

Lot No.: **09G**

Lot-specific information

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Olerup SSP[®] DQB1 high resolution for frequent alleles

Product number: 101.221-12 – including *Taq* polymerase
Lot number: 09G
Expiry date: 2011-August-01
Number of tests: 12 tests
Number of wells per test: 47 + 1
Storage - pre-aliquoted primers: dark at -20°C
- PCR Master Mix: -20°C
- Adhesive PCR seals RT
- Product Insert RT

This Product Description is only valid for Lot No. 09G.

CHANGES COMPARED TO THE PREVIOUS *OLERUP SSP[®]* DQB1 HIGH RESOLUTION LOT (12F)

The DQB1 high resolution specificity and interpretation tables have been updated with the DQB1 alleles described since the previous *Olerup SSP[®]* DQB1 high resolution lot was made (**Lot No. 12F**).

The primers of the wells detailed below have been exchanged, added or modified compared to the previous lot.

Well	5'-primer	3'-primer	rationale
4	Moved	Moved	Primer pair moved to well 43.
18	Added	Added	Primer pair added for the DQB1*0324 allele.
35	Added	-	Primer added for the DQB1*0635 allele.
38	-	Added	Primer added for the DQB1*061402 allele.
43	Added	Added	Primer pair from well 4.

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Well **48** contains Negative Control primer pairs, that will amplify more than 95% of the *Olerup* SSP[®] HLA Class I, DRB, DQB1 and DPB1 amplicons as well as the amplicons generated by control primer pairs.

PCR product sizes range from 75 to 430 base pairs.
The PCR product generated by the control primer pair is 430 base pairs.

Length of PCR product	105	200	105	80	75	80
5'-primer¹	164	340	440	45	45	43
	5'-CAC ^{3'}	5'-Agg ^{3'}	5'-TTA ^{3'}	5'-Tg g ^{3'}	5'-Tg g ^{3'}	5'-Tg g ^{3'}
3'-primer²	231	2nd I	507	59	58	57
	5'-TgC ^{3'}	5'-AAA ^{3'}	5'-TTg ^{3'}	5'-CTC ^{3'}	5'-ggC ^{3'}	5'-CTC ^{3'}
A*	+	+	+			
B*	+	+	+			
Cw*	+	+	+			
DRB1				+	+	
DRB3				+	+	
DRB5				+		
DQB1					+	
DPB1						+

¹The nucleotide position for HLA class I genes and the codon for HLA class II genes, in the 2nd or 3rd exon, matching the specificity-determining 3'-end of the primer is given. Nucleotide and codon numbering as on the www.ebi.ac.uk/imgt/hla web site. The sequence of the 3 terminal nucleotides of the primer is given.

²The nucleotide position for HLA class I genes and the codon for HLA class II genes, in the 2nd or 3rd exon or the 2nd intron, matching the specificity-determining 3'-end of the primer is given in the anti-sense direction. Nucleotide and codon numbering as on the www.ebi.ac.uk/imgt/hla web site. The sequence of the 3 terminal nucleotides of the primer is given.

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PRODUCT DESCRIPTION

DQB1 SSP typing

CONTENT

The primer set contains 5'- and 3'-primers for identifying the DQB1*0201 to DQB1*0635 alleles.

Please note that DQB1 amplifications usually are somewhat less pronounced than e.g. DRB and DQA1 amplifications even when using the same DNA preparation and exactly the same experimental procedures.

PLATE LAYOUT

Each test consists of 48 PCR reactions in a 48 well cut PCR plate.

1	2	3	4	5	6	7	8
9	10	11	12	13	14	15	16
17	18	19	20	21	22	23	24
25	26	27	28	29	30	31	32
33	34	35	36	37	38	39	40
41	42	43	44	45	46	47	48

Wells 1 to 47 – DQB1 primers.

Well 48 – Negative Control.

The 48 well cut PCR plate is marked with ‘DQB1 “HIGH”’ in silver/gray ink.

Well No. 1 is marked with the Lot No. ‘09G’.

The PCR plates are covered with a PCR-compatible foil.

Please note: When removing each 48 well PCR plate, make sure that the remaining plates stay covered. Use a scalpel or a similar instrument to carefully cut the foil between the plates.

INTERPRETATION

Only DQB1 alleles will be amplified by the DQB1 high resolution for frequent alleles typing kit. Thus, the interpretation of DQB1 high resolution for frequent alleles typings is not influenced by the DQB2 or DQB3 genes or other HLA class II genes.

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UNIQUELY IDENTIFIED ALLELES

All the phenotypically different DQB1 alleles, i.e. DQB1*0201 to 0205, DQB1*0301 to 0325, DQB1*0401 to 0403, DQB1*0501 to 0505 and DQB1*0601 to 0635, recognized by the HLA Nomenclature Committee in July 2009¹ will give rise to unique amplification patterns by the primers in the DQB1 high resolution for frequent alleles typing kit.

The DQB1 typing kit cannot distinguish the DQB1*020101 and 020102 alleles, the DQB1*030101, 030102 and 030104 alleles, the DQB1*030201 to 030204 alleles, the DQB1*030302 and 030303 alleles, the DQB1*030501 to 030504 alleles, the 040101 and 040102, the DQB1*050101 and 050102 alleles, the DQB1*050301 and 050302 alleles, the DQB1*060101 to 060104 alleles and the DQB1*060401 and 060403 alleles.

¹DQB1 alleles listed on the IMGT/HLA web page 2009-July-17, release 2.26.0, www.ebi.ac.uk/imgt/hla.

RESOLUTION IN HOMO- AND HETEROZYGOTES

The 73 DQB1 alleles generate 81 amplification patterns, the DQB1*030101/030102/030104 and 030103, the DQB1*050201 and 050202, the DQB1*060201 and 060202, the DQB1*060301 and 060302, the DQB1*060401/060403 and 060402, the DQB1*060501 and 060502, the DQB1*060801 and 060802, the DQB1*061101 and 061102 and the DQB1*061401 and 061402 alleles generate different amplification patterns. These patterns can be combined in 3321 homozygous and heterozygous combinations. 438 of these genotypes do not give rise to unique amplification patterns. (The different lengths of the specific PCR products in lanes 5, 9, 11, 13, 14, 17, 18, 35, 38, 40, 41, 42, 43 and 46 were not considered in these calculations, nor the differences in product length between the DQB1*317 and DQB1*0320 alleles.)

```
+++----- +----- +----- +----- +----- +----- 0201,0505 = 0204,050201 = 0204,0505
+++----- +----- +----- +----- +----- +----- 0201,0204 = 0204,0204
+++----- +----- +----- +----- +----- +----- 0201,0633 = 0205,060201 = 0205,0633
+++----- +----- +----- +----- +----- +----- 0201,0626N = 0205,0626N
+++----- +----- +----- +----- +----- +----- 0201,0205 = 0205,0205
++++----- +----- +----- +----- +----- +----- 0203,0301 = 0203,030103
---+----- +----- +----- +----- +----- +----- 050202,0505 = 0503,0505
---+----- +----- +----- +----- +----- +----- 050201,0505 = 0505,0505
---+----- +----- +----- +----- +----- +----- 0301,0308 = 0308,0319
---+----- +----- +----- +----- +----- +----- 0307,0309 = 0311,0316
---+----- +----- +----- +----- +----- +----- 0307,0324 = 0316,0318
---+----- +----- +----- +----- +----- +----- 0301,0307 = 0302,0316 = 0307,0316 =
0307,0319
---+----- +----- +----- +----- +----- +----- 0309,0318 = 0311,0324
---+----- +----- +----- +----- +----- +----- 0301,0311 = 0302,0309 = 0309,0311 =
0311,0319
---+----- +----- +----- +----- +----- +----- 0301,0318 = 0302,0324 = 0318,0319 =
0318,0324
---+----- +----- +----- +----- +----- +----- 0304,0315 = 0304,0316
---+----- +----- +----- +----- +----- +----- 0301,0314 = 0304,0312 = 0314,0319
---+----- +----- +----- +----- +----- +----- 030103,0314 = 0304,0314
---+----- +----- +----- +----- +----- +----- 0303,0304 = 0304,0319
---+----- +----- +----- +----- +----- +----- 0305,0316 = 0316,0317
---+----- +----- +----- +----- +----- +----- 0305,0309 = 0309,0317
---+----- +----- +----- +----- +----- +----- 0305,0313 = 0313,0317
---+----- +----- +----- +----- +----- +----- 0305,0324 = 0317,0324
---+----- +----- +----- +----- +----- +----- 0305,0321 = 0317,0321
---+----- +----- +----- +----- +----- +----- 0305,0322 = 0317,0322
---+----- +----- +----- +----- +----- +----- 0301,0305 = 0301,0317 = 030103,0317
= 0317,0319
---+----- +----- +----- +----- +----- +----- 0306,0309 = 0309,0325
---+----- +----- +----- +----- +----- +----- 0306,0313 = 0313,0325
---+----- +----- +----- +----- +----- +----- 0306,0324 = 0324,0325
```



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---+---+ +---+--- -+---+ -+---+ -+---+ 0306,0321 = 0321,0325
---+---+ +---+--- -+---+ -+---+ +---+--- 0306,0322 = 0322,0325
---+---+ +---+--- -+---+ -+---+ -+---+ 0301,0306 = 0301,0325 = 030103,0306
= 0306,0319
---+---+ +---+--- -+---+ -+---+ -+---+ 0309,0315 = 0309,0316
---+---+ +---+--- -+---+ -+---+ -+---+ 0310,0316 = 0312,0316
---+---+ +---+--- +---+--- -+---+ -+---+ 0313,0315 = 0313,0316
---+---+ +---+--- -+---+ -+---+ -+---+ 0315,0324 = 0316,0324
---+---+ +---+--- -+---+ -+---+ -+---+ 0315,0321 = 0316,0321
---+---+ +---+--- -+---+ -+---+ +---+--- 0315,0322 = 0316,0322
---+---+ +---+--- -+---+ -+---+ -+---+ 0301,0315 = 0301,0316 = 030103,0315
= 030103,0316 = 0303,0316 =
0315,0316 = 0315,0319 = 0316,0319
---+---+ +---+--- -+---+ -+---+ -+---+ 0303,0309 = 0309,0319
---+---+ +---+--- +---+--- -+---+ -+---+ 0310,0313 = 0312,0313
---+---+ +---+--- -+---+ -+---+ -+---+ 0301,0312 = 030103,0312 = 0310,0319
= 0312,0319
---+---+ +---+--- +---+--- -+---+ +---+--- -+---+ 0313,0604 = 0313,0634
---+---+ +---+--- +---+--- -+---+ -+---+ -+---+ 0301,0313 = 030103,0313 = 0303,0313
= 0313,0313 = 0313,0319
---+---+ +---+--- -+---+ -+---+ -+---+ 0303,0324 = 0319,0324
---+---+ +---+--- -+---+ -+---+ -+---+ 0301,0323 = 030103,0323 = 0319,0323
---+---+ +---+--- -+---+ -+---+ -+---+ 0303,0321 = 0319,0321
---+---+ +---+--- -+---+ -+---+ +---+--- 0303,0322 = 0319,0322
---+---+ +---+--- -+---+ -+---+ -+---+ 0301,0303 = 0301,0319 = 030103,0303
= 030103,0319
---+---+ +---+--- -+---+ -+---+ -+---+ 0303,0319 = 0319,0319
---+---+ +---+--- -+---+ -+---+ -+---+ 030103,0304 = 0304,0304
---+---+ +---+--- -+---+ -+---+ -+---+ 0301,0309 = 030103,0309 = 0309,0309
---+---+ +---+--- -+---+ -+---+ -+---+ 0301,0310 = 030103,0310
---+---+ +---+--- -+---+ -+---+ -+---+ 0301,0628 = 0324,060301 = 0324,0628
---+---+ +---+--- -+---+ -+---+ -+---+ 0301,0324 = 030103,0324
---+---+ +---+--- -+---+ -+---+ +---+--- 0301,0624 = 0322,060201 = 0322,0624
---+---+ +---+--- -+---+ -+---+ -+---+ 0301,0623 = 0321,0623
---+---+ +---+--- -+---+ -+---+ +---+--- 0301,0630 = 0322,0630
---+---+ +---+--- -+---+ -+---+ -+---+ 0301,0617 = 0322,0617
---+---+ +---+--- -+---+ -+---+ -+---+ 0301,0321 = 030103,0321 = 0321,0321
---+---+ +---+--- -+---+ -+---+ +---+--- 0301,0322 = 030103,0322 = 0322,0322
---+---+ +---+--- -+---+ -+---+ -+---+ 0301,0301 = 0301,030103
---+---+ +---+--- -+---+ -+---+ -+---+ 0303,0314 = 0310,0314 = 0312,0314
---+---+ +---+--- -+---+ -+---+ -+---+ 0305,0310 = 0310,0317
---+---+ +---+--- -+---+ -+---+ -+---+ 0306,0310 = 0310,0325
---+---+ +---+--- -+---+ -+---+ -+---+ 0303,0310 = 0310,0312
---+---+ +---+--- -+---+ -+---+ -+---+ 0308,0628 = 0318,060302
---+---+ +---+--- -+---+ -+---+ -+---+ 0302,0306 = 0306,0308
---+---+ +---+--- -+---+ -+---+ -+---+ 0308,060201 = 0308,060202
---+---+ +---+--- -+---+ -+---+ -+---+ 0308,060301 = 0308,060302
---+---+ +---+--- -+---+ -+---+ -+---+ 0302,0308 = 0308,0308
---+---+ +---+--- -+---+ -+---+ -+---+ 0302,0315 = 0303,0307 = 0307,0315
---+---+ +---+--- -+---+ -+---+ -+---+ 0302,0307 = 0307,0307
---+---+ +---+--- -+---+ -+---+ -+---+ 0302,0311 = 0311,0311
---+---+ +---+--- -+---+ -+---+ -+---+ 0318,060301 = 0318,0628
---+---+ +---+--- -+---+ -+---+ -+---+ 0302,0318 = 0318,0318
---+---+ +---+--- -+---+ -+---+ -+---+ 0305,0306 = 0306,0317 = 0317,0325
---+---+ +---+--- -+---+ -+---+ -+---+ 0305,0315 = 0315,0317
---+---+ +---+--- -+---+ -+---+ -+---+ 0305,0312 = 0312,0317
---+---+ +---+--- -+---+ -+---+ -+---+ 0305,0323 = 0317,0323
---+---+ +---+--- -+---+ -+---+ -+---+ 0303,0317 = 0305,0317 = 0317,0317
---+---+ +---+--- -+---+ -+---+ -+---+ 0306,0315 = 0315,0325
---+---+ +---+--- -+---+ -+---+ -+---+ 0306,0312 = 0312,0325
---+---+ +---+--- -+---+ -+---+ -+---+ 0306,0402 = 0306,0403
---+---+ +---+--- -+---+ -+---+ -+---+ 0306,060201 = 0306,060202
---+---+ +---+--- -+---+ -+---+ -+---+ 0306,060301 = 0306,060302
---+---+ +---+--- -+---+ -+---+ -+---+ 0303,0306 = 0306,0306 = 0306,0325
---+---+ +---+--- -+---+ -+---+ -+---+ 0325,0402 = 0325,0403
---+---+ +---+--- -+---+ -+---+ -+---+ 0325,060201 = 0325,060202
---+---+ +---+--- -+---+ -+---+ -+---+ 0325,060301 = 0325,060302
---+---+ +---+--- -+---+ -+---+ -+---+ 0303,0315 = 0315,0315
---+---+ +---+--- -+---+ -+---+ -+---+ 0303,0312 = 0312,0312
---+---+ +---+--- -+---+ -+---+ -+---+ 0323,060801 = 0323,060802
---+---+ +---+--- -+---+ -+---+ -+---+ 0323,0604 = 0323,060402
---+---+ +---+--- -+---+ -+---+ -+---+ 0402,0610 = 0403,0610
---+---+ +---+--- -+---+ -+---+ -+---+ 0402,0625 = 0403,0625
---+---+ +---+--- -+---+ -+---+ -+---+ 0402,0403 = 0403,0403



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---+----- ---+----- ----- ---++++- ---+----- ---+---+ 060202,0631 = 060302,0620
---+----- ---+----- ----- ---++++- ---+----- ---+---+ 060201,060302 = 060202,060301 =
060202,060302
---+----- ---+----- ----- ---+---+ ---+----- ---+---+ 060202,0620 = 060202,0635
---+----- ---+----- ----- ---+---+ ---+----- ---+---+ 060202,0613 = 060202,0616
---+----- ---+----- ----- ---+---+ ---+----- ---+---+ 060201,060202 = 060202,060202
---+----- ---+----- ----- ---+---+ ---+----- ---+---+ 060302,061101 = 060302,061102
---+----- ---+----- ----- ---++++- ---+----- ---+---+ 060302,061401 = 060302,061402
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060302,0604 = 060302,060402
---+----- ---+----- ----- ---++++- ---+----- ---+---+ 060302,060801 = 060302,060802
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060301,060302 = 060302,060302
---+----- ---+----- +----- ---+++++ ---+----- ---+---+ 060802,0634 = 0621,0634
---+----- ---+----- +----- ---+++++ ---+----- ---+---+ 0604,0634 = 0634,0634
---+----- ---+----- -+----- ---+++++ ---+----- ---+---+ 061101,0628 = 061102,0628
---+----- ---+----- -+----- ---+++++ ---+----- ---+---+ 061401,0628 = 061402,0628
---+----- ---+----- -+----- ---+++++ ---+----- ---+---+ 0604,0628 = 060402,0628
---+----- ---+----- -+----- ---+++++ ---+----- ---+---+ 060801,0628 = 060802,0628
---+----- ---+----- -+----- ---+++++ ---+----- ---+---+ 060301,0628 = 0628,0628
---+----- ---+----- +----- ---+++++ ---+----- ---+---+ 050201,050202 = 050201,0503 =
050202,050202 = 050202,0503
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060201,0631 = 060301,0620 =
0620,0631
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 061101,061401 = 061102,061401 =
061102,061402
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060201,0626N = 061102,0633 =
0626N,0633
---+----- ---+----- ----- ---+++++ ---+----- +---+---+ 060201,0630 = 0624,0630
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 061401,0620 = 061401,0635
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 0613,061401 = 061401,0616
---+----- ---+----- ----- ---+++++ ---+----- +---+---+ 060201,0617 = 0617,0624
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060201,0606 = 0606,0620
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060201,060501 = 060501,0620 =
0609,0620
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060201,060502 = 060502,0620
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 0615,0620 = 0615,0635
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 061402,0620 = 061402,0635
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 0613,0620 = 0616,0620 = 0616,0635
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 0620,0623 = 0623,0635
---+----- ---+----- ----- ---+++++ ---+----- +---+---+ 0620,0624 = 0624,0635
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 0620,0633 = 0633,0635
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060201,0620 = 060201,0635 =
0620,0620 = 0620,0635
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 0613,0615 = 0615,0616
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060201,0622 = 0616,0622
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 0613,061402 = 061402,0616
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 0613,0623 = 0616,0623
---+----- ---+----- ----- ---+++++ ---+----- +---+---+ 0613,0624 = 0616,0624
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 0613,0633 = 0616,0633
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 0613,0619 = 0616,0619
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060201,0613 = 060201,0616 =
0613,0616 = 0616,0616
---+----- ---+----- ----- ---+++++ ---+----- +---+---+ 060201,0624 = 0624,0624
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060201,0633 = 0633,0633
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 0604,061102 = 060402,061102
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060301,0606 = 0606,0631
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060301,0618 = 061102,0627
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060301,060502 = 060502,0631
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060801,061102 = 060802,061102
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060301,0635 = 061101,0631 =
061102,0631 = 0631,0635
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 061101,0626N = 061102,0626N
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060301,061101 = 060301,061102
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 061101,061102 = 061102,061102
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060801,061101 = 060802,061101
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 0604,061101 = 060402,061101 =
060402,0618
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060501,0606 = 060502,0606 =
0606,0606 = 0606,0609 = 0606,0618 =
0606,0635
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060501,060502 = 060501,0618 =
060501,0635 = 060502,0609 =
060502,0618 = 0609,0635
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 060502,060502 = 060502,0635
---+----- ---+----- ----- ---+++++ ---+----- ---+---+ 061401,0631 = 061402,0631



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--+----- ----- ----- ---++++- -----+-- ---+++- 061401,0626N = 061402,0626N
--+----- ----- ----- ---++++- -----+-- ---+++- 060301,061401 = 060301,061402
--+----- ----- ----- ---++++- -----+-- ---+++- 061401,061401 = 061401,061402
--+----- ----- ----- ---+++++ +-----+-- -----+ 0604,0631 = 060402,0631
--+----- ----- ----- ---+++++ +-----+-- -----+ 0604,0626N = 060402,0626N
--+----- ----- ----- ---+++++ +-----+-- -----+ 0607,060801 = 0607,060802 =
0607,0621
--+----- ----- ----- ---+++++ +-----+-- -----+ 060301,0604 = 060301,060402 =
0621,0632
--+----- ----- ----- ---+++++ -+-----+ +-----+ 060301,0617 = 0621,0630
--+----- ----- ----- ---+++++ +-----+-- -----+ 060301,060501 = 060501,0631 =
0609,0631
--+----- ----- ----- ---+++++ +-----+-- -----+ 060301,0609 = 0612,0632
--+----- ----- ----- ---+++++ +-----+-- -----+ 060301,0627 = 060801,0632 =
060802,0632
--+----- ----- ----- ---+++++ -+-----+ -----+ 060801,0631 = 060802,0631
--+----- ----- ----- ---+++++ -+-----+ +-----+ 060801,0630 = 060802,0630
--+----- ----- ----- ---+++++ -+-----+ -----+ 060801,0626N = 060802,0626N
--+----- ----- ----- ---+++++ -+-----+ -----+ 060301,060801 = 060301,060802
--+----- ----- ----- ---+++++ -+-----+ -----+ 060301,0631 = 0631,0631
--+----- ----- ----- ---+++++ +-----+-- +-----+ 0604,0630 = 060402,0630 = 0617,0632
--+----- ----- ----- ---+++++ +-----+-- -----+ 0604,0607 = 060402,0607
--+----- ----- ----- ---+++++ +-----+-- -----+ 0604,0632 = 060402,0632
--+----- ----- ----- ---+++++ +-----+-- -----+ 0604,060801 = 060402,060801 =
060402,060802 = 060402,0621
--+----- ----- ----- ---+++++ -+-----+ -----+ 060801,060801 = 060801,060802
--+----- ----- ----- ---+++++ +-----+-- -----+ 0604,060402 = 060402,060402
--+----- ----- ----- ---+++++ +-----+-- +-----+ 060802,0625 = 0621,0625
--+----- ----- ----- ---+++++ +-----+-- -----+ 0604,060802 = 0604,0621 = 0621,0627
--+----- ----- ----- ---+++++ -+-----+ +-----+ 060802,0617 = 0617,0621
--+----- ----- ----- ---+++++ -+-----+ -----+ 060802,0621 = 0621,0621
--+----- ----- ----- ---+++++ +-----+-- -----+ 060802,0609 = 0612,0627
--+----- ----- ----- ---+++++ +-----+-- -----+ 060501,060501 = 060501,0609
```

0201 = 020101-020102
0301 = 030101, 030102 and 030104
0302 = 030201-030204
0303 = 030302-030303
0305 = 030501-030504
0401 = 040101-040102
0501 = 050101-050102
0503 = 050301-050302
0601 = 060101-060104
0604 = 060401 and 060403



SPECIFICITY TABLE

DQB1 high resolution for frequent alleles SSP typing

Specificities and sizes of the PCR products of the 47 primer mixes used for DQB1 high resolution for frequent alleles SSP typing

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	Amplified DQB1 alleles
1³	120 bp	515 bp	*020101-0202, 0204, 0205
2³	85 bp	430 bp	*0203
3	145 bp	515 bp	*020101, 020102, 0204, 0205, 030101-0323, 0325, 040101-0403, 050101-0505, 060101-0635
4⁵	140 bp	430 bp	*0202, 0203
5^{4,7}	145 bp, 185 bp	430 bp	*0204, 0505
6⁴	165 bp	515 bp	*030101-030104, 0304, 0309, 0313, 0316, 0319, 0321, 0322, 0324
7⁴	220 bp	515 bp	*030101-030104, 0304, 0309, 0310, 0313, 0314, 0316, 0319, 0321, 0322, 0324
8^{3,4}	135 bp	515 bp	*020101-0202, 0204, 0205, 030201-030204, 0307, 0308, 0311, 0318, 0629
9^{4,8}	175 bp, 220 bp	515 bp	*020101-0205, 030201-030303, 030501-0308, 0311, 0312, 0313 [?] , 0314 [?] , 0315, 0316 [?] , 0317 [?] , 0318-0320, 0323, 0325, 040101-0403
10⁶	135 bp	430 bp	*0304, 0314
11^{3,9}	95 bp, 130 bp	430 bp	*030501-030504, 0317, 0320
12³	125 bp	430 bp	*0306, 0308, 0325, 060202, 060302
13^{4,10}	110 bp, 140 bp	430 bp	*0307, 0315, 0316
14¹¹	135 bp, 260 bp	430 bp	*0309, 0311
15⁴	135 bp	430 bp	*0310, 0312, 0314, 060101-060104
16⁴	260 bp	515 bp	*030101-030104, 0304, 0309, 0310, 0312-0314, 0316, 0319, 0321, 0322, 0324
17¹²	165 bp, 300 bp	430 bp	*0313, 0634
18^{5,13}	130 bp, 190 bp, 220 bp	430 bp	*0318, 0324, 0628
19	135 bp	515 bp	*030101-0315, 0317-0322, 0324
20⁶	130 bp	430 bp	*0203, 030101, 030102, 030104, 030302, 030303, 0306, 0309, 0310, 0312, 0313, 0315-0317, 0319-0324
21	205 bp	515 bp	*040101, 040102
22⁴	205 bp	430 bp	*0402, 0403

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23	135 bp	430 bp	*050101, 050102
24³	120 bp	430 bp	*050201, 050202, 0505
25³	95 bp	515 bp	*050202-050302
26³	120 bp	430 bp	*0504
27	210 bp	430 bp	*060101-060202, 060502 [?] , 0606 [?] , 0610-061102, 0613, 0616, 0618-0620, 0624, 0629, 0633, 0635
28	185 bp	430 bp	*060201, 060202, 061401-0616, 0619, 0620, 0623, 0624, 0633
29	130 bp	430 bp	*060301, 060302, 0607, 061102, 061401, 0626N, 0628, 0630-0632
30	160 bp	430 bp	*0323, 060301, 060302, 060402, 0607, 060801, 061101, 061102, 0626N, 0628, 0630-0632
31	170 bp	515 bp	*060301, 060302, 060801, 060802, 061102, 0612, 061401, 0621, 0626N, 0628, 0631
32	210 bp	515 bp	*060401-060403, 0607, 0617, 0621, 0625, 0634
33⁶	170 bp	430 bp	*060401-0607, 0609, 0618, 0625, 0627, 0632, 0634
34	130 bp	430 bp	*060401-060501, 0606, 060801-0609, 0612, 0617, 0618, 0621, 0627, 0634
35¹⁴	165 bp, 260 bp	515 bp	*060501, 060502 [?] , 0606 [?] , 0620, 0631, 0635
36	210 bp	430 bp	*060501, 060502 [?] , 0606 [?] , 0609, 0612, 0615, 0622
37	180 bp	430 bp	*030101-030104, 0304, 0306 [?] -0308 [?] , 0309, 0310, 0311 [?] -0318 [?] , 0320 [?] , 0321, 0322, 0323 [?] , 0324, 0606
38^{3,15}	100 bp, 185 bp	430 bp	*0610, 061401, 061402
39	190 bp	430 bp	*0613, 0616, 0622
40^{3,16}	110 bp, 150 bp	430 bp	*0321, 0623
41^{5,17}	175 bp, 205 bp	430 bp	*0322, 0617, 0624, 0630
42¹⁸	135 bp, 195 bp	430 bp	*0306, 0325, 0403, 0610, 0625
43¹⁹	155 bp, 185 bp, 215 bp, 250 bp	515 bp	*0205, 0626N, 0633
44	160 bp	430 bp	*060201, 060202, 0610, 0613-0616, 0620, 0623, 0624, 0629, 0633
45	130 bp	515 bp	*0607, 0615
46²⁰	160 bp, 210 bp	515 bp	*060301, 060302, 060801, 060802, 061401, 061402, 0621, 0627, 0628, 0630-0632
47⁴	155 bp	430 bp	*040101-0403, 0619

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¹ Alleles are assigned by the presence of specific PCR product(s). However, the sizes of the specific PCR products may be helpful in the interpretation of DQB1 high resolution typings.

When the primers in a primer mix can give rise to specific PCR products of more than one length this is indicated if the size difference is 20 base pairs or more. Size differences shorter than 20 base pairs are not given. For high resolution SSP kits the respective lengths of the specific PCR product(s) of the alleles amplified by these primer mixes are given.

Nonspecific amplifications, i.e. a ladder or a smear of bands, may sometimes be seen. GC-rich primers have a higher tendency of giving rise to nonspecific amplifications than other primers.

PCR fragments longer than the control bands may sometimes be observed. Such bands should be disregarded and do not influence the interpretation of the SSP typings.

PCR fragments migrating faster than the control bands, but slower than a 400 bp fragment may be seen in some gel read-outs. Such bands can be disregarded and do not influence the interpretation of the SSP typings.

Some primers may give rise to primer oligomer artifacts. Sometimes this phenomenon is an inherent feature of the primer pair(s) of a primer mix. More often it is due to other factors such as too low amount of DNA in the PCR reactions, taking too long time in setting up the PCR reactions, working at elevated room temperature or using thermal cyclers that are not pre-heated.

² The internal positive control primer pairs amplify segments of the human growth hormone gene. The two different control primer pairs give rise to either an internal positive control band of 430 base pairs, for most wells, or a band of 515 base pairs, for some wells.

Well number 1 contains the primer pair giving rise to the longer, 515 bp, internal positive control band in order to help in the correct orientation of the DQB1 high resolution for frequent alleles typing.

In addition, wells number 3, 6, 7, 8, 9, 16, 19, 21, 25, 31, 32, 35, 43, 45 and 46 contain the primer pair giving rise to the longer, 515 bp, internal positive control band in order to allow kit identification.

In the presence of a specific amplification the intensity of the control band often decreases.

³ Specific PCR fragments shorter than 125 base pairs are less intense and not as sharp as longer specific bands.

⁴ Primer mixes 6, 7, 8, 9, 15, 16, 22 and 47 may yield less specific PCR products than the other DQB1 primer mixes.

⁵ Primer mixes 4, 18 and 41 may give rise to primer oligomer formation.

⁶ Primer mixes 10, 20 and 33 may have a weak tendency of nonspecific amplifications.

⁷ Primer mix 5: Specific PCR fragment of 145 bp in the DQB1*0204 allele. Specific PCR fragment of 185 bp in the DQB1*0505 and alleles.

⁸ Primer mix 9: Specific PCR fragment of 175 bp in DQB1*030501-030504, 0313?, 0314?, 0316?, 0317?, 0319 and 040101-0402 alleles. Specific PCR fragment of 220 bp in the DQB1*020101, 0202 and 0204 alleles. Specific PCR fragments of 175 and 220 bp (both specific PCR fragments may not be obtained) in the DQB1*020102, 0203, 0205, 030201-030303, 0306-0308, 0311, 0312, 0315, 0318, 0320, 0323, 0325 and 0403 alleles.

⁹ Primer mix 11: Specific PCR fragment of 95 bp in the DQB1*0320 allele. Specific PCR fragment of 130 bp in the DQB1*030501-030504 and 0317 alleles.

¹⁰ Primer mix 13: Specific PCR fragment of 110 bp in the DQB1*0315 allele. Specific PCR fragment of 140 bp in the DQB1*0307 and 0316 alleles.

¹¹ Primer mix 14: Specific PCR fragment of 135 bp in the DQB1*0309 allele. Specific PCR fragment of 260 bp in the DQB1*0311 allele.

¹² Primer mix 17: Specific PCR fragment of 165 bp in the DQB1*0313 allele. Specific PCR fragment of 300 bp in the DQB1*0634 allele.

¹³ Primer mix 18: Specific PCR fragment of 130 bp in the DQB1*0324 allele. Specific PCR fragment of 190 bp in the DQB1*0628 allele. Specific PCR fragment of 220 bp in the DQB1*0318 allele.

¹⁴ Primer mix 35: Specific PCR fragment of 165 bp in DQB1*0635 allele. Specific PCR fragment of 260 bp in the DQB1*060501, 060502?, 0606?, 0620 and 0631 alleles.

¹⁵ Primer mix 38: Specific PCR fragment of 100 bp in DQB1*061401 and 061402 alleles. Specific PCR fragment of 185 bp in the DQB1*0610 allele.

¹⁶ Primer mix 40: Specific PCR fragment of 110 bp in the DQB1*0623 allele. Specific PCR fragment of 150 bp in the DQB1*0321 allele.

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¹⁷Primer mix 41: Specific PCR fragment of 175 bp in the DQB1*0617, 0624 and 0630 alleles. Specific PCR fragment of 205 bp in the DQB1*0322 allele.

¹⁸Primer mix 42: Specific PCR fragment of 135 bp in the DQB1*0610 and 0625 alleles. Specific PCR fragment of 195 bp in the DQB1*0306, 0325 and 0403 alleles.

¹⁹Primer mix 43: Specific PCR fragment of 155 bp in the DQB1*0205 allele. Specific PCR fragments of 185 and 215 bp in the DQB1*0626N allele. Specific PCR fragment of 265 bp in the DQB1*0633 allele.

²⁰Primer mix 46: Specific PCR fragment of 160 bp in the DQB1*0621 allele. Specific PCR fragment of 210 bp in the DQB1*0627, 0630 and 0632 alleles. Specific PCR fragment of 160 and 210 bp (both specific PCR fragments may not be obtained) in the *060301-060302, 060801-060802, 061401-061402, 0628 and 0631 alleles.

‘?’ nucleotide sequence information not available for the primer matching sequence.

‘w’, might be weakly amplified.

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INTERPRETATION TABLE																									
DQB1 high resolution SSP typing																									
Amplification patterns of the DQB1*0201 to DQB1*0635 alleles																									
	Well ⁵																								
	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	
Length of spec.	120	85	145	140	145	165	220	135	175	135	95	125	110	135	135	260	165	130	135	130	205	205	135	120	
PCR product(s)					185				220		130		140	260			300	190		220					
Length of int.	515	430	515	430	430	515	515	515	515	430	430	430	430	430	430	515	430	430	515	430	515	430	430	430	
pos. control ¹																									
5'-primer(s) ²	30(185) 5'-AAG ^{3'}	57(266) 5'-TgA ^{3'}	135(500) 5'-TgA ^{3'}	102(400) 5'-TCT ^{3'}	38(209) 5'-CgC ^{3'}	45(230) 5'-ggA ^{3'}	26(173) 5'-TTA ^{3'}	26(173) 5'-TCT ^{3'}	26(173) 5'-TCT ^{3'}	26(173) 5'-TTA ^{3'}	26(173) 5'-ggg ^{3'}	38(210) 5'-gCA ^{3'}	49(242) 5'-ggT ^{3'}	14(136) 5'-gCC ^{3'}	13(134) 5'-ggC ^{3'}	13(134) 5'-ggC ^{3'}	26(173) 5'-TTA ^{3'}	27(175) 5'-TTC ^{3'}	55(260) 5'-gCC ^{3'}	57(266) 5'-TgA ^{3'}	23(164) 5'-gCT ^{3'}	23(164) 5'-gCg ^{3'}	26(173) 5'-ggg ^{3'}	30(184) 5'-gAC ^{3'}	
					136(502) 5'-ACg ^{3'}				140(514) 5'-CAA ^{3'}		38(208) 5'-ACA ^{3'}		55(260) 5'-gCA ^{3'}	135(500) 5'-TgA ^{3'}			102(400) 5'-TCT ^{3'}	38(209) 5'-CgT ^{3'}							
													63(285) 5'-Agg ^{3'}					140(514) 5'-CAA ^{3'}							
3'-primer(s) ³	57(266) 5'-Cgg ^{3'}	71(309) 5'-CgT ^{3'}	170(604) 5'-gAC ^{3'}	135(500) 5'-ggC ^{3'}	86(353) 5'-ACg ^{3'}	86(353) 5'-gCT ^{3'}	86(353) 5'-gCT ^{3'}	57(266) 5'-Cgg ^{3'}	86(353) 5'-gCT ^{3'}	57(266) 5'-Cgg ^{3'}	55(260) 5'-gCg ^{3'}	62(282) 5'-CTA ^{3'}	86(353) 5'-gCT ^{3'}	86(353) 5'-gCT ^{3'}	45(230) 5'-CCC ^{3'}	86(353) 5'-gCT ^{3'}	67(296) 5'-ggT ^{3'}	86(353) 5'-gCT ^{3'}	86(353) 5'-gCT ^{3'}	86(353) 5'-gCT ^{3'}	86(353) 5'-gCT ^{3'}	77(327) 5'-ACg ^{3'}	77(327) 5'-ACg ^{3'}	57(266) 5'-CAA ^{3'}	57(265) 5'-gCT ^{3'}
					170(604) 5'-gAC ^{3'}				185(650) 5'-CgA ^{3'}			70(304) 5'-CCC ^{3'}		167(596) 5'-CAT ^{3'}			189(661) 5'-CCA ^{3'}	87(356) 5'-ggA ^{3'}							
																		170(604) 5'-gAT ^{3'}							
Well No.	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	



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INTERPRETATION TABLE																							
DQB1 high resolution SSP typing																							
Amplification patterns of the DQB1*0201 to DQB1*0635 alleles																							
Well ⁵																							
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	
95	120	210	185	130	160	170	210	170	130	165	210	180	100	190	110	175	135	155	160	130	160	155	Length of spec. PCR product(s)
										260			185		150	205	195	185				210	
																		215					
																		250					
515	430	430	430	430	430	515	515	430	430	515	430	430	430	430	430	430	430	515	430	515	515	430	Length of int. pos. control ¹
30(184)	30(184)	30(184)	9(122)	27(177)	9(122)	27(177)	30(184)	27(177)	27(177)	14(136)	30(184)	27(177)	9(122)	9(122)	26(173)	26(173)	26(173)	11(129)	9(122)	57(266)	30(184)	9(122)	5'-primer(s) ²
5'-gAC 3'	5'-gAT 3'	5'-gAT 3'	5'-gTT 3'	5'-gTA 3'	5'-gTA 3'	5'-gTA 3'	5'-gAC 3'	5'-gTA 3'	5'-gTA 3'	5'-gCC 3'	5'-gAT 3'	5'-gTA 3'	5'-gTT 3'	5'-gTT 3'	5'-ggg 3'	5'-TCT 3'	5'-TCT 3'	5'-TTA 3'	5'-gTT 3'	5'-TgA 3'	5'-gAC 3'	5'-gTT 3'	
		30(184)								45(230)		140(514)			133(493)	130(485)		29(181)					
		5'-gAT 3'								5'-ggA 3'		5'-CAA 3'			5'-TTT 3'	5'-CCA 3'		5'-CCT 3'					
																		38(210)					
																		5'-gTg 3'					
47(237)	57(265)	86(353)	57(266)	57(266)	48(240)	70(304)	86(353)	70(304)	57(266)	86(353)	86(353)	74(317)	27(177)	57(266)	48(240)	71(307)	57(265)	77(326)	48(240)	86(353)	70(304)	47(237)	3'-primer(s) ³
5'-CgA 3'	5'-gCT 3'	5'-ACg 3'	5'-CAT 3'	5'-CAT 3'	5'-gCg 3'	5'-CCC 3'	5'-ACC 3'	5'-CCT 3'	5'-CAA 3'	5'-ACC 3'	5'-ACC 3'	5'-CCg 3'	5'-gTT 3'	5'-CAA 3'	5'-gCg 3'	5'-ggC 3'	5'-gCT 3'	5'-CCg 3'	5'-gCg 3'	5'-ACC 3'	5'-CCC 3'	5'-CgA 3'	
										87(356)		185(650)	30(184)	60(274)	170(604)	185(650)	77(327)	86(353)			86(353)		
										5'-ggA 3'		5'-Cgg 3'	5'-gTg 3'	5'-gTT 3'	5'-gAC 3'	5'-Cgg 3'	5'-ACg 3'	5'-ACg 3'			5'-ACg 3'		
													57(265)										
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	Well No.



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Length of spec.	120	85	145	140	145	165	220	135	175	135	95	125	110	135	135	260	165	130	135	130	205	205	135	120
PCR product(s)					185				220		130		140	260			300	190						
Well No.	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24
DQB1 allele ⁴																								
*020101, 020102	1		3					8	9															
*0202	1			4				8	9															
*0203		2		4					9											20				
*0204	1		3		5			8	9															
*0205	1		3					8	9															
*030101, 030102, 030104			3			6	7									16			19	20				
*030103			3			6	7									16			19					
*030201-030204			3					8	9										19					
*030302, 030303			3						9										19	20				
*0304			3			6	7			10						16			19					
*030501-030504			3						9		11								19					
*0306			3						9			12							19	20				
*0307			3					8	9				13						19					
*0308			3					8	9			12							19					
*0309			3			6	7							14		16			19	20				
*0310			3				7								15	16			19	20				
*0311			3					8	9					14					19					
*0312			3						9						15	16			19	20				
*0313			3			6	7		?							16	17		19	20				
*0314			3				7		?	10					15	16			19					
*0315			3						9				13						19	20				
*0316			3			6	7		?				13			16				20				
*0317			3						?		11								19	20				
*0318			3					8	9										18	19				
*0319			3			6	7		9							16			19	20				
*0320			3						9		11								19	20				
*0321			3			6	7									16			19	20				
*0322			3			6	7									16			19	20				
*0323			3						9											20				
*0324						6	7									16			18	19	20			
*0325			3						9			12												
*040101, 040102			3						9												21			
*0402			3						9													22		
*0403			3						9														22	
Well No.	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24



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95	120	210	185	130	160	170	210	170	130	165	210	180	100	190	110	175	135	155	160	130	160	155	Length of spec. PCR product(s)	
										260			185		150	205	195	185			210			
																		215						
																		250						
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	Well No.	
																								DQB1 allele ⁴
																								*020101, 020102
																								*0202
																								*0203
																								*0204
																		43						*0205
												37												*030101, 030102, 030104
												37												*030103
																								*030201-030204
																								*030302, 030303
												37												*0304
																	42							*030501-030504
												?												*0306
												?												*0307
												?												*0308
												37												*0309
												37												*0310
												?												*0311
												?												*0312
												?												*0313
												?												*0314
												?												*0315
												?												*0316
												?												*0317
												?												*0318
												?												*0319
												?												*0320
												37			40									*0321
												37				41								*0322
					30							?												*0323
												37												*0324
																	42							*0325
																						47		*040101, 040102
																						47		*0402
																42						47		*0403
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	Well No.	



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Length of spec.	120	85	145	140	145	165	220	135	175	135	95	125	110	135	135	260	165	130	135	130	205	205	135	120
PCR product(s)					185				220		130		140	260			300	190						
																		220						
Well No.	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24
*050101, 050102			3																				23	
*050201			3																					24
*050202			3																					24
*050301, 050302			3																					
*0504			3																					
*0505			3		5																			24
*060101-060104			3												15									
*060201			3																					
*060202			3									12												
*060301			3																					
*060302			3									12												
*060401, 060403			3																					
*060402			3																					
*060501			3																					
*060502			3																					
*0606			3																					
*0607			3																					
*060801			3																					
*060802			3																					
*0609			3																					
*0610			3																					
*061101			3																					
*061102			3																					
*0612			3																					
*0613			3																					
*061401			3																					
*061402			3																					
*0615			3																					
*0616			3																					
*0617			3																					
*0618			3																					
*0619			3																					
*0620			3																					
*0621			3																					
*0622			3																					
*0623			3																					
*0624			3																					
Well No.	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24

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95	120	210	185	130	160	170	210	170	130	165	210	180	100	190	110	175	135	155	160	130	160	155	Length of spec. PCR product(s)		
										260			185		150	205	195	185			210				
																		215							
																		250							
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	Well No.		
																								*050101, 050102	
																								*050201	
25																								*050202	
25																								*050301, 050302	
	26																							*0504	
																								*0505	
		27																						*060101-060104	
		27	28																	44				*060201	
		27	28																	44				*060202	
				29	30	31																46		*060301	
				29	30	31																46		*060302	
							32	33	34															*060401, 060403	
					30		32	33	34															*060402	
								33	34	35	36													*060501	
		?						33		?	?													*060502	
		?						33	34	?	?	37												*0606	
				29	30		32	33														45		*0607	
					30	31			34														46	*060801	
						31			34														46	*060802	
								33	34		36													*0609	
		27										38					42		44					*0610	
		27			30																			*061101	
		27		29	30	31																		*061102	
						31			34		36													*0612	
		27												39						44				*0613	
			28	29		31							38							44		46		*061401	
			28										38							44		46		*061402	
			28								36									44		45		*0615	
		27	28											39						44				*0616	
							32		34							41								*0617	
		27						33	34															*0618	
		27	28																				47	*0619	
		27	28							35										44				*0620	
					31	32			34													46		*0621	
										36			39											*0622	
			28											40						44				*0623	
		27	28												41					44				*0624	
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	Well No.		



Lot No.: **09G**

Lot-specific information

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Length of spec.	120	85	145	140	145	165	220	135	175	135	95	125	110	135	135	260	165	130	135	130	205	205	135	120
PCR product(s)					185				220		130		140	260			300	190						
																		220						
Well No.	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24
*0625			3																					
*0626N			3																					
*0627			3																					
*0628			3															18						
*0629			3					8																
*0630			3																					
*0631			3																					
*0632			3																					
*0633			3																					
*0634			3															17						
*0635			3																					
DQB1 allele ⁴																								
Well No.	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24

¹The internal positive control primer pairs amplify segments of the human growth hormone gene. The two different control primer pairs give rise to either an internal positive control band of 430 base pairs, for most wells, or a band of 515 base pairs, for some wells.

Well number 1 contains the primer pair giving rise to the longer, 515 bp, internal positive control band in order to help in the correct orientation of the DQB1 high resolution for frequent alleles typing.

In addition, wells number 3, 6, 7, 8, 9, 16, 19, 21, 25, 31, 32, 35, 43, 45 and 46 contain the primer pair giving rise to the longer, 515 bp, internal positive control band in order to allow kit identification.

²The codon, in the 2nd or 3rd exon, matching the specificity-determining 3'-end of the primer is given. Codon numbering as on the www.ebi.ac.uk/imgt/hla web site. The sequence of the 3 terminal nucleotides of the primer is given. Empty spaces indicate codon boundaries.

³The codon, in the 2nd or 3rd exon, matching the specificity-determining 3'-end of the primer is given in the anti-sense direction. Codon numbering as on the www.ebi.ac.uk/imgt/hla web site. The sequence of the 3 terminal nucleotides of the primer is given. Empty spaces indicate codon boundaries.

⁴The sequence of the DQB1*03031 allele has been shown to be identical to DQB1*030302.

⁵Primer mix 5: Specific PCR fragment of 145 bp in the DQB1*0204 allele. Specific PCR fragment of 185 bp in the DQB1*0505 and alleles.

Primer mix 9: Specific PCR fragment of 175 bp in DQB1*030501-030504, 0313?, 0314?, 0316?, 0317?, 0319 and 040101-0402 alleles. Specific PCR fragment of 220 bp in the DQB1*020101, 0202 and 0204 alleles. Specific PCR fragments of 175 and 220 bp (both specific PCR fragments may not be obtained) in the DQB1*020102, 0203, 0205, 030201-030303, 0306-0308, 0311, 0312, 0315, 0318, 0320, 0323, 0325 and 0403 alleles.

Primer mix 11: Specific PCR fragment of 95 bp in the DQB1*0320 allele. Specific PCR fragment of 130 bp in the DQB1*030501-030504 and 0317 alleles.

Primer mix 13: Specific PCR fragment of 110 bp in the DQB1*0315 allele. Specific PCR fragment of 140 bp in the DQB1*0307 and 0316 alleles.

Lot No.: **09G**

Lot-specific information

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95	120	210	185	130	160	170	210	170	130	165	210	180	100	190	110	175	135	155	160	130	160	155	Length of spec. PCR product(s)	
										260			185		150	205	195	185			210			
																		215						
																		250						
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	Well No.	
							32	33									42							*0625
				29	30	31												43						*0626N
								33	34													46		*0627
				29	30	31																46		*0628
		27																	44					*0629
				29	30											41						46		*0630
				29	30	31				35												46		*0631
				29	30			33														46		*0632
		27	28																43	44				*0633
							32	33	34															*0634
		27								35														*0635
																								DQB1 allele ⁴
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	Well No.	

Primer mix 14: Specific PCR fragment of 135 bp in the DQB1*0309 allele. Specific PCR fragment of 260 bp in the DQB1*0311 allele.

Primer mix 17: Specific PCR fragment of 165 bp in the DQB1*0313 allele. Specific PCR fragment of 300 bp in the DQB1*0634 allele.

Primer mix 18: Specific PCR fragment of 130 bp in the DQB1*0324 allele. Specific PCR fragment of 190 bp in the DQB1*0628 allele. Specific PCR fragment of 220 bp in the DQB1*0318 allele.

Primer mix 35: Specific PCR fragment of 165 bp in DQB1*0635 allele. Specific PCR fragment of 260 bp in the DQB1*060501, 060502[?], 0606[?], 0620 and 0631 alleles.

Primer mix 38: Specific PCR fragment of 100 bp in DQB1*061401 and 061402 alleles. Specific PCR fragment of 185 bp in the DQB1*0610 allele.

Primer mix 40: Specific PCR fragment of 110 bp in the DQB1*0623 allele. Specific PCR fragment of 150 bp in the DQB1*0321 allele.

Primer mix 41: Specific PCR fragment of 175 bp in the DQB1*0617, 0624 and 0630 alleles. Specific PCR fragment of 205 bp in the DQB1*0322 allele.

Primer mix 42: Specific PCR fragment of 135 bp in the DQB1*0610 and 0625 alleles. Specific PCR fragment of 195 bp in the DQB1*0306, 0325 and 0403 alleles.

Primer mix 43: Specific PCR fragment of 155 bp in the DQB1*0205 allele. Specific PCR fragments of 185 and 215 bp in the DQB1*0626N allele. Specific PCR fragment of 265 bp in the DQB1*0633 allele.

Primer mix 46: Specific PCR fragment of 160 bp in the DQB1*0621 allele. Specific PCR fragment of 210 bp in the DQB1*0627, 0630 and 0632 alleles. Specific PCR fragment of 160 and 210 bp (both specific PCR fragments may not be obtained) in the *060301-060302, 060801-060802, 061401-061402, 0628 and 0631 alleles.

‘w’, might be weakly amplified.

‘?’ , nucleotide sequence information not available for the primer matching sequence.

Lot No.: **09G**

Lot-specific information

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CELL LINE VALIDATION SHEET																			
DQB1 high resolution primer set																			
			Production No.	Well															
				1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16
				200851401	200851402	200851403	200961804	200851405	200851406	200851407	200851408	200851409	200851410	200851411	200851412	200851413	200851414	200851415	200851416
	IHWC cell line	DQB1																	
1	9001 SA	*0501		-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-
2	9280 LK707	*0601	*0202	+	-	+	+	-	-	-	+	+	-	-	-	-	-	+	-
3	9011 E4181324	*0601		-	-	+	-	-	-	-	-	-	-	-	-	-	-	+	-
4	9275 GU373	*0201		+	-	+	-	-	-	-	+	+	-	-	-	-	-	-	-
5	9009 KAS011	*0502		-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-
6	9353 SM	*0302	*0601	-	-	+	-	-	-	-	+	+	-	-	-	-	-	+	-
7	9020 QBL	*0201		+	-	+	-	-	-	-	+	+	-	-	-	-	-	-	-
8	9025 DEU	*0301		-	-	+	-	-	+	+	-	-	-	-	-	-	-	-	+
9	9026 YAR	*0302		-	-	+	-	-	-	-	+	+	-	-	-	-	-	-	-
10	9107 LKT3	*0401		-	-	+	-	-	-	-	-	+	-	-	-	-	-	-	-
11	9051 PITOUT	*0202		+	-	-	+	-	-	-	+	+	-	-	-	-	-	-	-
12	9052 DBB	*0303		-	-	+	-	-	-	-	-	+	-	-	-	-	-	-	-
13	9004 JESTHOM	*0501		-	-	+	-	-	-	-	-	+	-	-	-	-	-	-	-
14	9071 OLGA	*0402		-	-	+	-	-	-	-	-	+	-	-	-	-	-	-	-
15	9075 DKB	*0303		-	-	+	-	-	-	-	-	+	-	-	-	-	-	-	-
16	9037 SWEIG007	*0301		-	-	+	-	-	+	+	-	-	-	-	-	-	-	-	+
17	9282 CTM3953540	*0201	*0603	+	-	+	-	-	-	-	+	+	-	-	-	-	-	-	-
18	9257 32367	*0602	*0202	+	-	+	+	-	-	-	+	+	-	-	-	-	-	-	-
19	9038 BM16	*0301		-	-	+	-	-	+	+	-	-	-	-	-	-	-	-	+
20	9059 SLE005	*0604		-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-
21	9064 AMALA	*0301		-	-	+	-	-	+	+	-	-	-	-	-	-	-	-	+
22	9056 KOSE	*0503	*0604	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-
23	9124 IHL	*0503	*0601	-	-	+	-	-	-	-	-	-	-	-	-	-	-	+	-
24	9035 JBUSH	*0301		-	-	+	-	-	+	+	-	-	-	-	-	-	-	-	+
25	9049 IBW9	*0202		+	-	-	+	-	-	-	+	+	-	-	-	-	-	-	-
26	9285 WT49	*0201		+	-	+	-	-	-	-	+	+	-	-	-	-	-	-	-
27	9191 CH1007	*0401	*0501	-	-	+	-	-	-	-	-	+	-	-	-	-	-	-	-
28	9320 BEL5GB	*0202	*0301	+	-	+	+	-	+	+	+	+	-	-	-	-	-	-	+
29	9050 MOU	*0202		+	-	-	+	-	-	-	+	+	-	-	-	-	-	-	-
30	9021 RSH	*0402		-	-	+	-	-	-	-	-	+	-	-	-	-	-	-	-
31	9019 DUCAF	*0201		+	-	+	-	-	-	-	+	+	-	-	-	-	-	-	-
32	9297 HAG	*0301		-	-	+	-	-	+	+	-	-	-	-	-	-	-	-	+
33	9098 MT14B	*0302		-	-	+	-	-	-	-	+	+	-	-	-	-	-	-	-
34	9104 DHIF	*0301		-	-	+	-	-	+	+	-	-	-	-	-	-	-	-	+
35	9302 SSTO	*0305		-	-	+	-	-	-	-	-	+	-	+	-	-	-	-	-
36	9024 KT17	*0302		-	-	+	-	-	-	-	+	+	-	-	-	-	-	-	-
37	9065 HHKB	*0603		-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-
38	9099 LZL	*0301		-	-	+	-	-	+	+	-	-	-	-	-	-	-	-	+
39	9315 CML	*0201	*0301	+	-	+	-	-	+	+	+	+	-	-	-	-	-	-	+
40	9134 WHONP199	*0202	*0303	+	-	+	+	-	-	-	+	+	-	-	-	-	-	-	-
41	9055 H0301	*0609		-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-
42	9066 TAB089	*0601		-	-	+	-	-	-	-	-	-	-	-	-	-	-	+	-
43	9076 T7526	*0303		-	-	+	-	-	-	-	-	+	-	-	-	-	-	-	-
44	9057 TEM	*0503		-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-
45	9239 SHJO	*0202		+	-	-	+	-	-	-	+	+	-	-	-	-	-	-	-
46	9013 SCHU	*0602		-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-
47	9045 TUBO	*0301		-	-	+	-	-	+	+	-	-	-	-	-	-	-	-	+
48	9303 TER-ND	*0501		-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-



Lot No.: **09G**

Lot-specific information

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CELL LINE VALIDATION SHEET																					
DQB1 high resolution primer set																					
				Production No.	Well																
					17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	
					200851417	200961818	200851419	200851420	200851421	200851422	200851423	200851424	200851425	200851426	200851427	200851428	200851429	200851430	200851431	200851432	
	IHWC cell line	DQB1																			
1	9001 SA	*0501			-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
2	9280 LK707	*0601	*0202		-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-
3	9011 E4181324	*0601			-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-
4	9275 GU373	*0201			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
5	9009 KAS011	*0502			-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-
6	9353 SM	*0302	*0601		-	-	+	-	-	-	-	-	-	-	+	-	-	-	-	-	-
7	9020 QBL	*0201			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
8	9025 DEU	*0301			-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
9	9026 YAR	*0302			-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
10	9107 LKT3	*0401			-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-
11	9051 PITOUT	*0202			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
12	9052 DBB	*0303			-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
13	9004 JESTHOM	*0501			-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
14	9071 OLGA	*0402			-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
15	9075 DKB	*0303			-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
16	9037 SWEIG007	*0301			-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
17	9282 CTM3953540	*0201	*0603		-	-	-	-	-	-	-	-	-	-	-	-	-	+	+	+	-
18	9257 32367	*0602	*0202		-	-	-	-	-	-	-	-	-	-	+	+	-	-	-	-	-
19	9038 BM16	*0301			-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
20	9059 SLE005	*0604			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
21	9064 AMALA	*0301			-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
22	9056 KOSE	*0503	*0604		-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	+
23	9124 IHL	*0503	*0601		-	-	-	-	-	-	-	-	+	-	+	-	-	-	-	-	-
24	9035 JBUSH	*0301			-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
25	9049 IBW9	*0202			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
26	9285 WT49	*0201			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
27	9191 CH1007	*0401	*0501		-	-	-	-	+	-	+	-	-	-	-	-	-	-	-	-	-
28	9320 BEL5GB	*0202	*0301		-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
29	9050 MOU	*0202			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
30	9021 RSH	*0402			-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
31	9019 DUCAF	*0201			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
32	9297 HAG	*0301			-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
33	9098 MT14B	*0302			-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
34	9104 DHIF	*0301			-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
35	9302 SSTO	*0305			-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
36	9024 KT17	*0302			-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
37	9065 HHKB	*0603			-	-	-	-	-	-	-	-	-	-	-	-	-	+	+	+	-
38	9099 LZL	*0301			-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
39	9315 CML	*0201	*0301		-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
40	9134 WHONP199	*0202	*0303		-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
41	9055 H0301	*0609			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
42	9066 TAB089	*0601			-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-
43	9076 T7526	*0303			-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
44	9057 TEM	*0503			-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-
45	9239 SHJO	*0202			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
46	9013 SCHU	*0602			-	-	-	-	-	-	-	-	-	-	+	+	-	-	-	-	-
47	9045 TUBO	*0301			-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
48	9303 TER-ND	*0501			-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-



Lot No.: **09G**

Lot-specific information

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CELL LINE VALIDATION SHEET																			
DQB1 high resolution primer set																			
				Production No.	Well														
					33	34	35	36	37	38	39	40	41	42	43	44	45	46	47
					200851433	200851434	200961835	200851436	200961837	200961838	200851439	200851440	200851441	200851442	200961843	200851444	200851445	200851446	200851447
	IHWC cell line	DQB1																	
1	9001 SA	*0501			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
2	9280 LK707	*0601	*0202		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
3	9011 E4181324	*0601			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
4	9275 GU373	*0201			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
5	9009 KAS011	*0502			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
6	9353 SM	*0302	*0601		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
7	9020 QBL	*0201			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
8	9025 DEU	*0301			-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
9	9026 YAR	*0302			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
10	9107 LKT3	*0401			-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
11	9051 PITOUT	*0202			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
12	9052 DBB	*0303			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
13	9004 JESTHOM	*0501			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
14	9071 OLGA	*0402			-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
15	9075 DKB	*0303			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
16	9037 SWEIG007	*0301			-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
17	9282 CTM3953540	*0201	*0603		-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
18	9257 32367	*0602	*0202		-	-	-	-	-	-	-	-	-	-	-	+	-	-	-
19	9038 BM16	*0301			-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
20	9059 SLE005	*0604			+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
21	9064 AMALA	*0301			-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
22	9056 KOSE	*0503	*0604		+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
23	9124 IHL	*0503	*0601		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
24	9035 JBUSH	*0301			-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
25	9049 IBW9	*0202			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
26	9285 WT49	*0201			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
27	9191 CH1007	*0401	*0501		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
28	9320 BEL5GB	*0202	*0301		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
29	9050 MOU	*0202			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
30	9021 RSH	*0402			-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
31	9019 DUCAF	*0201			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
32	9297 HAG	*0301			-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
33	9098 MT14B	*0302			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
34	9104 DHIF	*0301			-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
35	9302 SSTO	*0305			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
36	9024 KT17	*0302			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
37	9065 HHKB	*0603			-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
38	9099 LZL	*0301			-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
39	9315 CML	*0201	*0301		-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
40	9134 WHONP199	*0202	*0303		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
41	9055 H0301	*0609			+	+	-	+	-	-	-	-	-	-	-	-	-	-	-
42	9066 TAB089	*0601			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
43	9076 T7526	*0303			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
44	9057 TEM	*0503			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
45	9239 SHJO	*0202			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
46	9013 SCHU	*0602			-	-	-	-	-	-	-	-	-	-	-	+	-	-	-
47	9045 TUBO	*0301			-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
48	9303 TER-ND	*0501			-	-	-	-	-	-	-	-	-	-	-	-	-	-	-



Lot No.: **09G**

Lot-specific information

www.olerup.com

CERTIFICATE OF ANALYSIS

Olerup SSP[®] DQB1 high resolution for frequent alleles SSP

Product number: 101.221-12 – including *Taq* polymerase
Lot number: 09G
Expiry date: 2011-August-01
Number of tests: 12 tests
Number of wells per test: 47 + 1

Well specifications:

Well No.	Production No.	Well No.	Production No.	Well No.	Production No.
1	2008-514-01	17	2008-514-17	33	2008-514-33
2	2008-514-02	18	2009-618-18	34	2008-514-34
3	2008-514-03	19	2008-514-19	35	2009-618-35
4	2009-618-04	20	2008-514-20	36	2008-514-36
5	2008-514-05	21	2008-514-21	37	2009-618-37
6	2008-514-06	22	2008-514-22	38	2009-618-38
7	2008-514-07	23	2008-514-23	39	2008-514-39
8	2008-514-08	24	2008-514-24	40	2008-514-40
9	2008-514-09	25	2008-514-25	41	2008-514-41
10	2008-514-10	26	2008-514-26	42	2008-514-42
11	2008-514-11	27	2008-514-27	42	2009-618-43
12	2008-514-12	28	2008-514-28	43	2008-514-44
13	2008-514-13	29	2008-514-29	45	2008-514-45
14	2008-514-14	30	2008-514-30	46	2008-514-46
15	2008-514-15	31	2008-514-31	47	2008-514-47
16	2008-514-16	32	2008-514-32		

The specificity of each primer solution of the kit has been tested against 48 well characterized IHWC cell line DNAs.

No DNAs carrying the alleles to be amplified by primer solutions 2, 5, 10, 12, 13, 14, 17, 18, 26, 35, 38-43 and 45 were available. The specificities of the primers in primer solutions 2, 5, 10, 12, 14, 18, 26, 35, 38, 39, 40, 41, 42 and 45 were tested by separately adding one additional 5'-primer, respectively one additional 3'-primer. In primer solution 17 it was only possible to test the 5'-primers, the 3'-primers were not possible to test. In primer solution 13 and 43 it was only possible to test the 3'-primers, the 5'-primers were not possible to test. In primer solutions 5, 18, 40 and 41 one 5'-primer was not possible to test. In primer solutions 14, 18, 39 and 41 one 3'-primer was not possible to test. Additional primers in primer solutions 37 and 46 were tested by separately adding one additional 5'-primer, respectively one additional 3'-primer.

The negative control primer pairs, **Production No. 2008-495-01**, can detect contamination with PCR products diluted 10^{-7} .

Results: No false positive or false negative amplifications were obtained.

Date of approval: 2009-August-28

Approved by:

Quality Control, Supervisor

Lot No.: **09G**

Lot-specific information

www.olerup.com

Declaration of Conformity

Product name: *Olerup* SSP[®] DQB1
Product number: 101.221-12
Lot number: 09G

Intended use: DQB1 high resolution histocompatibility testing

Manufacturer: *Olerup* SSP AB
Hasselstigen 1
SE-133 33 Saltsjöbaden, Sweden
Phone: +46-8-717 88 27
Fax: +46-8-717 88 18

We, *Olerup* SSP AB, hereby declare that this product, to which this Declaration of Conformity relates is in conformity with the following Standard(s) and other normative document(s) ISO 9001:2000, ISO 17025:1999 and ISO 13485:2000, following the provisions of the 98/79/EC Directive on *in vitro* diagnostic medical devices, Annex III.

The Technical Construction File is maintained at *Olerup* SSP AB, Hasselstigen 1, SE-133 33 Saltsjöbaden, Sweden.

The Authorized Representative located within the Community is: *Olerup* SSP AB.

Saltsjöbaden, Sweden
2009-August-28

Olle Olerup
Managing Director

Lot No.: **09G**

Lot-specific information

www.olerup.com

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