

UK NEQAS FOR H&I SCHEME 9 – KIR GENOTYPING

9/01/2019

Result summary	2DL1	2DL2	2DL3	2DL4	2DL5	3DL1	3DL2	3DL3	3DS1	2DS1	2DS2	2DS3	2DS4	2DS5	2DP1	3DP1			
Total distributed	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13			
Total submitted	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13			
Positive	12	13	13	13	-	13	13	13	-	-	13	-	13	-	13	13			
Negative	1	-	-	-	13	-	-	-	13	13	-	13	-	13	-	-			
% Positive	92.3	100	100	100	-	100	100	100	-	-	100	-	100	-	100	100			
% Negative	7.7	-	-	-	-	100	-	-	100	100	-	100	-	100	-	-			
Consensus	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Positive	Negative	Positive	Positive	Positive	Positive			
Reference	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-			
Expected	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Positive	Negative	Positive	Positive	Positive	Positive			
Unacceptable	Negative	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-			
Number acceptable	12	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13			
Number unacceptable	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0			
Assessed <input checked="" type="checkbox"/>																			
Comments																			

Lab	2DL1	2DL2	2DL3	2DL4	2DL5	3DL1	3DL2	3DL3	3DS1	2DS1	2DS2	2DS3	2DS4	2DS5	2DP1	3DP1	Haplotype		Comments	Received	Tested	Assessment
																	Subtypes for 2DS4 and 3DP1 not ascertained.	A	B			
38	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Positive	Negative	Positive	Negative	Positive	Positive	No	No	24/07/2019	29/07/2019	Acceptable	
122	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Negative	Positive	Negative	Positive	Positive	Yes	Yes	30/07/2019	31/07/2019	Acceptable	
133	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Negative	Positive	Negative	Positive	Positive	Yes	Yes	25/07/2019	02/08/2019	Acceptable	
185	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Negative	Positive	Negative	Positive	Positive	Yes	Yes	24/07/2019	01/08/2019	Acceptable	
227	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Negative	Positive	Negative	Positive	Positive	Yes	Yes	24/07/2019	24/07/2019	Unacceptable	
267	Negative	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Negative	Positive	Negative	Positive	Positive	Yes	Yes	24/07/2019	24/07/2019	Unacceptable	
276	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Negative	Positive	Negative	Positive	Positive	Yes	Yes	24/07/2019	26/07/2019	Acceptable	
281	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Negative	Positive	Negative	Positive	Positive	Yes	Yes	24/07/2019	25/07/2019	Acceptable	
332	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Negative	Positive	Negative	Positive	Positive	No	No	27/07/2019	01/08/2019	Acceptable	
346	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Negative	Positive	Negative	Positive	Positive	Yes	Yes	29/07/2019	01/08/2019	Acceptable	
385	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Negative	Positive	Negative	Positive	Positive	Yes	Yes	24/07/2019	29/07/2019	Acceptable	
393	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Negative	Positive	Negative	Positive	Positive	Yes	Yes	24/07/2019	01/08/2019	Acceptable	
403	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Negative	Positive	Negative	Positive	Positive	Yes	Yes	28/08/2019	29/08/2019	Acceptable	

UK NEQAS FOR H&I SCHEME 9 – KIR GENOTYPING

9/02/2019

Result summary	2DL1	2DL2	2DL3	2DL4	2DL5	3DL1	3DL2	3DL3	3DS1	2DS1	2DS2	2DS3	2DS4	2DS5	2DP1	3DP1			
Total distributed	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13			
Total submitted	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13			
Positive	12	-	13	13	-	13	13	13	-	-	-	-	13	-	13	13			
Negative	1	13	-	-	13	-	-	-	13	13	13	13	-	13	-	-			
% Positive	92.3	-	100	100	-	100	100	100	-	-	-	-	100	-	100	100			
% Negative	7.7	100	-	-	100	-	-	-	100	100	100	100	-	100	-	-			
Consensus	Positive	Negative	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Negative	Positive	Negative	Positive	Positive			
Reference	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-			
Expected	Positive	Negative	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Negative	Positive	Negative	Positive	Positive			
Unacceptable	Negative	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-			
Number acceptable	12	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13			
Number unacceptable	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0			
Assessed <input checked="" type="checkbox"/>																			
Comments																			

Lab	2DL1	2DL2	2DL3	2DL4	2DL5	3DL1	3DL2	3DL3	3DS1	2DS1	2DS2	2DS3	2DS4	2DS5	2DP1	3DP1	2DS4 report	3DP1 report	Haplotype	Haplotype	Comments	Received	Tested	Assessment
																			A	B				
38	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Negative	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Positive	Subtypes for 2DS4 and	Yes	No	24/07/2019	01/08/2019	Acceptable	
122	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Negative	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Positive		No	No	24/07/2019	29/07/2019	Acceptable	
133	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Negative	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Positive	Deletion	Yes	No	30/07/2019	31/07/2019	Acceptable	
185	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Positive		Yes	No	25/07/2019	02/08/2019	Acceptable	
227	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Positive	Deletion	Yes	No	24/07/2019	01/08/2019	Acceptable	
267	Negative	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Negative	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Positive	Full Length	Yes	No	24/07/2019	24/07/2019	Unacceptable	
276	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Negative	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Positive		Yes	No	24/07/2019	26/07/2019	Acceptable	
281	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Positive		Yes	No	24/07/2019	25/07/2019	Acceptable	
332	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Positive		No	No	27/07/2019	01/08/2019	Acceptable	
346	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Positive		Yes	No	29/07/2019	01/08/2019	Acceptable	
385	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Negative	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Positive		Yes	No	24/07/2019	29/07/2019	Acceptable	
393	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Negative	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Positive	Deletion	Yes	No	24/07/2019	01/08/2019	Acceptable	
403	Positive	Negative	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Positive	Full Length	Yes	No	28/08/2019	29/08/2019	Acceptable	

UK NEQAS FOR H&I SCHEME 9 – KIR GENOTYPING

9 03/2019

Result summary	2DL1	2DL2	2DL3	2DL4	2DL5	3DL1	3DL2	3DL3	3DS1	2DS1	2DS2	2DS3	2DS4	2DS5	2DP1	3DP1			
Total distributed	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13			
Total submitted	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13			
Positive	12	13	13	13	13	13	13	13	13	13	13	-	13	13	13	13			
Negative	1	-	-	-	-	-	-	-	-	-	-	13	-	-	-	-			
% Positive	92.3	100	100	100	100	100	100	100	100	100	100	-	100	100	100	100			
% Negative	7.7	-	-	-	-	-	-	-	-	-	-	100	-	-	-	-			
Consensus	Positive	Negative	Positive	Positive	Positive	Positive													
Reference	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-			
Expected	Positive	Negative	Positive	Positive	Positive	Positive													
Unacceptable	Negative	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-			
Number acceptable	12	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13			
Number unacceptable	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0			
Assessed <input checked="" type="checkbox"/>																			
Comments																			

Lab	2DL1	2DL2	2DL3	2DL4	2DL5	3DL1	3DL2	3DL3	3DS1	2DS1	2DS2	2DS3	2DS4	2DS5	2DP1	3DP1	Haplotype		Comments	Received	Tested	Assessment	
																	A	B					
38	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Subtypes for 2DS4 and 3DP1 not ascertained.		No	Yes	24/07/2019	08/01/2019	Acceptable
122	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Full Length		No	No	24/07/2019	29/07/2019	Acceptable
133	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Full Length		Yes	Yes	30/07/2019	31/08/2019	Acceptable
185	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Full Length		Yes	Yes	25/07/2019	02/08/2019	Acceptable
227	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Full Length		Yes	Yes	24/07/2019	01/08/2019	Acceptable
267	Negative	Positive	Negative	Positive	Positive	Positive	Positive	Deletion		Yes	Yes	24/07/2019	24/07/2019	Unacceptable									
276	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Haplotype A/B or B/B		Yes	Yes	24/07/2019	26/07/2019	Acceptable
281	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Haplotype A/B or B/B		Yes	Yes	24/07/2019	25/07/2019	Acceptable
332	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Haplotype A/B or B/B		No	No	27/07/2019	01/08/2019	Acceptable
346	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Haplotype A/B or B/B		Yes	Yes	29/07/2019	01/08/2019	Acceptable
385	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Haplotype A/B or B/B		Yes	Yes	24/07/2019	29/07/2019	Acceptable
393	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Haplotype A/B or B/B		Yes	Yes	24/07/2019	01/08/2019	Acceptable
403	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Haplotype A/B or B/B		Yes	Yes	28/08/2019	29/08/2019	Acceptable

UK NEQAS FOR H&I SCHEME 9 – KIR GENOTYPING

9/04/2019

Result summary	2DL1	2DL2	2DL3	2DL4	2DL5	3DL1	3DL2	3DL3	3DS1	2DS1	2DS2	2DS3	2DS4	2DS5	2DP1	3DP1			
Total distributed	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13			
Total submitted	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13			
Positive	13	13	13	13	13	13	13	13	-	-	13	13	13	-	13	13			
Negative	-	-	-	-	-	-	-	-	13	13	-	-	13	-	-	-			
% Positive	100	100	100	100	100	100	100	100	-	-	100	100	100	-	100	100			
% Negative	-	-	-	-	-	-	-	-	100	100	-	-	100	-	-	-			
Consensus	Positive	Negative	Negative	Positive	Positive	Positive	Negative	Positive	Positive										
Reference	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-			
Expected	Positive	Negative	Negative	Positive	Positive	Positive	Negative	Positive	Positive										
Unacceptable	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-			
Number acceptable	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13			
Number unacceptable	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0			
Assessed <input checked="" type="checkbox"/>																			
Comments																			

Lab	2DL1	2DL2	2DL3	2DL4	2DL5	3DL1	3DL2	3DL3	3DS1	2DS1	2DS2	2DS3	2DS4	2DS5	2DP1	3DP1	Haplotype		Comments	Received	Tested	Assessment		
																	2DS4 report	3DP1 report						
38	Positive	Negative	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Subtypes for 2DS4 and 3DP1 not ascertained.		No	Yes		24/07/2019	01/08/2019	Acceptable							
122	Positive	Negative	Negative	Positive	Positive	Positive	Positive	Positive	Positive	Deletion		No	No		24/07/2019	29/07/2019	Acceptable							
133	Positive	Negative	Positive	Positive	Positive	Positive	Positive	Positive	Deletion		Yes	Yes		30/07/2019	31/07/2019	Acceptable								
185	Positive	Negative	Positive	Positive	Positive	Positive	Positive	Positive	Full Length		Yes	Yes		25/07/2019	02/08/2019	Acceptable								
227	Positive	Negative	Positive	Positive	Positive	Positive	Positive	Positive	Deletion		Yes	Yes		24/07/2019	01/08/2019	Acceptable								
267	Positive	Negative	Positive	Positive	Positive	Positive	Positive	Positive	Full Length		Yes	Yes		24/07/2019	24/07/2019	Acceptable								
276	Positive	Negative	Positive	Positive	Positive	Positive	Positive	Positive	Full Length		Yes	Yes		24/07/2019	26/07/2019	Acceptable								
281	Positive	Negative	Positive	Positive	Positive	Positive	Positive	Positive	Full Length		Yes	Yes		24/07/2019	25/07/2019	Acceptable								
332	Positive	Negative	Positive	Positive	Positive	Positive	Positive	Positive	Full Length		No	No		27/07/2019	01/08/2019	Acceptable								
346	Positive	Negative	Positive	Positive	Positive	Positive	Positive	Positive	Full Length		Yes	Yes		29/07/2019	01/08/2019	Acceptable								
385	Positive	Negative	Positive	Positive	Positive	Positive	Positive	Positive	Full Length		Yes	Yes		24/07/2019	29/07/2019	Acceptable								
393	Positive	Negative	Positive	Positive	Positive	Positive	Positive	Positive	Full Length		Yes	Yes		24/07/2019	01/08/2019	Acceptable								
403	Positive	Negative	Positive	Positive	Positive	Positive	Positive	Positive	Full Length		Yes	Yes		28/08/2019	29/08/2019	Acceptable								

UK NEQAS FOR H&I SCHEME 9 – KIR GENOTYPING

9 05/2019																								
Result summary	2DL1	2DL2	2DL3	2DL4	2DL5	3DL1	3DL2	3DL3	3DS1	2DS1	2DS2	2DS3	2DS4	2DS5	2DP1	3DP1								
Total distributed	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13								
Total submitted	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13								
Positive	13	-	13	13	-	13	13	13	-	-	-	-	-	13	-	13								
Negative	-	13	-	-	13	-	-	-	13	13	13	13	-	13	-	13								
% Positive	100	-	100	100	-	100	100	100	-	-	-	-	100	-	100	100								
% Negative	-	100	-	-	100	-	-	-	100	100	100	100	-	100	-	100								
Consensus	Positive	Negative	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive								
Reference	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-								
Expected	Positive	Negative	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Negative	Positive	Negative	Positive	Positive								
Unacceptable	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-								
Number acceptable	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13								
Number unacceptable	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0								
Assessed <input checked="" type="checkbox"/>																								
Comments																								
Lab	2DL1	2DL2	2DL3	2DL4	2DL5	3DL1	3DL2	3DL3	3DS1	2DS1	2DS2	2DS3	2DS4	2DS5	2DP1	3DP1	2DS4 report	3DP1 report	Haplotype A	Haplotype B	Comments	Received	Tested	Assessment
38	Positive	Negative	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Subtypes for 2DS4 and 3DP1 not ascertained.		Yes	No	24/07/2019	01/08/2019	Acceptable	
122	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Full Length		No	No	24/07/2019	29/07/2019	Acceptable	
133	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Positive	Positive	Positive	Full Length		Yes	No	30/07/2019	31/07/2019	Acceptable	
185	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Full Length		Yes	No	25/07/2019	02/08/2019	Acceptable	
227	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Full Length		Yes	No	24/07/2019	01/08/2019	Acceptable	
267	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Full Length		Yes	No	24/07/2019	24/07/2019	Acceptable	
276	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Full Length		Yes	No	24/07/2019	26/07/2019	Acceptable	
281	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Full Length		Yes	No	24/07/2019	25/07/2019	Acceptable	
332	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Full Length		No	No	27/07/2019	01/08/2019	Acceptable	
346	Positive	Negative	Positive	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Full Length		Yes	No	29/07/2019	01/08/2019	Acceptable	
385	Positive	Negative	Positive	Positive	Negative	Positive	Positive	Positive	Positive	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Full Length		Yes	No	24/07/2019	29/07/2019	Acceptable	
393	Positive	Negative	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Full Length		Yes	No	KIR 2DS4 gene is present as full length as well as del 22bp form			
403	Positive	Negative	Positive	Positive	Negative	Positive	Positive	Positive	Negative	Negative	Negative	Negative	Positive	Negative	Positive	Positive	Full Length		Yes	No	24/07/2019	01/08/2019	Acceptable	
																					28/08/2019	29/08/2019	Acceptable	