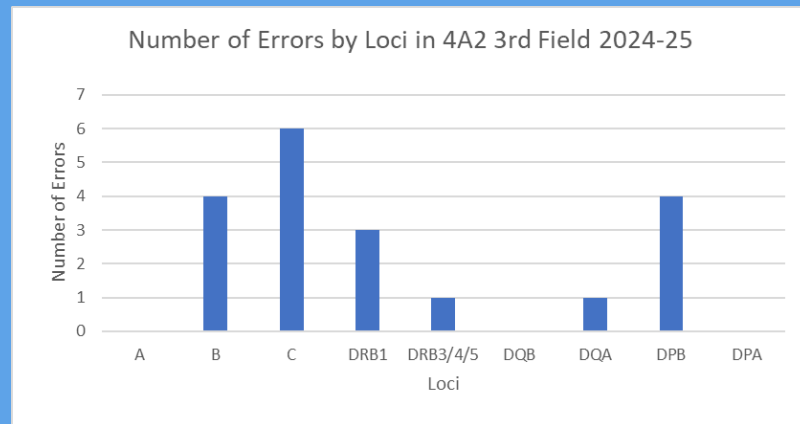
# Scheme 4A2: Incorrect Assignments: 3<sup>rd</sup> Field

19/3560 (0.53%) incorrect HLA alleles reported by 7 labs (1 UK&I) – last year (0.81%)

- 5 errors at 2<sup>nd</sup> field e.g. C\*03:03:01 rather than 03:321:XX
- 5 incorrect assignments e.g. C\*04:01:01 rather than C\*05:01:01
- 3 errors at 3<sup>rd</sup> field e.g. DPB1\*03:01:03 rather than DPB1\*03:01:01
- 1 report of unresolved ambiguities e.g. reporting G groups

4 (57%) labs made 1 error  
2 (29%) labs made 2 errors  
1 (14%) labs made 3+ errors

4 (29%) HLA types with multiple errors  
10 (71%) HLA types with a single error



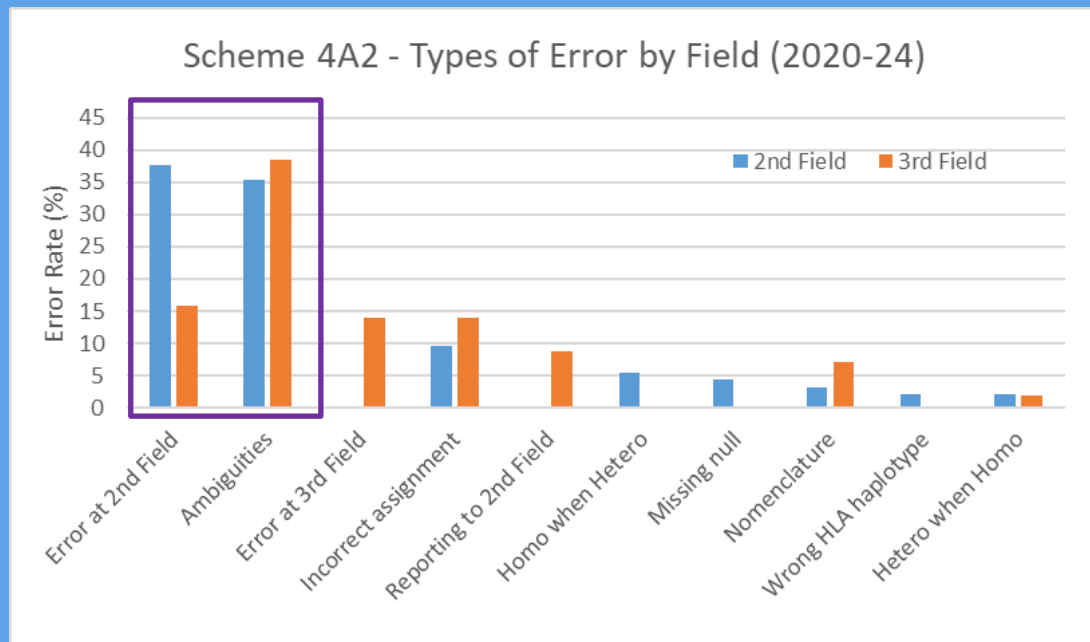
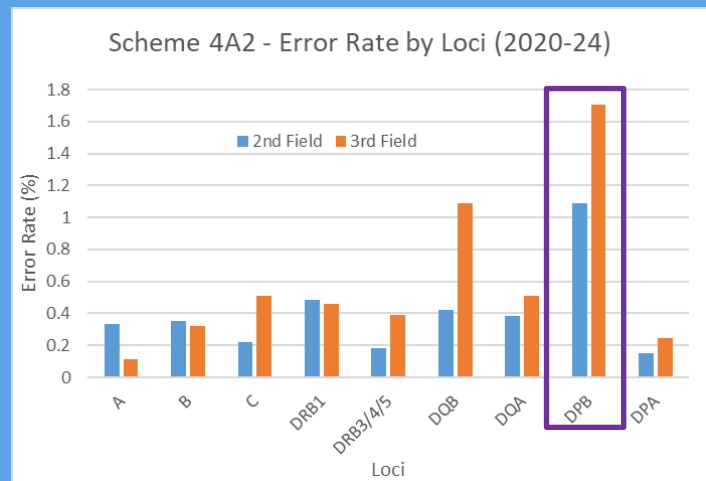
# Scheme 4A2: Unacceptable Performers 2024



| Lab  | Sample      | Error                                                                                                                                                                                                                                                                                                                                                               | Field           | CAPA Response                                      |
|------|-------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------|----------------------------------------------------|
| 284  | 01-03       | Reported G groups                                                                                                                                                                                                                                                                                                                                                   | 2 <sup>nd</sup> | EQA Specific Reporting issue                       |
| 29   | 04+05       | 3rd Field:<br>4A2 04/2024: Reported <b>B*40:01:01</b> homozygous, consensus B*40:01:02 homozygous<br>4A2 05/2024: Reported <b>DPB1*03:01:03</b> , consensus DPB1*03:01:01                                                                                                                                                                                           | 3 <sup>rd</sup> | Transcription error                                |
| 112  | 02+05+08+10 | 2nd Field:<br>4A2 02/2024: Reported <b>A*01:01, 01:03</b> , consensus A*01:01 homozygous<br>4A2 05/2024: Reported <b>C*03:03</b> , consensus C*03:321<br>4A2 08/2024: Reported <b>DQA1*03:02</b> , consensus DQA1*03:03<br>4A2 10/2024: Reported <b>DQA1*04:01</b> , consensus DQA1*04:02                                                                           | 2 <sup>nd</sup> | Kit resolution issue                               |
| 134  | 02+07+08+09 | 3rd Field:<br>4A2 02/2024: Reported <b>DPB1*02:01:01</b> , consensus DPB1*02:01:02<br>4A2 07/2024: Reported <b>C*04:01:01</b> , consensus C*05:01:01<br>4A2 08/2024: Reported <b>DPB1*01:03:01, 02:01:02</b> , consensus DPB1*01:01:01, 04:01:01<br>4A2 09/2024: Reported incomplete allele ( <b>C*07:02:0</b> ), consensus C*07:02:01, reported G groups for DRB1* | 3 <sup>rd</sup> | No reply                                           |
| 309  | 03+07+08+09 | 3rd Field:<br>4A2 03/2024: Reported <b>DRB3*01:01:02 homozygous</b> , consensus DRB3*01:01:02, 03:01:01<br>4A2 07/2024: Reported <b>B*15:01:01</b> , consensus B*35:01:01<br>4A2 08/2024: Reported <b>DRB1*01:01:01 and DQA1*03:01:01</b> , consensus DRB1*03:01:01 and DQA1*03:03:01<br>4A2 09/2024: Reported <b>B*07:01:01</b> , consensus B*07:02:01             | 3 <sup>rd</sup> | Transcription error                                |
| 1433 | 05+06+09    | 2nd Field:<br>4A2 05/2024: Reported <b>C*03:03</b> , consensus C*03:321<br>4A2 06/2024: Reported <b>B*42:02</b> , consensus B*44:02<br>4A2 09/2024: Reported <b>DRB1*11:04</b> , consensus DRB1*11:01                                                                                                                                                               | 2 <sup>nd</sup> | Transcription error /<br>Kit resolution issue      |
| 185  | 08-10       | 2nd Field:<br>4A2 08/2024: Reported Unacceptable ambiguities DPB1*677:01/875:01N/1086:01<br>4A2 09/2024: Reported Unacceptable ambiguities DPB1*677:01/875:01N/1086:01<br>4A2 10/2024: Reported Unacceptable ambiguities DPB1*727:01/1285:01N, 677:01/875:01N/1086:01                                                                                               | 2 <sup>nd</sup> | Unacceptable ambiguities /<br>Kit resolution issue |
| 223  | 07+09+10    | 2nd Field:<br>4A2 07/2024: Reported <b>DRB1*15:02</b> , consensus DRB1*15:01<br>4A2 09/2024: Reported <b>DQA1*04:01</b> , consensus DQA1*04:05<br>4A2 10/2024: Reported <b>DQA1*04:01</b> , consensus DQA1*04:02                                                                                                                                                    | 2 <sup>nd</sup> | Kit resolution issue                               |



# Scheme 4A2: Types of Errors Over 5 Years





Scheme

9

# KIR Genotyping



# Scheme 9: KIR Genotyping

## Purpose

Assess participants ability to correctly determine the presence or absence of specific KIR genes.

## Satisfactory Performance

9 or more full KIR genotypes in agreement with consensus/reference genotype in a distribution year.



## Consensus

At least 75% agreement on the presence/absence of each gene.  
Reference type used where consensus is not met

*10 blood samples over 2 distributions*





# Scheme 9: KIR Genotyping

- Participants able to report any of the following: *KIR2DL1, KIR2DL2, KIR2DL3, KIR2DL4, KIR2DL5, KIR3DL1, KIR3DL2, KIR3DL3, KIR3DS1, KIR2DS1, KIR2DS2, KIR2DS3, KIR2DS4, KIR2DS5, KIR2DP1, KIR3DP1.*
- Also able to report any other KIR polymorphisms they detected for information
- Participants can also report an 'A' or 'B' haplotype for each sample based on the gene content of the sample





# Scheme 9: Performance

- 1 lab with unsatisfactory performance

|                                                  | 2018  | 2019   | 2020   | 2021   | 2022   | 2023   | 2024   |
|--------------------------------------------------|-------|--------|--------|--------|--------|--------|--------|
| Number of Participants<br>(UK&I)                 | 9 (1) | 12 (1) | 12 (1) | 15 (1) | 15 (1) | 15 (1) | 13 (1) |
| Number with Unsatisfactory<br>Performance (UK&I) | 1 (0) | 3 (0)  | 0 (0)  | 1 (0)  | 0 (0)  | 1 (0)  | 1 (0)  |
| % Unsatisfactory<br>Performance                  | 11.1% | 25%    | 0%     | 6.7%   | 0%     | 6.7%   | 7.7%   |





## Scheme 9: Unacceptable Performers 2024

| Lab | Polymorphism        | Error                  | CAPA Response                             |
|-----|---------------------|------------------------|-------------------------------------------|
| 332 | 3DS1 & 2DS5<br>2DP1 | False Pos<br>False Neg | Interpretation issues<br>(staff training) |





Scheme

10

# HPA Genotyping



# Scheme 10: HPA Genotyping

## Purpose

Assess participants ability to correctly determine HPA polymorphisms.

## Satisfactory Performance

9 or more full HPA types in agreement with consensus/reference genotype in a distribution year.



## Consensus

At least 75% agreement on the presence/absence of each allele. Reference type used where consensus is not met

*10 blood samples over 2 distributions*





# Scheme 10: HPA Genotyping

- Participants able to report any of the following: *HPA-1, HPA-2, HPA-3, HPA-4, HPA-5, HPA-6, HPA-15*
  - 35/38 reported HPA-1, 2 , 3, 4, 5 and 15
  - 30/38 labs reported HPA-6
- Also able to report any other HPA polymorphisms detected, for information





# Scheme 10: HPA Genotyping

- 0 lab with unsatisfactory performance

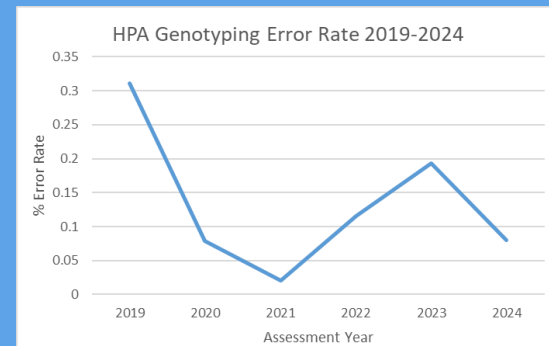
|                                                        | 2017   | 2018   | 2019   | 2020   | 2021   | 2022   | 2023   | 2024   |
|--------------------------------------------------------|--------|--------|--------|--------|--------|--------|--------|--------|
| Number of Participants (UK&I)                          | 15 (5) | 37 (6) | 38 (6) | 40 (0) | 38 (6) | 39 (6) | 39 (6) | 38 (6) |
| Number with Unsatisfactory Performance (< 100%) (UK&I) | 1 (0)  | 1 (0)  | 3 (0)  | 0 (0)  | 0 (0)  | 1 (0)  | 1 (0)  | 0 (0)  |
| % Unsatisfactory Performance                           | 6.7%   | 2.7%   | 7.9%   | 0%     | 0%     | 2.6%   | 2.6%   | 0%     |

# Scheme 10: Errors in HPA Genotypes 2024



- 3 labs made 1 error
- Error rate extremely low 0.08% but errors at some clinically relevant polymorphisms.
- Errors found at HPA-3b (n=2), HPA-1b (n=1), HPA-4b (n=1)

| Errors 2024  | HPA-1 a | HPA-1 b | HPA-2 a | HPA-2 b | HPA-3 a | HPA-3 b | HPA-4 a | HPA-4 b | HPA-5 a | HPA-5 b | HPA-6 a | HPA-6 b | HPA-15 a | HPA-15 b | Total |
|--------------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|----------|----------|-------|
| False Neg    | 0       | 1       | 0       | 0       | 0       | 0       | 0       | 1       | 0       | 0       | 0       | 0       | 0        | 0        | 2     |
| False Pos    | 0       | 0       | 0       | 0       | 0       | 2       | 0       | 0       | 0       | 0       | 0       | 0       | 0        | 0        | 2     |
| Total Error  | 0       | 1       | 0       | 0       | 0       | 2       | 0       | 1       | 0       | 0       | 0       | 0       | 0        | 0        | 4     |
| % Error      | 0.0     | 0.3     | 0.0     | 0.0     | 0.0     | 0.5     | 0.0     | 0.3     | 0.0     | 0.0     | 0.0     | 0.0     | 0.0      | 0.0      | 0.08  |
| Total Tested | 375     | 375     | 375     | 375     | 375     | 375     | 345     | 345     | 375     | 375     | 295     | 295     | 375      | 375      | 5030  |



- Even split of false positive (n=2) and false negative (n=2) errors.



Scheme



1B

HLA-B27 Testing



# Scheme 1B: HLA-B27 Testing

## Purpose

Assess participants ability to correctly determine HLA-B27/2708/B\*27 status.

## Satisfactory Performance

Making 10/10 reports that are in agreement with consensus in a distribution year.



## Consensus

At least 75% agreement on B27 status. Reference type used where consensus is not met

*10 blood samples sent over 5 distributions*





# Scheme 1B: Performance

- 16 labs with unsatisfactory performance (3 UK&I)

|                                                                      | 2018           | 2019           | 2020           | 2021         | 2022         | 2023         | 2024          |
|----------------------------------------------------------------------|----------------|----------------|----------------|--------------|--------------|--------------|---------------|
| Number of Participants<br>(UK&I)                                     | 133<br>(54)    | 133<br>(53)    | 141<br>(52)    | 141<br>(50)  | 139<br>(49)  | 134<br>(50)  | 131<br>(49)   |
| Number with<br>Unsatisfactory<br>Performance<br>( $< 100\%$ ) (UK&I) | 10<br>(3)      | 4<br>(1)       | 12<br>(2)      | 3<br>(0)     | 8<br>(0)     | 11<br>(3)    | 16<br>(3)     |
| % Unsatisfactory<br>Performance (UK&I)                               | 7.5%<br>(5.5%) | 3.0%<br>(1.9%) | 8.5%<br>(3.8%) | 2.1%<br>(0%) | 5.7%<br>(0%) | 8.2%<br>(6%) | 12.2%<br>(6%) |

- 5/10 samples distributed were HLA-B27 positive

# Scheme 1B: 2024 Incorrect Assignments



| Sample     | Result              | Lab Number                                | Technique                                                                            | HLA Type              | Lab Identified Cause                                                                                                              |
|------------|---------------------|-------------------------------------------|--------------------------------------------------------------------------------------|-----------------------|-----------------------------------------------------------------------------------------------------------------------------------|
| 1B 02      | False neg           | 1392<br>1435                              | Molecular<br>Serological                                                             | B27 B44               | No reply<br>No reply                                                                                                              |
| 1B 03      | False neg           | 40                                        | Serological                                                                          | B27 B40               | Procedural/processing errors                                                                                                      |
| 1B 05 & 06 | No results          | 1441                                      | Serological                                                                          | B27 B40<br>B27 B40    | No reply                                                                                                                          |
| 1B 06      | False neg           | 1435<br>31<br>225<br>357                  | Serological<br>Serological<br>Molecular<br>Serological                               | B27 B40               | No reply<br>Technical/testing issue<br>Kit/interpretation/reagent issue<br>No reply                                               |
| 1B 07 & 08 | False pos/false neg | 32                                        | Molecular                                                                            | B*27:08 B35<br>B7 B40 | Sample mix up                                                                                                                     |
| 1B 07      | False neg           | 1431<br>106<br>324<br>409<br>1308<br>1312 | Molecular<br>Serological<br>Serological<br>Serological<br>Serological<br>Serological | B*27:08 B35           | Reporting error<br>Interpretation/reagent issue<br>Interpretation/cut off values<br>Interpretation issues<br>No reply<br>No reply |
| 1B 09      | False pos           | 324                                       | Serological                                                                          | B7 B35                | Interpretation/cut off values                                                                                                     |
| 1B 09 & 10 | No results          | 1402                                      | Unknown                                                                              | B7 B35<br>B7 B15      | No reply                                                                                                                          |

- 88% false negative, 12% false positive
- 70.5% errors involved serological techniques
- Overall 73% use molecular methods, 27% use serological methods

7/11 labs with unsatisfactory performance completed CAPA



Scheme



5A

HFE Typing



# Scheme 5A: HFE Testing

## Purpose

Assess participants ability to correctly determine HFE mutations.

3 mutations assessed:

Codon 63: Histidine63Aspartic acid (H63D)

Codon 282: cysteine282tyrosine (C282Y)

Codon 65: Serine63Cysteine (S65C)

## Satisfactory Performance

10 reports in agreement with consensus/reference result in a distribution year.



## Consensus

At least 75% agreement on each HFE mutation. Reference type used where consensus is not met

*10 donor samples sent over 3 distributions*



# Scheme 5A: Performance



- 3 labs with unsatisfactory performance (1 UK&I)

|                                                        | 2018       | 2019           | 2020           | 2021           | 2022             | 2023           | 2024         |
|--------------------------------------------------------|------------|----------------|----------------|----------------|------------------|----------------|--------------|
| Number of Participants (UK&I)                          | 58<br>(44) | 51<br>(38)     | 49<br>(36)     | 45<br>(32)     | 37<br>(27)       | 38<br>(26)     | 37<br>(25)   |
| Number with Unsatisfactory Performance (< 100%) (UK&I) | 0<br>(0)   | 2<br>(1)       | 1<br>(1)       | 1<br>(1)       | 4<br>(3)         | 1<br>(1)       | 3<br>(1)     |
| % Unsatisfactory Performance                           | 0%         | 3.9%<br>(2.6%) | 2.0%<br>(2.8%) | 2.2%<br>(3.1%) | 10.8%<br>(11.1%) | 2.6%<br>(3.8%) | 8.1%<br>(4%) |

## CAPA responses (n=2/3)

- Sample mix up – 33.3%
- Transcription error – 33.3%
- No reply – 33.3%





Scheme



7

HLA-B\*57:01 Typing for Drug Hypersensitivity



# Scheme 7: HLA-B\*57:01 Typing for Drug Hypersensitivity



## Purpose

Assess participants ability to correctly determine HLA-B\*57:01 status

## Satisfactory Performance

Making 10 sample reports in agreement with the consensus/reference result in a distribution year.



## Consensus

At least 75% agreement on the status of HLA-B\*57:01. Reference result used when consensus not met.

*10 blood samples over 3 distributions*



# Scheme 7: Performance

- 1 lab with unacceptable performance

|                                                      | 2018       | 2019       | 2020       | 2021       | 2022       | 2023       | 2024       |
|------------------------------------------------------|------------|------------|------------|------------|------------|------------|------------|
| Number of Participants (UK&I)                        | 67<br>(27) | 67<br>(27) | 67<br>(27) | 64<br>(25) | 52<br>(18) | 50<br>(18) | 46<br>(18) |
| Number with Unacceptable Performance (< 100%) (UK&I) | 2<br>(0)   | 0<br>(0)   | 2<br>(0)   | 1<br>(1)   | 3<br>(0)   | 2<br>(0)   | 1<br>(0)   |
| % Unsatisfactory Performance                         | 3.0%       | 0.0%       | 3.1%       | 1.6%       | 5.8%       | 4.0%       | 2.2%       |

- 5/10 samples distributed were HLA-B\*57:01 positive



## Scheme 7: Unacceptable Performers 2024



| Lab | Sample | Error     | CAPA Response |
|-----|--------|-----------|---------------|
| 308 | 03     | False neg | No reply      |





## Scheme 7: **UPDATE**

~~HLA B\*57:01 Typing for Drug Hypersensitivity~~



### **HLA-related Pharmacogenetics**

#### **NEW ASSESSMENT OPTIONS**

(NO ADDITIONAL CHARGES)



*Abacavir Hypersensitivity - B\*57:01*

*Allopurinol Hypersensitivity - B\*58:01*

*Carbamazepine Hypersensitivity - A\*31:01, B\*15:02*

*Oxcarbazepine Hypersensitivity - B\*15:02*

*Lamotrigine Hypersensitivity - B\*15:02*

*Flucloxacillin Hypersensitivity - B\*57:01*

*Phenytoin Hypersensitivity - B\*15:02*

*Tebentafusp Suitability - A\*02:01*



All scheme 7 material will now be previously frozen whole blood



# Scheme



8

HLA Genotyping for Coeliac and other HLA Associated Disease



# Scheme 8: HLA Genotyping for Coeliac and other HLA Associated Disease.



## Purpose

Assess participants ability to correctly determine HLA type associated with various diseases e.g. coeliac disease, narcolepsy.

## Satisfactory Performance

Making 10 sample reports in agreement with the reference genotype in a distribution year.



## Assessment

Lab results reported in format identical to clinical report. Reference HLA result used for assesment.

*10 blood samples over 3 distributions*



# Scheme 8: Performance

- 10 Unsatisfactory Performers (1 UK&I)

|                                                                | 2018         | 2019         | 2020         | 2021         | 2022           | 2023           | 2024            |
|----------------------------------------------------------------|--------------|--------------|--------------|--------------|----------------|----------------|-----------------|
| Number of Participants (UK&I)                                  | 52<br>(10)   | 50<br>(11)   | 55<br>(12)   | 55<br>(10)   | 54<br>(11)     | 57<br>(11)     | 55<br>(11)      |
| Number with Unsatisfactory Performance<br>( $< 100\%$ ) (UK&I) | 14<br>(4)    | 13<br>(2)    | 17<br>(5)    | 12<br>(2)    | 25<br>(5)      | 18<br>(2)      | 10<br>(1)       |
| % Unsatisfactory Performance                                   | 27%<br>(40%) | 26%<br>(18%) | 31%<br>(42%) | 22%<br>(20%) | 46.3%<br>(45%) | 31.6%<br>(18%) | 18.2%<br>(9.1%) |

## CAPA responses (n=8/10)

- Transcription errors – 46%
- Kit interpretation error – 17%
- Reporting error – 7%
- Procedural error – 7%
- Unknown – 23%



# Scheme 8: Unacceptable Performance by Disease



| Disease                      | HLA Association   | Number of Participants | No. of Participants with Unacceptable Performance |
|------------------------------|-------------------|------------------------|---------------------------------------------------|
| Coeliac                      | DQ2.5, DQ8, DQ2.2 | 51                     | 9 (18%)                                           |
| Narcolepsy                   | DQB1*06:02        | 24                     | 0                                                 |
| Actinic Prurigo              | DRB1*04:07        | 5                      | 0                                                 |
| Birdshot Retinopathy         | A*29              | 14                     | 0                                                 |
| Behçet's                     | B*51              | 21                     | 0                                                 |
| Rheumatoid Arthritis         | DRB1*04           | 6                      | 0                                                 |
| Diabetes                     | DR3, DR4          | 7                      | 1 (14%)                                           |
| Psoriasis                    | C*06              | 6                      | 0                                                 |
| Allopurinol Hypersensitivity | B*58              | 8                      | 0                                                 |
| Carbamazepine                | A*31:01           | 9                      | 0                                                 |
| Phenytoin                    | B*15:02           | 3                      | 0                                                 |
| Tebentafusp                  | A*02:01           | 3                      | 0                                                 |

New for 2025-26



## Scheme 8: **UPDATE**

### HLA Genotyping for Coeliac and other HLA Associated Diseases



Coeliac Disease  
Narcolepsy  
Actinic Prurigo  
Birdshot Retinopathy  
Behcet's Disease  
Rheumatoid Arthritis  
Diabetes  
Psoriasis



Allopurinol hypersensitivity  
Carbamazepine hypersensitivity  
Phenytoin hypersensitivity  
Tebentafusp suitability



NOW PART OF **SCHEME 7**



# Scheme 8: Interpretative Comments

- Interpretation of the genotype in terms of predisposition to CD not currently assessed

## iv. DQ2.5 heterozygous (cis or trans)

### HLA genotype result:

HLA-DQA1\*05, DQB1\*02

HLA-DQA1\*X, DQB1\*X

OR

HLA-DQA1\*05, DQB1\*X

HLA-DQA1\*X, DQB1\*02

### HLA CD heterodimer result:

HLA-DQB1\*02, DQA1\*05 (DQ2.5) positive

HLA-DQB1\*02, DQA1\*02 (DQ2.2) negative

HLA-DQB1\*03:02 (DQ8) negative

**Genotype comment:** Positive for DQ2.5 (heterozygous)

**Interpretative comment < 1 >:** This individual has a genotype which is associated with coeliac disease

**Interpretative comment < 2 >:** This presence of DQA1\*05, DQB1\*02 (HLA-DQ2.5) has a strong association with coeliac disease in patients where laboratory tests or symptoms or endoscopic features suggest coeliac disease.



# Scheme 8: Assessment of Interpretative Comments

- Pilot assessment based on points:

| Coeliac Disease                                      | Outcome                                               | Improvement Point | Assessment   |
|------------------------------------------------------|-------------------------------------------------------|-------------------|--------------|
| HLA Genotype                                         | HLA genotype aligned to reference type                | N/A               | Acceptable   |
|                                                      | Result not reported                                   | N/A               | Unacceptable |
|                                                      | HLA genotype not aligned to reference type            | N/A               | Unacceptable |
| Interpretation (>1 improvement point = unacceptable) | *HLA Comments / Correct Nomenclature Used             | 0                 | Acceptable   |
|                                                      | *Incorrect HLA Comments / Incorrect Nomenclature Used | 1                 | Unacceptable |
|                                                      | Risk of CD Present/Absent Correctly Identified        | 0                 | Acceptable   |
|                                                      | Risk of CD Present/Absent Incorrectly Identified      | 1                 | Unacceptable |
|                                                      | *Stratification of Risk Identified                    | 0                 | Acceptable   |
|                                                      | *Stratification of Risk Incorrectly Identified        | 1                 | Unacceptable |
|                                                      | Diagnostic Disclaimer Applied Correctly               | 0                 | Acceptable   |
|                                                      | Diagnostic Disclaimer Not Applied or Incorrect        | 0.5               | Acceptable   |



# Scheme 8: Coeliac Disease – examples

DRB1\*07:01, 15:01; DQB1\*02:02, 06:02; DQA1\*01:02, 02:01  
(DQ2.2, DQ6.1)

| ID | Example of Interpretative Comment                                                                                                                                                                                          | Assessment |
|----|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------|
| 1  | Positive for DQ2.2. This individual carries HLA-DQA1*02, DQB1*02:02 (DQ2.2) that has a weak association with coeliac disease in patients where laboratory test or symptoms or endoscopic features suggest coeliac disease. | ✓ / ✗      |
| 2  | 90-95% of coeliac patients are HLA DQ2 and DQ8 positive.                                                                                                                                                                   | ✓ / ✗      |
| 3  | The presence of HLA-DQ2 is associated with, but not diagnostic for, coeliac disease. HLA-DQ2 is present in about 21% of Caucasians in the normal population.                                                               | ✓ / ✗      |
| 4  | Positive                                                                                                                                                                                                                   | ✓ / ✗      |

Patient genotype associated with coeliac disease.

Weak association with coeliac disease.

In patients where laboratory tests or symptoms or endoscopic features suggest coeliac disease.

The presence of an associated HLA genotype does not confer a diagnosis of coeliac disease and has a low positive predictive value for coeliac disease.



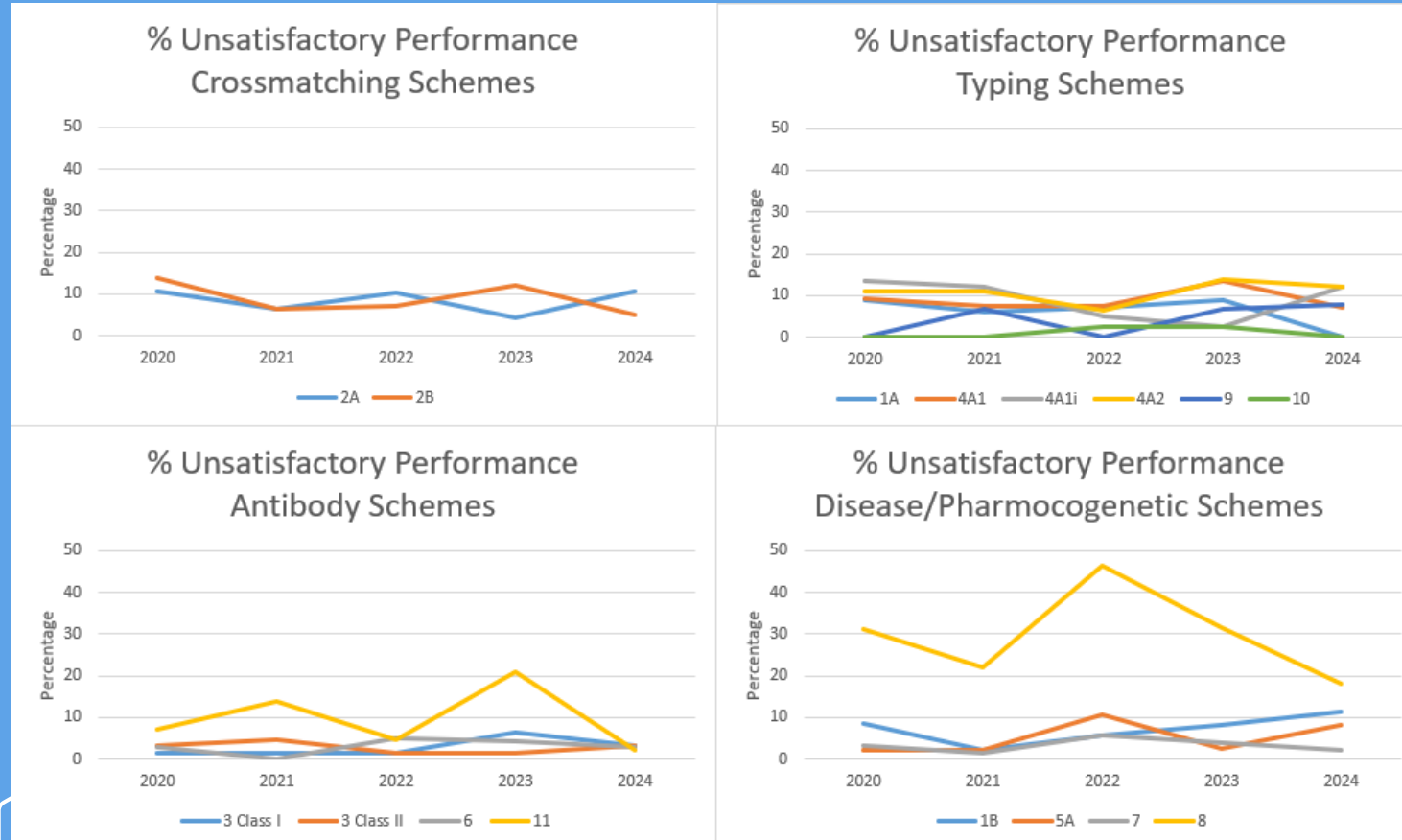


# Scheme Summary

Performance Summary for all Schemes



# 5 Year Trends in Unsatisfactory Performance



UK NEQAS for H&I  
Educational Crossmatch Scenario (EDXM)  
**Amy De'Ath**  
UK NEQAS for H&I Manager





"Schemes should relate more closely to clinical scenarios rather than testing individual test assays."



# Whole Process 'EQA'



## Assessed Schemes

- 1A, 4A1, 4A2 – HLA Typing
- 6 – HLA Antibody Detection
- 3 – HLA Antibody Specification
- 2A, 2B – Crossmatching



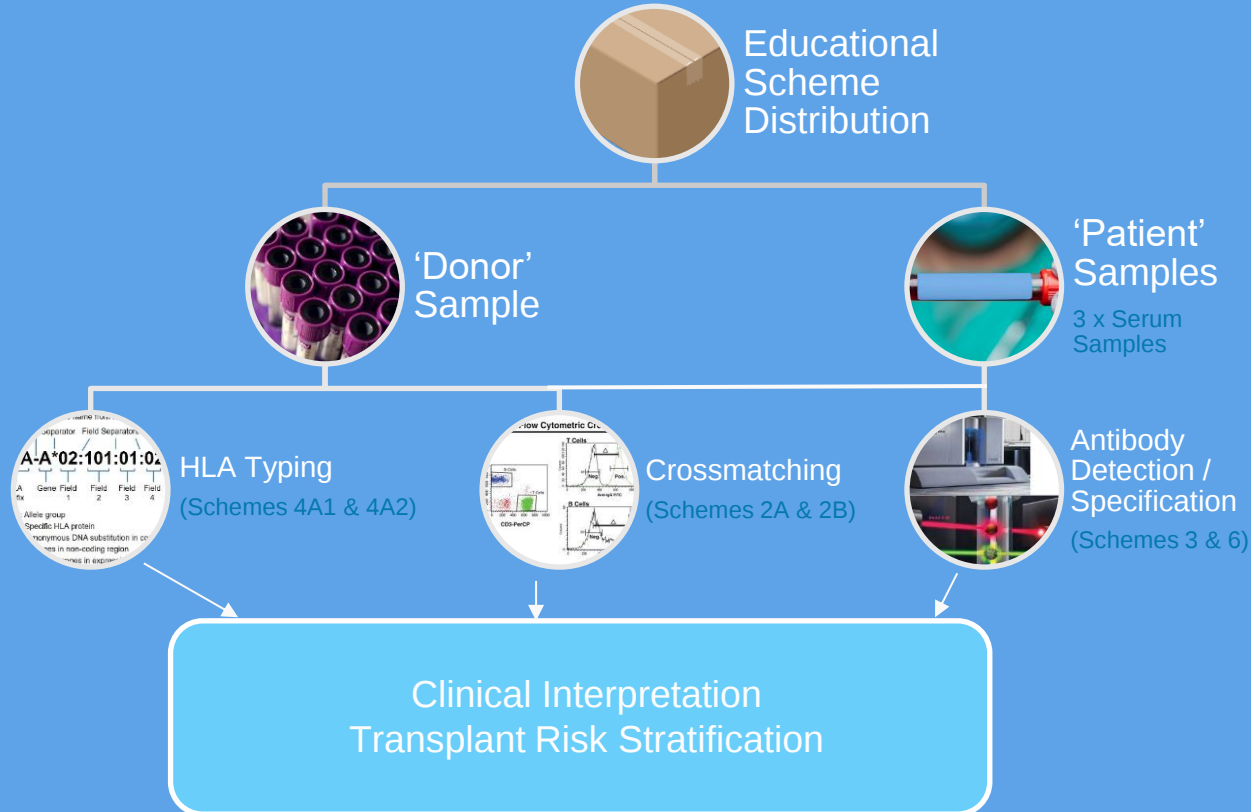
## Educational Schemes

- Interpretative Educational Scenarios
- Educational Crossmatch Scheme
  - Clinical decision making based on results from multiple assays
  - Each assay only gives part of the picture
  - Results from one assay can influence the interpretation of another
  - Variation between centres (repertoires, cut-offs)





# Educational Scheme Distribution





- 33 participants submitted results
- Not all labs reported results for all tests
- HLA genotype:

[illegible]



# Serum 1

Results





# Serum 1 Results

|                         | Result                                                                                           | % Consensus                                 | Comments                                                                                         |
|-------------------------|--------------------------------------------------------------------------------------------------|---------------------------------------------|--------------------------------------------------------------------------------------------------|
| HLA Class I Antibodies  | No Consensus                                                                                     | 58% (19/33)                                 |                                                                                                  |
| HLA Class II Antibodies | Negative                                                                                         | 76% (25/33)                                 |                                                                                                  |
| DSA                     | None                                                                                             | 100% (31/31)                                |                                                                                                  |
| CDC XM                  | PBL Negative<br>T cell Negative<br><i>B cell No Consensus</i>                                    | 100% (4/4)<br>100% (10/10)<br>71% (5/7 Neg) | CDC Negative – B cell XM with DTT 100% negative, without DTT 71% (5/7) negative<br>FCXM Negative |
| FCXM T Cell             | Negative                                                                                         | 96% (27/28)                                 |                                                                                                  |
| FCXM B Cell             | Negative                                                                                         | 96% (25/26)                                 |                                                                                                  |
| Transplant Risk         | Low/Standard<br>Intermediate                                                                     | 97% (29/30)<br>3% (1/30)                    |                                                                                                  |
| Immunological Advice    | Suitable for direct transplantation.<br>Low level HLA antibodies present but not donor specific. |                                             |                                                                                                  |
| Recommendations         | Proceed to transplant.                                                                           |                                             |                                                                                                  |



# Serum 2

Results



# Serum 2 Results

|                                | Result                                                                                                                                                                                             | % Consensus                            | Comments                                                          |
|--------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------|-------------------------------------------------------------------|
| <b>HLA Class I Antibodies</b>  | Positive                                                                                                                                                                                           | 100% (33/33)                           | Multiple A, B and Cw ab >10,000                                   |
| <b>HLA Class II Antibodies</b> | Negative                                                                                                                                                                                           | 85% (28/33)                            |                                                                   |
| <b>DSA</b>                     | Present                                                                                                                                                                                            | 100% (31/31)                           | 100% B antibody reported at 6-20,000<br>39% Cw antibody 600-1,750 |
| <b>CDC XM</b>                  | PBL Negative<br>T cell Negative<br>B cell Negative                                                                                                                                                 | 75% (3/4)<br>100% (10/10)<br>86% (6/7) | CDCXM Negative<br>FCXM Positive                                   |
| <b>FCXM T Cell</b>             | Positive                                                                                                                                                                                           | 89% (25/28)                            |                                                                   |
| <b>FCXM B Cell</b>             | Positive                                                                                                                                                                                           | 81% (21/26)                            |                                                                   |
| <b>Transplant Risk</b>         | Intermediate<br>High/Contraindication                                                                                                                                                              | 16% (5/31)<br>84% (26/31)              |                                                                   |
| <b>Immunological Advice</b>    | Not suitable for direct transplantation. High risk of AMR.<br>If transplant proceeds use enhanced immunosuppression and post-transplant monitoring.<br>Test for non-HLA and autologous antibodies. |                                        |                                                                   |
| <b>Recommendations</b>         | Seek alternative donor.<br>Consider de-sensitisation. Monitor antibodies over time to consider de-listing.<br>Discuss risk with patient.                                                           |                                        |                                                                   |



# Serum 3

Results





# Serum 3 Results

|                                | Result                                                                                                                                                                                                                   | % Consensus                             | Comments                                                |
|--------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------|---------------------------------------------------------|
| <b>HLA Class I Antibodies</b>  | Positive                                                                                                                                                                                                                 | 100% (33/33)                            | B and Cw antibodies 2,000 - >10,000 MFI                 |
| <b>HLA Class II Antibodies</b> | Positive                                                                                                                                                                                                                 | 100% (33/33)                            | DQ and DP antibodies 2,000 - >10,000 MFI                |
| <b>DSA</b>                     | Present                                                                                                                                                                                                                  | 100% (31/31)                            | Multiple DSA to CI and CII ranging from 600-33,000      |
| <b>CDC XM</b>                  | PBL No consensus<br>T cell Negative<br>B cell Positive                                                                                                                                                                   | 50% (2/4 Pos)<br>78% (7/9)<br>86% (6/7) | CDCXM T cell Negative, B cell Positive<br>FCXM Positive |
| <b>FCXM T Cell</b>             | Positive                                                                                                                                                                                                                 | 100% (28/28)                            |                                                         |
| <b>FCXM B Cell</b>             | Positive                                                                                                                                                                                                                 | 92% (24/26)                             |                                                         |
| <b>Transplant Risk</b>         | High/Contraindication                                                                                                                                                                                                    | 100% (31/31)                            |                                                         |
| <b>Immunological Advice</b>    | Not suitable for direct transplantation. Risk of AMR. Consider de-sensitisation.<br>If transplant proceeds use enhanced immunosuppression and post-transplant monitoring.<br>Test for non-HLA and autologous antibodies. |                                         |                                                         |
| <b>Recommendations</b>         | Seek alternative donor.<br>Consider de-sensitisation. Monitor antibodies over time to consider de-listing.<br>Discuss risk with patient.                                                                                 |                                         |                                                         |

# Summary of Crossmatch and DSA Detection Results



The table shows the percentage of participants identifying a DSA and the most common MFI range it was reported in.

| 2024 Results           |        | Serum 1                        |          | Serum 2                          |          | Serum 3                  |                              |
|------------------------|--------|--------------------------------|----------|----------------------------------|----------|--------------------------|------------------------------|
| DSA Defined by Luminex |        | Class I                        | Class II | Class I                          | Class II | Class I                  | Class II                     |
| MFI >10,000            |        | N/A                            | N/A      | B60 (100%)                       | N/A      | B51 (100%)<br>Cw10 (97%) | DR4 (100%)                   |
| MFI 5,001-9,999        |        | N/A                            | N/A      | N/A                              | N/A      | Cw15 (84%)               | DQ8 (97%)<br>DQ9 (97%)       |
| MFI 2,501-5,000        |        | N/A                            | N/A      | N/A                              | N/A      | N/A                      | DR9 (45%)<br>DQA1*03:02 (3%) |
| MFI <2,500             |        | N/A                            | N/A      | Cw10 (39%)<br>Cw3 (3%)           | N/A      | N/A                      | DR53 (10%)                   |
| CDCXM<br>B CELL        | No DTT | Negative                       |          | Negative                         |          | Positive                 |                              |
|                        | DTT    | Negative                       |          | Negative                         |          | Positive                 |                              |
| FCXM<br>B CELL         | T Cell | Negative                       |          | Positive                         |          | Positive                 |                              |
|                        | B Cell | Negative                       |          | Positive                         |          | Positive                 |                              |
| Risk                   |        | Low (97%)<br>Intermediate (3%) |          | High (84%)<br>Intermediate (16%) |          | High (100%)              |                              |





# Benefits



## Benchmarking

- Monitor performance of multiple techniques
- Make clinical interpretations on own results
- Compare local policies for clinical assessment



## Education

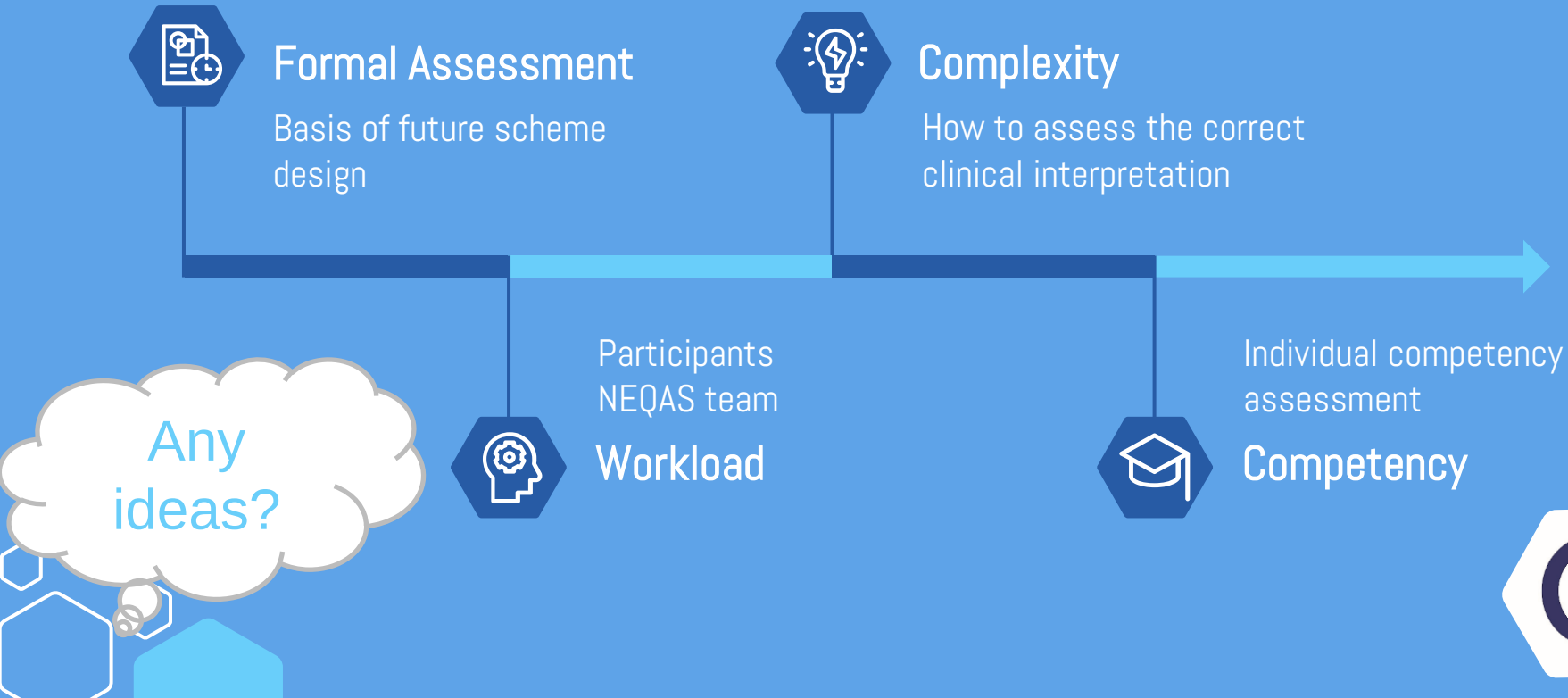
- Monitor concordances
- Review variations
- Staff training



## Competency

- Laboratory staff
- Clinical staff

# Future Considerations



The background is a solid blue color. In the corners, there are decorative geometric patterns consisting of white and light blue hexagons and lines. A large, dark blue hexagon with rounded corners is centered on the slide.

Do you have any  
questions?

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