

101.201-48/12 – including *Taq* polymerase
101.201-48u/12u – without *Taq* polymerase

Visit www.caredx.com for
“Instructions for Use” (IFU)

Lot No.: **2S4**

Lot-specific information

Olerup SSP® DQ low

Product number: 101.201-48/12 – including *Taq* pol.
101.201-48u/12u – without *Taq* pol.

Lot number: 2S4

Expiry date: 2027-06-01

Number of tests: 48 tests – Product No. 101.201-48/48u
12 tests – Product No. 101.201-12/12u

Number of wells per test: 15+1

Storage - pre-aliquoted primers: dark, between -15°C and -25°C

- PCR Master Mix: between -15°C and -25°C
- Adhesive PCR seals RT

This Product Description is only valid for Lot No. 2S4.

Complete product documentation consists of generic Instructions for Use (IFU), lot specific Product Insert, Worksheet and Certificate.

CHANGES COMPARED TO THE PREVIOUS OLERUP SSP® DQ LOW LOT (5N9)

The product documentation has been updated for new alleles of IMGT 3.52.0.

The DQ low primer set, specificity and interpretation tables have been updated for the HLA-DQB1 alleles described since the previous *Olerup SSP*® DQ low lot was made (**Lot No 5N9**).

The primers of the wells detailed below have been exchanged, added or modified compared to the previous lot (**Lot No. 5N9**).

Well	5'-primer	3'-primer	rationale
8	Added	-	5'-primer added for the *04:03:03 allele.



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Well **16** contains Negative Control primer pairs, that will amplify the majority of the *Olerup SSP*® HLA Class I, DRB, DQB1, DPB1 and DQA1 amplicons as well as all the amplicons generated by the control primer pairs matching the human growth hormone gene.

HLA-specific PCR product sizes range from 75 to 200 base pairs.

The PCR product generated by the positive control primer pair is 200 base pairs.

Length of PCR product	105	200	105	80	75	80	85
5'-primer¹	164	340	440	45	45	43	36
	5'-CAC ^{3'}	5'-Agg ^{3'}	5'-TTA ^{3'}	5'-Tgg ^{3'}	5'-Tgg ^{3'}	5'-Tgg ^{3'}	5'-TAC ^{3'}
							36
							5'-TAT ^{3'}
3'-primer²	231	2nd I	507	59	58	57	47
	5'-TgC ^{3'}	5'-AAA ^{3'}	5'-TTg ^{3'}	5'-CTC ^{3'}	5'-ggC ^{3'}	5'-CTC ^{3'}	5'-ACA ^{3'}
							48
							5'-gCA ^{3'}
							48
							5'-gCC ^{3'}
							52
							5'-TgT ^{3'}
A*	+	+	+				
B*	+	+	+				
C*	+	+	+				
DRB1				+	+		
DRB3				+	+		
DRB5				+			
DQB1					+		
DPB1						+	
DQA1							+

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

DQ low SSP typing

CONTENT

The primer set contains 5'- and 3'-primers for grouping the DQB1 alleles into the serological groups DQ2 to DQ9.

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 16 PCR reactions in a 16 well PCR plate.

1	2	3	4	5	6	7	8
9	10	11	12	13	14	15	NC

The 16 well cut PCR plate is marked with 'DQ LOW' in silver/gray ink.

Well No. 1 is marked with the Lot No. '2S4'.

Wells 1 to 15 – DQ low primers.

Well 16 – Negative Control (NC).

A faint row of numbers is seen between wells 1 and 2 or wells 7 and 8 of the PCR trays. These stem from the manufacture of the trays, and should be disregarded. The PCR plates are covered with a PCR-compatible foil.

Please note: When removing each 16 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 the DQB1 alleles will be amplified by the DQ low typing kit. Thus, the interpretation of DQ low typings is not influenced by the DQB2 and DQB3 genes.

UNIQUELY IDENTIFIED ALLELES

All the DQB1 alleles, i.e. **DQB1*05:01:01:01 to 05:321**, **DQB1*06:01:01:01 to 06:471**, **DQB1*02:01:01 to 02:212**, **DQB1*03:01:01:01 to 03:510** and **DQB1*04*01:01:01 to 04:96**, recognized by the HLA Nomenclature Committee in April 2023^{1,2,4} will be amplified by the primers in the DQ low SSP kit. The DQB1 alleles will be grouped into their corresponding serological specificities, i.e.:

DQ5(1) = DQB1*05:01-05:05³
 DQ6(1) = DQB1*06:01-06:33³
 DQ2 = DQB1*02:01-02:05³
 DQ3 = DQB1*03:01-03:20³
 DQ4 = DQB1*04:01-04:02³



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The DQ3 alleles may be further subdivided into the DQ3, DQ7, DQ8 and DQ9 based upon serology and expert assignment. Thus:

DQ3	=	DQB1*03:06, 03:10, 03:14
DQ7	=	DQB1*03:01:01-03:01:04, 03:04, 03:09, 03:13, 03:16, 03:19
DQ8	=	DQB1*03:02, 03:05, 03:07, 03:08, 03:11, 03:18
DQ9	=	DQB1*03:03, 03:12, 03:15, 03:17, 03:20

¹HLA-DQB1 alleles listed on the IMGT/HLA web page 2023-April-17, release 3.52.0, www.ebi.ac.uk/imgt/hla.

²Alleles that have been deleted from or renamed in the official WHO HLA Nomenclature up to and including the last IMGT/HLA database release can be retrieved from web page <http://hla.alleles.org/alleles/deleted.html>.

³The serological split of the DQB1*05:05-05:321 alleles, the DQB1*06:06-06:07, DQB1*06:10, DQB1*06:13, DQB1*06:15-06:24 and DQB1*06:27-06:510 alleles, the DQB1*02:04-02:212 alleles, the DQB1*03:07-03:09 and DQB1*03:11 to 03:510, and the DQB1*04:03 to 04:96 alleles is not known. The grouping of not serologically defined alleles is taken from the expert-assigned serological grouping in Tissue Antigens (2009) 73:95-170.

⁴The DQ low alleles will be grouped into their corresponding serological specificities, except for the following alleles that will give rise to identical amplification patterns:

Alleles

DQB1*05:01:01-05:01:15, 05:01:17-05:02:26, 05:03:01:01-05:03:23, 05:03:26-05:03:30, 05:05:01-05:43:02, 05:45-05:51, 05:53, 05:55-05:59, 05:61-05:71, 05:73-05:76, 05:78-05:81, 05:84-05:97, 05:99-05:104, 05:106-05:113, 05:115, 05:117-05:127, 05:129-05:131, 05:133-05:145, 05:147-05:174, 05:177-05:206N, 05:208N-05:217, 05:219-05:243, 05:245-05:257, 05:259-05:261, 05:263-05:299, 05:301-05:303, 05:305-05:321, DQB1*06:325



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Lot-specific information
SPECIFICITY TABLE

DQ low SSP typing

Specificities and sizes of the PCR products of the 15+1 primer mixes used for DQ low SSP typing

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	DQ serology ³	Amplified DQB1 alleles ⁴
1	135 bp, 230 bp	515 bp	5	*05:01:01:01-05:01:15, 05:01:17-05:03:24, 05:03:26-05:59, 05:61-05:81, 05:84-05:97, 05:99-05:115, 05:117-05:174, 05:176-05:217, 05:219-05:257, 05:259-05:321, 06:325, 06:389
2⁸	135 bp, 185 bp, 220 bp, 270 bp	515 bp	1, 5, 6	*02:03:02, 03:23:01-03:23:02, 03:217, 03:259, 03:355, 04:10, 05:176, 06:01:01:01-06:155, 06:157-06:161, 06:163-06:168, 06:170-06:219, 06:221-06:324, 06:326-06:382, 06:384-06:398, 06:400-06:455, 06:457-06:471
3	210 bp	430 bp	2	*02:01:01:01-02:01:09, 02:01:11-02:01:13, 02:01:15-02:212
4	130 bp, 220 bp	515 bp	3, 7	*03:01:01:01-03:01:01:12, 03:01:01:14-03:01:58, 03:04:01:01-03:04:04, 03:09-03:10:03, 03:13-03:14:02, 03:16, 03:19:01:01-03:19:06, 03:21-03:22:02, 03:24, 03:27-03:29, 03:35-03:36, 03:42, 03:44, 03:46-03:60, 03:69, 03:71, 03:73, 03:75-03:77, 03:80, 03:82-03:84N, 03:92-03:94, 03:101-03:103, 03:108-03:109, 03:114-03:116, 03:118N-03:122, 03:127-03:131, 03:133-03:135, 03:138-03:140, 03:142-03:144, 03:147-03:148, 03:150, 03:152, 03:154, 03:157-03:160, 03:162-03:167, 03:169-03:173, 03:180, 03:182-03:183, 03:186-03:188, 03:191-03:198:02, 03:201-03:202, 03:206-03:208, 03:216, 03:218-03:219, 03:231-03:232, 03:235-03:236, 03:241-03:243, 03:246, 03:252-03:257, 03:260, 03:264, 03:266-03:268, 03:271, 03:275-03:276N, 03:281, 03:284-03:286, 03:288, 03:290-03:294, 03:297, 03:302-03:303N, 03:305-03:307, 03:309:01-03:309:02, 03:311-03:312, 03:317:01-03:318, 03:326-03:331, 03:335, 03:338N, 03:340N-03:342, 03:347, 03:350, 03:353-03:355, 03:358N, 03:360-03:361, 03:366, 03:370, 03:372-03:373, 03:376N-03:378, 03:380-03:381, 03:385N, 03:387, 03:389-03:391, 03:394, 03:396, 03:399N-03:401, 03:404, 03:407N-03:408, 03:417-03:421, 03:423-03:428, 03:430-03:432, 03:434-03:436, 03:438-03:439, 03:443, 03:448-03:449, 03:451, 03:454-03:455, 03:458, 03:460, 03:465, 03:467-03:470, 03:472-03:476, 03:480Q, 03:482-03:483, 03:485-03:486, 03:488N, 03:491-03:492, 03:496-03:497, 03:499N, 03:503, 03:506, 03:508, 06:209



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5⁶	130 bp, 220 bp	515 bp	6, 8	*03:02:01:01-03:02:37, 03:05:01-03:05:05, 03:07-03:08, 03:11, 03:18, 03:32, 03:37, 03:45:01-03:45:02, 03:61, 03:63-03:64, 03:66N-03:68, 03:70, 03:85, 03:104, 03:106-03:107, 03:125, 03:132, 03:146, 03:153, 03:161, 03:174-03:175, 03:178-03:179, 03:181, 03:184-03:185, 03:189-03:190, 03:199, 03:203-03:205, 03:210-03:211, 03:213N-03:215, 03:220-03:221, 03:223-03:224, 03:226, 03:228-03:229, 03:233, 03:237N, 03:240, 03:245, 03:247, 03:250-03:251, 03:261-03:263:01:02, 03:265, 03:269N, 03:273-03:274, 03:277-03:279, 03:287, 03:289, 03:295-03:296, 03:298-03:301, 03:310N, 03:315, 03:320-03:324, 03:333-03:334N, 03:339N, 03:343-03:346, 03:348-03:349, 03:352, 03:362, 03:364, 03:367-03:369, 03:371, 03:379, 03:383, 03:386, 03:388, 03:392, 03:403N, 03:409-03:410, 03:412, 03:415-03:416, 03:422N, 03:429, 03:433, 03:437, 03:440N-03:442, 03:444, 03:446-03:447, 03:450, 03:452, 03:456-03:457, 03:459, 03:462-03:464, 03:466, 03:471, 03:479, 03:481, 03:484, 03:490, 03:493, 03:495, 03:498, 03:500-03:502, 03:504, 04:75, 06:29, 06:123, 06:139, 06:337
6⁶	135 bp	515 bp	2, 3, 9	*02:77, 02:180, 03:03:02:01-03:03:25, 03:03:27-03:03:29, 03:06, 03:12, 03:15, 03:20, 03:23:03, 03:25:01-03:26, 03:30-03:31, 03:33-03:34, 03:38:01-03:41, 03:43, 03:65, 03:74, 03:79, 03:86-03:91Q, 03:95N-03:99Q, 03:104-03:105, 03:111-03:113, 03:117, 03:123-03:124, 03:126, 03:136-03:137, 03:141, 03:145, 03:149, 03:155-03:156, 03:168, 03:176-03:177, 03:200, 03:209, 03:212, 03:222, 03:227, 03:230, 03:234, 03:238-03:239, 03:248-03:249, 03:258, 03:270, 03:280, 03:282N-03:283, 03:304, 03:313, 03:316, 03:319, 03:332, 03:336-03:337, 03:351, 03:356N-03:357N, 03:359, 03:363, 03:365, 03:374-03:375N, 03:382, 03:384, 03:393, 03:395, 03:397-03:398, 03:402, 03:405-03:406, 03:411N, 03:414, 03:445, 03:453, 03:461, 03:477-03:478, 03:487, 03:489, 03:494, 03:505, 03:507, 03:509N-03:510, 04:03:01-04:03:03, 06:02:43, 06:03:10, 06:03:33, 06:51:01, 06:66, 06:96:01, 06:118:04, 06:168, 06:172, 06:322:01-06:322:02, 06:377
7^{5,7}	80 bp	515 bp	3, 7, 8, 9	*03:01:01:01-03:01:01:12, 03:01:01:14-03:01:06, 03:01:07 ^w , 03:01:08-03:02:05, 03:02:07-03:02:17, 03:02:19-03:02:29, 03:02:31-03:05:05, 03:07, 03:08 ^w , 03:09-03:12, 03:13 ^w , 03:14:01-03:24, 03:26-03:57, 03:58 ^w , 03:59-03:64, 03:65 ^w , 03:66N-03:106, 03:107 ^w , 03:108-03:124, 03:126-03:136, 03:137 ^w , 03:138-03:146, 03:148-03:193, 03:194 ^w , 03:195-03:227, 03:229-03:231, 03:232 ^w , 03:233-03:261, 03:262 ^w , 03:263:01:01-03:282N, 03:283 ^w , 03:284-03:324, 03:326-03:410, 03:412-03:417, 03:419-03:448, 03:450-03:510



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8⁶	135 bp, 160 bp, 185 bp, 210 bp, 235 bp	430 bp	4	*03:132, 04:01:01:01-04:02:01:01, 04:02:01:04-04:96
9	225 bp	430 bp	4	*04:01:01:01-04:02:01:01, 04:02:01:04-04:02:18, 04:02:20-04:03:03, 04:06-04:21, 04:22 ^w , 04:23, 04:24 ^w , 04:25N-04:32, 04:34-04:37, 04:38 ^w , 04:39-04:48, 04:50-04:96
10	185 bp, 215 bp	430 bp	5	*05:01:01:01-05:02:26, 05:03:01:01-05:03:23, 05:03:25-05:03:30, 05:05:01-05:43:02, 05:45-05:51, 05:53, 05:55-05:71, 05:73-05:76, 05:78-05:104, 05:106-05:113, 05:115-05:127, 05:129-05:131, 05:133-05:145, 05:147-05:175, 05:177-05:206N, 05:208N-05:243, 05:245-05:261, 05:263-05:299, 05:301-05:303, 05:305-05:321, 06:156, 06:162, 06:169, 06:325
11⁶	185 bp	430 bp	8, 9	*03:03:11, 03:05:01, 03:05:03, 03:05:05, 03:17:01, 03:61, 03:72, 03:100, 03:181, 03:250, 03:346
12⁶	185 bp	430 bp	4, 7, 8, 9	*03:01:01:01-03:01:01:12, 03:01:01:14-03:01:01:58, 03:01:03-03:01:07, 03:01:09-03:02:02:02, 03:02:04-03:02:12, 03:02:14-03:03:02:11, 03:03:04-03:03:15, 03:03:17-03:04:04, 03:05:03-03:05:04, 03:06-03:17:01, 03:18-03:19:06, 03:21-03:22:02, 03:23:02-03:36, 03:38:01, 03:39-03:60, 03:62-03:71, 03:74, 03:76-03:98, 03:101-03:103, 03:106-03:108, 03:110-03:111, 03:113-03:117, 03:119-03:131, 03:133-03:153, 03:155, 03:157-03:161, 03:163-03:180, 03:182, 03:184-03:188, 03:190-03:203, 03:205-03:222, 03:224-03:225, 03:227-03:232, 03:234-03:236, 03:239-03:249, 03:251, 03:253-03:257, 03:259-03:261, 03:263:01:01-03:290, 03:292-03:324, 03:326-03:334N, 03:337-03:345, 03:347-03:350, 03:353, 03:355-03:370, 03:372-03:390, 03:392-03:442, 03:444-03:510, 04:01:03, 04:02:16, 04:03:03
13⁷	185 bp	515 bp	6	*04:10, 06:02:01:01-06:02:42, 06:02:44-06:02:59, 06:14:01-06:16, 06:19:01-06:20, 06:23-06:24, 06:33, 06:37, 06:46-06:50, 06:51:02, 06:68, 06:70-06:84:01:02, 06:95, 06:96:02-06:97, 06:107, 06:109, 06:111-06:117, 06:122, 06:124-06:127, 06:136-06:138, 06:146:01-06:147, 06:150-06:152, 06:156, 06:159, 06:161-06:163, 06:166, 06:173-06:175, 06:178-06:179N, 06:182-06:183, 06:188, 06:192, 06:197-06:198, 06:200-06:201, 06:208:01-06:208:02, 06:211, 06:213, 06:215-06:216N, 06:219, 06:224-06:228, 06:232, 06:235-06:237, 06:240, 06:242, 06:249, 06:255-06:256, 06:262, 06:264, 06:270:02-06:271, 06:273, 06:284, 06:286, 06:289-06:290, 06:293-06:298, 06:300, 06:304N, 06:306N, 06:308N, 06:311, 06:314-06:315, 06:317N, 06:322:03, 06:324, 06:326, 06:333, 06:335, 06:338, 06:341N, 06:344, 06:347,



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				06:354-06:357, 06:363-06:364, 06:366, 06:370, 06:372, 06:374, 06:376, 06:379N-06:380, 06:383N-06:384, 06:386, 06:388, 06:390, 06:395, 06:397N, 06:401-06:402, 06:404-06:406, 06:409, 06:411-06:413, 06:416Q, 06:422N, 06:430-06:431, 06:436-06:438, 06:445-06:448, 06:451, 06:456N-06:457, 06:461, 06:468
14	185 bp	430 bp	1, 5, 6	*06:02:01-01-06:02:12, 06:02:14-06:03:10, 06:03:12-06:03:47, 06:05:02 [?] -06:06 [?] , 06:08:01-06:08:03, 06:10-06:11:04, 06:13:01-06:14:03, 06:16, 06:18:01-06:20, 06:23-06:24, 06:26N-06:27:02, 06:29-06:33, 06:37, 06:40-06:41:01:03, 06:44, 06:47-06:51:02, 06:59-06:65, 06:67-06:68, 06:70-06:78, 06:80-06:84:01:02, 06:87, 06:90-06:91, 06:95-06:97, 06:99:01-06:99:03, 06:106-06:107, 06:109-06:117, 06:122-06:128, 06:130-06:131, 06:133-06:134, 06:136-06:139, 06:141, 06:143, 06:145:01-06:145:02, 06:147-06:148, 06:150-06:152, 06:154, 06:156, 06:159, 06:161-06:163, 06:165-06:166, 06:169-06:170, 06:173-06:176, 06:178-06:179N, 06:182-06:185, 06:187-06:188, 06:190:01-06:192, 06:195-06:198, 06:200-06:201, 06:206:01-06:206:02, 06:208:01-06:208:02, 06:211, 06:213, 06:215-06:216N, 06:218-06:219, 06:221-06:228, 06:230, 06:232-06:234, 06:236-06:238, 06:240, 06:242, 06:244, 06:248-06:250, 06:253, 06:255-06:256, 06:259, 06:262, 06:264, 06:269-06:273, 06:276, 06:278-06:279, 06:284, 06:286, 06:289-06:290, 06:293-06:298, 06:300, 06:304N, 06:306N, 06:308N, 06:311, 06:314-06:317N, 06:319:01-06:319:02, 06:322:01-06:322:03, 06:324, 06:326-06:329, 06:331, 06:333-06:338, 06:341N, 06:344-06:347, 06:350, 06:352, 06:354-06:357, 06:360, 06:362-06:368, 06:370-06:374, 06:376, 06:378-06:380, 06:383N-06:384, 06:388-06:397N, 06:399-06:406, 06:408-06:413, 06:416Q-06:417, 06:422N-06:425, 06:428, 06:430-06:431, 06:433, 06:437-06:438, 06:440, 06:442-06:443, 06:445-06:448, 06:450, 06:453-06:457, 06:459-06:461, 06:468, 06:470
15	135 bp	430 bp	3, 5, 6, 9	*03:10:01-03:10:02:02, 03:12, 03:14:01-03:14:02, 03:70, 03:179, 03:183, 03:195, 03:474, 06:01:01:01-06:01:06, 06:01:08-06:01:31, 06:01:33-06:01:34, 06:06 [?] , 06:43, 06:54N-06:58, 06:98, 06:99:02-06:105, 06:108, 06:120, 06:132, 06:140, 06:153:01-06:153:02, 06:157, 06:167-06:168, 06:177, 06:181, 06:194, 06:205, 06:209, 06:214, 06:229, 06:239, 06:243, 06:245-06:247, 06:251, 06:257-06:258, 06:260, 06:263, 06:268, 06:274, 06:277, 06:285, 06:302, 06:305, 06:307, 06:309-06:310, 06:312, 06:321, 06:323, 06:330N, 06:342, 06:359, 06:377, 06:382, 06:415, 06:418-06:419, 06:421, 06:427, 06:435, 06:462-06:464, 06:466, 06:469, 06:471
16⁹	-	-	-	Negative Control



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Lot-specific information

¹Alleles 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 DQ low SSP typings. When the primers in a primer mix can give rise to HLA-specific PCR products of more than one length this is indicated if the size difference is more than 20 base pairs. Size differences of 20 base pairs or less are not given. For high resolution SSP kits, the alleles listed are specified according to amplicon length.

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.

²The internal positive control primer pairs amplify segments of the human growth hormone gene. The internal positive control bands are 430 or 515 base pairs respectively, well distribution as outlined in the table. Well number 1 contains the longer, 515 bp, internal positive control band. The well distribution of the internal controls can help in orientation of the kit on gel photo, as well as allow for kit identification. In the presence of a specific amplification the intensity of the control band often decreases.

³The serological split of the DQB1*05:05-05:282 alleles, the DQB1*06:06-06:07, DQB1*06:10, DQB1*06:13, DQB1*06:15-06:24 and DQB1*06:27-06:394N alleles, the DQB1*02:04-02:186 alleles, the DQB1*03:07-03:09 and DQB1*03:11 to 03:466, and the DQB1*04:03 to 04:86 alleles is not known. The grouping of not serologically defined alleles is taken from the expert-assigned serological grouping in Tissue Antigens (2009) 73:95-170.

⁴For several DQB1 alleles 1st and/or 3rd exon(s) and beyond, as well as intron nucleotide sequences, are not available. In these instances it is not known whether some of the primers of the SSP sets are completely matched with the target sequences or not. Assumption is made that unknown sequences in these regions are conserved within allelic groups.

⁵HLA-Specific PCR products shorter than 125 base pairs have a lower intensity and are less sharp than longer PCR products.

⁶Primer mixes 5, 6, 8, 11 and 12 may give a lower yield of HLA-specific PCR products than the other DQ low primer mixes.

⁷Primer mixes 7 and 13 may have tendencies of unspecific amplification.

⁸Primer mix 2 may have tendencies for primer oligomer formation.

⁹Primer mix 16 contains a negative control, which will amplify the majority of the HLA amplicons as well as the amplicons generated by the control primer pairs matching the human growth hormone gene. HLA-specific PCR product sizes range from 75 to 200 base pairs and the PCR product generated by the HGH positive control primer pair is 200 base pairs.

Abbreviations

ser: serological HLA specificity

w: might be weakly amplified.

?: nucleotide sequence information not available for the primer matching sequence.



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Lot-specific information

PRIMER SPECIFICATION

Well No.	1	2	3	4	5	6	7	8	9	10	11	12
Length of spec.	135	135	210	130	130	135	80	135	225	185	185	185
PCR product	230	185		220	220			160		215		
		220						185				
		270						210				
								235				
Length of int.	515	515	430	515	515	515	515	430	430	430	430	430
pos. control ¹												
5'-primer(s) ²	26(173) 5'-ggg 3'	9(122) 5'-gCT 3'	28(178) 5'-TgC 3'	26(173) 5'-TTA 3'	28(179) 5'-gAC 3'	26(173) 5'-TCT 3'	72(312) 5'-Cgg 3'	9(122) 5'-gTT 3'	9(122) 5'-gTT 3'	30(184) 5'-gAC 3'	21(159) 5'-ACC 3'	38(210) 5'-gCA 3'
	26(173) 5'-ggA 3'	25(169) 5'-TgT 3'	30(184) 5'-gAg 3'		28(179) 5'-gAC 3'			21(159) 5'-ACC 3'				
	26(173) 5'-ggg 3'	26(172) 5'-ATC 3'	30(185) 5'-AAg 3'		28(179) 5'-gAC 3'			23(164) 5'-gCT 3'				
	26(173) 5'-gTg 3'	26(173) 5'-TTA 3'	30(185) 5'-AAA 3'					38(210) 5'-gCg 3'				
		26(173) 5'-TCT 3'										
		30(184) 5'-gAT 3'										
3'-primer(s) ³	57(266) 5'-CAA 3'	57(266) 5'-CAA 3'	86(353) 5'-gCT 3'	55(260) 5'-gCg 3'	57(266) 5'-Cgg 3'	57(266) 5'-CgT 3'	86(353) 5'-gCT 3'	70(304) 5'-CTC 3'	70(304) 5'-CTC 3'	77(327) 5'-ACT 3'	70(304) 5'-CCT 3'	86(353) 5'-gCT 3'
	87(356) 5'-ggT 3'	57(266) 5'-CAT 3'		86(353) 5'-gCT 3'	57(266) 5'-CAg 3'	57(266) 5'-CgT 3'	89(361) 5'-CgT 3'	77(327) 5'-ACg 3'		87(356) 5'-ggT 3'		87(357) 5'-CgT 3'
	87(356) 5'-ggT 3'	86(353) 5'-ACg 3'		86(354) 5'-AgT 3'	57(266) 5'-Cgg 3'					87(356) 5'-ggA 3'		
	89(361) 5'-CCT 3'	86(353) 5'-ACC 3'			87(356) 5'-ggg 3'							
		86(354) 5'-AAT 3'										
Well No.	1	2	3	4	5	6	7	8	9	10	11	12

Well No.	13	14	15
Length of spec.	185	185	135
PCR product			
Length of int.	515	430	430
pos. control ¹			
5'-primer(s) ²	9(122) 5'-gTT 3'	38(209) 5'-CgC 3'	13(134) 5'-ggC 3'
3'-primer(s) ³	57(266) 5'-CAT 3'	86(353) 5'-ACg 3'	45(230) 5'-CCC 3'
	58(270) 5'-TCC 3'		
Well No.	13	14	15

¹The internal positive control primer pairs amplify segments of the human growth hormone gene. The internal positive control bands are 430 or 515 base pairs respectively, well distribution as outlined in the table. Well number 1 contains the longer, 515 bp, internal positive control band. The well distribution of the internal controls can help in orientation of the kit on gel photo, as well as allow for kit identification. In the presence of a specific amplification the intensity of the control band often decreases.



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Lot-specific information

²The nucleotide position matching the specificity-determining 3'-end of the primer is given. Nucleotide numbering as on the www.ebi.ac.uk/imgt/hla web site. The sequence of the 3 terminal nucleotides of the primer is given.

³The nucleotide position matching the specificity-determining 3'-end of the primer is given in the anti-sense direction. Nucleotide 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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Lot-specific information

CELL LINE VALIDATION SHEET																			
DQ low resolution primer set ²																			
				Well															
				1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	
				Production No.	202349101	202349102	202349103	202349104	202349105	202349106	202349107	202351408	202349109	202349110	202349111	202349112	202349113	202349114	202349115
	IHWC cell line ¹			DQB1															
1	9001	SA	*05:01		+	-	-	-	-	-	-	-	-	+	-	-	-	-	
2	9280	LK707	*06:01	*02:02	-	+	+	-	-	-	-	-	-	-	-	-	-	+	
3	9011	E4181324	*06:01		-	+	-	-	-	-	-	-	-	-	-	-	-	+	
4	9275	GU373	*02:01		-	-	+	-	-	-	-	-	-	-	-	-	-	-	
5	9009	KAS011	*05:02		+	-	-	-	-	-	-	-	-	+	-	-	-	-	
6	9353	SM	*03:02	*06:01	-	+	-	-	+	-	+	-	-	-	-	+	-	+	
7	9020	QBL	*02:01		-	-	+	-	-	-	-	-	-	-	-	-	-	-	
8	9025	DEU	*03:01		-	-	-	+	-	-	+	-	-	-	+	-	-	-	
9	9026	YAR	*03:02		-	-	-	-	+	-	+	-	-	-	+	-	-	-	
10	9107	LKT3	*04:01		-	-	-	-	-	-	-	+	+	-	-	-	-	-	
11	9051	PITOUT	*02:02		-	-	+	-	-	-	-	-	-	-	-	-	-	-	
12	9052	DBB	*03:03		-	-	-	-	-	+	+	-	-	-	-	+	-	-	
13	9004	JESTHOM	*05:01		+	-	-	-	-	-	-	-	-	+	-	-	-	-	
14	9071	OLGA	*04:02		-	-	-	-	-	-	-	+	+	-	-	-	-	-	
15	9075	DKB	*03:03		-	-	-	-	-	+	+	-	-	-	+	-	-	-	
16	9037	SWEIG007	*03:01		-	-	-	+	-	-	+	-	-	-	+	-	-	-	
17	9282	CTM3953540	*02:01	*06:03	-	+	+	-	-	-	-	-	-	-	-	-	+	-	
18	9257	32367	*06:02	*02:02	-	+	+	-	-	-	-	-	-	-	-	+	+	-	
19	9038	BM16	*03:01		-	-	-	+	-	-	+	-	-	-	-	+	-	-	
20	9059	SLE005	*06:04		-	+	-	-	-	-	-	-	-	-	-	-	-	-	
21	9064	AMALA	*03:01		-	-	-	+	-	-	+	-	-	-	+	-	-	-	
22	9056	KOSE	*05:03	*06:04	+	+	-	-	-	-	-	-	-	+	-	-	-	-	
23	9124	IHL	*05:03	*06:01	+	+	-	-	-	-	-	-	-	+	-	-	-	+	
24	9035	JBUSH	*03:01		-	-	-	+	-	-	+	-	-	-	+	-	-	-	
25	9049	IBW9	*02:02		-	-	+	-	-	-	-	-	-	-	-	-	-	-	
26	9285	WT49	*02:01		-	-	+	-	-	-	-	-	-	-	-	-	-	-	
27	9191	CH1007	*04:01	*05:01	+	-	-	-	-	-	-	+	+	+	-	-	-	-	
28	9320	BEL5GB	*02:02	*03:01	-	-	+	+	-	-	+	-	-	-	+	-	-	-	
29	9050	MOU	*02:02		-	-	+	-	-	-	-	-	-	-	-	-	-	-	
30	9021	RSH	*04:02		-	-	-	-	-	-	-	+	+	-	-	-	-	-	
31	9019	DUCAF	*02:01		-	-	+	-	-	-	-	-	-	-	-	-	-	-	
32	9297	HAG	*03:01		-	-	-	+	-	-	+	-	-	-	+	-	-	-	
33	9098	MT14B	*03:02		-	-	-	-	+	-	+	-	-	-	+	-	-	-	
34	9104	DHIF	*03:01		-	-	-	+	-	-	+	-	-	-	-	-	-	-	
35	9302	SSTO	*03:05		-	-	-	-	+	-	+	-	-	+	-	-	-	-	
36	9024	KT17	*03:02		-	-	-	-	+	-	+	-	-	-	+	-	-	-	
37	9065	HHKB	*06:03		-	+	-	-	-	-	-	-	-	-	-	-	+	-	
38	9099	LZL	*03:01		-	-	-	+	-	-	+	-	-	-	+	-	-	-	
39	9315	CML	*02:01	*03:01	-	-	+	+	-	-	+	-	-	-	+	-	-	-	
40	9134	WHONP199	*02:02	*03:03	-	-	+	-	-	+	+	-	-	-	+	-	-	-	
41	9055	H0301	*06:09		-	+	-	-	-	-	-	-	-	-	-	-	-	-	
42	9066	TAB089	*06:01		-	+	-	-	-	-	-	-	-	-	-	-	-	+	
43	9076	T7526	*03:03		-	-	-	-	-	+	+	-	-	-	+	-	-	-	
44	9057	TEM	*05:03		+	-	-	-	-	-	-	-	-	+	-	-	-	-	
45	9239	SHJO	*02:02		-	-	+	-	-	-	-	-	-	-	-	-	-	-	
46	9013	SCHU	*06:02		-	+	-	-	-	-	-	-	-	-	-	+	+	-	
47	9045	TUBO	*03:01		-	-	-	+	-	-	+	-	-	-	+	-	-	-	
48	9303	TER-ND	*05:01		+	-	-	-	-	-	-	-	-	+	-	-	-	-	



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¹The provided cell line HLA specificities are retrieved from the <http://www.ihwg.org/hla> web site. The specificity of an individual cell line may thus be subject to change.

²The specificity of each primer solution in the kit has been tested against 48 well characterized cell line DNAs and where applicable, additional cell line DNAs.

In primer solutions 1 to 3 and 5 one, two or three 5'-primers were not possible to be tested, and in primer solutions 1, 2, 4, 5, 12 and 13 one or more of the 3'-primers were not possible to be tested.

The specificities of the primers in primer solution 2 was tested by separately adding one additional 5'-primer, and one additional 3'-primer accordingly. In addition, one 3'-primer in primer solutions 6 and 10 was tested by separately adding one 5'-primer.



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Lot-specific information

ADDRESSES:

Manufacturer:

CareDx AB, Franzengatan 5, SE-112 51 Stockholm, Sweden.

Tel: +46-8-508 939 00

Fax: +46-8-717 88 18

E-mail: orders-se@caredx.com

Web page: www.caredx.com

CareDx Lab Solutions Inc., 901 S. Bolmar St., Suite R, West Chester, PA 19382

Tel: 1-877-653-78171

Fax: 610-344-7989

E-mail: orders-us@caredx.com

Web page: www.caredx.com

For information on CareDx distributors worldwide, contact **CareDx AB**.

