

101.516-24/04 – including *Taq* polymerase
101.516-24u/04u – without *Taq* polymerase

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

Lot No.: 3S0

Lot-specific information
***Olerup SSP[®]* HLA-B*15**

Product number:	101.516-24/04 – including <i>Taq</i> pol. 101.516-24u/04u – without <i>Taq</i> pol.
Lot number:	3S0
Expiry date:	2027-07-01
Number of tests:	24 tests – Product No. 101.516-24/24u 4 tests – Product No. 101.516-04/04u
Number of wells per test:	95+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. 3S0.

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

CHANGES COMPARED TO THE PREVIOUS OLERUP SSP[®]
HLA-B*15 LOT (9N3)

- The product documentation has been updated for new alleles of IMGT 3.52.0.
- The kit resolution focuses on common and well documented (CWD) alleles¹.

¹As described in section Uniquely Identified Alleles.

The HLA-B*15 specificity and interpretation tables have been updated for the HLA-B alleles described since the previous *Olerup SSP[®]* HLA-B*15 lot was made (**Lot No. 9N3**).

The HLA-B*15 primer set is unchanged compared to the previous *Olerup SSP[®]* HLA-B*15 (**Lot No. 9N3**).

¹S. J. Mack, P. Cano, J. A. Hollenbach et al.
 Common and well-documented HLA alleles: 2012 update to
 the CWD catalogue. *Tissue Antigens*, 2013, 81, 194–203



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Well **96** 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

HLA-B*15 SSP subtyping

CONTENT

The primer set contains 5'- and 3'-primers for identifying the B*15:01 to B*15:665 alleles.

PLATE LAYOUT

Each test consists of 96 PCR reactions in a 96 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
49	50	51	52	53	54	55	56
57	58	59	60	61	62	63	64
65	66	67	68	69	70	71	72
73	74	75	76	77	78	79	80
81	82	83	84	85	86	87	88
89	90	91	92	93	94	95	NC

The 96 well cut PCR plate is marked with ‘HLA-B*15’ in silver/gray ink.

Well No. 1 is marked with the Lot No. ‘3S0’.

Wells 1 to 95 – HLA-B*15 high resolution primers.

Well 96 – 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.

INTERPRETATION

Due to the sharing of sequence motifs between HLA-B alleles, non-HLA-B*15 alleles will be amplified by some primer mixes. For further details see Specificity Table.

UNIQUELY IDENTIFIED ALLELES

All the HLA-B*15 alleles, i.e. **B*15:01 to B*15:665**, recognized by the HLA Nomenclature Committee in April 2023^{1,2} will be amplified by the primers in the HLA-B*15 subtyping kit.

The HLA-B*15 kit enables separation of the confirmed HLA-B*15 alleles as listed in the IMGT/HLA database 3.25.0. An HLA allele is listed as confirmed by IMGT/HLA if it has been sequenced by more than a single laboratory or from multiple sources. Current allele confirmation status for HLA-B*15 alleles is listed below.

The HLA-B*15 kit also enables identification of many null and alternatively expressed alleles.



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The following HLA-B*15 alleles can be distinguished by the different sizes of the HLA-specific PCR product:

Alleles	Primer mix	Alleles	Primer mix
B*15:81, 15:430	82	B*15:166, 15:193	86
B*15:82, 15:256	36	B*15:169, 15:225	87
B*15:92, 15:104, 15:591	52	B*15:202, 15:239, 15:477	90
B*15:120, 15:258N	93	B*15:206, 15:343	50
B*15:142, 15:147, 15:429	78	B*15:208, 15:230	70
B*15:146, 15:441	75	B*15:301, 15:302N	64

The HLA-B*15 primer set cannot separate the following alleles. These alleles can be distinguished by the HLA-B low resolution kit and/or the respective high resolution kits:

Alleles

B*15:180, 46:71, 46:89
 B*15:515, 35:43:01, 35:135, 35:185, 51:220

¹HLA-B 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>.



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ALLELE CONFIRMATION STATUS

Allele	Status ¹	Allele	Status ¹	Allele	Status ¹	Allele	Status ¹
B*15:01:01:01	Confirmed	B*15:02:09	Unconfirmed	B*15:18:01:01	Confirmed	B*15:54	Confirmed
B*15:01:01:02N	Confirmed	B*15:02:10	Unconfirmed	B*15:18:01:02	Unconfirmed	B*15:55	Confirmed
B*15:01:01:03	Unconfirmed	B*15:02:11	Unconfirmed	B*15:18:02	Confirmed	B*15:56	Confirmed
B*15:01:01:04	Confirmed	B*15:03:01:01	Confirmed	B*15:18:03	Confirmed	B*15:57	Unconfirmed
B*15:01:01:05	Confirmed	B*15:03:01:02	Confirmed	B*15:18:04	Confirmed	B*15:58	Confirmed
B*15:01:02	Confirmed	B*15:03:02	Confirmed	B*15:18:05	Unconfirmed	B*15:60	Unconfirmed
B*15:01:03	Unconfirmed	B*15:03:03	Confirmed	B*15:18:06	Unconfirmed	B*15:61	Confirmed
B*15:01:04	Unconfirmed	B*15:03:04	Confirmed	B*15:19	Confirmed	B*15:62	Confirmed
B*15:01:06	Confirmed	B*15:03:05	Confirmed	B*15:20	Confirmed	B*15:63	Confirmed
B*15:01:07	Unconfirmed	B*15:03:06	Unconfirmed	B*15:21	Confirmed	B*15:64:01	Confirmed
B*15:01:08	Unconfirmed	B*15:04:01:01	Confirmed	B*15:23	Confirmed	B*15:64:02	Confirmed
B*15:01:09	Confirmed	B*15:04:01:02	Unconfirmed	B*15:24:01	Confirmed	B*15:65	Unconfirmed
B*15:01:10	Confirmed	B*15:04:02	Confirmed	B*15:24:02	Unconfirmed	B*15:66	Unconfirmed
B*15:01:11	Confirmed	B*15:04:03	Unconfirmed	B*15:25:01	Confirmed	B*15:67	Confirmed
B*15:01:12	Unconfirmed	B*15:04:04	Confirmed	B*15:25:02	Confirmed	B*15:68	Confirmed
B*15:01:13	Confirmed	B*15:05:01	Confirmed	B*15:25:03	Unconfirmed	B*15:69	Confirmed
B*15:01:14	Confirmed	B*15:05:02	Confirmed	B*15:26N	Confirmed	B*15:70	Confirmed
B*15:01:15	Confirmed	B*15:06	Confirmed	B*15:27:01	Confirmed	B*15:71	Confirmed
B*15:01:16	Confirmed	B*15:07:01	Confirmed	B*15:27:02	Unconfirmed	B*15:72	Unconfirmed
B*15:01:17	Unconfirmed	B*15:07:02	Unconfirmed	B*15:27:03	Unconfirmed	B*15:73	Confirmed
B*15:01:18	Unconfirmed	B*15:07:03	Confirmed	B*15:28	Unconfirmed	B*15:74	Confirmed
B*15:01:19	Unconfirmed	B*15:08:01	Confirmed	B*15:29	Confirmed	B*15:75	Confirmed
B*15:01:20	Unconfirmed	B*15:08:02	Unconfirmed	B*15:30	Confirmed	B*15:76	Unconfirmed
B*15:01:21	Unconfirmed	B*15:09	Confirmed	B*15:31	Confirmed	B*15:77	Unconfirmed
B*15:01:22	Unconfirmed	B*15:10:01	Confirmed	B*15:32:01	Confirmed	B*15:78:01	Confirmed
B*15:01:23	Unconfirmed	B*15:10:02	Confirmed	B*15:32:02	Unconfirmed	B*15:78:02	Unconfirmed
B*15:01:24	Unconfirmed	B*15:10:03	Unconfirmed	B*15:33	Confirmed	B*15:78:03	Unconfirmed
B*15:01:25	Unconfirmed	B*15:10:04	Unconfirmed	B*15:34	Confirmed	B*15:79N	Unconfirmed
B*15:01:26	Unconfirmed	B*15:11:01	Confirmed	B*15:35	Confirmed	B*15:80	Unconfirmed
B*15:01:27	Unconfirmed	B*15:11:02	Confirmed	B*15:36	Unconfirmed	B*15:81	Unconfirmed
B*15:01:28	Unconfirmed	B*15:11:03	Unconfirmed	B*15:37	Confirmed	B*15:82	Unconfirmed
B*15:01:29	Unconfirmed	B*15:11:04	Unconfirmed	B*15:38:01	Confirmed	B*15:83	Unconfirmed
B*15:01:30	Unconfirmed	B*15:11:05	Unconfirmed	B*15:38:02	Confirmed	B*15:84	Unconfirmed
B*15:01:31	Confirmed	B*15:11:06	Unconfirmed	B*15:39:01	Confirmed	B*15:85	Unconfirmed
B*15:01:32	Unconfirmed	B*15:11:07	Unconfirmed	B*15:39:02	Unconfirmed	B*15:86	Confirmed
B*15:01:33	Unconfirmed	B*15:12	Confirmed	B*15:40:01	Confirmed	B*15:87	Unconfirmed
B*15:01:34	Unconfirmed	B*15:13:01	Confirmed	B*15:40:02	Unconfirmed	B*15:88	Unconfirmed
B*15:01:35	Unconfirmed	B*15:13:02	Unconfirmed	B*15:42	Confirmed	B*15:89	Confirmed
B*15:01:36	Confirmed	B*15:14	Confirmed	B*15:43	Unconfirmed	B*15:90	Unconfirmed
B*15:01:37	Unconfirmed	B*15:15	Confirmed	B*15:44	Unconfirmed	B*15:91	Unconfirmed
B*15:01:38	Unconfirmed	B*15:16:01:01	Confirmed	B*15:45	Confirmed	B*15:92	Unconfirmed
B*15:01:39	Unconfirmed	B*15:16:01:02	Confirmed	B*15:46	Confirmed	B*15:93	Unconfirmed
B*15:02:01	Confirmed	B*15:16:02	Confirmed	B*15:47:01	Confirmed	B*15:94N	Unconfirmed
B*15:02:02	Unconfirmed	B*15:16:03	Unconfirmed	B*15:47:02	Confirmed	B*15:95	Unconfirmed
B*15:02:03	Unconfirmed	B*15:17:01:01	Confirmed	B*15:48	Confirmed	B*15:96	Unconfirmed
B*15:02:04	Confirmed	B*15:17:01:02	Unconfirmed	B*15:49	Unconfirmed	B*15:97	Unconfirmed
B*15:02:05	Unconfirmed	B*15:17:02	Confirmed	B*15:50	Confirmed	B*15:98	Unconfirmed
B*15:02:06	Unconfirmed	B*15:17:03	Unconfirmed	B*15:51	Unconfirmed	B*15:99	Unconfirmed
B*15:02:07	Unconfirmed	B*15:17:04	Unconfirmed	B*15:52	Confirmed	B*15:101	Unconfirmed
B*15:02:08	Unconfirmed	B*15:17:05	Unconfirmed	B*15:53	Confirmed	B*15:102	Unconfirmed



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B*15:103	Unconfirmed	B*15:154	Confirmed	B*15:202	Unconfirmed	B*15:251	Unconfirmed
B*15:104	Unconfirmed	B*15:155	Confirmed	B*15:203	Unconfirmed	B*15:252	Unconfirmed
B*15:105	Unconfirmed	B*15:156	Unconfirmed	B*15:204	Confirmed	B*15:253	Unconfirmed
B*15:106	Confirmed	B*15:157	Confirmed	B*15:205	Unconfirmed	B*15:254	Unconfirmed
B*15:107	Unconfirmed	B*15:158	Confirmed	B*15:206	Confirmed	B*15:255	Unconfirmed
B*15:108	Confirmed	B*15:159	Confirmed	B*15:207	Confirmed	B*15:256	Unconfirmed
B*15:109	Unconfirmed	B*15:160	Confirmed	B*15:208	Confirmed	B*15:257	Unconfirmed
B*15:110	Unconfirmed	B*15:161	Confirmed	B*15:209N	Unconfirmed	B*15:258N	Unconfirmed
B*15:111N	Unconfirmed	B*15:162	Confirmed	B*15:210	Unconfirmed	B*15:259	Unconfirmed
B*15:112	Unconfirmed	B*15:163	Unconfirmed	B*15:211	Confirmed	B*15:260	Unconfirmed
B*15:113	Confirmed	B*15:164	Unconfirmed	B*15:212	Unconfirmed	B*15:261	Unconfirmed
B*15:114	Unconfirmed	B*15:165	Unconfirmed	B*15:213	Unconfirmed	B*15:262N	Unconfirmed
B*15:115	Unconfirmed	B*15:166	Unconfirmed	B*15:214	Unconfirmed	B*15:263	Confirmed
B*15:116	Unconfirmed	B*15:167	Confirmed	B*15:215	Unconfirmed	B*15:264	Unconfirmed
B*15:117	Unconfirmed	B*15:168	Unconfirmed	B*15:216	Unconfirmed	B*15:265	Unconfirmed
B*15:118	Unconfirmed	B*15:169	Unconfirmed	B*15:217	Unconfirmed	B*15:266	Confirmed
B*15:119	Unconfirmed	B*15:170	Confirmed	B*15:218Q	Unconfirmed	B*15:267	Unconfirmed
B*15:120	Unconfirmed	B*15:171	Confirmed	B*15:219	Unconfirmed	B*15:268	Confirmed
B*15:121	Unconfirmed	B*15:172	Confirmed	B*15:220	Confirmed	B*15:269	Unconfirmed
B*15:122	Unconfirmed	B*15:173	Confirmed	B*15:221	Unconfirmed	B*15:270	Unconfirmed
B*15:123	Unconfirmed	B*15:174	Unconfirmed	B*15:222	Unconfirmed	B*15:271	Confirmed
B*15:124	Unconfirmed	B*15:175	Unconfirmed	B*15:223	Unconfirmed	B*15:272N	Confirmed
B*15:125	Confirmed	B*15:176	Confirmed	B*15:224	Unconfirmed	B*15:273	Unconfirmed
B*15:126	Unconfirmed	B*15:177	Unconfirmed	B*15:225	Unconfirmed	B*15:274	Unconfirmed
B*15:127	Unconfirmed	B*15:178	Unconfirmed	B*15:226N	Unconfirmed	B*15:275:01	Unconfirmed
B*15:128	Unconfirmed	B*15:179:01	Unconfirmed	B*15:227	Unconfirmed	B*15:275:02	Unconfirmed
B*15:129	Unconfirmed	B*15:179:02	Unconfirmed	B*15:228	Confirmed	B*15:276	Unconfirmed
B*15:131	Unconfirmed	B*15:180	Confirmed	B*15:229	Confirmed	B*15:277	Confirmed
B*15:132	Confirmed	B*15:181N	Confirmed	B*15:230	Unconfirmed	B*15:278	Confirmed
B*15:133	Unconfirmed	B*15:182N	Unconfirmed	B*15:231	Unconfirmed	B*15:279	Unconfirmed
B*15:134	Unconfirmed	B*15:183	Confirmed	B*15:232	Confirmed	B*15:280	Unconfirmed
B*15:135	Confirmed	B*15:184	Unconfirmed	B*15:233	Unconfirmed	B*15:281	Unconfirmed
B*15:136	Unconfirmed	B*15:185	Unconfirmed	B*15:234	Unconfirmed	B*15:282	Confirmed
B*15:137	Unconfirmed	B*15:186	Unconfirmed	B*15:235	Confirmed	B*15:283	Unconfirmed
B*15:138	Unconfirmed	B*15:187	Unconfirmed	B*15:236	Confirmed	B*15:284	Unconfirmed
B*15:139	Unconfirmed	B*15:188	Unconfirmed	B*15:237	Confirmed	B*15:285	Confirmed
B*15:140	Unconfirmed	B*15:189	Confirmed	B*15:238	Unconfirmed	B*15:286	Unconfirmed
B*15:141	Confirmed	B*15:190N	Confirmed	B*15:239	Confirmed	B*15:287	Unconfirmed
B*15:142	Confirmed	B*15:191	Unconfirmed	B*15:240	Confirmed	B*15:288	Unconfirmed
B*15:143	Unconfirmed	B*15:192	Unconfirmed	B*15:241	Confirmed	B*15:289	Unconfirmed
B*15:144	Unconfirmed	B*15:193	Unconfirmed	B*15:242:01	Unconfirmed	B*15:290	Confirmed
B*15:145	Unconfirmed	B*15:194	Unconfirmed	B*15:242:02	Unconfirmed	B*15:291	Unconfirmed
B*15:146	Confirmed	B*15:195	Unconfirmed	B*15:243	Unconfirmed	B*15:292	Unconfirmed
B*15:147	Confirmed	B*15:196	Unconfirmed	B*15:244	Unconfirmed	B*15:293	Unconfirmed
B*15:148	Unconfirmed	B*15:197:01	Unconfirmed	B*15:245Q	Unconfirmed	B*15:294N	Unconfirmed
B*15:149N	Unconfirmed	B*15:197:02	Unconfirmed	B*15:246N	Unconfirmed	B*15:295	Unconfirmed
B*15:150	Confirmed	B*15:198	Unconfirmed	B*15:247	Unconfirmed	B*15:296	Unconfirmed
B*15:151	Confirmed	B*15:199	Unconfirmed	B*15:248	Confirmed	B*15:297	Unconfirmed
B*15:152	Unconfirmed	B*15:200	Unconfirmed	B*15:249	Unconfirmed	B*15:298	Unconfirmed
B*15:153	Confirmed	B*15:201	Unconfirmed	B*15:250	Unconfirmed	B*15:299	Unconfirmed



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B*15:300	Confirmed	B*15:330	Unconfirmed	B*15:360	Unconfirmed	B*15:390	Unconfirmed
B*15:301	Confirmed	B*15:331	Unconfirmed	B*15:361	Unconfirmed	B*15:391	Unconfirmed
B*15:302N	Unconfirmed	B*15:332	Confirmed	B*15:362	Unconfirmed	B*15:392	Unconfirmed
B*15:303	Unconfirmed	B*15:333	Unconfirmed	B*15:363:01	Unconfirmed	B*15:393	Unconfirmed
B*15:304N	Unconfirmed	B*15:334	Unconfirmed	B*15:364	Unconfirmed	B*15:394	Unconfirmed
B*15:305	Unconfirmed	B*15:335	Confirmed	B*15:365	Unconfirmed	B*15:395	Unconfirmed
B*15:306	Unconfirmed	B*15:336	Unconfirmed	B*15:366	Unconfirmed	B*15:396	Unconfirmed
B*15:307	Unconfirmed	B*15:337	Unconfirmed	B*15:367	Unconfirmed	B*15:397	Unconfirmed
B*15:308	Unconfirmed	B*15:338	Unconfirmed	B*15:368	Unconfirmed	B*15:398	Unconfirmed
B*15:309	Unconfirmed	B*15:339	Unconfirmed	B*15:369	Confirmed		
B*15:310	Unconfirmed	B*15:340	Unconfirmed	B*15:370	Unconfirmed		
B*15:311	Unconfirmed	B*15:341	Unconfirmed	B*15:371	Unconfirmed		
B*15:312	Unconfirmed	B*15:342	Confirmed	B*15:372	Unconfirmed		
B*15:313	Unconfirmed	B*15:343	Unconfirmed	B*15:373	Unconfirmed		
B*15:314	Unconfirmed	B*15:344	Unconfirmed	B*15:374	Unconfirmed		
B*15:315	Confirmed	B*15:345	Unconfirmed	B*15:375N	Unconfirmed		
B*15:316	Unconfirmed	B*15:346	Unconfirmed	B*15:376	Unconfirmed		
B*15:317	Unconfirmed	B*15:347	Confirmed	B*15:377Q	Unconfirmed		
B*15:318	Unconfirmed	B*15:348	Unconfirmed	B*15:378	Confirmed		
B*15:319	Confirmed	B*15:349	Unconfirmed	B*15:379	Confirmed		
B*15:320	Unconfirmed	B*15:350	Unconfirmed	B*15:380N	Confirmed		
B*15:321Q	Unconfirmed	B*15:351	Unconfirmed	B*15:381	Unconfirmed		
B*15:322	Unconfirmed	B*15:352	Unconfirmed	B*15:382	Unconfirmed		
B*15:323	Unconfirmed	B*15:353	Unconfirmed	B*15:383	Unconfirmed		
B*15:324	Unconfirmed	B*15:354	Unconfirmed	B*15:384	Unconfirmed		
B*15:325	Unconfirmed	B*15:355	Unconfirmed	B*15:385	Unconfirmed		
B*15:326	Unconfirmed	B*15:356	Unconfirmed	B*15:386	Unconfirmed		
B*15:327	Unconfirmed	B*15:357	Unconfirmed	B*15:387	Unconfirmed		
B*15:328	Unconfirmed	B*15:358	Unconfirmed	B*15:388	Unconfirmed		
B*15:329	Unconfirmed	B*15:359	Unconfirmed	B*15:389	Unconfirmed		

¹Allele status “confirmed” or “unconfirmed” as listed on the IMGT/HLA web page 2016-July-14, release 3.25.0, www.ebi.ac.uk/imgt/hla.

RESOLUTION IN HOMO- AND HETEROZYGOTES

Results file with resolution in HLA-B*15 homo- and heterozygotes is available upon request.



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SPECIFICITY TABLE

HLA-B*15 SSP subtyping

Specificities and sizes of the PCR products of the 95+1 primer mixes used for HLA-B*15 SSP subtyping

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	Amplified HLA-B*15 alleles ³	Other amplified Class I alleles
1 ⁹	400 bp	800 bp	*15:01:01:01-15:01:04, 15:01:06-15:03:11, 15:03:13-15:07:03, 15:12:01-15:15:01:02, 15:19-15:20, 15:24:01-15:28, 15:30:01:01-15:36, 15:38:01-15:40:02, 15:42-15:43, 15:45-15:50, 15:53-15:55, 15:57-15:58, 15:60-15:65, 15:69-15:71, 15:73:01:01-15:73:01:02, 15:75, 15:77-15:79N, 15:81-15:89:02, 15:91-15:92, 15:94N, 15:96, 15:98, 15:102-15:107, 15:109-15:113, 15:116-15:118, 15:120-15:123, 15:125:01-15:129, 15:131-15:132, 15:135-15:138, 15:140-15:142, 15:144-15:147, 15:150-15:151, 15:154-15:160:01:02, 15:163-15:167, 15:169-15:175, 15:178-15:179:02, 15:181N-15:185, 15:187-15:188, 15:190N, 15:192-15:195, 15:199, 15:201-15:207, 15:210-15:214, 15:217-15:220:04, 15:223-15:225, 15:227-15:228, 15:231-15:235, 15:237, 15:239-15:242:02, 15:244-15:251:03, 15:253, 15:256-15:262:02N, 15:264-15:267, 15:269-15:272N, 15:274, 15:276-15:283, 15:285-15:289, 15:291, 15:295-15:302N, 15:304N, 15:308-15:310, 15:315-15:322, 15:324-15:328, 15:330-15:334, 15:336, 15:339-15:350, 15:352-15:353, 15:355, 15:357-15:360, 15:363:01-15:366, 15:368-15:375N, 15:377Q-15:379, 15:381, 15:383-15:387, 15:389-15:393, 15:395, 15:397-15:402, 15:405-15:407, 15:409-15:410, 15:412-15:413, 15:415-15:419, 15:421-15:422, 15:425, 15:428-15:432, 15:437-15:438, 15:440-15:444, 15:448-15:450, 15:452-15:454N, 15:456-15:461, 15:463N-15:478, 15:480-15:485, 15:488-15:497, 15:499, 15:501-15:502, 15:505, 15:507, 15:510-15:512, 15:514, 15:517-15:522, 15:524-15:525, 15:527-15:529, 15:531, 15:533:01-15:545, 15:547, 15:549N, 15:551-15:552, 15:554, 15:556-15:562N, 15:564-15:569, 15:571-15:572, 15:574, 15:576-15:593, 15:595-15:601,	*40:12:01-40:12:02, 40:26, 40:40, 40:516, 44:02:68, C*12:301, C*16:85



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2⁶	290 bp	1070 bp	*15:01:01:01-15:01:03, 15:01:06-15:01:53, 15:01:55-15:02:13, 15:03:03, 15:04:01:01-15:08:02, 15:10:02, 15:11:01-15:11:03, 15:11:05-15:15:01:02, 15:17:01:01-15:17:07, 15:19-15:21:01:02, 15:24:01-15:28, 15:30:01:01-15:36, 15:38:01-15:40:02, 15:42-15:46, 15:48, 15:50, 15:55-15:58, 15:60, 15:63, 15:65-15:66, 15:70-15:71, 15:73:01:01-15:73:01:02, 15:75-15:79N, 15:81-15:89:02, 15:92, 15:94N, 15:96-15:97, 15:101-15:102, 15:104-15:107, 15:109-15:113, 15:116-15:118, 15:120-15:122, 15:125:01-15:126, 15:128-15:129, 15:135-15:150, 15:152, 15:154-15:155, 15:157, 15:159-15:160:01:02, 15:162-15:172, 15:174-15:175, 15:177-15:179:02, 15:181N-15:185, 15:187-15:196, 15:199, 15:201-15:209N, 15:211, 15:213-15:219, 15:223-15:225, 15:227-15:228, 15:230-15:234, 15:236-15:237, 15:239-15:241, 15:244-15:251:02, 15:256-15:262:02N, 15:264-15:265, 15:267-15:273, 15:276-15:280, 15:283-15:289, 15:291, 15:295-15:305, 15:308-15:310, 15:315-15:322, 15:324-15:328, 15:330-15:334, 15:339-15:346, 15:348-15:350, 15:352-15:353, 15:355-15:361, 15:363:01-15:368, 15:370-15:375N, 15:377Q-15:378, 15:381, 15:383-15:387, 15:390-15:396, 15:398-15:407, 15:409-15:413, 15:415-15:417, 15:419-15:425, 15:428-15:432, 15:437-15:438, 15:440-15:443, 15:446, 15:449-15:450, 15:452-15:454N, 15:456-15:474, 15:476-15:486, 15:488-15:497, 15:499-15:501, 15:504:02-15:505, 15:507-15:508, 15:510, 15:512-15:514, 15:516, 15:518-15:525, 15:528N-15:529, 15:531-15:538, 15:540N-15:545, 15:547, 15:549N, 15:551-15:552, 15:554-15:558, 15:561-15:562N, 15:564-15:569, 15:572, 15:574-15:581, 15:583-15:587, 15:589-15:601, 15:603-15:606, 15:608-15:609, 15:615, 15:618-15:619, 15:621-15:627, 15:629-15:633, 15:635-15:642, 15:644-15:649, 15:651-15:653, 15:655-15:665
			*46:01:01:01-46:70, 46:72-46:76, 46:78-46:88, 46:90-46:98N, 51:367, C*01:73, C*03:02:23, C*05:01:28, C*06:02:93, C*07:02:30, C*08:16:02, C*14:02:27



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3 ⁴	120 bp	1070 bp	*15:01:01:01-15:01:04, 15:01:06-15:01:16, 15:01:17 ^w , 15:01:18-15:01:30, 15:01:32-15:01:80, 15:04:01:01-15:04:04, 15:07:01:01-15:08:02, 15:11:01-15:12:03, 15:14-15:15:01:02, 15:19, 15:24:01-15:24:02, 15:26N-15:28, 15:30:01:01-15:30:01:03, 15:32:01-15:32:02, 15:34:01-15:35:02, 15:38:01-15:38:02, 15:43, 15:45-15:46, 15:50, 15:53-15:54, 15:56-15:58, 15:60, 15:63, 15:66, 15:68, 15:70-15:71, 15:73:01:01-15:73:01:02, 15:75-15:77, 15:79N, 15:81-15:82, 15:85, 15:87, 15:90, 15:92, 15:94N, 15:96-15:97, 15:101-15:102, 15:104-15:105, 15:109-15:111N, 15:113, 15:117-15:118, 15:120, 15:122, 15:125:01-15:126, 15:128-15:129, 15:135, 15:137, 15:140, 15:142-15:149N, 15:152, 15:154, 15:157, 15:159-15:160:01:02, 15:163-15:167, 15:169, 15:171-15:172, 15:174-15:175, 15:178, 15:180-15:184, 15:187, 15:189:01-15:193, 15:201-15:203, 15:205-15:207, 15:209N, 15:211-15:212, 15:215, 15:217, 15:225, 15:227-15:228, 15:231-15:234, 15:236, 15:239, 15:241, 15:244-15:247, 15:251:01-15:251:03, 15:256-15:262:02N, 15:264, 15:267, 15:269-15:270, 15:272N, 15:276-15:280, 15:284-15:286, 15:296, 15:298-15:299, 15:303-15:305, 15:309, 15:315-15:318, 15:320-15:321Q, 15:324, 15:326-15:327, 15:331, 15:333-15:334, 15:336, 15:340, 15:342-15:344, 15:346-15:347, 15:349:01-15:350, 15:353, 15:355, 15:359, 15:363:01-15:363:02, 15:365, 15:367-15:368, 15:370-15:373, 15:375N, 15:377Q, 15:381, 15:385-15:387, 15:390-15:392, 15:398, 15:400N-15:401, 15:405-15:406, 15:409-15:410, 15:412-15:413, 15:415-15:417, 15:419, 15:422, 15:428-15:432, 15:440-15:443, 15:450, 15:452-15:453, 15:456-15:457, 15:459, 15:461, 15:464-15:467, 15:469-15:474, 15:476-15:484, 15:486, 15:488-15:490, 15:492-15:493, 15:496N, 15:498-15:499, 15:501, 15:507, 15:510, 15:512-15:515, 15:520Q-15:522, 15:525, 15:528N-15:529, 15:531, 15:533:01-15:536, 15:538, 15:540N-15:542, 15:545, 15:547, 15:549N, 15:551, 15:554, 15:556, 15:558-15:559, 15:561-15:562N, 15:565, 15:568, 15:572, 15:576, 15:579-15:581, 15:583, 15:585-15:586, 15:590-15:591, 15:593, 15:595, 15:597-15:598,	*07:100, 13:31, 13:41, 13:131, 13:145, 13:164, 18:19, 18:164, 27:25, 27:115, 27:161, 27:245, 35:14:01-35:14:03, 35:43:01, 35:43:03-35:44, 35:62, 35:67, 35:79, 35:86, 35:102, 35:117, 35:135, 35:163, 35:185, 35:213, 35:265, 35:343, 35:390N, 35:413, 35:475-35:476, 35:512, 35:549, 37:66, 39:36, 44:146, 44:275, 44:299, 45:05, 46:01:01:01-46:02, 46:04-46:05, 46:07N, 46:09-46:10, 46:12, 46:14-46:17, 46:20, 46:22-46:24, 46:26-46:32:02, 46:34-46:42, 46:44, 46:46-46:53, 46:55N-46:57, 46:60-46:61, 46:63-46:82, 46:84-46:92, 46:94-46:98N, 50:47, 51:61:01-51:61:02, 51:220, 52:21:01-52:21:02, 52:97, 53:70, 54:06, 54:33, 54:40, 55:21, 56:03, 56:43, 57:147, 57:161, 58:78, A*25:66, A*26:68, A*26:100, A*26:200, A*68:56:01-68:56:02, C*02:56, C*06:20, C*12:50
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4	180 bp 390 bp	800 bp	*15:01:01:02N *15:285, 15:378
5⁶	385 bp	1070 bp	*15:02:01:01-15:02:13, 15:08:01:01-15:11:07, 15:13:01-15:13:03, 15:15:01:01-15:15:01:02, 15:18:01:01-15:18:10, 15:21:01:01-15:21:01:02, 15:23, 15:29:01:01-15:29:01:02, 15:31:01:01-15:31:01:03, 15:37, 15:44, 15:51-15:52, 15:55, 15:64:01-15:64:02, 15:72, 15:76 ^w , 15:80, 15:88-15:90, 15:93, 15:99, 15:108, 15:112, 15:114-15:115, 15:119, 15:121, 15:124, 15:133-15:134, 15:139, 15:143-15:144, 15:148, 15:153, 15:161, 15:170, 15:176, 15:186, 15:189:01-15:189:02, 15:191, 15:194, 15:197:01-15:198, 15:200, 15:209N, 15:213-15:215, 15:221, 15:223, 15:226N, 15:229, 15:238, 15:252, 15:263, 15:265, 15:275:01-15:275:02, 15:283-15:284, 15:290-15:294N, 15:297, 15:301-15:303, 15:305-15:308, 15:311-15:314, 15:319, 15:323, 15:325, 15:328-15:330, 15:335, 15:337-15:338, 15:341, 15:345, 15:350-15:351, 15:354, 15:357-15:358, 15:367, 15:374, 15:378, 15:380N, 15:382, 15:384:01-15:384:02, 15:388, 15:394, 15:399, 15:404, 15:414, 15:418, 15:420, 15:425-15:427, 15:433-15:439, 15:445, 15:447, 15:451, 15:454N-15:455, 15:460, 15:479, 15:486-15:487N, 15:491, 15:495, 15:498, 15:503-15:505, 15:508, 15:509 ^w , 15:513, 15:526, 15:530, 15:537, 15:548, 15:553, 15:557, 15:563, 15:567, 15:569-15:570, 15:573-15:574, 15:592, 15:596N, 15:607, 15:611-15:612, 15:615, 15:620, 15:622-15:623, 15:627, 15:629-15:630, 15:637, 15:643, 15:646, 15:648, 15:659, 15:662, 15:664-15:665
6	525 bp	1070 bp	*15:02:01:01-15:02:13, 15:13:01-15:13:03, 15:21:01:01-15:21:01:02, 15:25:01:01-15:25:04, 15:36, 15:44, 15:62, 15:77, 15:80, 15:85, 15:88-15:89:02, 15:106, 15:112, 15:121, 15:139, 15:144, 15:154, 15:165, 15:170, 15:194, 15:204, 15:213-15:214, 15:223, 15:240, 15:250, 15:265, 15:271, 15:283, 15:289, 15:291, 15:297, 15:301-15:302N, 15:308, 15:319, 15:325, 15:328, 15:330, 15:341, 15:345,
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7	330 bp	1070 bp	*15:03:01:01-15:03:13, 15:09:01- 15:10:08, 15:18:01:01-15:18:10, 15:23, 15:29:01:01-15:29:01:02, 15:37, 15:46- 15:47:02, 15:49, 15:51-15:54, 15:61- 15:62, 15:64:01-15:64:02, 15:69, 15:72, 15:74, 15:80, 15:90-15:91, 15:93, 15:98- 15:99, 15:103, 15:106, 15:108, 15:114- 15:115, 15:119, 15:123-15:124, 15:127, 15:131-15:134, 15:143, 15:151, 15:153, 15:156, 15:158, 15:161, 15:173, 15:176, 15:186, 15:197:01-15:198, 15:200, 15:210, 15:212, 15:220:01:01-15:221, 15:226N, 15:229, 15:235, 15:238, 15:242:01-15:243, 15:252-15:253, 15:255, 15:259, 15:263, 15:266, 15:274- 15:275:02, 15:281-15:282, 15:290, 15:292-15:294N, 15:306-15:307, 15:311- 15:314, 15:323, 15:329, 15:335, 15:337- 15:338, 15:351, 15:354, 15:369, 15:379- 15:380N, 15:382, 15:388-15:389, 15:397, 15:414, 15:426-15:427, 15:433, 15:435, 15:439, 15:444-15:445, 15:447- 15:448, 15:451, 15:455, 15:475, 15:487N, 15:498, 15:502-15:504:02, 15:509, 15:511, 15:517, 15:525-15:527, 15:530, 15:539, 15:548, 15:553, 15:560, 15:563, 15:570-15:571, 15:582, 15:588, 15:607, 15:610-15:612, 15:614:01:01- 15:614:01:02, 15:616Q-15:617, 15:620, 15:628, 15:643, 15:650, 15:654	*27:195, 40:12:01-40:12:02, 40:26, 40:40, 40:516, 44:02:68, C*03:342, C*07:46, C*07:356, C*07:546, C*12:184, C*15:99, C*16:85
8 ⁴	105 bp	1070 bp	*15:04:01:01-15:04:04, 15:16:01:01- 15:16:05, 15:67, 15:95, 15:137, 15:155, 15:222, 15:254, 15:293, 15:310, 15:362, 15:408, 15:499, 15:517, 15:531, 15:550, 15:602, 15:609, 15:613	*13:04, 13:46, 13:72, 13:127, 13:140, 13:160, 35:37, 35:235, 35:265, 35:276, 39:06:01:01-39:06:08, 39:33- 39:34, 39:50, 39:57, 39:62, 39:64, 39:83:01-39:83:03, 39:90, 39:102, 39:109, 39:127, 39:129, 39:132, 39:140-39:141, 39:154, 39:178, 39:186-39:187, 40:86, 40:159, 40:330, 40:537, 41:30, 41:50, 45:02, 45:09, 46:32:01-46:32:02, 49:04:01-49:05:02, 49:50, 50:09, 50:33, 51:13:01- 51:13:02, 51:37, 51:63:01-51:63:02, 51:92:01-51:92:02, 51:97, 51:172,



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9	365 bp	1070 bp	*15:05:01:01-15:05:04, 15:31:01:01-15:31:01:03, 15:48, 15:51 ^w , 15:52, 15:55, 15:84, 15:86, 15:91, 15:107, 15:114, 15:123-15:124, 15:136, 15:151, 15:155, 15:179:01 ^w -15:179:02 ^w , 15:185-15:186, 15:188, 15:218Q ^w , 15:222, 15:224, 15:235, 15:237, 15:275:01-15:275:02, 15:283, 15:332, 15:348, 15:383, 15:393, 15:396, 15:407, 15:415, 15:421, 15:449, 15:485, 15:499, 15:507, 15:518 ^w , 15:539, 15:552, 15:577, 15:578 ^w , 15:587, 15:599, 15:623, 15:628, 15:633, 15:636
10	300 bp	1070 bp	*15:37-15:38:02, 15:185, 15:323, 15:335, 15:364, 15:368, 15:435, 15:610, 15:645, 15:658
	515 bp		*15:06, 15:27:01:01-15:27:04, 15:84, 15:109, 15:112, 15:195, 15:327, 15:344, 15:398, 15:656
11	140 bp	800 bp	*15:07:01:01-15:07:03, 15:45, 15:55, 15:68, 15:126, 15:197:02, 15:207, 15:324, 15:329, 15:331, 15:360, 15:383, 15:405, 15:431, 15:450, 15:661
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12	400 bp	800 bp	*15:08:01:01-15:08:02, 15:29:01:01-15:29:01:02, 15:49, 15:56, 15:191, 15:236, 15:479, 15:508, 15:553, 15:622
			*51:367, C*06:147
13	395 bp	800 bp	*15:09:01-15:10:08, 15:18:01:01-15:18:10, 15:21:01:01-15:21:01:02, 15:23, 15:37, 15:44, 15:51-15:52, 15:66, 15:72, 15:80, 15:90, 15:93, 15:99, 15:108, 15:114-15:115, 15:119, 15:124,



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28	195 bp 240 bp	1070 bp	*15:400N *15:35:01:01-15:35:02, 15:141, 15:290, 15:381, 15:597, 15:641		
29	255 bp	1070 bp	*15:37-15:38:02, 15:185, 15:323, 15:335, 15:364, 15:368, 15:435, 15:610	*07:218, 07:220, 08:118, 14:01:12, 14:05-14:06:01, 14:08:02, 14:13, 14:53, 14:56, 14:96, 18:01:01:01- 18:01:05, 18:01:07-18:01:40, 18:01:42-18:01:49, 18:01:51-18:01:53, 18:03:01:01-18:07:03:02, 18:09, 18:12:01-18:15, 18:17N, 18:19-18:20, 18:23N-18:26, 18:28-18:34, 18:36- 18:44:01, 18:45-18:51, 18:53-18:55, 18:57:01-18:57:02, 18:59-18:100, 18:102-18:104, 18:106Q-18:109, 18:111-18:140, 18:142-18:148, 18:151-18:163, 18:166-18:167, 18:169, 18:172-18:178, 18:180- 18:188, 18:190-18:203, 18:205- 18:209, 18:211-18:228, 18:230- 18:231Q, 35:519, 38:106, 39:32, 39:43, 39:48, 40:51, 40:113, 40:163, 40:174, 51:06:01:01-51:06:01:03, 51:06:03-51:06:04, 51:45, 51:62, 51:64, 51:135, 51:194, 51:293, 51:319, 51:341, 51:378, 56:31, 67:06, C*02:142	
30 ⁶	320 bp 355 bp 405 bp	1070 bp	*15:40:01-15:40:02, 15:47:01-15:47:02, 15:49, 15:52, 15:114, 15:117, 15:124, 15:138, 15:238, 15:241, 15:389, 15:421, 15:519, 15:532, 15:656 *15:75, 15:312 *15:375N	*46:06, 46:25, 46:30	
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33	200 bp 255 bp 340 bp	1070 bp	*15:118 *15:18:03, 15:42, 15:73:01:01- 15:73:01:02, 15:86, 15:224, 15:303, 15:485-15:486, 15:552 *15:192, 15:281, 15:424 *46:11, 46:18
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48	420 bp 1070 bp	*15:01:01:01-15:01:04, 15:01:06- 15:01:16, 15:01:17 ^w , 15:01:18-15:19, 15:21:01:01-15:21:01:02, 15:23- 15:32:02, 15:34:01-15:40:02, 15:43- 15:58, 15:60-15:71, 15:73:01:01-15:77, 15:79N-15:99, 15:101-15:106, 15:108- 15:115, 15:117-15:129, 15:131-15:140, 15:142-15:149N, 15:151-15:187, 15:189:01-15:227, 15:229-15:247, 15:249-15:286, 15:288-15:294N, 15:296- 15:351, 15:353-15:382, 15:384:01- 15:420, 15:422-15:523, 15:525-15:613, 15:615-15:665	*35:43:01, 35:135, 35:185, 46:01:01:01-46:05, 46:07N-46:44, 46:46-46:53, 46:55N-46:57, 46:59- 46:98N, 51:220, 56:03
49	135 bp 1070 bp	*15:57, 15:135, 15:648	*13:39, 13:67
50⁴	85 bp 1070 bp	*15:67, 15:343	*35:110, 44:497, 58:77, A*02:598, A*02:695, A*68:56:01-68:56:02, A*69:05
	185 bp	*15:206	
51	135 bp 1070 bp 175 bp 200 bp	*15:98, 15:109, 15:163 *15:277-15:278 *15:246N	*55:104 *35:122, 40:339, 56:29
52	200 bp 800 bp	*15:104, 15:307, 15:321Q, 15:560, 15:591 *15:92, 15:213	*35:364, 35:379, 35:462, 39:187, 42:31, 51:142, 52:80, 57:116
53	345 bp 210 bp	800 bp *15:96	*35:537, C*03:175, C*03:199, C*03:245
54	250 bp 800 bp 410 bp	*15:187 *15:97	C*12:36
55⁶	175 bp 1070 bp	*15:214	*07:120, 08:301, 14:61, 18:81, 35:250, 37:93, 39:145, 39:158, 40:150, 40:427, 44:311, 48:45, 51:165, 55:108, 57:29
	210 bp 255 bp	*15:321Q *15:102, 15:284	*35:474, 37:71, 39:179, 44:278, 51:30, 58:105, 73:01:01:01-73:03
56⁴	70 bp 1070 bp 260 bp	*15:103 *15:227, 15:630, 15:641	*07:329, 13:151, 18:183, 27:69, 27:192, 37:23, 40:213, 40:485, 42:28, 44:338
57	340 bp 1070 bp 480 bp	*15:111N, 15:300 *15:369	
58	175 bp 1070 bp	*15:113, 15:319, 15:633	*44:500, 52:17



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59 ⁵	230 bp 370 bp 420 bp	800 bp	*15:191, 15:234 *15:198, 15:270 *15:94N, 15:294N	C*06:33 *35:161
60	160 bp	800 bp	*15:20, 15:228, 15:271, 15:305, 15:435, 15:599, 15:658	*27:197, 35:01:01:01-35:01:27, 35:01:29-35:01:46, 35:01:48- 35:30:01:02, 35:32:01-35:32:02, 35:33-35:34, 35:36-35:42:02, 35:45, 35:47-35:56, 35:58-35:59:02, 35:61:01-35:72, 35:74-35:78, 35:80- 35:86, 35:88-35:134N, 35:136-35:184, 35:186-35:208:01, 35:209-35:278, 35:280-35:358, 35:360-35:466, 35:468-35:470, 35:472-35:474, 35:477-35:482, 35:484:01-35:533, 35:535-35:544, 35:546-35:548, 35:550-35:578, 37:80, 48:02:01- 48:02:03, 49:66, 51:01:01:01- 51:01:55, 51:01:57-51:24:05, 51:26- 51:41N, 51:43-51:46, 51:48-51:156, 51:158:01-51:163, 51:165-51:219, 51:221-51:236, 51:238-51:294, 51:296-51:299, 51:301-51:306N, 51:308-51:358, 51:360-51:381, 52:01:01:01-52:01:50, 52:01:52- 52:06:03, 52:08-52:107, 52:109- 52:113, 53:01:01:01-53:01:27, 53:01:29-53:58, 53:60-53:72, 56:05:01-56:05:02, 56:21, 56:36, 58:01:01:01-58:01:21, 58:01:23- 58:01:44, 58:02:01:01-58:02:02, 58:04-58:10N, 58:12-58:19, 58:21- 58:29, 58:31N-58:104, 58:107-58:143, 67:05, 78:01:01:02-78:10, 81:03
61	160 bp 205 bp	1070 bp	*15:175 *15:125:01	*40:331 *13:124, 14:73, 37:85, 40:342, 40:470, 44:22, 44:105, 48:57, 49:24, C*12:301
62 ^{4,5}	105 bp	1070 bp	*15:126	*07:175
63	175 bp 280 bp	1070 bp	*15:127 *15:315	*46:77
64 ⁴	90 bp 135 bp 240 bp 365 bp	800 bp	*15:302N *15:270 *15:301 *15:122	*08:246 C*12:318, C*14:134 C*03:48, C*05:126, C*06:85, C*07:540
65	405 bp 470 bp 520 bp	1070 bp	*15:375N *15:232 *15:129, 15:342, 15:395, 15:629	*46:64
66	435 bp	1070 bp	*15:02:01:01-15:02:13, 15:13:01- 15:13:03, 15:20-15:21:01:02, 15:25:01:01-15:25:04, 15:36, 15:44, 15:77, 15:85, 15:88-15:89:02, 15:118, 15:121, 15:139, 15:144, 15:154, 15:194, 15:204, 15:213-15:214, 15:223, 15:240,	*13:01:01:01-13:01:22, 13:06-13:07N, 13:12:01-13:13:02, 13:17, 13:20- 13:23, 13:25-13:26:02, 13:28-13:29, 13:36, 13:39, 13:43, 13:50-13:52, 13:57, 13:60-13:61, 13:63N, 13:73, 13:77-13:78, 13:80, 13:83,



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67	265 bp 800 bp	*15:53, 15:183, 15:212, 15:336, 15:345, 15:511, 15:559	*27:195, 40:26, 40:40, 40:516, 44:02:68, 46:10, C*02:167, C*03:504, C*04:459, C*05:205, C*06:284, C*07:828, C*16:85, C*07:02:92
	300 bp	*15:132	
68^{5,6}	215 bp 1070 bp 380 bp	*15:134 *15:247	*44:284, C*03:490, C*03:548, C*05:03
69	130 bp 1070 bp	*15:57	
70⁴	90 bp 1070 bp 185 bp	*15:208 *15:138, 15:230, 15:241, 15:297	*51:252 *07:68:01-07:68:03, 07:214, 07:237, 07:262:01-07:262:02, 07:424, 18:35, 18:110, 35:66, 40:77, 40:87:01-40:87:02, 40:121, 40:158, 40:222, 40:237, 40:368, 40:422, 44:150, 46:87, 48:12, 48:14, 48:26, 48:29
71	290 bp 1070 bp 315 bp 400 bp	*15:53, 15:212, 15:336, 15:347, 15:559, 15:588 *15:205 *15:139	*27:195, 40:26, 40:40, 40:516, 44:02:68, C*01:124, C*16:85
72⁴	110 bp 1070 bp 200 bp 295 bp 390 bp	*15:226N *15:380N *15:140 *15:201	*35:216N, 38:165N, 39:87N C*15:199 *35:543, 51:166
73	155 bp 1070 bp	*15:09:01-15:09:02, 15:30:01:01 ^w -15:30:01:03 ^w , 15:42, 15:58 ^w , 15:73:01:01 ^w -15:73:01:02 ^w , 15:150 ^w , 15:229, 15:243, 15:252, 15:303 ^w , 15:324 ^w , 15:329, 15:361, 15:435, 15:453 ^w , 15:460, 15:486 ^w , 15:541 ^w , 15:637 ^w	*07:05:01:01-07:06:05, 07:32, 07:34, 07:40, 07:53, 07:69, 07:78, 07:80, 07:90, 07:97, 07:105, 07:112, 07:123, 07:137-07:138, 07:140, 07:176, 07:182N, 07:201N, 07:206-07:207, 07:210, 07:213-07:214, 07:222, 07:249, 07:258, 07:262:01-07:262:02, 07:264, 07:269-07:270, 07:278, 07:283, 07:287, 07:293, 07:304, 07:317, 07:324, 07:327, 07:333, 07:340, 07:352-07:353, 07:356-07:358, 07:368, 07:378, 07:387, 07:389, 07:395, 07:399, 07:426Q, 07:434, 07:443, 07:474,



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 35:413^w, 38:10, 38:32, 38:98, 38:106,
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74⁴	75 bp	800 bp	*15:145, 15:176, 15:411	*35:237, 35:352, 51:228, 52:28, 53:68, C*03:420
	230 bp		*15:262:01N	
75	280 bp 315 bp	1070 bp	*15:146 *15:367, 15:441	*35:552, 51:48, C*02:83, C*05:224, C*06:327, C*08:24, C*12:323, C*16:90, C*16:180
	390 bp		*15:201	
76	475 bp	1070 bp	*15:02:01:01-15:02:13, 15:06, 15:13:01-15:13:03, 15:17:01:01-15:17:07, 15:21:01:01-15:21:01:02, 15:25:01:01-15:25:04, 15:31:01:01-15:31:01:03, 15:36, 15:44, 15:55, 15:77, 15:88-15:89:02, 15:106, 15:112, 15:121, 15:128, 15:137, 15:139, 15:144, 15:156, 15:162, 15:168, 15:170, 15:177, 15:196, 15:204, 15:208, 15:213-15:214, 15:216, 15:230, 15:233, 15:240, 15:250, 15:265, 15:271, 15:273, 15:283, 15:289, 15:291, 15:297, 15:301-15:302N, 15:308, 15:319, 15:325, 15:328, 15:330, 15:339, 15:341-15:342, 15:345, 15:356-15:358, 15:360, 15:366, 15:374, 15:378, 15:384:01-15:384:02, 15:394-15:396, 15:399, 15:403-15:405, 15:411, 15:418, 15:420, 15:423-15:425, 15:437-15:438, 15:446, 15:454N, 15:458, 15:462-15:463N, 15:468, 15:488, 15:491, 15:495, 15:500, 15:505, 15:516, 15:523, 15:532, 15:537, 15:543-15:544N, 15:546Q, 15:553, 15:555, 15:557, 15:564, 15:566-15:567, 15:569, 15:574-15:575N, 15:578, 15:584N, 15:589, 15:592, 15:594, 15:596N, 15:603, 15:613, 15:615, 15:619, 15:625, 15:627, 15:629-15:630, 15:642, 15:644-15:645, 15:648, 15:655, 15:657, 15:659, 15:664-15:665	
77^{6,8}	185 bp 330 bp	1070 bp	*15:174, 15:655 *15:123, 15:220:01:01-15:220:04, 15:388, 15:475, 15:517, 15:539, 15:560, 15:571, 15:610, 15:650	*27:179, 35:549, 44:312, 51:299
78⁴	110 bp	1070 bp	*15:142, 15:429	*07:29, 07:186, 07:383, 08:01:11, 08:56:01, 08:133, 08:176, 08:180, 13:93, 35:218, 35:256, 37:62, 51:68,



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	385 bp 475 bp		*15:147 *15:232, 15:369	51:176, 52:77, 57:49, C*06:02:93, C*07:02:30 *46:42 *46:64
79	215 bp 365 bp 420 bp	800 bp	*15:182N *15:282 *15:148, 15:330, 15:636	C*03:84, C*03:281, C*07:275
80^{4,5}	85 bp 200 bp 225 bp 340 bp	1070 bp	*15:149N *15:380N *15:263 *15:320	C*15:199 C*03:602
81⁶	395 bp	1070 bp	*15:11:01-15:11:07, 15:143, 15:148- 15:149N, 15:152, 15:209N, 15:215, 15:284, 15:303, 15:305, 15:350, 15:367, 15:486, 15:513, 15:637, 15:662	
82^{4,5}	80 bp 200 bp 245 bp	1070 bp	*15:81 *15:246N *15:277, 15:430	*53:35, 58:47 *35:222, 35:525, 35:570, 40:76, C*05:30, C*06:125, C*16:144
83⁶	255 bp	1070 bp	*15:159, 15:260, 15:491	*35:302, 35:391, 35:437, 35:539, 40:79, 41:14, 46:52
84	155 bp 195 bp 235 bp	1070 bp	*15:167 *15:211, 15:400N *15:197:01, 15:215	*35:408, 44:496 *35:560, C*03:168 *35:117
85⁵	340 bp 395 bp	1070 bp	*15:157, 15:500 *15:76, 15:101, 15:255, 15:509	C*01:240, C*03:542, C*07:335, C*07:1059, C*14:62
86	420 bp 230 bp 260 bp 400 bp	1070 bp	*15:164 *15:166 *15:272N *15:193	
87⁴	85 bp	1070 bp	*15:71, 15:175, 15:204, 15:225, 15:344, 15:381	*08:29, 18:98:01-18:98:02, 27:214, 35:50, 35:231, 35:280, 35:323:01- 35:323:02, 37:22:01-37:22:03, 40:248, 40:282, 40:351, 40:364, 40:526, 41:12, 41:36, 42:27, 46:14, 49:03, 49:48, 51:151, 52:51, 54:02, 55:16, 56:35, 57:39, 57:73, 58:41, C*03:278, C*16:86
88	295 bp 240 bp 450 bp	800 bp	*15:169 *15:171 *15:158, 15:240, 15:377Q	C*04:492, C*07:756, C*14:67 C*05:80, C*12:308, C*15:173
89	180 bp 240 bp	1070 bp	*15:160:01:01-15:160:01:02 *15:94N, 15:294N, 15:377Q	*08:72N
90⁴	95 bp	1070 bp	*15:202, 15:239, 15:376	*07:400, 13:58, 18:12:01-18:12:02, 18:195, 35:10, 35:13, 35:16, 35:28:01:01-35:28:01:03, 35:69, 35:80, 35:226, 35:270, 35:318, 35:477, 35:534, 35:541, 37:01:01:01- 37:01:05, 37:01:07-37:05, 37:07, 37:09-37:27, 37:29-37:59, 37:62- 37:66, 37:68-37:81, 37:83-37:107,



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91 ^{6,7}	135 bp 325 bp	1070 bp	*15:184, 15:203 *15:170, 15:356	*18:109, 38:62
92 ⁴	95 bp 155 bp	1070 bp	*15:266, 15:332 *15:172, 15:328	*13:37, 38:49, 51:75 *50:78
93 ⁹	255 bp 295 bp 385 bp	1070 bp	*15:258N, 15:304N *15:53, 15:177, 15:212, 15:336, 15:347, 15:559, 15:588 *15:120	*27:195, 40:26, 40:40, 40:516, 44:02:68, C*01:124, C*16:85
94	260 bp 285 bp 325 bp	1070 bp	*15:272N *15:123 *15:178	
95 ^{5,6}	160 bp 340 bp 395 bp	1070 bp	*15:358 *15:181N, 15:584N *15:40:01-15:40:02, 15:47:01-15:47:02, 15:49, 15:52, 15:114, 15:117, 15:124, 15:138, 15:238, 15:241, 15:389, 15:421, 15:519, 15:532, 15:656	*46:06, 46:25, 46:30
96 ¹⁰	-	-	Negative Control	

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 HLA-B*15 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 1070 or 800 base pairs respectively, well distribution as outlined in the table. Well number 1 contains the shorter, 800 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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For *In Vitro* Diagnostic Use
MA123 v02 SSP PI Template
Date: July 2023, Rev. No: 00

101.516-24/04 – including *Taq* polymerase
 101.516-24u/04u – without *Taq* polymerase

Visit www.caredx.com for
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Lot No.: 3S0

Lot-specific information

³For several HLA Class I alleles 1st and/or 4th 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.

⁴HLA-specific PCR products shorter than 125 base pairs have a lower intensity and are less sharp than longer PCR products.

⁵Primer mixes 40, 59, 62, 68, 80, 82, 85 and 95 have a tendency to giving rise to primer oligomer formation.

⁶Primer mixes 2, 5, 15, 30, 36, 55, 68, 77, 81, 83 and 95 may have tendencies of unspecific amplifications.

⁷Primer mix 91 may give rise to a long unspecific amplification product of approximately 800 bp. This should be disregarded when interpreting the B*15 typings.

⁸Primer mixes 23 and 77 may give rise to a lower yield of HLA-specific PCR product than the other HLA-B*15 resolution primer mixes.

⁹In primer mixes 1 and 93 the positive control band may be weaker than for other HLA-B*15 primer mixes.

¹⁰Primer mix 96 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

'w', might be weakly amplified.



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For *In Vitro* Diagnostic Use
 MA123 v02 SSP PI Template
 Date: July 2023, Rev. No: 00

101.516-24/04 – including *Taq* polymerase
 101.516-24u/04u – without *Taq* polymerase

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

Lot No.: **3S0**

Lot-specific information
PRIMER SPECIFICATION

Well No.	1	2	3	4	5	6	7	8	9	10	11	12
Length of spec.	400	290	120	180	385	525	330	105	365	300	140	400
PCR product				390						515		
Length of int. pos. control ¹	800	1070	1070	800	1070	1070	1070	1070	1070	1070	800	800
5'-primer(s) ²	45	45	463	45	45	355	45	357	2 nd	360	363	45
	5'-ggA 3' 5'	5'-ggA 3' 5'	5'-TgA 3' 5'	5'-ggA 3' 5'	5'-ggA 3' 5'	5'-TCA 3' 5'	5'-ggA 3' 5'	5'-Tgg 3' 5'	5'-CAA 3' 5'	5'-CAC 3' 5'	5'-AgC 3' 5'	5'-ggA 3' 5'
										368		
										5'-gTT 3'		
										583		
										5'-gAC 3'		
3'-primer(s) ³	272	165	538	1 st	259	3 rd	206	419	527	3 rd	463	272
	5'-Tgg 3' 5'	5'-Tgg 3' 5'	5'-CCA 3' 5'	5'-gAC 3' 5'	5'-gTT 3' 5'	5'-ATC 3' 5'	5'-CCT 3' 5'	5'-Cgg 3' 5'	5'-CCA 3' 5'	5'-ATC 3' 5'	5'-gCT 3' 5'	5'-TgA 3' 5'
	283		545	267				419	527			287
	5'-Tgg 3'		5'-AgA 3' 5'	5'-ATg 3'				5'-CgA 3' 5'	5'-CCA 3'			5'-TCC 3'
Well No.	1	2	3	4	5	6	7	8	9	10	11	12

Well No.	13	14	15	16	17	18	19	20	21	22	23	24
Length of spec.	395	465	105	120	435	195	395	360	205	190	230	205
PCR product	435	515		190				505	425			305
Length of int. pos. control ¹	800	1070	1070	1070	1070	1070	800	1070	1070	800	800	1070
5'-primer(s) ²	45	368	369	419	45	419	45	379	45	409	821	2 nd
	5'-ggA 3' 5'	5'-gTg 3' 5'	5'-TAC 3' 5'	5'-gTC 3' 5'	5'-ggA 3' 5'	5'-gTC 3' 5'	5'-ggA 3' 5'	5'-ACC 3' 5'	5'-ggA 3' 5'	5'-ggC 3' 5'	5'-gCT 3' 5'	5'-CAA 3' 5'
		419						523		410		
		5'-gTA 3'						5'-CCT 3'		5'-CCA 3'		
3'-primer(s) ³	272	3 rd	435	499	309	572	266	3 rd	83	559	916	369
	5'-TgC 3' 5'	5'-ATC 3' 5'	5'-TCT 3' 5'	5'-ggA 3' 5'	5'-ATC 3' 5'	5'-gCg 3' 5'	5'-TCC 3' 5'	5'-ATC 3' 5'	5'-TgA 3' 5'	5'-CAg 3' 5'	5'-gAT 3' 5'	5'-CCT 3' 5'
	311			570			278		301			463
	5'-ggT 3'			5'-CCg 3'			5'-TgA 3'		5'-gTC 3'			5'-gCg 3'
												471
												5'-gTT 3'
Well No.	13	14	15	16	17	18	19	20	21	22	23	24



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For *In Vitro* Diagnostic Use
 MA123 v02 SSP PI Template
 Date: July 2023, Rev. No: 00

101.516-24/04 – including *Taq* polymerase
 101.516-24u/04u – without *Taq* polymerase

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Lot No.: **3S0**

Lot-specific information

Well No.	25	26	27	28	29	30	31	32	33	34	35	36
Length of spec.	315	230	115	240	255	320	320	275	200	160	165	360
PCR product	385		195			355	395	350	255			435
						405		380	340			
Length of int. control ¹	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	800
Primer(s) ²	45	368	404	362	369	479	488	2 nd I	2 nd I	419	412	45
	5'-ggA 3' 5'	-gTC 3' 5'	-g 3' 5'	-gAC 3' 5'	-TAC 3' 5'	-C 3' 5'	-ggg 3' 5'	-CAA 3' 5'	-CAA 3' 5'	-gTC 3' 5'	-ATA 3' 5'	-ggA 3' 5'
			485			526	560					
			5'-CAA 3'			5'-gTC 3' 5'	-CAC 3'					
						560						
						5'-CgA 3'						
Primer(s) ³	193	558	559	559	583	3 rd I	3 rd I	437	362	538	538	238
	5'-CgT 3' 5'	-Agg 3' 5'	-CAg 3' 5'	-CAg 3' 5'	-gTg 3' 5'	-ATC 3' 5'	-ATC 3' 5'	-AAA 3' 5'	-TCA 3' 5'	-gTC 3' 5'	-CCA 3' 5'	-CCT 3' 5'
	263							513	420			309
	5'-gTA 3'							5'-TCT 3' 5'	-gCT 3'			5'-gTg 3'
								544	505			
								5'-ggT 3' 5'	-gCT 3'			
Well No.	25	26	27	28	29	30	31	32	33	34	35	36

Well No.	37	38	39	40	41	42	43	44	45	46	47	48
Length of spec.	170	225	315	220	465	350	525	515	475	360	320	420
PCR product			385	425		460						
Length of int. control ¹	1070	1070	800	1070	1070	1070	1070	1070	1070	800	1070	1070
Primer(s) ²	409	106	45	45	419	420	357	369	409	522	558	463
	5'-ggC 3' 5'	-CCA 3' 5'	-ggA 3' 5'	-ggA 3' 5'	-gTC 3' 5'	-TTA 3' 5'	-Tgg 3' 5'	-TAC 3' 5'	-ggC 3' 5'	-gCT 3' 5'	-ggA 3' 5'	-TgA 3' 5'
						522			410	527	560	
						5'-gCT 3'			5'-CCA 3' 5'	-TgA 3' 5'	-CCT 3'	
						539						
						5'-gCT 3'						
Primer(s) ³	538	292	181	97	3 rd I	3 rd I	3 rd I	3 rd I	3 rd I	3 rd I	3 rd I	3 rd I
	5'-CCg 3' 5'	-gTC 3' 5'	-gTT 3' 5'	-gTg 3' 5'	-ATC 3' 5'	-ATC 3' 5'	-ATC 3' 5'	-ATC 3' 5'	-ATC 3' 5'	-ATC 3' 5'	-ATC 3' 5'	-ATC 3' 5'
			203	302								
			5'-CTg 3' 5'	-ggT 3'								
			257									
			5'-CCT 3'									
			267									
			5'-ATg 3'									
Well No.	37	38	39	40	41	42	43	44	45	46	47	48



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For *In Vitro* Diagnostic Use
 MA123 v02 SSP PI Template
 Date: July 2023, Rev. No: 00

101.516-24/04 – including *Taq* polymerase
 101.516-24u/04u – without *Taq* polymerase

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Lot No.: **3S0**

Lot-specific information

Well No.	49	50	51	52	53	54	55	56	57	58	59	60
Length of spec.	135	85	135	200	210	250	175	70	340	175	230	160
PCR product		185	175	345		410	210	260	480		370	
			200				255				420	
Length of int. pos. control ¹	1070	1070	1070	800	800	800	1070	1070	1070	1070	800	800
5'-primer(s) ²	209	463	106	45	419	45	3 rd I	25	400	133	45	652
	5'-ggC 3' 5'	-TgA 3' 5'	-CCA 3' 5'	-ggA 3' 5'	-gTC 3' 5'	-ggA 3' 5'	-Agg 3' 5'	-CCg 3' 5'	-TCT 3' 5'	-CCA 3' 5'	-ggA 3' 5'	-CCg 3' 5'
				3 rd I				652	539	145	673	661
				5'-Agg 3'				5'-CCA 3' 5'	-gTg 3' 5'	-CAA 3' 5'	-CCA 3' 5'	-ACg 3'
								540			679	
								5'-TgC 3'			5'-ggC 3'	
3'-primer(s) ³	302	506	200	221	589	127	704	56	3 rd I	272	107	774
	5'-ggT 3' 5'	-TgT 3' 5'	-TCA 3' 5'	-ATA 3' 5'	-CTT 3' 5'	-CTg 3' 5'	-CCA 3' 5'	-TCg 3' 5'	-ATC 3' 5'	-Tgg 3' 5'	-ACT 3' 5'	-ggT 3' 5'
		608	200	723		288	735	872			295	
		5'-gCC 3' 5'	-TCA 3' 5'	-Tgg 3'		5'-gTg 3' 5'	-TCg 3' 5'	-TCA 3'			5'-TCA 3'	
			239	727			779				916	
			5'-gCT 3' 5'	-CCA 3'			5'-CTA 3'				5'-gAT 3'	
			263	735			787					
			5'-g 3' 5'	-TCg 3'			5'-TCC 3'					
Well No.	49	50	51	52	53	54	55	56	57	58	59	60

Well No.	61	62	63	64	65	66	67	68	69	70	71	72
Length of spec.	160	105	175	90	405	435	265	215	130	90	290	110
PCR product	205		280	135	470		300	380		185	315	200
				240	520						400	295
				365								390
Length of int. pos. control ¹	1070	1070	1070	800	1070	1070	800	1070	1070	1070	1070	1070
5'-primer(s) ²	106	363	25	45	363	209	45	419	209	463	45	261
	5'-CCg 3' 5'	-AgC 3' 5'	-CCg 3' 5'	-ggA 3' 5'	-AAT 3' 5'	-ggC 3' 5'	-ggA 3' 5'	-gTC 3' 5'	-ggC 3' 5'	-TgA 3' 5'	-ggA 3' 5'	-AAC 3' 5'
	151		419	679	410	209		668				662
	5'-gCC 3'		5'-gTC 3' 5'	-ggC 3' 5'	-gCg 3' 5'	-ggC 3'		5'-ggT 3'				5'-CCT 3'
				724	479							754
				5'-g.C 3' 5'	-..C 3'							5'-ACg 3'
												852
												5'-CAA 3'
3'-primer(s) ³	272	425	134	115	3 rd I	353	142	593	299	510	167	331
	5'-Tgg 3' 5'	-CgC 3' 5'	-AgT 3' 5'	-ggA 3' 5'	-ATC 3' 5'	-TgA 3' 5'	-ggT 3' 5'	-CgC 3' 5'	-TCA 3' 5'	-CAA 3' 5'	-ACA 3' 5'	-CTA 3' 5'
			554	242		353	177	916		605	189	916
			5'-CCg 3' 5'	-CCA 3'		5'-TgA 3' 5'	-AAA 3' 5'	-gAT 3'		5'-gCT 3' 5'	-gCT 3' 5'	-gAT 3'
				774		362					277	
				5'-ggT 3'		5'-TCA 3'					5'-ggC 3'	
Well No.	61	62	63	64	65	66	67	68	69	70	71	72



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For *In Vitro* Diagnostic Use
 MA123 v02 SSP PI Template
 Date: July 2023, Rev. No: 00

101.516-24/04 – including *Taq* polymerase
 101.516-24u/04u – without *Taq* polymerase

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Lot No.: **3S0**

Lot-specific information

Well No.	73	74	75	76	77	78	79	80	81	82	83	84
Length of spec.	155	75	280	475	185	110	215	85	395	80	255	155
PCR product		230	315		330	385	365	200		200		195
			390			475	420	225		245		235
								340				
Length of int. pos. control ¹	1070	800	1070	1070	1070	1070	800	1070	1070	1070	1070	1070
5'-primer(s) ²	412	368	662	409	47	97	45	267	45	106	106	363
	5' -ATA 3' 5'	-gTA 3' 5'	-CCT 3' 5'	-ggT 3' 5'	-AgT 3' 5'	-TCg 3' 5'	-ggA 3' 5'	-Ag 3' 5'	-ggA 3' 5'	-CCA 3' 5'	-CCA 3' 5'	-AgT 3' 5'
		523	736		463	400		709				404
		5' -CCg 3' 5'	-gCA 3'		5' -TgA 3' 5'	-TCT 3'		5' -AgT 3'				5' -CCT 3'
			769			410		824				404
			5' -Agg 3'			5' -gCg 3'		5' -TTg 3'				5' -g 3'
						495		852				446
						5' -CAC 3'		5' -CAA 3'				5' -CgT 3'
3'-primer(s) ³	527	559	916	3 rd I	206	165	93	311	272	146	322	559
	5' -CCT 3' 5'	-CAg 3' 5'	-gAT 3' 5'	-ATC 3' 5'	-CCT 3' 5'	-Tgg 3' 5'	-AAC 3' 5'	-ggT 3' 5'	-TgT 3' 5'	-CCg 3' 5'	-gTg 3' 5'	-CAg 3' 5'
					605	3 rd I	242	916		263		
					5' -gCA 3' 5'	-ATC 3' 5'	-CCT 3' 5'	-gAT 3'		5' -g 3'		
							296			312		
							5' -CTT 3'			5' -AgT 3'		
Well No.	73	74	75	76	77	78	79	80	81	82	83	84

Well No.	85	86	87	88	89	90	91	92	93	94	95
Length of spec.	340	230	85	240	180	95	135	95	255	260	160
PCR product	395	260	295	450	240	195	325	155	295	285	340
	420	400							385	325	395
Length of int. pos. control ¹	1070	1070	1070	800	1070	1070	1070	1070	1070	1070	1070
5'-primer(s) ²	45	2 nd I	97	45	106	206	45	463	45	2 nd I	2 nd I
	5' -ggA 3' 5'	-CAA 3' 5'	-TCC 3' 5'	-ggA 3' 5'	-CCA 3' 5'	-gAC 3' 5'	-ggA 3' 5'	-TgA 3' 5'	-ggA 3' 5'	-CAA 3' 5'	-CAA 3' 5'
			2 nd I			463	365		369	767	3 rd I
			5' -CAA 3'			5' -TgA 3' 5'	-gAC 3'		5' -TAC 3' 5'	-ggg 3' 5'	-Agg 3'
							376				
							5' -gCA 3'				
3'-primer(s) ³	218	394	142	118	247	259	203	514	135	426	502
	5' -gCA 3' 5'	-gCT 3' 5'	-TgC 3' 5'	-gCT 3' 5'	-ATg 3' 5'	-CTC 3' 5'	-TCT 3' 5'	-CTg 3' 5'	-Ag 3' 5'	-TCC 3' 5'	-CTA 3' 5'
	272	426	457	320	295	616	463	515	167	487	559
	5' -TgT 3' 5'	-TCC 3' 5'	-gTA 3' 5'	-AgC 3' 5'	-TCA 3' 5'	-CgT 3' 5'	-gCT 3' 5'	-CCC 3' 5'	-ACA 3' 5'	-CgT 3' 5'	-CTC 3' 5'
	296	565		323	320			578	173	916	685
	5' -CTg 3' 5'	-CAT 3'		5' -AgA 3' 5'	-AgC 3'			5' -TgT 3' 5'	-TCC 3' 5'	-gAT 3' 5'	-ggT 3'
				337					261		
				5' -CTT 3'					5' -gTA 3'		
									585		
									5' -AgT 3'		
Well No.	85	86	87	88	89	90	91	92	93	94	95



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For *In Vitro* Diagnostic Use
 MA123 v02 SSP PI Template
 Date: July 2023, Rev. No: 00

101.516-24/04 – including *Taq* polymerase
101.516-24u/04u – without *Taq* polymerase

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Lot No.: 3S0

Lot-specific information

¹The internal positive control primer pairs amplify segments of the human growth hormone gene. The internal positive control bands are 1070 or 800 base pairs respectively, well distribution as outlined in the table. Well number 1 contains the shorter, 800 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.

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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For *In Vitro* Diagnostic Use
 MA123 v02 SSP PI Template
 Date: July 2023, Rev. No: 00

101.516-24/04 – including *Taq* polymerase
101.516-24u/04u – without *Taq* polymerase

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

Lot No.: **3S0**

Lot-specific information

CELL LINE VALIDATION SHEET																				
HLA-B*15 SSP subtyping kit ²																				
				Well																
				1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	
				Prod. No.:	202239501	202239502	202239503	202239504	202239505	202239506	202239507	202239508	202239509	202239510	202239511	202239512	202239513	202239514	202239515	202239516
IHC cell line ¹		B*																		
1	9001 SA	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
2	9280 LK707	*52:01	*73:01	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-
3	9011 E4181324	*52:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
4	9275 GU373	*15:10	*53:01	-	-	-	-	+	-	+	-	-	-	-	-	+	+	-	-	-
5	9009 KAS011	*37:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
6	9353 SM	*39:01	*51:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
7	9020 QBL	*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
8	9025 DEU	*35:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
9	9026 YAR	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
10	9107 LKT3	*54:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
11	9051 PITOUT	*44:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
12	9052 DBB	*57:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
13	9004 JESTHOM	*27:05		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
14	9071 OLGA	*15:01	*15:20	+	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
15	9075 DKB	*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
16	9037 SWEIG007	*40:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
17	9282 CTM3953540	*08:01	*55:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
18	9257 32367	*14:01	*56:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
19	9038 BM16	*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
20	9059 SLE005	*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
21	9064 AMALA	*15:01		+	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
22	9056 KOSE	*35:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
23	9124 IHL	*40:02	*56:02	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
24	9035 JBUSH	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
25	9049 IBW9	*14:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
26	9285 WT49	*58:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
27	9191 CH1007	*07:05	*51:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
28	9320 BEL5GB	*44:02	*44:03	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
29	9050 MOU	*44:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
30	9021 RSH	*42:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
31	9019 DUCAF	*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
32	9297 HAG	*41:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
33	9098 MT14B	*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
34	9104 DHIF	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
35	9302 SSTO	*44:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
36	9024 KT17	*15:01	*35:01	+	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
37	9065 HHKB	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
38	9099 LZL	*15:01		+	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
39	9315 CML	*08:01	*27:05	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
40	9134 WHONP199	*13:02	*46:01	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
41	9055 H0301	*14:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
42	9066 TAB089	*46:01		-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
43	9076 T7526	*46:01		-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
44	9057 TEM	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
45	9239 SHJO	*42:01	*50:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
46	9013 SCHU	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
47	9045 TUBO	*51:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
48	9303 TER-ND	*35:01	*44:03	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-



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For *In Vitro* Diagnostic Use
MA123 v02 SSP PI Template
Date: July 2023, Rev. No: 00

101.516-24/04 – including Taq polymerase
 101.516-24u/04u – without Taq polymerase

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Lot No.: **3S0**

Lot-specific information

CELL LINE VALIDATION SHEET																				
HLA-B*15 SSP subtyping kit²																				
				Well																
				Prod. No.:																
				202239517	202239518	202239519	202239520	202239521	202239522	202239523	202239524	202239525	202239526	202239527	202239528	202239529	202239530	202239531	202239532	
IHC cell line ¹		B*																		
1	9001 SA	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
2	9280 LK707	*52:01	*73:01	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
3	9011 E4181324	*52:01		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
4	9275 GU373	*15:10	*53:01	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
5	9009 KAS011	*37:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
6	9353 SM	*39:01	*51:01	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
7	9020 QBL	*18:01		-	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-
8	9025 DEU	*35:01		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
9	9026 YAR	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
10	9107 LKT3	*54:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
11	9051 PITOUT	*44:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
12	9052 DBB	*57:01		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
13	9004 JESTHOM	*27:05		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
14	9071 OLGA	*15:01	*15:20	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
15	9075 DKB	*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
16	9037 SWEIG007	*40:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
17	9282 CTM3953540	*08:01	*55:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
18	9257 32367	*14:01	*56:01	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
19	9038 BM16	*18:01		-	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-
20	9059 SLE005	*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
21	9064 AMALA	*15:01		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
22	9056 KOSE	*35:03		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
23	9124 IHL	*40:02	*56:02	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
24	9035 JBUSH	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
25	9049 IBW9	*14:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
26	9285 WT49	*58:01		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
27	9191 CH1007	*07:05	*51:01	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
28	9320 BEL5GB	*44:02	*44:03	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
29	9050 MOU	*44:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
30	9021 RSH	*42:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
31	9019 DUCAF	*18:01		-	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-
32	9297 HAG	*41:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
33	9098 MT14B	*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
34	9104 DHIF	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
35	9302 SSTO	*44:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
36	9024 KT17	*15:01	*35:01	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
37	9065 HHKB	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
38	9099 LZL	*15:01		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
39	9315 CML	*08:01	*27:05	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
40	9134 WHONP199	*13:02	*46:01	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
41	9055 H0301	*14:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
42	9066 TAB089	*46:01		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
43	9076 T7526	*46:01		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
44	9057 TEM	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
45	9239 SHJO	*42:01	*50:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
46	9013 SCHU	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
47	9045 TUBO	*51:01		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
48	9303 TER-ND	*35:01	*44:03	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-



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For *In Vitro* Diagnostic Use
 MA123 v02 SSP PI Template
 Date: July 2023, Rev. No: 00

101.516-24/04 – including Taq polymerase
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Lot No.: **3S0**

Lot-specific information

CELL LINE VALIDATION SHEET																				
HLA-B*15 SSP subtyping kit ²																				
					Well															
					33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48
					Prod. No.:															
					202239533	202239534	202239535	202239536	202239537	202239538	202239539	202239540	202239541	202239542	202239543	202239544	202239545	202239546	202239547	202239548
IHCW cell line ¹		B*																		
1	9001	SA	*07:02		-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
2	9280	LK707	*52:01	*73:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
3	9011	E4181324	*52:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
4	9275	GU373	*15:10	*53:01	-	-	-	-	-	-	-	-	-	+	-	+	+	+	+	+
5	9009	KAS011	*37:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
6	9353	SM	*39:01	*51:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
7	9020	QBL	*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
8	9025	DEU	*35:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
9	9026	YAR	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
10	9107	LKT3	*54:01		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
11	9051	PITOUT	*44:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
12	9052	DBB	*57:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
13	9004	JESTHOM	*27:05		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
14	9071	OLGA	*15:01	*15:20	-	-	-	-	-	-	-	-	+	-	-	+	+	+	+	+
15	9075	DKB	*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
16	9037	SWEIG007	*40:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
17	9282	CTM3953540	*08:01	*55:01	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
18	9257	32367	*14:01	*56:01	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
19	9038	BM16	*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
20	9059	SLE005	*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
21	9064	AMALA	*15:01		-	-	-	-	-	-	-	-	+	-	-	+	+	+	+	+
22	9056	KOSE	*35:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
23	9124	IHL	*40:02	*56:02	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
24	9035	JBUSH	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
25	9049	IBW9	*14:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
26	9285	WT49	*58:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
27	9191	CH1007	*07:05	*51:01	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
28	9320	BEL.5GB	*44:02	*44:03	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
29	9050	MOU	*44:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
30	9021	RSH	*42:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
31	9019	DUCAF	*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
32	9297	HAG	*41:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
33	9098	MT14B	*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
34	9104	DHIF	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
35	9302	SSTO	*44:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
36	9024	KT17	*15:01	*35:01	-	-	-	-	-	-	-	-	+	-	-	+	+	+	+	+
37	9065	HHKB	*07:02		-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
38	9099	LZL	*15:01		-	-	-	-	-	-	-	-	+	-	-	+	+	+	+	+
39	9315	CML	*08:01	*27:05	-	-	-	-	-	W	-	-	-	-	-	-	-	-	-	-
40	9134	WHONP199	*13:02	*46:01	-	-	-	-	-	+	-	-	+	-	-	+	+	+	+	+
41	9055	H0301	*14:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
42	9066	TAB089	*46:01		-	-	-	-	-	+	-	-	+	-	-	+	+	+	+	+
43	9076	T7526	*46:01		-	-	-	-	-	+	-	-	+	-	-	+	+	+	+	+
44	9057	TEM	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
45	9239	SHJO	*42:01	*50:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
46	9013	SCHU	*07:02		-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
47	9045	TUBO	*51:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
48	9303	TER-ND	*35:01	*44:03	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-



0197

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Lot-specific information

CELL LINE VALIDATION SHEET																				
HLA-B*15 SSP subtyping kit ²																				
				Well																
				49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	
				Prod. No.:	202239549	202239550	202239551	202239552	202239553	202239554	202239555	202239556	202239557	202239558	202239559	202239560	202239561	202239562	202239563	202239564
	IHWC cell line ¹		B*																	
1	9001 SA		*07:02	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
2	9280 LK707		*52:01	*73:01	-	-	-	-	-	-	+	-	-	-	-	+	-	-	-	-
3	9011 E4181324		*52:01		-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-
4	9275 GU373		*15:10	*53:01	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-
5	9009 KAS011		*37:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
6	9353 SM		*39:01	*51:01	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-
7	9020 QBL		*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
8	9025 DEU		*35:01		-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-
9	9026 YAR		*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
10	9107 LKT3		*54:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
11	9051 PITOUT		*44:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
12	9052 DBB		*57:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
13	9004 JESTHOM		*27:05		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
14	9071 OLGA		*15:01	*15:20	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-
15	9075 DKB		*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
16	9037 SWEIG007		*40:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
17	9282 CTM3953540		*08:01	*55:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
18	9257 32367		*14:01	*56:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
19	9038 BM16		*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
20	9059 SLE005		*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
21	9064 AMALA		*15:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
22	9056 KOSE		*35:03		-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-
23	9124 IHL		*40:02	*56:02	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
24	9035 JBUSH		*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
25	9049 IBW9		*14:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
26	9285 WT49		*58:01		-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-
27	9191 CH1007		*07:05	*51:01	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-
28	9320 BEL5GB		*44:02	*44:03	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
29	9050 MOU		*44:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
30	9021 RSH		*42:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
31	9019 DUCAF		*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
32	9297 HAG		*41:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
33	9098 MT14B		*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
34	9104 DHIF		*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
35	9302 SSTO		*44:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
36	9024 KT17		*15:01	*35:01	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-
37	9065 HHKB		*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
38	9099 LZL		*15:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
39	9315 CML		*08:01	*27:05	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
40	9134 WHONP199		*13:02	*46:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
41	9055 H0301		*14:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
42	9066 TAB089		*46:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
43	9076 T7526		*46:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
44	9057 TEM		*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
45	9239 SHJO		*42:01	*50:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
46	9013 SCHU		*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
47	9045 TUBO		*51:01		-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-
48	9303 TER-ND		*35:01	*44:03	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-



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For *In Vitro* Diagnostic Use
MA123 v02 SSP PI Template
Date: July 2023, Rev. No: 00

101.516-24/04 – including Taq polymerase
101.516-24u/04u – without Taq polymerase

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

Lot No.: **3S0**

Lot-specific information

CELL LINE VALIDATION SHEET					Well															
HLA-B*15 SSP subtyping kit ²					65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80
Prod. No.:					202239565	202239566	202239567	202239568	202239569	202239570	202239571	202239572	202239573	202239574	202239575	202239576	202239577	202239578	202239579	202239580
	IHWC cell line ¹		B*																	
1	9001 SA		*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
2	9280 LK707		*52:01	*73:01	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-
3	9011 E4181324		*52:01		-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-
4	9275 GU373		*15:10	*53:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
5	9009 KAS011		*37:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
6	9353 SM		*39:01	*51:01	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-
7	9020 QBL		*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
8	9025 DEU		*35:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
9	9026 YAR		*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
10	9107 LKT3		*54:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
11	9051 PITOUT		*44:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
12	9052 DBB		*57:01		-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
13	9004 JESTHOM		*27:05		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
14	9071 OLGA		*15:01	*15:20	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
15	9075 DKB		*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
16	9037 SWEIG007		*40:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
17	9282 CTM3953540		*08:01	*55:01	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-
18	9257 32367		*14:01	*56:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
19	9038 BM16		*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
20	9059 SLE005		*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
21	9064 AMALA		*15:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
22	9056 KOSE		*35:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
23	9124 IHL		*40:02	*56:02	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
24	9035 JBUSH		*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
25	9049 IBW9		*14:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
26	9285 WT49		*58:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
27	9191 CH1007		*07:05	*51:01	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-
28	9320 BEL5GB		*44:02	*44:03	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
29	9050 MOU		*44:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
30	9021 RSH		*42:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
31	9019 DUCAF		*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
32	9297 HAG		*41:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
33	9098 MT14B		*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
34	9104 DHIF		*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
35	9302 SSTO		*44:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
36	9024 KT17		*15:01	*35:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
37	9065 HHKB		*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
38	9099 LZL		*15:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
39	9315 CML		*08:01	*27:05	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
40	9134 WHONP199		*13:02	*46:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
41	9055 H0301		*14:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
42	9066 TAB089		*46:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
43	9076 T7526		*46:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
44	9057 TEM		*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
45	9239 SHJO		*42:01	*50:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
46	9013 SCHU		*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
47	9045 TUBO		*51:01		-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-
48	9303 TER-ND		*35:01	*44:03	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-



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For *In Vitro* Diagnostic Use
MA123 v02 SSP PI Template
Date: July 2023, Rev. No: 00

101.516-24/04 – including Taq polymerase
 101.516-24u/04u – without Taq polymerase

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 “Instructions for Use” (IFU)

Lot No.: **3S0**

Lot-specific information

CELL LINE VALIDATION SHEET																		
HLA-B*15 SSP subtyping kit ²																		
			Prod. No.:	Well														
				81	82	83	84	85	86	87	88	89	90	91	92	93	94	95
				202239581	202239582	202239583	202239584	202239585	202239586	202239587	202239588	202239589	202239590	202239591	202239592	202239593	202239594	202239595
	IHWC cell line ¹	B*																
1	9001 SA	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
2	9280 LK707	*52:01	*73:01	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-
3	9011 E4181324	*52:01		-	-	-	-	-	-	-	-	-	+	-	-	-	-	-
4	9275 GU373	*15:10	*53:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
5	9009 KAS011	*37:01		-	-	-	-	-	-	-	-	-	+	-	-	-	-	-
6	9353 SM	*39:01	*51:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
7	9020 QBL	*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
8	9025 DEU	*35:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
9	9026 YAR	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
10	9107 LKT3	*54:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
11	9051 PITOUT	*44:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
12	9052 DBB	*57:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
13	9004 JESTHOM	*27:05		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
14	9071 OLGGA	*15:01	*15:20	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
15	9075 DKB	*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
16	9037 SWEIG007	*40:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
17	9282 CTM3953540	*08:01	*55:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
18	9257 32367	*14:01	*56:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
19	9038 BM16	*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
20	9059 SLE005	*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
21	9064 AMALA	*15:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
22	9056 KOSE	*35:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
23	9124 IHL	*40:02	*56:02	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
24	9035 JBUSH	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
25	9049 IBW9	*14:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
26	9285 WT49	*58:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
27	9191 CH1007	*07:05	*51:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
28	9320 BEL5GB	*44:02	*44:03	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
29	9050 MOU	*44:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
30	9021 RSH	*42:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
31	9019 DUCAF	*18:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
32	9297 HAG	*41:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
33	9098 MT14B	*40:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
34	9104 DHIF	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
35	9302 SSTO	*44:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
36	9024 KT17	*15:01	*35:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
37	9065 HHKB	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
38	9099 LZL	*15:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
39	9315 CML	*08:01	*27:05	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
40	9134 WHONP199	*13:02	*46:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
41	9055 H0301	*14:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
42	9066 TAB089	*46:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
43	9076 T7526	*46:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
44	9057 TEM	*38:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
45	9239 SHJO	*42:01	*50:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
46	9013 SCHU	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
47	9045 TUBO	*51:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
48	9303 TER-ND	*35:01	*44:03	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-



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For *In Vitro* Diagnostic Use
 MA123 v02 SSP PI Template
 Date: July 2023, Rev. No: 00

101.516-24/04 – including *Taq* polymerase
101.516-24u/04u – without *Taq* polymerase

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Lot No.: 3S0**Lot-specific information**

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.

No DNAs carrying the alleles to be amplified by primer solutions 4, 9, 10, 18, 24 to 27, 30 to 34, 36, 39, 49 to 54, 56 to 59, 61 to 65, 67 to 72, 74, 75, 78 to 80, 82 to 89 and 91 to 95 were available. The specificities of the primers in primer solutions 9, 10, 18, 24 to 27, 30, 31, 33, 34, 36, 49, 53, 59, 61, 63 to 65, 67, 68, 70 to 72, 78, 82, 85, 87, 91 and 93 to 95 were tested by separately adding one or two additional 5'-primers, respectively one to three additional 3'-primers. In primer solutions 4, 32, 39, 50 to 52, 54, 56, 62, 69, 79, 83, 86, 88, 89 and 92 it was only possible to test the 5'-primers, the 3'-primers were not possible to test. In primer solutions 57, 58, 74, 75, 80 and 84 it was only possible to test the 3'-primers, the 5'-primers were not possible to test. In primer solutions 1, 3, 12, 13, 19, 21, 24, 25, 33, 36, 55, 59, 63, 64, 67, 68, 70 to 72, 77, 82, 85, 87, 90, 91, 93 to 95 one or more 3'-primers were not possible to test. In primer solutions 10, 14, 20, 22, 27, 30, 31, 42, 45 to 47, 59 to 61, 64, 65, 72, 77, 91 and 94 one, two or three 5'-primers were not possible to test. In addition, the primers in primer solutions 16, 40, 42, 66, 77 and 90 were tested by separately adding one additional 5'-primer or one additional 3'-primer.

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For *In Vitro* Diagnostic Use
MA123 v02 SSP PI Template
Date: July 2023, Rev. No: 00

101.516-24/04 – including *Taq* polymerase
101.516-24u/04u – without *Taq* polymerase

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“Instructions for Use” (IFU)

Lot No.: **3S0**

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

Distributed by:

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**.



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For *In Vitro* Diagnostic Use
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