

101.615-24/04 – including *Taq* pol., IFU-01
101.615-24u/04u – without *Taq* pol., IFU-02

Visit www.olerup-ssp.com for
“Instructions for Use” (IFU)

Lot No.: 66N

Lot-specific information
Olerup SSP® HLA-C*07

Product number:	101.615-24/04 – including <i>Taq</i> pol. 101.615-24u/04u – without <i>Taq</i> pol.
Lot number:	66N
Expiry date:	2014-October-01
Number of tests:	24 tests – Product No. 101.615-24 4 tests – Product No. 101.615-04
Number of wells per test:	64
Storage - pre-aliquoted primers:	dark at -20°C
- PCR Master Mix:	-20°C
- Adhesive PCR seals	RT
- Product Insert	RT

This Product Description is only valid for Lot No. 66N.

**CHANGES COMPARED TO THE PREVIOUS OLERUP SSP®
HLA-C*07 LOT (62M)**

The HLA-C*07 kit is updated for new alleles to enable separation of:

- Confirmed¹ alleles as listed in the IMGT/HLA database
- Polymorphisms in exons outside of the region encoding the peptide binding domain
- Null and Alternatively expressed alleles

Sixteen wells have been added to the HLA-C*07 kit, wells **49 to 64**.

The Lot-specific information for HLA-C*07 including and without *Taq* polymerase is now described in one common Product Insert.

¹As described in section Uniquely Identified Alleles.

The HLA-C*07 specificity and interpretation tables have been updated for the HLA-C alleles described since the previous *Olerup SSP® HLA-C*07* lot was made (**Lot No. 62M**).

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

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Well	5'-primer	3'-primer	rationale
11	Added	-	5'-primer added for the C*07:112 allele.
15	-	-	Exchanged positive control primer pair, to decrease primer oligomer formation.
20	Modified	-	Increased yield of specific PCR product.
31	-	Modified	Increased yield of specific PCR product.
34	-	-	Exchanged positive control primer pair, to decrease primer oligomer formation.
35	Moved	Moved	Exchanged positive control primer pair, primer pair moved to well 43, to decrease primer oligomer formation.
42	Added	-	5'-primers added for the C*07:123, 07:173 and 07:207 alleles.
43	Added	Added	Primer pair from well 35.
46	-	Added	3'-primers added for the C*07:152N, and 07:202 alleles.
47	-	Added	3'-primer added for the C*07:152N allele.
49	New	New	New primer pairs for the C*07:100, 07:101 and 07:148 alleles.
50	New	New	New primer pairs for the C*07:104N, 07:119, 07:124, 07:155, and 07:122 alleles.
51	New	New	New primer pairs for the C*07:117, 07:121Q and 07:126 alleles.
52	New	New	New primer pairs for the C*07:118, 07:203 and 07:106 alleles.
53	New	New	New primer pairs for the C*07:140, 07:151 and 07:154 alleles.
54	New	New	New primer pairs for the C*07:125, 07:129, 07:153 and 07:172 alleles.
55	New	New	New primer pairs for the C*07:132 and 07:179 alleles.
56	New	New	New primer pairs for the C*07:143, 07:150Q, 07:123 and 07:173 alleles.
57	New	New	New primer pair for the C*07:130 allele.
58	New	New	New primer pairs for the C*07:137:02, 07:164N and 07:202 alleles.
59	New	New	New primer pairs for the C*07:141, 07:145 and 07:191N alleles.
60	New	New	New primer pairs for the C*07:147, 07:165 and 07:180 alleles.
61	New	New	New primer pairs for the C*07:120 and 07:218 alleles.
62	New	New	New primer pairs for the C*07:106, 07:127 and 07:205 alleles.
63	New	New	New primer pairs for the C*07:198N, 07:24 and 07:202 alleles.
64	New	New	New primer pairs for the C*07:176 and 07:192 alleles.

Change in revision R01 compared to R00:

- The positive control primer pair in primer mix 63 has been corrected to 800 base pairs in the specificity and amplification tables.

Change in revision R02 compared to R01:

- The HLA-C*07:24 and 07:218 and the C*03:125, 12:45 and 16:13 alleles are not amplified by primer mix 61. This has been corrected in the Specificity and Interpretation Tables. Thus, this lot of the HLA-C*07 subtyping kit cannot distinguish the C*07:198N and the C*07:218 alleles.

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PRODUCT DESCRIPTION

HLA-C*07 SSP subtyping

CONTENT

The primer set contains 5'- and 3'-primers for identifying the HLA-C*07:01 to HLA-C*07:220 alleles.

PLATE LAYOUT

Each test consists of 64 PCR reactions in a 64 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	54

The 64 well cut PCR plate is marked with ‘HLA-C*07’ in silver/gray ink.

Well No. 1 is marked with the Lot No. ‘66N’.

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 heat-sealed with a PCR-compatible foil.

INTERPRETATION

The interpretation of HLA-C*07 SSP subtypings will be influenced by the C*01:60 allele, two C*02, twelve C*03, ten C*04, the C*05, most C*06, most C*08, five C*12, the C*14:17 allele, two C*15, the C*16:13, the C*17 and the C*18 alleles when present on the other haplotype.

In addition, the A*29:10 allele will be amplified by primer mix 12, the A*30:01:03 allele by primer mix 20, the A*33:03:09 allele will be amplified by primer mixes 20 and 24, the B*07:77 allele will be amplified by primer mixes 18 and 37, the B*07:87 and B*27:55 will be amplified by primer mix 50, the B*08:17, B*08:39 and B*08:47 alleles by primer mix 32, the B*27:36, B*48:21 and B*48:26 alleles by primer mix 7, the B*35:08:04 allele by primer mixes 2, 11, 20, 27, 42, 56 and 59, the B*35:178 and B*73:01-73:02 alleles will be amplified by primer mix 45, the B*37:04:02 allele by primer mixes 11, 14, 20, 53 and 59, the B*38:23 allele by primer mix 8, the B*40:60 allele by primer mixes 4, 18 and 37, the B*40:100 allele by primer mix 10, the B*46:06 allele by primer mix 16 and the B*56:08 and B*56:14 alleles by primer mix 25.

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UNIQUELY IDENTIFIED ALLELES

All the HLA-C*07 alleles, i.e. C*07:01 to C*07:220, recognized by the HLA Nomenclature Committee in January 2012¹ will be amplified by the primers in the HLA-C*07 SSP kit.

The HLA-C*07 kit enables separation of the confirmed HLA-C*07 alleles as listed in the IMGT/HLA database. 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-C*07 alleles is listed below.

The HLA-C*07 kit also enables identification of polymorphisms in exons outside of the region encoding the peptide binding domain and of null and alternatively expressed alleles.

The C*07:93 and 07:207 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 42.

The C*07:100 and 07:161 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 49.

The C*07:104N, 07:119 and 07:124 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 50.

The C*07:117, 07:121Q and 07:126 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 51.

The C*07:118 and 07:203 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 52.

The C*07:122 and 07:151 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 53.

The C*07:129 and 07:153 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 54.

The C*07:132 and 07:179 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 55.

The C*07:164N and 07:212 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 58.

The C*07:165 and 07:180 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 60.

The HLA-C*07 subtyping kit cannot distinguish the following silent mutations: the C*07:01:01-07:01:04 and 07:01:06-07:01:24 alleles, the C*07:02:01:01-07:02:29 alleles, the C*07:04:01-07:04:07, the C*07:27:01-07:27:02 or the C*07:56:01-07:56:02 alleles.

¹HLA-C alleles listed on the IMGT/HLA web page 2012-January-12, release 3.7.0, www.ebi.ac.uk/imgt/hla.

²This lot of the HLA-C*07 subtyping kit cannot distinguish the C*07:198N and the C*07:218 alleles.

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ALLELIC CONFIRMATION STATUS

Allele	Status ¹	Allele	Status ¹	Allele	Status ¹	Allele	Status ¹	Allele	Status ¹
C*07:01:01	Confirmed	C*07:04:05	Unconfirmed	C*07:61N	Confirmed	C*07:121Q	Confirmed	C*07:180	Confirmed
C*07:01:02	Confirmed	C*07:04:06	Unconfirmed	C*07:62	Unconfirmed	C*07:122	Confirmed	C*07:181	Unconfirmed
C*07:01:03	Unconfirmed	C*07:04:07	Unconfirmed	C*07:63	Unconfirmed	C*07:123	Confirmed	C*07:182	Unconfirmed
C*07:01:04	Unconfirmed	C*07:05	Confirmed	C*07:64	Unconfirmed	C*07:124	Confirmed	C*07:183	Unconfirmed
C*07:01:05	Confirmed	C*07:06	Confirmed	C*07:65	Confirmed	C*07:125	Confirmed	C*07:184	Confirmed
C*07:01:06	Unconfirmed	C*07:07	Unconfirmed	C*07:66	Confirmed	C*07:126	Confirmed	C*07:185	Unconfirmed
C*07:01:07	Confirmed	C*07:08	Unconfirmed	C*07:67	Confirmed	C*07:127	Confirmed	C*07:186	Unconfirmed
C*07:01:08	Confirmed	C*07:09	Unconfirmed	C*07:68	Confirmed	C*07:128	Unconfirmed	C*07:187	Unconfirmed
C*07:01:09	Confirmed	C*07:10	Confirmed	C*07:69	Confirmed	C*07:129	Confirmed	C*07:188	Unconfirmed
C*07:01:10	Confirmed	C*07:11	Unconfirmed	C*07:70	Confirmed	C*07:130	Confirmed	C*07:189	Unconfirmed
C*07:01:11	Unconfirmed	C*07:12	Confirmed	C*07:71	Unconfirmed	C*07:131	Unconfirmed	C*07:190	Unconfirmed
C*07:01:12	Unconfirmed	C*07:13	Confirmed	C*07:72	Confirmed	C*07:132	Confirmed	C*07:191N	Unconfirmed
C*07:01:13	Unconfirmed	C*07:14	Unconfirmed	C*07:73	Confirmed	C*07:133	Unconfirmed	C*07:192	Confirmed
C*07:01:14	Unconfirmed	C*07:15	Confirmed	C*07:74	Unconfirmed	C*07:134	Unconfirmed	C*07:193	Unconfirmed
C*07:01:15	Confirmed	C*07:16	Confirmed	C*07:75	Confirmed	C*07:135	Unconfirmed	C*07:194	Unconfirmed
C*07:01:16	Unconfirmed	C*07:17	Confirmed	C*07:76	Confirmed	C*07:136	Unconfirmed	C*07:195	Unconfirmed
C*07:01:17	Unconfirmed	C*07:18	Confirmed	C*07:77	Unconfirmed	C*07:137:01	Unconfirmed	C*07:196	Confirmed
C*07:01:18	Confirmed	C*07:19	Confirmed	C*07:78	Unconfirmed	C*07:137:02	Confirmed	C*07:197	Unconfirmed
C*07:01:19	Unconfirmed	C*07:20	Confirmed	C*07:79	Confirmed	C*07:138	Unconfirmed	C*07:198N	Unconfirmed
C*07:01:20	Unconfirmed	C*07:21	Unconfirmed	C*07:80	Confirmed	C*07:139	Unconfirmed	C*07:199	Unconfirmed
C*07:01:21	Unconfirmed	C*07:22	Confirmed	C*07:81	Unconfirmed	C*07:140	Confirmed	C*07:200	Unconfirmed
C*07:01:22	Unconfirmed	C*07:23	Unconfirmed	C*07:82	Unconfirmed	C*07:141	Confirmed	C*07:201	Unconfirmed
C*07:01:23	Unconfirmed	C*07:24	Unconfirmed	C*07:83	Confirmed	C*07:142	Unconfirmed	C*07:202	Confirmed
C*07:01:24	Unconfirmed	C*07:25	Unconfirmed	C*07:84	Unconfirmed	C*07:143	Confirmed	C*07:203	Confirmed
C*07:02:01:01	Confirmed	C*07:26	Confirmed	C*07:85	Confirmed	C*07:144	Unconfirmed	C*07:204	Unconfirmed
C*07:02:01:02	Unconfirmed	C*07:27:01	Unconfirmed	C*07:86	Unconfirmed	C*07:145	Confirmed	C*07:205	Confirmed
C*07:02:01:03	Confirmed	C*07:27:02	Confirmed	C*07:87	Unconfirmed	C*07:146	Unconfirmed	C*07:206	Unconfirmed
C*07:02:02	Unconfirmed	C*07:28	Confirmed	C*07:88	Confirmed	C*07:147	Confirmed	C*07:207	Confirmed
C*07:02:03	Unconfirmed	C*07:29	Unconfirmed	C*07:89	Confirmed	C*07:148	Confirmed	C*07:208	Unconfirmed
C*07:02:04	Confirmed	C*07:30	Unconfirmed	C*07:90	Confirmed	C*07:149	Unconfirmed	C*07:209	Unconfirmed
C*07:02:05	Unconfirmed	C*07:31	Confirmed	C*07:91	Confirmed	C*07:150Q	Unconfirmed	C*07:210	Unconfirmed
C*07:02:06	Confirmed	C*07:32N	Confirmed	C*07:92	Unconfirmed	C*07:151	Confirmed	C*07:211	Unconfirmed
C*07:02:07	Confirmed	C*07:33N	Confirmed	C*07:93	Confirmed	C*07:152N	Unconfirmed	C*07:212	Confirmed
C*07:02:08	Confirmed	C*07:35	Confirmed	C*07:94	Unconfirmed	C*07:153	Confirmed	C*07:213	Unconfirmed
C*07:02:09	Unconfirmed	C*07:36	Confirmed	C*07:95	Confirmed	C*07:154	Confirmed	C*07:214	Unconfirmed
C*07:02:10	Confirmed	C*07:37	Unconfirmed	C*07:96	Confirmed	C*07:155	Confirmed	C*07:215	Unconfirmed
C*07:02:11	Confirmed	C*07:38	Unconfirmed	C*07:97	Confirmed	C*07:156	Unconfirmed	C*07:216	Unconfirmed
C*07:02:12	Unconfirmed	C*07:39	Unconfirmed	C*07:98N	Unconfirmed	C*07:157	Unconfirmed	C*07:217	Unconfirmed
C*07:02:13	Unconfirmed	C*07:40	Confirmed	C*07:99	Unconfirmed	C*07:158	Unconfirmed	C*07:218	Confirmed
C*07:02:14	Unconfirmed	C*07:41	Unconfirmed	C*07:100	Confirmed	C*07:159	Unconfirmed	C*07:219	Unconfirmed
C*07:02:15	Unconfirmed	C*07:42	Unconfirmed	C*07:101	Confirmed	C*07:160	Unconfirmed	C*07:220	Unconfirmed
C*07:02:16	Confirmed	C*07:43	Confirmed	C*07:102	Unconfirmed	C*07:161	Unconfirmed		
C*07:02:17	Confirmed	C*07:44	Unconfirmed	C*07:103	Unconfirmed	C*07:162	Unconfirmed		
C*07:02:18	Confirmed	C*07:45	Unconfirmed	C*07:104N	Confirmed	C*07:163	Unconfirmed		
C*07:02:19	Unconfirmed	C*07:46	Confirmed	C*07:105	Unconfirmed	C*07:164N	Unconfirmed		
C*07:02:20	Confirmed	C*07:47	Unconfirmed	C*07:106	Confirmed	C*07:165	Confirmed		
C*07:02:21	Unconfirmed	C*07:48	Unconfirmed	C*07:107	Unconfirmed	C*07:166	Unconfirmed		
C*07:02:22	Unconfirmed	C*07:49	Confirmed	C*07:108	Confirmed	C*07:167	Unconfirmed		
C*07:02:23	Unconfirmed	C*07:50	Unconfirmed	C*07:109	Confirmed	C*07:168	Unconfirmed		
C*07:02:24	Confirmed	C*07:51	Confirmed	C*07:110	Unconfirmed	C*07:169	Unconfirmed		
C*07:02:25	Unconfirmed	C*07:52	Unconfirmed	C*07:111	Unconfirmed	C*07:170	Unconfirmed		
C*07:02:26	Unconfirmed	C*07:53	Unconfirmed	C*07:112	Confirmed	C*07:171	Unconfirmed		
C*07:02:27	Unconfirmed	C*07:54	Confirmed	C*07:113	Unconfirmed	C*07:172	Confirmed		
C*07:02:28	Unconfirmed	C*07:55N	Confirmed	C*07:114	Unconfirmed	C*07:173	Confirmed		
C*07:02:29	Unconfirmed	C*07:56:01	Unconfirmed	C*07:115	Unconfirmed	C*07:174	Unconfirmed		
C*07:03	Unconfirmed	C*07:56:02	Confirmed	C*07:116	Unconfirmed	C*07:175	Unconfirmed		
C*07:04:01	Confirmed	C*07:57	Unconfirmed	C*07:117	Confirmed	C*07:176	Confirmed		
C*07:04:02	Unconfirmed	C*07:58	Unconfirmed	C*07:118	Confirmed	C*07:177	Unconfirmed		
C*07:04:03	Unconfirmed	C*07:59	Confirmed	C*07:119	Confirmed	C*07:178	Unconfirmed		
C*07:04:04	Unconfirmed	C*07:60	Confirmed	C*07:120	Confirmed	C*07:179	Confirmed		



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¹Allele status “confirmed” or “unconfirmed” as listed on the IMGT/HLA web page 2012-January-12, release 3.7.0, www.ebi.ac.uk/imgt/hla.

RESOLUTION IN HOMO- AND HETEROZYGOTES

A total of 281 alleles generate 147 amplification patterns that can be combined in 10878 homozygous and heterozygous combinations. 3312 of these genotypes do not give rise to unique amplification patterns. The different lengths of the specific PCR products were not considered in these calculations.

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Lot-specific information
SPECIFICITY TABLE

HLA-C*07 SSP subtyping

Specificities and sizes of the PCR products of the 64 primer mixes used for HLA-C*07 SSP subtyping

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	Amplified HLA-C*07 alleles ³	Other amplified HLA-Class I alleles ⁴
1 ¹⁰	245 bp, 425 bp	800 bp	*07:01:01-07:33N, 07:35-07:220	
2	290 bp	800 bp	*07:01:01-07:01:04, 07:01:06-07:01:24, 07:05-07:07, 07:09, 07:14, 07:16, 07:18, 07:20-07:22, 07:24, 07:26-07:28, 07:30-07:31, 07:35-07:36, 07:40, 07:43-07:44, 07:52-07:53, 07:55N, 07:58-07:60, 07:65, 07:69-07:71, 07:73, 07:77-07:78, 07:81-07:83, 07:85-07:86, 07:89, 07:91-07:96, 07:98N, 07:103-07:104N, 07:106, 07:109, 07:111, 07:113, 07:115-07:116, 07:118-07:120, 07:122, 07:124, 07:128-07:129, 07:131-07:132, 07:134, 07:140-07:141, 07:148, 07:150Q-07:151, 07:153, 07:156, 07:162-07:166, 07:170, 07:173, 07:176, 07:179-07:180, 07:182, 07:184, 07:188-07:191N, 07:196-07:197, 07:200-07:201, 07:203-07:207, 07:210, 07:212, 07:214-07:215, 07:219	*03:04:19, 03:113, 05:01:17, 05:09:02, 05:17, 08:15:02, 12:03:10, B*35:08:04
3 ⁵	110 bp	800 bp	*07:01:01-07:01:24, 07:06, 07:09, 07:18-07:22, 07:24, 07:26, 07:28, 07:30, 07:35-07:36, 07:40, 07:44, 07:52-07:53, 07:55N, 07:57-07:59, 07:65, 07:69-07:71, 07:73, 07:77-07:78, 07:81-07:83,	*06:44, 18:05



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			07:85-07:86, 07:89, 07:91- 07:96, 07:98N, 07:103- 07:104N, 07:106, 07:108, 07:110-07:113, 07:115- 07:116, 07:118-07:120, 07:122, 07:124, 07:128- 07:129, 07:131-07:132, 07:134, 07:140-07:141, 07:148, 07:150Q-07:151, 07:153, 07:156, 07:162- 07:166, 07:170, 07:173, 07:176, 07:179-07:180, 07:182, 07:188-07:191N, 07:196-07:197, 07:200- 07:201, 07:203-07:207, 07:210, 07:212, 07:214- 07:215, 07:219
4	185 bp	1070 bp	*07:02:01:01-07:02:29, 07:10, B*40:60 07:13, 07:15, 07:19, 07:23, 07:25, 07:29, 07:32N-07:33N, 07:37-07:39, 07:42, 07:46- 07:51, 07:54, 07:56:01- 07:56:02, 07:61N-07:62, 07:64, 07:66-07:68, 07:72, 07:74-07:76, 07:79-07:80, 07:84, 07:87-07:88, 07:90, 07:97, 07:99-07:100, 07:102, 07:105, 07:107, 07:114, 07:117, 07:121Q, 07:123, 07:125-07:127, 07:130, 07:133, 07:135-07:138, 07:143-07:147, 07:149, 07:152N, 07:154-07:155, 07:157-07:161, 07:167- 07:169, 07:172, 07:174- 07:175, 07:178, 07:183, 07:185-07:187, 07:192- 07:195, 07:198N, 07:202, 07:208-07:209, 07:211, 07:213, 07:216-07:218, 07:220
5^{5,8,11}	70 bp, 145 bp, 245 bp	800 bp	*07:03, 07:53, 07:57, 07:216
6⁵	100 bp	1070 bp	*07:04:01-07:04:07, 07:11- *05:62 07:12, 07:63, 07:68, 07:101, 07:139, 07:142, 07:181, 07:199



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7	190 bp	1070 bp	*07:05 B*27:36, B*48:21, 48:26
8 ¹²	285 bp, 505 bp	1070 bp	*07:06, 07:45 *04:40, B*38:23
9 ^{6,13}	215 bp, 240 bp, 425 bp	800 bp	*07:07, 07:09, 07:29, 07:33N, 07:49, 07:76, 07:177, 07:184, 07:210 *03:04:19, 03:113, 06:17, 18:01-18:05
10	185 bp	1070 bp	*07:08, 07:108 *04:58, B*40:100
11 ¹⁴	275 bp, 300 bp	800 bp	*07:10, 07:24, 07:43, 07:112, 07:184, 07:196, 07:218 *03:04:19, 03:113, B*35:08:04, B*37:04:02
12	130 bp	1070 bp	*07:12, 07:41 *05:01:01:01-05:01:18, 05:03-05:08, 05:10-05:16, 05:18:01-05:22:02, 05:24-05:48N, 05:50-05:51Q, 05:53-05:61, 05:63-05:72, 08:01:01-08:06, 08:08-08:14, 08:16-08:46, 08:48-08:50, 08:52N-08:56, A*29:10
13 ^{5,7}	90 bp	1070 bp	*07:11, 07:56:01-07:56:02, 07:60, 07:79, 07:109 *03:69, 04:27, 04:52, 06:53, 12:58
14	240 bp	800 bp	*07:13, 07:28 *05:01:17, 05:09:02, 05:17, 08:15:02, B*37:04:02
15 ^{5,15}	80 bp, 290 bp, 460 bp	1070 bp	*07:14, 07:50, 07:89
16	535 bp	1070 bp	*07:02:01:01-07:02:29, 07:04:01-07:05, 07:08, 07:10-07:11, 07:13-07:15, 07:23, 07:25, 07:27:01-07:27:02, 07:29, 07:31-07:33N, 07:37-07:39, 07:42-07:43, 07:45-07:51, 07:54, 07:56:01-07:56:02, 07:61N-07:64, 07:66-07:68, 07:72, 07:74-07:76, 07:79-07:80, 07:84, 07:87-07:88, 07:90, 07:97, 07:99-07:102, 07:105, 07:107, 07:114, 07:117, 07:121Q, 07:123, 07:125-07:127, 07:130, 07:133, 07:135-07:139, *01:60, 03:14, 04:58, 05:23, 05:62, 08:07, 08:47, 12:14:01-12:14:02, 14:17, 17:01:01-17:11, B*46:06



101.615-24/04 – including *Taq pol.*, IFU-01
 101. 615-24u/04u – without *Taq pol.*, IFU-02

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Lot No.: 66N

Lot-specific information

			07:142-07:147, 07:149, 07:152N, 07:154-07:155, 07:157-07:161, 07:167- 07:169, 07:172, 07:174- 07:175, 07:177-07:178, 07:181, 07:183-07:187, 07:192-07:195, 07:198N- 07:199, 07:202, 07:208- 07:209, 07:211, 07:213, 07:216-07:218, 07:220
17 ⁷	130 bp	1070 bp	*07:04:01-07:04:07, 07:11, 07:63, 07:68, 07:101, 07:139, 07:142, 07:181, 07:199
18 ^{5,16}	120 bp, 230 bp	800 bp	*07:15, 07:52
			*05:03, B*07:77, B*40:60
19 ¹⁷	220 bp, 355 bp, 465 bp	800 bp	*07:07, 07:16, 07:26, 07:33N, 07:51, 07:80, 07:92, 07:96, 07:177, 07:181
20 ¹⁸	145 bp, 170 bp, 195 bp	1070 bp	*07:03, 07:17, 07:44, 07:48, 07:81, 07:140-07:141, 07:168
			*03:04:19, 03:113, 05:01:17, 05:09:02, 05:17, 08:15:02, 12:03:10, A*30:01:03, A*33:03:09, B*35:08:04, B*37:04:02
21 ^{5,19}	100 bp, 225 bp	800 bp	*07:06, 07:18-07:19, 07:54
22 ²⁰	205 bp, 250 bp	800 bp	*07:20, 07:84, 07:96, 07:127
			*02:49, 03:15, 03:32, 03:45, 03:60, 03:136, 04:03, 04:06, 04:16, 04:80, 15:25
23 ²¹	155 bp, 185 bp	1070 bp	*07:21, 07:23
24	130 bp	1070 bp	*07:22, 07:25, 07:137:01, 07:138
			*12:03:10, A*33:03:09
25 ^{5,22}	105 bp, 150 bp	1070 bp	*07:30, 07:46
26 ^{5,23}	100 bp, 250 bp, 300 bp,	1070 bp	*07:31-07:32N, 07:55N, 07:177
27 ⁵	125 bp	1070 bp	*07:37, 07:69, 07:176
			*05:01:17, 05:09:02, 08:15:02, B*35:08:04
28 ²⁴	175 bp, 260 bp	800 bp	*07:35, 07:44, 07:47
29	255 bp	1070 bp	*07:36, 07:42
30 ⁵	60 bp	1070 bp	*07:75



101.615-24/04 – including *Taq* pol., IFU-01
 101. 615-24u/04u – without *Taq* pol., IFU-02

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Lot No.: 66N		Lot-specific information		
31	415 bp	1070 bp	*07:39-07:40, 07:177, 07:210 ^w	
32 ²⁵	245 bp, 285 bp	1070 bp	*07:07, 07:09, 07:49, 07:60, 07:76, 07:79, 07:210	*02:14, 03:39, 03:67, 04:42, 05:43, 06:02:01:01- 06:02:01:02, 06:02:03- 06:02:05, 06:02:07- 06:02:11, 06:02:13- 06:02:19, 06:04-06:10, 06:12-06:51, 06:53- 06:69, 15:23, 18:01- 18:05, B*08:17, B*08:39, B*08:47
33 ²⁶	205 bp, 230 bp	1070 bp	*07:66, 07:82	
34 ^{5,27}	105 bp, 270 bp	1070 bp	*07:67, 07:70	
35 ^{6,28}	320 bp, 360 bp	1070 bp	*07:71-07:72	
36 ⁶	265 bp	1070 bp	*07:20, 07:64, 07:73, 07:92, 07:96, 07:172	*18:04
37	265 bp	1070 bp	*07:38, 07:63, 07:68	*08:38, B*07:77, B*40:60
38 ^{9,29}	135 bp, 195 bp, 405 bp	1070 bp	*07:58, 07:61N, 07:65	
39 ^{5,30}	100 bp, 255 bp	800 bp	*07:59, 07:74	
40 ³¹	205 bp, 440 bp	1070 bp	*07:07, 07:09, 07:62, 07:76, 07:78	*06:17, 18:01-18:05
41 ³²	140 bp, 255 bp	1070 bp	*07:88, 07:91	
42 ^{5,33}	100 bp, 205 bp, 250 bp	1070 bp	*07:15, 07:77, 07:93, 07:123, 07:173, 07:207	B*35:08:04
43 ^{5,6,34}	105 bp, 140 bp, 175 bp	1070 bp	*07:80, 07:94, 07:99, 07:183	
44 ^{6,35}	165 bp, 195 bp	1070 bp	*07:65, 07:77, 07:98N	
45	475 bp	1070 bp	*07:95	*04:34, B*35:178, B*73:01-73:02
46 ³⁶	200 bp, 315 bp, 350 bp, 420 bp	1070 bp	*07:83, 07:90, 07:152N, 07:202	
47 ³⁷	200 bp, 420 bp, 455 bp	1070 bp	*07:86, 07:97, 07:152N	
48 ³⁸	220 bp, 400 bp	1070 bp	*07:85, 07:87	

101.615-24/04 – including *Taq* pol., IFU-01
 101. 615-24u/04u – without *Taq* pol., IFU-02

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Lot No.: 66N		Lot-specific information		
49 ³⁹	155 bp, 220 bp	800 bp	*07:100-07:101, 07:148, 07:161	*05:42, 05:46, 06:67, 08:05, 08:21, 17:05
50 ^{5,40}	105 bp, 170 bp, 425 bp	1070 bp	*07:104N, 07:119, 07:124, 07:155	B*07:87, B*27:55
51 ^{5,41}	65 bp, 235 bp, 285 bp	1070 bp	*07:117, 07:121Q, 07:126	
52 ^{5,42}	65 bp, 205 bp, 280 bp	1070 bp	*07:106, 07:118, 07:203	
53 ^{5,43}	110 bp, 155 bp	1070 bp	*07:122, 07:140, 07:151, 07:154	B*37:04:02
54 ^{5,44}	120 bp, 235 bp, 340 bp, 370 bp	1070 bp	*07:64, 07:125, 07:129, 07:153	
55 ⁴⁵	150 bp, 290 bp	1070 bp	*07:132, 07:179	*06:07
56 ⁴⁶	210 bp, 255 bp	1070 bp	*07:123, 07:143, 07:150Q, 07:173	B*35:08:04
57	175 bp	800 bp	*07:130	*05:01:17, 05:09:02, 05:17, 08:15:02
58 ^{5,47}	80 bp, 450 bp,	1070 bp	*07:137:01-07:137:02, 07:164N, 07:186, 07:212	
59 ⁶	200 bp	1070 bp	*07:81, 07:145, 07:168, 07:191N	B*35:08:04, B*37:04:02
60 ⁴⁸	130 bp, 305 bp, 370 bp	1070 bp	*07:147, 07:165, 07:180	*03:04:19, 03:113, 05:01:17, 05:09:02, 08:15:02, 12:03:10
61	245 bp	1070 bp	*07:120	
62 ⁴⁹	200 bp, 280 bp	1070 bp	*07:84, 07:106, 07:205	
63 ⁵⁰	275 bp, 320 bp	800 bp	*07:24, 07:198N, 07:202, 07:218	
64 ⁵¹	135 bp, 225 bp	1070 bp	*07:69, 07:192	

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-C*07 SSP typings.

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

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

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

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

101.615-24/04 – including Taq pol., IFU-01
101. 615-24u/04u – without Taq pol., IFU-02

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Lot No.: 66N

Lot-specific information

Some primers may give rise to primer oligomer artifacts. Sometimes this phenomenon is an inherit 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 two different control primer pairs give rise to either an internal positive control band of 1070 base pairs, for most wells, or a band of 800 base pairs, for some wells.

Well number 1 contains the primer pair giving rise to the shorter, 800 bp, internal positive control band in order to help in the correct orientation of the HLA-C*07 subtyping.

In addition, wells number 2, 3, 5, 9, 11, 14, 18, 19, 21, 22, 28, 39, 49, 57 and 63 contain the primer pair giving rise to the shorter, 800 bp, internal positive control band in order to allow kit identification.

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

³For several HLA-C*07 alleles 1st, 4th, 5th, 6th and 7th exon or 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. We assume that unknown sequences in the 1st, 4th, 5th, 6th and 7th exons and in the introns are conserved within allelic groups.

⁴Due to the sharing of sequence motifs between HLA-C alleles non-HLA-C*07 alleles will be amplified by primer mixes 2, 3, 6, 8 to 14, 16 to 18, 20, 22, 24, 26, 27, 32, 36, 37, 40, 45, 49, 55, 57, 60 and 61. In addition, the A*29:10 allele will be amplified by primer mix 12, the A*30:01:03 allele by primer mix 20, the A*33:03:09 allele will be amplified by primer mixes 20 and 24, the B*07:77 allele will be amplified by primer mixes 18 and 37, the B*07:87 and B*27:55 will be amplified by primer mix 50, the B*08:17, B*08:39 and B*08:47 alleles by primer mix 32, the B*27:36, B*48:21 and B*48:26 alleles by primer mix 7, the B*35:08:04 allele by primer mixes 2, 11, 20, 27, 42, 56 and 59, the B*35:178 and B*73:01-73:02 alleles will be amplified by primer mix 45, the B*37:04:02 allele by primer mixes 11, 14, 20, 53 and 59, the B*38:23 allele by primer mix 8, the B*40:60 allele by primer mixes 4, 18 and 37, the B*40:100 allele by primer mix 10, the B*46:06 allele by primer mix 16 and the B*56:08 and B*56:14 alleles by primer mix 25.

⁵Short specific PCR fragments are less intense and not as sharp as longer specific bands.

⁶Primer mixes 9, 35, 36, 43, 44 and 59 have a tendency of primer oligomer formation.

⁷Primer mixes 13 and 17 have a tendency of giving rise to nonspecific amplifications.

⁸In primer mix 5 the positive control band may be weaker than for other HLA-C*07 primer mixes.

⁹Primer mix 38 may generate a false band of about 900 base pairs. This band should be disregarded when interpreting HLA-C*07 SSP typings.

¹⁰Primer mix 1: Specific PCR fragment of 245 bp in the C*07:07, 07:09, 07:49 and 07:76, 07:115, and 07:210 alleles. Specific PCR fragment of 425 bp in the C*07:02:21, 07:41 and 07:166 alleles. Specific PCR fragment of 245 and 425 bp in the C*07:01:01-07:06, 07:08, 07:10-07:33N, *07:35-07:40, 07:42-07:48, 07:50-07:75, 07:77-07:114 and 07:211-07:220 alleles.

¹¹Primer mix 5: Specific PCR fragment of 70 bp in the C*07:53 and 07:216 alleles. Specific PCR fragment of 145 bp in the C*07:57 allele. Specific PCR fragment of 245 bp in the C*07:03 allele.

¹²Primer mix 8: Specific PCR fragment of 285 bp in the C*07:45 and the C*04:40 and in the B*38:23 alleles. Specific PCR fragment of 505 bp in the C*07:06 allele.

¹³Primer mix 9: Specific PCR fragment of 215 bp in the C*07:33N allele. Specific PCR fragment of 240 bp in the C*07:29, 07:177 and 07:184 and in the C*03:04:19 and 03:113 alleles. Specific PCR fragment of 425 bp in the C*07:07, 07:09, 07:49, 07:76 and 07:210 and in the C*06:17 and 18:01-18:05 alleles.

¹⁴Primer mix 11: Specific PCR fragment of 275 bp in the C*07:24 and 07:218 alleles. Specific PCR fragment of 300 bp in the C*07:10, 07:43, 07:112, 07:184 and 07:196 and the C*03:04:19 and 03:113 and in the B*35:08:04 and B*37:04:02 alleles.

¹⁵Primer mix 15: Specific PCR fragment of 80 bp in the C*07:50 allele. Specific PCR fragment of 290 bp in the C*07:14 allele. Specific PCR fragment of 460 bp in the C*07:89 allele.

¹⁶Primer mix 18: Specific PCR fragment of 120 bp in the C*07:52 and the C*05:03 alleles. Specific PCR fragment of 230 bp in the C*07:15 and in the B*07:77 and B*40:60 alleles.

¹⁷Primer mix 19: Specific PCR fragment of 220 bp in the C*07:26, 07:33N, 07:92 and 07:96 alleles. Specific PCR fragment of 355 bp in the C*07:80 allele. Specific PCR fragment of 465 bp in the C*07:07, 07:16, 07:51, 07:177 and 07:181 alleles.

**101.615-24/04 – including Taq pol., IFU-01
101. 615-24u/04u – without Taq pol., IFU-02**

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Lot No.: 66N

Lot-specific information

- ¹⁸Primer mix 20: Specific PCR fragment of 145 bp in the C*07:03, 07:17 and 07:140-07:141 and the C*03:04:19, 03:113, 05:01:17, 05:09:02, 05:17, 08:15:02 and 12:03:10 and in the A*30:01:03 and A*33:03:09 alleles. Specific PCR fragment of 170 bp in the 07:44 and 07:48 alleles. Specific PCR fragment of 195 bp in the C*07:81 allele. Specific PCR fragment of 145 bp and 195 bp in the B*35:08:04 and B*37:04:02 alleles.
- ¹⁹Primer mix 21: Specific PCR fragment of 100 bp in the C*07:54 allele. Specific PCR fragment of 225 bp in the C*07:06, 07:18 and 07:19 alleles.
- ²⁰Primer mix 22: Specific PCR fragment of 205 bp in the C*07:84 allele. Specific PCR fragment of 250 bp in the C*07:20, 07:96 and 07:127 and in the C*02:49, 03:15, 03:32, 03:45, 03:60, 03:136, 04:03, 04:06, 04:16, 04:80, 15:25 alleles.
- ²¹Primer mix 23: Specific PCR fragment of 155 bp in the C*07:23 allele. Specific PCR fragment of 185 bp in the C*07:21 allele.
- ²²Primer mix 25: Specific PCR fragment of 105 bp in the C*07:46 and B*56:06^w and B*56:14 alleles. Specific PCR fragment of 150 bp in the C*07:30 allele.
- ²³Primer mix 26: Specific PCR fragment of 100 bp in the C*07:32N allele. Specific PCR fragment of 250 bp in the C*07:55N allele. Specific PCR fragment of 300 bp in the C*07:31 and 07:177 and the C*12:03:10 alleles.
- ²⁴Primer mix 28: Specific PCR fragment of 175 bp in the C*07:44 allele. Specific PCR fragment of 260 bp in the C*07:35 and 07:47 alleles.
- ²⁵Primer mix 32: Specific PCR fragment of 245 bp in the C*07:07, 07:09, 07:49, 07:76 and 07:210 and the C*02:14, 04:42, 05:43, 06:02:01:01-06:02:01:02, 06:02:03-06:02:05, 06:02:07-06:02:11, 06:02:13-06:02:19, 06:04-06:10, 06:12-06:51, 06:53-06:69, 15:23 and 18:01-18:05 and in the B*08:17 and B*08:47 alleles. Specific PCR fragment of 285 bp in the C*07:60 and 07:79 and the C*03:39 and 03:67 and the B*08:39 alleles.
- ²⁶Primer mix 33: Specific PCR fragment of 205 bp in the C*07:66 allele. Specific PCR fragment of 230 bp in the C*07:82 allele.
- ²⁷Primer mix 34: Specific PCR fragment of 105 bp in the C*07:67 allele. Specific PCR fragment of 270 bp in the C*07:70 allele.
- ²⁸Primer mix 35: Specific PCR fragment of 320 bp in the C*07:72 allele. Specific PCR fragment of 360 bp in the C*07:71 allele.
- ²⁹Primer mix 38: Specific PCR fragment of 135 bp in the C*07:58 allele. Specific PCR fragment of 195 bp in the C*07:65 allele. Specific PCR fragment of 405 bp in the C*07:61N allele.
- ³⁰Primer mix 39: Specific PCR fragment of 100 bp in the C*07:74 allele. Specific PCR fragment of 255 bp in the C*07:59 allele.
- ³¹Primer mix 40: Specific PCR fragment of 205 bp in the C*07:62 allele. Specific PCR fragment of 440 bp in the C*07:07, 07:09, 07:76 and 07:78 and in the C*06:17 and 18:01-18:05 alleles.
- ³²Primer mix 41: Specific PCR fragment of 140 bp in the C*07:91 allele. Specific PCR fragment of 255 bp in the C*07:88 allele.
- ³³Primer mix 42: Specific PCR fragment of 100 bp in the C*07:15 and 07:207 alleles. Specific PCR fragment of 205 bp in the C*07:77 and 07:93 alleles. Specific PCR fragment of 250 bp in the C*07:123 and 07:173 and the B*35:08:04 alleles.
- ³⁴Primer mix 43: Specific PCR fragment of 105 bp in the C*07:94 and 07:183 alleles. Specific PCR fragment of 140 bp in the C*07:99 allele. Specific PCR fragment of 175 bp in the C*07:80 allele.
- ³⁵Primer mix 44: Specific PCR fragment of 165 bp in the C*07:98N allele. Specific PCR fragment of 195 bp in the C*07:65 and 07:77 alleles.
- ³⁶Primer mix 46: Specific PCR fragment of 200 bp in the C*07:83 allele. Specific PCR fragment of 315 bp in the C*07:202 allele. Specific PCR fragment of 350 bp in the C*07:90 allele. Specific PCR fragment of 420 bp in the C*07:152N allele.
- ³⁷Primer mix 47: Specific PCR fragment of 200 bp in the C*07:86 allele. Specific PCR fragment of 420 bp in the C*07:152N allele. Specific PCR fragment of 455 bp in the C*07:97 allele.
- ³⁸Primer mix 48: Specific PCR fragment of 220 bp in the C*07:87 allele. Specific PCR fragment of 400 bp in the C*07:85 allele.
- ³⁹Primer mix 49: Specific PCR fragment of 155 bp in the C*07:101, 07:148, 07:161, 05:42, 05:46, 06:67, 08:05, 08:21, 17:05 allele. Specific PCR fragment of 220 bp in the C*07:100 allele.



101.615-24/04 – including *Taq pol.*, IFU-01
101. 615-24u/04u – without *Taq pol.*, IFU-02

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Lot No.: 66N**Lot-specific information**

⁴⁰Primer mix 50: Specific PCR fragment of 105 bp in the C*07:104N allele. Specific PCR fragment of 170 bp in the C*07:119 and in the B*07:87 and B*27:55 alleles. Specific PCR fragment of 425 bp in the C*07:124 and 07:155 alleles.

⁴¹Primer mix 51: Specific PCR fragment of 65 bp in the C*07:126 allele. Specific PCR fragment of 235 bp in the C*07:121Q allele. Specific PCR fragment of 285 bp in the C*07:117 allele.

⁴²Primer mix 52: Specific PCR fragment of 65 bp in the C*07:203 allele. Specific PCR fragment of 205 bp in the C*07:118 allele. Specific PCR fragment of 280 bp in the C*07:106 allele.

⁴³Primer mix 53: Specific PCR fragment of 110 bp in the C*07:140, 07:151 and 07:154 and in the B*37:04:02 alleles. Specific PCR fragment of 155 bp in the C*07:122 allele.

⁴⁴Primer mix 54: Specific PCR fragment of 120 bp in the C*07:153 allele. Specific PCR fragment of 235 bp in the C*07:64 allele. Specific PCR fragment of 340 bp in the C*07:125 allele. Specific PCR fragment of 370 bp in the C*07:129 allele.

⁴⁵Primer mix 55: Specific PCR fragment of 150 bp in the C*07:179 and the C*06:07 alleles. Specific PCR fragment of 290 bp in the C*07:132 allele.

⁴⁶Primer mix 56: Specific PCR fragment of 210 bp in the C*07:150Q allele. Specific PCR fragment of 255 bp in the C*07:123, 07:143 and 07:173 and the B*35:08:04 alleles.

⁴⁷Primer mix 58: Specific PCR fragment of 80 bp in the C*07:137:01-07:137:02, 07:186 and 07:212 alleles. Specific PCR fragment of 450 bp in the C*07:164N allele.

⁴⁸Primer mix 60: Specific PCR fragment of 130 bp in the C*07:165 and the C*03:04:19, 03:113, 05:01:17, 05:09:02, 08:15:02 and 12:03:10 alleles. Specific PCR fragment of 305 bp in the C*07:147 allele. Specific PCR fragment of 370 bp in the C*07:180 allele.

⁴⁹Primer mix 62: Specific PCR fragment of 200 bp in the C*07:84 and 07:205 alleles. Specific PCR fragment of 280 bp in the C*07:106 allele.

⁵⁰Primer mix 63: Specific PCR fragment of 275 bp in the C*07:24 and 07:218 alleles. Specific PCR fragment of 320 bp in the C*07:198N and 07:202 alleles.

⁵¹Primer mix 64: Specific PCR fragment of 135 bp in the C*07:69 allele. Specific PCR fragment of 225 bp in the C*07:192 allele.

‘w’, may be weakly amplified.

101.615-24/04 – including Taq pol., IFU-01
101.615-24u/04u – without Taq pol., IFU-02

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Lot No.: 66N

Lot-specific information

Length of spec. PCR product(s)	Length of int. pos. control ¹	5'-primer(s) ²	Well ¹⁶											
			1	2	3	4	5	6	7	8	9	10	11	12
5' -CAT ³ 853	5' -ggC ³ 302	5' -CAC ³ 648	5' -Agg ³ 47	800	425	245								
5' -CT ³ 618		5' -gTA ³ 368	800	290										
5' -Cgt ³ 341		5' -AAC ³ 270	800	110	3									
5' -CCA ³ 512		5' -gTC ³ 368	5' -gTC ³ 368	1070	185									
5' -TCT ³ 601	5' -ggC ³ 121	5' -gTT ³ 512	5' -Agg ³ 47	800	245	145	70	5	6					
5' -CCA ³ 512	5' -CTT ³ 601		5' -gAC ³ 540	1070	100									
5' -CAC ³ 1016	5' -CTC ³ 601		5' -AAT ³ 363	1070	190	7								
5' -CT ³ 618	5' -ggT ³ 302	5' -TA ³ 992	5' -CCT ³ 355	1070	505	285	8							
	5' -Ag ³ 90	5' -gTA ³ 419	5' -Agg ³ 47	800	425	240	215	9	10					
5' -CCA ³ 512			5' -gTT ³ 368	1070	185									
5' -CT ³ 618	5' -ggT ³ 385	5' -gAC ³ 365	5' -TCA ³ 355	800	300	275		11	12					
5' -CTT ³ 601			5' -TCA ³ 355	800	300	275								
5' -AgC ³ 1087	5' -g ³ 343	5' -T ³ 343	5' -g ³ 1049	5' -Agg ³ 289	1070	90								
5' -AgT ³ 1087	5' -TgT ³ 616	5' -CT ³ 618	5' -C ³ 1049	5' -TAC ³ 369	5' -Agg ³ 47	1070	240	14						
5' -TgA ³ 668	5' -CTC ³ 559			5' -Agg ³ 270	1070	535								
5' -Cgg ³ 341	5' -CCC ³ 233	5' -gTA ³ 97	5' -Ag ³ 90	5' -CT ³ 618	5' -gTg ³ 512	5' -C ³ 512	5' -gTT ³ 512	17	18					
						5' -Cg ³ 3 rd I	5' -gTC ³ 368	800	230	120				
						5' -gTC ³ 368	5' -Agg ³ 47	800	465	355				
							5' -AgA ³ 463	1070	195	170	145			
								800	465	355	220			
									21	22	23	24	25	100
5' -CAA ³ 1043	5' -gTg ³ 427						5' -gTC ³ 368	800	225	205	185	155	125	
5' -CT ³ 618	5' -Cgt ³ 341						5' -gTC ³ 368	800	250	205	185	155	125	
5' -CCC ³ 527	5' -TAC ³ 270	5' -gTg ³ 512	5' -gTg ³ 427	5' -gTC ³ 419	5' -AgA ³ 206	1070	1070	1070	1070	1070	1070	1070	1070	255
5' -CT ³ 618	5' -Cgt ³ 341	5' -AgC ³ 289	5' -AgT ³ 289	5' -CAC ³ 560	5' -TAC ³ 411	5' -AgT ³ 361	1070	300	250	100				
5' -Tgg ³ 341	5' -ggT ³ 302			5' -AgT ³ 289	5' -gCg ³ 539	5' -CCA ³ 193	1070	1070	1070	1070	1070	1070	1070	60
							5' -AgA ³ 390	800	260	175				
								800	260	175				
									29	30	31	32	31	32



101.615-24/04 – including *Taq* pol., IFU-01
 101. 615-24u/04u – without *Taq* pol., IFU-02

Visit www.olerup-ssp.com for
 “Instructions for Use” (IFU)

Lot No.: 66N

Lot-specific information

INTERPRETATION TABLE																
HLA-C*07 SSP subtyping																
Amplification patterns of the C*07:01 to C*07:220 alleles																
33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	Well ¹⁶
33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48
33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49 50 51 52
33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49 50 51 52 53 54 55 56
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
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
33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	Well No.



**101.615-24/04 – including Taq pol., IFU-01
101.615-24u/04u – without Taq pol., IFU-02**

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Lot No.: 66N

Lot-specific information

Lot-specific information																	Length of spec.
																	PCR product(s)
230	205	270	105	360	320	265	265	405	195	135	255	100	440	205	255	140	
250	205	100	175	140	105	195	165	420	350	315	200	455	420	200	400	220	
49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64		Well No.
41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	HLA-C allele^{4,5}
33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	*07:01:01-07:01:04, 07:01:06-07:01:24, 07:103, 07:111, 07:113, 07:115-07:116, 07:128, 07:131, 07:134, 07:156, 07:162-07:163, 07:166, 07:170, 07:182, 07:188- 07:190, 07:197, 07:200- 07:201, 07:204, 07:206, 07:214-07:215, 07:219 *07:01:05, 07:110
37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	*07:02:01-01-07:02:29, 07:102, 07:105, 07:107, 07:114, 07:133, 07:135- 07:136, 07:144, 07:146, 07:149, 07:157-07:160, 07:167, 07:169, 07:174- 07:175, 07:178, 07:185, 07:187, 07:193-07:195, 07:208-07:209, 07:211, 07:213, 07:217, 07:220 *07:03
38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	*07:04:01-07:04:07, 07:139, 07:142, 07:199 *07:05
39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	*07:06
40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	*07:07
41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	*07:08
42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	*07:09
43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	*07:10
44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	*07:11
45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	*07:12
46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	*07:13
47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	*07:14
48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	*07:15
49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	*07:16
50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	*07:17
51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	*07:18
52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	*07:19
53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	*07:20
54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	*07:21
55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	*07:22
56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	*07:23
57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	*07:24
58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	*07:25, 07:138
59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	*07:26
60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	



101.615-24/04 – including Taq pol., IFU-01
101. 615-24u/04u – without Taq pol., IFU-02

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Lot No.: 66N

Lot-specific information

¹The internal positive control primer pairs amplify segments of the human growth hormone gene. The two different control primer pairs give rise to either an internal positive control band of 1070 base pairs, for most wells, or a band of 800 base pairs, for some wells.

Well number 1 contains the primer pair giving rise to the shorter, 800 bp, internal positive control band in order to help in the correct orientation of the HLA-C*07 subtyping.

In addition, wells number 2, 3, 5, 9, 11, 14, 18, 19, 21, 22, 28, 39, 49, 57 and 63 contain the primer pair giving rise to the shorter, 800 bp, internal positive control band in order to allow kit identification. In the presence of a specific amplification the intensity of the control band often decreases.

²The nucleotide position, in the 1st, 2nd, 3rd, 4th, 5th or 7th exon or the 3rd or 6th intron, 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, in the 2nd, 3rd, 4th, 6th or 7th exon or the 3rd intron, 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.

⁴The HLA-Cw*0734 allele has been renamed to HLA-C*07:27:02.

⁵HLA-C*07 alleles in bold lettering are listed as confirmed alleles on the IMGT/HLA web page www.ebi.ac.uk/imgt/hla, release 3.7.0, January 2012.

⁶The C*07:93 and 07:207 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 42.

⁷The C*07:100 and 07:161 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 49.

⁸The C*07:104N, 07:119 and 07:124 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 50.

⁹The C*07:117, 07:121Q and 07:126 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 51.

¹⁰The C*07:118 and 07:203 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 52.

¹¹The C*07:122 and 07:151 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 53.

¹²The C*07:129 and 07:153 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 54.

¹³The C*07:132 and 07:179 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 55.

¹⁴The C*07:164N and 07:212 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 58.

¹⁵The C*07:165 and 07:180 alleles can be distinguished by the different sizes of the specific PCR products generated by primer mix 60.

¹⁶Primer mix 1: Specific PCR fragment of 245 bp in the C*07:07, 07:09, 07:49 and 07:76, 07:115, and 07:210 alleles. Specific PCR fragment of 425 bp in the C*07:02:21, 07:41 and 07:166 alleles. Specific PCR fragment of 245 and 425 bp in the C*07:01:01-07:06, 07:08, 07:10-07:33N, *07:35-07:40, 07:42-07:48, 07:50-07:75, 07:77-07:114 and 07:211-07:220 alleles.

Primer mix 5: Specific PCR fragment of 70 bp in the C*07:53 and 07:216 alleles. Specific PCR fragment of 145 bp in the C*07:57 allele. Specific PCR fragment of 245 bp in the C*07:03 allele.

Primer mix 8: Specific PCR fragment of 285 bp in the C*07:45 and the C*04:40 and in the B*38:23alleles. Specific PCR fragment of 505 bp in the C*07:06 allele.

Primer mix 9: Specific PCR fragment of 215 bp in the C*07:33N allele. Specific PCR fragment of 240 bp in the C*07:29, 07:177 and 07:184 and in the C*03:04:19 and 03:113 alleles. Specific PCR fragment of 425 bp in the C*07:07, 07:09, 07:49, 07:76 and 07:210 and in the C*06:17 and 18:01-18:05 alleles.

Primer mix 11: Specific PCR fragment of 275 bp in the C*07:24 and 07:218 alleles. Specific PCR fragment of 300 bp in the C*07:10, 07:43, 07:112, 07:184 and 07:196 and the C*03:04:19 and 03:113 and in the B*35:08:04 and B*37:04:02 alleles.

Primer mix 15: Specific PCR fragment of 80 bp in the C* 07:50 allele. Specific PCR fragment of 290 bp in the C*07:14 allele. Specific PCR fragment of 460 bp in the C*07:89 allele.

Primer mix 18: Specific PCR fragment of 120 bp in the C*07:52 and the C*05:03 alleles. Specific PCR fragment of 230 bp in the C*07:15 and in the B*07:77 and B*40:60 alleles.

**101.615-24/04 – including Taq pol., IFU-01
101. 615-24u/04u – without Taq pol., IFU-02**

Visit www.olerup-ssp.com for
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Lot No.: 66N

Lot-specific information

Primer mix 19: Specific PCR fragment of 220 bp in the C*07:26, 07:33N, 07:92 and 07:96 alleles. Specific PCR fragment of 355 bp in the C*07:80 allele. Specific PCR fragment of 465 bp in the C*07:07, 07:16, 07:51, 07:177 and 07:181 alleles.

Primer mix 20: Specific PCR fragment of 145 bp in the C*07:03, 07:17 and 07:140-07:141 and the C*03:04:19, 03:113, 05:01:17, 05:09:02, 05:17, 08:15:02 and 12:03:10 and in the A*30:01:03 and A*33:03:09 alleles. Specific PCR fragment of 170 bp in the 07:44 and 07:48 alleles. Specific PCR fragment of 195 bp in the C*07:81 allele. Specific PCR fragment of 145 bp and 195 bp in the B*35:08:04 and B*37:04:02 alleles.

Primer mix 21: Specific PCR fragment of 100 bp in the C*07:54 allele. Specific PCR fragment of 225 bp in the C*07:06, 07:18 and 07:19 alleles.

Primer mix 22: Specific PCR fragment of 205 bp in the C*07:84 allele. Specific PCR fragment of 250 bp in the C*07:20, 07:96 and 07:127 and in the C*02:49, 03:15, 03:32, 03:45, 03:60, 03:136, 04:03, 04:06, 04:16, 04:80, 15:25 alleles.

Primer mix 23: Specific PCR fragment of 155 bp in the C*07:23 allele. Specific PCR fragment of 185 bp in the C*07:21 allele.

Primer mix 25: Specific PCR fragment of 105 bp in the C*07:46 and B*56:06^w and B*56:14 alleles. Specific PCR fragment of 150 bp in the C*07:30 allele.

Primer mix 26: Specific PCR fragment of 100 bp in the C*07:32N allele. Specific PCR fragment of 250 bp in the C*07:55N allele. Specific PCR fragment of 300 bp in the C*07:31 and 07:177 and the C*12:03:10 alleles.

Primer mix 28: Specific PCR fragment of 175 bp in the C*07:44 allele. Specific PCR fragment of 260 bp in the C*07:35 and 07:47 alleles.

Primer mix 32: Specific PCR fragment of 245 bp in the C*07:07, 07:09, 07:49, 07:76 and 07:210 and the C*02:14, 04:42, 05:43, 06:02:01:01-06:02:01:02, 06:02:03-06:02:05, 06:02:07-06:02:11, 06:02:13-06:02:19, 06:04-06:10, 06:12-06:51, 06:53-06:69, 15:23 and 18:01-18:05 and in the B*08:17 and B*08:47 alleles. Specific PCR fragment of 285 bp in the C*07:60 and 07:79 and the C*03:39 and 03:67 and the B*08:39 alleles.

Primer mix 33: Specific PCR fragment of 205 bp in the C*07:66 allele. Specific PCR fragment of 230 bp in the C*07:82 allele.

Primer mix 34: Specific PCR fragment of 105 bp in the C*07:67 allele. Specific PCR fragment of 270 bp in the C*07:70 allele.

Primer mix 35: Specific PCR fragment of 320 bp in the C*07:72 allele. Specific PCR fragment of 360 bp in the C*07:71 allele.

Primer mix 38: Specific PCR fragment of 135 bp in the C*07:58 allele. Specific PCR fragment of 195 bp in the C*07:65 allele. Specific PCR fragment of 405 bp in the C*07:61N allele.

Primer mix 39: Specific PCR fragment of 100 bp in the C*07:74 allele. Specific PCR fragment of 255 bp in the C*07:59 allele.

Primer mix 40: Specific PCR fragment of 205 bp in the C*07:62 allele. Specific PCR fragment of 440 bp in the C*07:07, 07:09, 07:76 and 07:78 and in the C*06:17 and 18:01-18:05 alleles.

Primer mix 41: Specific PCR fragment of 140 bp in the C*07:91 allele. Specific PCR fragment of 255 bp in the C*07:88 allele.

Primer mix 42: Specific PCR fragment of 100 bp in the C*07:15 and 07:207 alleles. Specific PCR fragment of 205 bp in the C*07:77 and 07:93 alleles. Specific PCR fragment of 250 bp in the C*07:123 and 07:173 and the B*35:08:04 alleles.

Primer mix 43: Specific PCR fragment of 105 bp in the C*07:94 and 07:183 alleles. Specific PCR fragment of 140 bp in the C*07:99 allele. Specific PCR fragment of 175 bp in the C*07:80 allele.

Primer mix 44: Specific PCR fragment of 165 bp in the C*07:98N allele. Specific PCR fragment of 195 bp in the C*07:65 and 07:77 alleles.

Primer mix 46: Specific PCR fragment of 200 bp in the C*07:83 allele. Specific PCR fragment of 315 bp in the C*07:202 allele. Specific PCR fragment of 350 bp in the C*07:90 allele. Specific PCR fragment of 420 bp in the C*07:152N allele.

Primer mix 47: Specific PCR fragment of 200 bp in the C*07:86 allele. Specific PCR fragment of 420 bp in the C*07:152N allele. Specific PCR fragment of 455 bp in the C*07:97 allele.

Primer mix 48: Specific PCR fragment of 220 bp in the C*07:87 allele. Specific PCR fragment of 400 bp in the C*07:85 allele.

Primer mix 49: Specific PCR fragment of 155 bp in the C*07:101, 07:148, 07:161, 05:42, 05:46, 06:67, 08:05, 08:21, 17:05 allele. Specific PCR fragment of 220 bp in the C*07:100 allele.



101.615-24/04 – including *Taq pol.*, IFU-01
101. 615-24u/04u – without *Taq pol.*, IFU-02

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Lot No.: 66N**Lot-specific information**

Primer mix 50: Specific PCR fragment of 105 bp in the C*07:104N allele. Specific PCR fragment of 170 bp in the C*07:119 and in the B*07:87 and B*27:55 alleles. Specific PCR fragment of 425 bp in the C*07:124 and 07:155 alleles.

Primer mix 51: Specific PCR fragment of 65 bp in the C*07:126 allele. Specific PCR fragment of 235 bp in the C*07:121Q allele. Specific PCR fragment of 285 bp in the C*07:117 allele.

Primer mix 52: Specific PCR fragment of 65 bp in the C*07:203 allele. Specific PCR fragment of 205 bp in the C*07:118 allele. Specific PCR fragment of 280 bp in the C*07:106 allele.

Primer mix 53: Specific PCR fragment of 110 bp in the C*07:140, 07:151 and 07:154 and in the B*37:04:02 alleles. Specific PCR fragment of 155 bp in the C*07:122 allele.

Primer mix 54: Specific PCR fragment of 120 bp in the C*07:153 allele. Specific PCR fragment of 235 bp in the C*07:64 allele. Specific PCR fragment of 340 bp in the C*07:125 allele. Specific PCR fragment of 370 bp in the C*07:129 allele.

Primer mix 55: Specific PCR fragment of 150 bp in the C*07:179 and the C*06:07 alleles. Specific PCR fragment of 290 bp in the C*07:132 allele.

Primer mix 56: Specific PCR fragment of 210 bp in the C*07:150Q allele. Specific PCR fragment of 255 bp in the C*07:123, 07:143 and 07:173 and the B*35:08:04 alleles.

Primer mix 58: Specific PCR fragment of 80 bp in the C*07:137:01-07:137:02, 07:186 and 07:212 alleles. Specific PCR fragment of 450 bp in the C*07:164N allele.

Primer mix 60: Specific PCR fragment of 130 bp in the C*07:165 and the C*03:04:19, 03:113, 05:01:17, 05:09:02, 08:15:02 and 12:03:10 alleles. Specific PCR fragment of 305 bp in the C*07:147 allele. Specific PCR fragment of 370 bp in the C*07:180 allele.

Primer mix 62: Specific PCR fragment of 200 bp in the C*07:84 and 07:205 alleles. Specific PCR fragment of 280 bp in the C*07:106 allele.

Primer mix 63: Specific PCR fragment of 275 bp in the C*07:24 and 07:218 alleles. Specific PCR fragment of 320 bp in the C*07:198N and 07:202 alleles.

Primer mix 64: Specific PCR fragment of 135 bp in the C*07:69 allele. Specific PCR fragment of 225 bp in the C*07:192 allele.

¹⁷This lot of the HLA-C*07 subtyping kit cannot distinguish the C*07:198N and the C*07:218 alleles.

'w', may be weakly amplified.

101.615-24/04 – including *Taq pol.*, IFU-01
101. 615-24u/04u – without *Taq pol.*, IFU-02

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Lot No.: 66N

Lot-specific information

CELL LINE VALIDATION SHEET																	
HLA-C*07 SSP subtyping kit																	
Prod. No.:		Well															
		17	18	19	20	21	22	23	24	25	26	27	28	29	30		
		201069817	201069818	201069819	201200520	201069821	201069822	201184123	201189524	201069825	201069826	201200527	201189528	201189529	201189530	201200531	201069832
IHWC cell line	C*																
1	9001	SA	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	
2	9280	LK707	*07:01	*15:05	-	-	-	-	-	-	-	-	-	-	-	-	
3	9011	E4181324	*12:02		-	-	-	-	-	-	-	-	-	-	-	-	
4	9275	GU373	*03:04	*04:01	-	-	-	-	-	-	-	-	-	-	-	-	
5	9009	KAS011	*06:02		-	-	-	-	-	-	-	-	-	-	-	+	
6	9353	SM	*03:04	*07:02	-	-	-	-	-	-	-	-	-	-	-	-	
7	9020	QBL	*05:01		-	-	-	-	-	-	-	-	-	-	-	-	
8	9025	DEU	*04:01		-	-	-	-	-	-	-	-	-	-	-	-	
9	9026	YAR	*12:03		-	-	-	-	-	-	-	-	-	-	-	-	
10	9107	LKT3	*01:02		-	-	-	-	-	-	-	-	-	-	-	-	
11	9051	PITOUT	*16:01		-	-	-	-	-	-	-	-	-	-	-	-	
12	9052	DBB	*06:02		-	-	-	-	-	-	-	-	-	-	-	+	
13	9004	JESTHOM	*01:02		-	-	-	-	-	-	-	-	-	-	-	-	
14	9071	OLGA	*01:02	*03:04	-	-	-	-	-	-	-	-	-	-	-	-	
15	9075	DKB	*03:04		-	-	-	-	-	-	-	-	-	-	-	-	
16	9037	SWEIG007	*02:02		-	-	-	-	-	-	-	-	-	-	-	-	
17	9282	CTM3953540	*03:03	*07:01	-	-	-	-	-	-	-	-	-	-	-	-	
18	9257	32367	*01:02	*07:05	-	-	-	-	-	-	-	-	-	-	-	-	
19	9038	BM16	*07:01		-	-	-	-	-	-	-	-	-	-	-	-	
20	9059	SLE005	*03:04		-	-	-	-	-	-	-	-	-	-	-	-	
21	9064	AMALA	*03:03		-	-	-	-	-	-	-	-	-	-	-	-	
22	9056	KOSE	*12:03		-	-	-	-	-	-	-	-	-	-	-	-	
23	9124	IHL	*01:02	*15:02	-	-	-	-	-	-	-	-	-	-	-	-	
24	9035	JBUSH	*12:03		-	-	-	-	-	-	-	-	-	-	-	-	
25	9049	IBW9	*08:02		-	-	-	-	-	-	-	-	-	-	-	-	
26	9285	WT49	*07:01		-	-	-	-	-	-	-	-	-	-	-	-	
27	9191	CH1007	*07:04	*15:05	+	-	-	-	-	-	-	-	-	-	-	-	
28	9320	BEL5GB	*05:01	*16:01	-	-	-	-	-	-	-	-	-	-	-	-	
29	9050	MOU	*16:01		-	-	-	-	-	-	-	-	-	-	-	-	
30	9021	RSH	*17:01		-	-	-	-	-	-	-	-	-	-	-	-	
31	9019	DUCAF	*05:01		-	-	-	-	-	-	-	-	-	-	-	-	
32	9297	HAG	*17:01	*17:03	-	-	-	-	-	-	-	-	-	-	-	-	
33	9098	MT14B	*03:04		-	-	-	-	-	-	-	-	-	-	-	-	
34	9104	DHIF	*12:03		-	-	-	-	-	-	-	-	-	-	-	-	
35	9302	SSTO	*05:01		-	-	-	-	-	-	-	-	-	-	-	-	
36	9024	KT17	*03:03	*04:01	-	-	-	-	-	-	-	-	-	-	-	-	
37	9065	HHKB	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	
38	9099	LZL	*03:03		-	-	-	-	-	-	-	-	-	-	-	-	
39	9315	CML	*02:02	*07:01	-	-	-	-	-	-	-	-	-	-	-	-	
40	9134	WHONP199	*01:02	*06:02	-	-	-	-	-	-	-	-	-	-	-	+	
41	9055	H0301	*08:02		-	-	-	-	-	-	-	-	-	-	-	-	
42	9066	TAB089	*01:02		-	-	-	-	-	-	-	-	-	-	-	-	
43	9076	T7526	*01:02	*08:01	-	-	-	-	-	-	-	-	-	-	-	-	
44	9057	TEM	*12:03		-	-	-	-	-	-	-	-	-	-	-	-	
45	9239	SHJO	*06:02	*17:01	-	-	-	-	-	-	-	-	-	-	-	+	
46	9013	SCHU	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	
47	9045	TUBO	*07:04	*15:02	+	-	-	-	-	-	-	-	-	-	-	-	
48	9303	TER-ND	*04:01	*16:01	-	-	-	-	-	-	-	-	-	-	-	-	

For *In Vitro Diagnostic Use*

101.615-24/04 – including *Taq* pol., IFU-01
101. 615-24u/04u – without *Taq* pol., IFU-02

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“Instructions for Use” (IFU)

Lot No.: 66N

Lot-specific information

CELL LINE VALIDATION SHEET																
HLA-C*07 SSP subtyping kit																
Prod. No.:	Well															
	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64
IHWG cell line	C*															
1	9001	SA	*07:02	-	-	-	-	-	-	-	-	-	-	-	-	-
2	9280	LK707	*07:01	*15:05	-	-	-	-	-	-	-	-	-	-	-	-
3	9011	E4181324	*12:02	-	-	-	-	-	-	-	-	-	-	-	-	-
4	9275	GU373	*03:04	*04:01	-	-	-	-	-	-	-	-	-	-	-	-
5	9009	KAS011	*06:02	-	-	-	-	-	-	-	-	-	-	-	-	-
6	9353	SM	*03:04	*07:02	-	-	-	-	-	-	-	-	-	-	-	-
7	9020	QBL	*05:01	-	-	-	-	-	-	-	-	-	-	-	-	-
8	9025	DEU	*04:01	-	-	-	-	-	-	-	-	-	-	-	-	-
9	9026	YAR	*12:03	-	-	-	-	-	-	-	-	-	-	-	-	-
10	9107	LKT3	*01:02	-	-	-	-	-	-	-	-	-	-	-	-	-
11	9051	PITOUT	*16:01	-	-	-	-	-	-	-	-	-	-	-	-	-
12	9052	DBB	*06:02	-	-	-	-	-	-	-	-	-	-	-	-	-
13	9004	JESTHOM	*01:02	-	-	-	-	-	-	-	-	-	-	-	-	-
14	9071	OLGA	*01:02	*03:04	-	-	-	-	-	-	-	-	-	-	-	-
15	9075	DKB	*03:04	-	-	-	-	-	-	-	-	-	-	-	-	-
16	9037	SWEIG007	*02:02	-	-	-	-	-	-	-	-	-	-	-	-	-
17	9282	CTM3953540	*03:03	*07:01	-	-	-	-	-	-	-	-	-	-	-	-
18	9257	32367	*01:02	*07:05	-	-	-	-	-	-	-	-	-	-	-	-
19	9038	BM16	*07:01	-	-	-	-	-	-	-	-	-	-	-	-	-
20	9059	SLE005	*03:04	-	-	-	-	-	-	-	-	-	-	-	-	-
21	9064	AMALA	*03:03	-	-	-	-	-	-	-	-	-	-	-	-	-
22	9056	KOSE	*12:03	-	-	-	-	-	-	-	-	-	-	-	-	-
23	9124	IHL	*01:02	*15:02	-	-	-	-	-	-	-	-	-	-	-	-
24	9035	JBUSH	*12:03	-	-	-	-	-	-	-	-	-	-	-	-	-
25	9049	IBW9	*08:02	-	-	-	-	-	-	-	-	-	-	-	-	-
26	9285	WT49	*07:01	-	-	-	-	-	-	-	-	-	-	-	-	-
27	9191	CH1007	*07:04	*15:05	-	-	-	-	-	-	-	-	-	-	-	-
28	9320	BEL5GB	*05:01	*16:01	-	-	-	-	-	-	-	-	-	-	-	-
29	9050	MOU	*16:01	-	-	-	-	-	-	-	-	-	-	-	-	-
30	9021	RSH	*17:01	-	-	-	-	-	-	-	-	-	-	-	-	-
31	9019	DUCAF	*05:01	-	-	-	-	-	-	-	-	-	-	-	-	-
32	9297	HAG	*17:01	*17:03	-	-	-	-	-	-	-	-	-	-	-	-
33	9098	MT14B	*03:04	-	-	-	-	-	-	-	-	-	-	-	-	-
34	9104	DHIF	*12:03	-	-	-	-	-	-	-	-	-	-	-	-	-
35	9302	SSTO	*05:01	-	-	-	-	-	-	-	-	-	-	-	-	-
36	9024	KT17	*03:03	*04:01	-	-	-	-	-	-	-	-	-	-	-	-
37	9065	HHKB	*07:02	-	-	-	-	-	-	-	-	-	-	-	-	-
38	9099	LZL	*03:03	-	-	-	-	-	-	-	-	-	-	-	-	-
39	9315	CML	*02:02	*07:01	-	-	-	-	-	-	-	-	-	-	-	-
40	9134	WHONP199	*01:02	*06:02	-	-	-	-	-	-	-	-	-	-	-	-
41	9055	H0301	*08:02	-	-	-	-	-	-	-	-	-	-	-	-	-
42	9066	TAB089	*01:02	-	-	-	-	-	-	-	-	-	-	-	-	-
43	9076	T7526	*01:02	*08:01	-	-	-	-	-	-	-	-	-	-	-	-
44	9057	TEM	*12:03	-	-	-	-	-	-	-	-	-	-	-	-	-
45	9239	SHJO	*06:02	*17:01	-	-	-	-	-	-	-	-	-	-	-	-
46	9013	SCHU	*07:02	-	-	-	-	-	-	-	-	-	-	-	-	-
47	9045	TUBO	*07:04	*15:02	-	-	-	-	-	-	-	-	-	-	-	-
48	9303	TER-ND	*04:01	*16:01	-	-	-	-	-	-	-	-	-	-	-	-



101.615-24/04 – including *Taq* pol., IFU-01
 101.615-24u/04u – without *Taq* pol., IFU-02

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 “Instructions for Use” (IFU)

Lot No.: 66N

Lot-specific information

CERTIFICATE OF ANALYSIS**Olerup SSP® HLA-C*07 SSP**

Product number: 101.615-24/04 – including *Taq* pol.
 101.615-24u/04u – without *Taq* pol.

Lot number: 66N

Expiry date: 2014-October-01

Number of tests: 24 tests – Product No. 101.615-24
 4 tests – Product No. 101.615-04

Number of wells per test: 64

Well specifications:

Well No.	Production No.	Well No.	Production No.	Well No.	Production No.
1	2010-698-01	25	2010-698-25	49	2012-005-49
2	2011-895-02	26	2010-698-26	50	2012-005-50
3	2011-895-03	27	2012-005-27	51	2012-005-51
4	2010-698-04	28	2011-895-28	52	2012-005-52
5	2012-005-05	29	2011-895-29	53	2012-005-53
6	2010-698-06	30	2011-895-30	54	2012-005-54
7	2010-698-07	31	2012-005-31	55	2012-005-55
8	2010-698-08	32	2010-698-32	56	2012-005-56
9	2010-698-09	33	2011-895-33	57	2012-005-57
10	2010-698-10	34	2012-005-34	58	2012-005-58
11	2012-005-11	35	2012-005-35	59	2012-005-59
12	2010-698-12	36	2010-698-36	60	2012-005-60
13	2010-698-13	37	2010-698-37	61	2012-005-61
14	2010-698-14	38	2010-698-38	62	2012-005-62
15	2012-005-15	39	2010-698-39	63	2012-005-63
16	2012-005-16	40	2010-698-40	64	2012-005-64
17	2010-698-17	41	2010-698-41		
18	2010-698-18	42	2012-005-42		
19	2010-698-19	43	2012-005-43		
20	2012-005-20	44	2012-005-44		
21	2010-698-21	45	2010-698-45		
22	2010-698-22	46	2012-005-46		
23	2011-841-23	47	2012-005-47		
24	2011-895-24	48	2010-698-48		

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

No DNAs carrying the alleles to be amplified by primer solutions 5, 10, 11, 13 to 15, 18, 19, 23 to 31, 34 to 39, 41 to 44 and 46 to 64 were available.

The specificities of the primers in primer solutions 5, 10, 11, 13 to 15, 18, 19, 24 to 27, 31, 36 to 38, 42 to 44, 46 to 50, 53, 54, 56 to 60, 63 and 64 were tested by separately adding one 5'-primer, respectively one 3'-primer. In primer solutions 35, 51 and 61 it was only possible to test the 5'-primer, the 3'-primer

101.615-24/04 – including *Taq pol.*, IFU-01
101. 615-24u/04u – without *Taq pol.*, IFU-02

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Lot No.: 66N**Lot-specific information**

were not possible to test. In primer solutions 23, 28 to 30, 34, 39, 41, 52, 55 and 62 it was only possible to test the 3'-primers, the 5'-primers were not possible to test. In primer solutions 8, 11, 20, 22, 26, 33, 37, 38, 42, 44, 46 to 49, 53, 56, 59 and 63 one or two 5'-primers were not possible to test, and in primer solutions 5, 9, 13, 15, 18, 19, 21, 25, 32, 38, 40, 43, 46 to 48, 50, 54, 58, 60, 63 and 64 one or two 3'-primers were not possible to test.
Additional primers in primer solutions 2, 8, 9, 20, 21 and 33 were tested by separately adding either one 5'-primer or one 3'-primer.

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

Date of approval: 2012-May-11

Approved by:

Production Quality Control

101.615-24/04 – including *Taq* pol., IFU-01
101.615-24u/04u – without *Taq* pol., IFU-02

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Lot No.: 66N

Lot-specific information

Declaration of Conformity

Product name: Olerup SSP® HLA-C*07

Product number: 101.615-24/24u, -04/04u

Lot number: 66N

Intended use: HLA-C*07 high resolution histocompatibility testing

Manufacturer: Olerup SSP AB
Franzengatan 5
SE-112 51 Stockholm, Sweden
Phone: +46-8-717 88 27
Fax: +46-8-717 88 18

We, Olerup SSP AB, hereby declare that this product, to which this Declaration of Conformity relates is in conformity with the following Standard(s) and other normative document(s) ISO 9001:2008 and ISO 13485:2003, following the provisions of the 98/79/EC Directive on *in vitro* diagnostic medical devices, Annex III, as transposed into the national laws of the Member States of the European Union.

The Technical Documentation File is maintained at Olerup SSP AB,
Franzengatan 5, SE-112 51 Stockholm, Sweden.

Stockholm, Sweden
2012-May-11

Ann-Cathrin Jareman
Head of QA and Regulatory Affairs

101.615-24/04 – including *Taq* pol., IFU-01
101. 615-24u/04u – without *Taq* pol., IFU-02

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101. 615-24u/04u – without *Taq* pol., IFU-02

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Lot No.: 66N

Lot-specific information

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