

101.512-24/03 – including *Taq* pol., IFU-01
101.512-24u/03u – without *Taq* pol., IFU-02

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

Lot No.: **3D9**

Lot-specific information

Olerup SSP® HLA-B*07

Product number:	101.512-24/03 – including <i>Taq</i> pol. 101.512-24u/03u – without <i>Taq</i> pol.
Lot number:	3D9
Expiry date:	2018-07-01
Number of tests:	24 tests – Product No. 101.512-24/24u 3 tests – Product No. 101.512-03/03u
Number of wells per test:	87+1
Storage - pre-aliquoted primers:	dark at -20°C
- PCR Master Mix:	-20°C
- Adhesive PCR seals	RT
- Product Insert	RT

This Product Description is only valid for Lot No. 3D9.

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*07 LOT (84V)

The HLA-B*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

A well containing Negative Control primer pairs has been added.

The format of the Product Insert and Worksheet have been changed.

Four wells have been added to HLA-B*07, wells **85 to 88**.

¹As described in section Uniquely Identified Alleles.

The HLA-B*07 primer set, specificity and interpretation tables have been updated for the HLA-B alleles described since the previous *Olerup SSP®* HLA-B*07 lot was made (**Lot No. 84V**). The kit design is based on IMGT/HLA database 3.22.0.

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As of lot series V, the Specificity Table is included in the lot-specific Product Insert, and the Interpretation Table is included in the Worksheet.

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

Well	5'-primer	3'-primer	rationale
3	Added	-	5'-primer added for the B*07:02:46 allele.
35	Modified	-	5'-primer modified for improved HLA-specific amplification.
54	-	Added	3'-primer added for the B*07:238 allele.
69	-	Added	3'-primer added for the B*07:136:02 allele.
76	-	Added	3'-primer added for the B*07:136:02 allele.
78	-	Added	3'-primer added for the B*07:231N allele.
81	Exchanged	Exchanged	Exchanged primer pair for increased amplification.
84	Added	Added	Updated negative control moved to well 88, primer pair added for the B*07:248 allele.
85	New	New	New primer pairs added for the B*07:208 and B*07:238 alleles.
86	New	New	New primer pair added for the B*07:219 allele.
87	New	New	New primer pair added for the B*07:251N allele.
88	-	-	Updated negative control added from well 84.

Changes in revision R01 compared to R00:

1. Primer mix 84 does not amplify the B*07:248 allele. Thus, this lot of the B*07 kit cannot distinguish the B*07:163 and the B*07:248 alleles. This primer mix may also give rise to a lower yield of HLA-specific PCR product than the other HLA-B*07 high resolution primer mixes. This has been corrected in the Specificity and Interpretation Tables.

Changes in revision R02 compared to R01:

1. Primer mix 22 does not amplify the B*07:23 and the B*40:79, 40:190, 40:289 and 41:14 alleles. Thus, this lot of the B*07 kit cannot distinguish the B*07:23 and the B*07:02:01-07:02:25, 07:02:27, 07:02:29-07:02:48, 07:124, 07:127, 07:133-07:134, 07:141-07:142, 07:144-07:145, 07:150, 07:152, 07:155, 07:157-07:159, 07:166, 07:170-07:171, 07:173, 07:177-07:179, 07:185, 07:187-07:190, 07:192, 07:194-07:195, 07:200, 07:203, 07:205, 07:211-07:212, 07:215-07:218, 07:221, 07:224, 07:230, 07:233-07:234, 07:241-07:245, 07:247, 07:250, 07:252, 07:254 and 07:256 alleles. This has been corrected in the Specificity and Interpretation Tables.

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Well **88** contains Negative Control primer pairs, that will amplify more than 95% 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 430 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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Lot-specific information

PRODUCT DESCRIPTION

HLA-B*07 SSP subtyping

CONTENT

The primer set contains 5'- and 3'-primers for identifying the B*07:02 to B*07:256 alleles.

PLATE LAYOUT

Each test consists of 88 PCR reactions in a 96 well cut PCR plate. Wells 89 to 96 are empty.

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	NC
empty	empty	empty	empty	empty	empty	empty	empty

The 96 well cut PCR plate is marked with 'HLA-B*07' in silver/gray ink.

Well No. 1 is marked with the Lot Number '3D9'.

Wells 1 to 87 – HLA-B*07 high resolution primers.

Well 88 – 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*07 alleles will be amplified by primer mixes 1 to 20, 22 to 27, 29 to 33, 35, 37 to 40, 43 to 53, 56, 59, 60, 62 to 69, 72 to 75, 77, 81 to 84 and 86. In addition, a few HLA-A and HLA-C alleles will be amplified by primer mixes 3 to 5, 7, 9, 10, 12 to 14, 18, 20, 23, 26, 27, 29 to 32, 48, 53, 56, 59, 63, 72, 81, 82 and 86.

For further details see Specificity Table.

UNIQUELY IDENTIFIED ALLELES

All the HLA-B*07 alleles, i.e. **B*07:02 to B*07:256**, recognized by the HLA Nomenclature Committee in October 2015^{1,2} will be amplified by the primers in the HLA-B*07 subtyping kit^{3,4,5}.

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The HLA-B*07 kit enables separation of the confirmed HLA-B*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-B*07 alleles is listed below.

The HLA-B*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 following HLA-B*07 alleles can be distinguished by the different sizes of the HLA-specific PCR product:

Alleles	Primer mix	Alleles	Primer mix
B*07:42, 07:118	26	B*07:94, 07:135	63
B*07:49N, 07:169	73	B*07:102, 07:106	69
B*07:61, 07:149	33	B*07:107, 07:181N	71
B*07:63, 07:143	35	B*07:121, 07:126	75
B*07:89, 07:246	59	B*07:129, 07:231N	78

The HLA-B*07 subtyping kit cannot distinguish the following silent mutations: the B*07:02:01-07:02:25, 07:02:27 and 07:02:29-07:02:48, the B*07:05:01-07:05:08, the B*07:22:01-07:22:02, the B*07:33:01-07:33:03, the B*07:56:01-B*07:56:02, the B*07:68:01-07:68:03, the B*07:96:01-07:96:02, the 07:136:01-07:136:02 or the B*07:151:01-07:151:02 alleles.

¹HLA-B alleles listed on the IMGT/HLA web page 2015-October-10, release 3.22.0, www.ebi.ac.uk/imgt/hla.

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

³The B*07:50 and B*48:08 alleles will give rise to identical amplification patterns. These alleles can e.g. be distinguished by the HLA-B low resolution kit and/or the HLA-B*48 high resolution kit.

⁴This lot of the B*07 kit cannot distinguish the B*07:163 and the B*07:248 alleles.

⁵This lot of the B*07 kit cannot distinguish the B*07:23 and the B*07:02:01-07:02:25, 07:02:27, 07:02:29-07:02:48, 07:124, 07:127, 07:133-07:134, 07:141-07:142, 07:144-07:145, 07:150, 07:152, 07:155, 07:157-07:159, 07:166, 07:170-07:171, 07:173, 07:177-07:179, 07:185, 07:187-07:190, 07:192, 07:194-07:195, 07:200, 07:203, 07:205, 07:211-07:212, 07:215-07:218, 07:221, 07:224, 07:230, 07:233-07:234, 07:241-07:245, 07:247, 07:250, 07:252, 07:254 and 07:256 alleles.

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ALLELE CONFIRMATION STATUS

Allele	Status ¹	Allele	Status ¹	Allele	Status ¹	Allele	Status ¹
B*07:02:01	Confirmed	B*07:05:01	Confirmed	B*07:44	Unconfirmed	B*07:91	Confirmed
B*07:02:02	Confirmed	B*07:05:02	Unconfirmed	B*07:45	Unconfirmed	B*07:92	Confirmed
B*07:02:03	Confirmed	B*07:05:03	Unconfirmed	B*07:46	Unconfirmed	B*07:93	Unconfirmed
B*07:02:04	Unconfirmed	B*07:05:04	Unconfirmed	B*07:47	Confirmed	B*07:94	Confirmed
B*07:02:05	Unconfirmed	B*07:05:05	Confirmed	B*07:48	Confirmed	B*07:95	Confirmed
B*07:02:06	Unconfirmed	B*07:05:06	Unconfirmed	B*07:49N	Unconfirmed	B*07:96:01	Confirmed
B*07:02:07	Confirmed	B*07:05:07	Unconfirmed	B*07:50	Unconfirmed	B*07:96:02	Unconfirmed
B*07:02:08	Confirmed	B*07:05:08	Confirmed	B*07:51	Confirmed	B*07:97	Unconfirmed
B*07:02:09	Unconfirmed	B*07:06	Confirmed	B*07:52	Unconfirmed	B*07:98	Confirmed
B*07:02:10	Confirmed	B*07:07	Confirmed	B*07:53	Confirmed	B*07:99	Unconfirmed
B*07:02:11	Unconfirmed	B*07:08	Confirmed	B*07:54	Confirmed	B*07:100	Unconfirmed
B*07:02:12	Unconfirmed	B*07:09	Confirmed	B*07:55	Unconfirmed	B*07:101	Unconfirmed
B*07:02:13	Confirmed	B*07:10	Confirmed	B*07:56:01	Unconfirmed	B*07:102	Confirmed
B*07:02:14	Confirmed	B*07:11	Unconfirmed	B*07:56:02	Confirmed	B*07:103	Confirmed
B*07:02:15	Confirmed	B*07:12	Confirmed	B*07:57	Confirmed	B*07:104	Unconfirmed
B*07:02:16	Unconfirmed	B*07:13	Confirmed	B*07:58	Unconfirmed	B*07:105	Confirmed
B*07:02:17	Confirmed	B*07:14	Confirmed	B*07:59	Unconfirmed	B*07:106	Confirmed
B*07:02:18	Unconfirmed	B*07:15	Confirmed	B*07:60	Unconfirmed	B*07:107	Unconfirmed
B*07:02:19	Unconfirmed	B*07:16	Confirmed	B*07:61	Unconfirmed	B*07:108	Unconfirmed
B*07:02:20	Unconfirmed	B*07:17	Confirmed	B*07:62	Confirmed	B*07:109	Confirmed
B*07:02:21	Confirmed	B*07:18:01	Unconfirmed	B*07:63	Confirmed	B*07:110	Confirmed
B*07:02:22	Unconfirmed	B*07:18:02	Confirmed	B*07:64	Confirmed	B*07:111N	Confirmed
B*07:02:23	Confirmed	B*07:19	Confirmed	B*07:65	Unconfirmed	B*07:112	Unconfirmed
B*07:02:24	Unconfirmed	B*07:20	Confirmed	B*07:66	Unconfirmed	B*07:113	Unconfirmed
B*07:02:25	Unconfirmed	B*07:21	Unconfirmed	B*07:67N	Unconfirmed	B*07:114	Unconfirmed
B*07:02:26	Unconfirmed	B*07:22:01	Unconfirmed	B*07:68:01	Confirmed	B*07:115	Unconfirmed
B*07:02:27	Confirmed	B*07:22:02	Confirmed	B*07:68:02	Confirmed	B*07:116	Unconfirmed
B*07:02:28	Confirmed	B*07:23	Unconfirmed	B*07:68:03	Confirmed	B*07:117	Unconfirmed
B*07:02:29	Unconfirmed	B*07:24	Confirmed	B*07:69	Confirmed	B*07:118	Unconfirmed
B*07:02:30	Unconfirmed	B*07:25	Unconfirmed	B*07:70	Unconfirmed	B*07:119	Confirmed
B*07:02:31	Unconfirmed	B*07:26	Confirmed	B*07:71	Confirmed	B*07:120	Unconfirmed
B*07:02:32	Unconfirmed	B*07:27	Unconfirmed	B*07:72	Confirmed	B*07:121	Unconfirmed
B*07:02:33	Confirmed	B*07:28	Unconfirmed	B*07:73	Confirmed	B*07:122	Confirmed
B*07:02:34	Confirmed	B*07:29	Unconfirmed	B*07:74	Confirmed	B*07:123	Unconfirmed
B*07:02:35	Confirmed	B*07:30	Unconfirmed	B*07:75	Unconfirmed	B*07:124	Unconfirmed
B*07:02:36	Confirmed	B*07:31	Confirmed	B*07:76	Confirmed	B*07:125	Confirmed
B*07:02:37	Unconfirmed	B*07:32	Confirmed	B*07:77	Confirmed	B*07:126	Confirmed
B*07:02:38	Unconfirmed	B*07:33:01	Confirmed	B*07:78	Unconfirmed	B*07:127	Unconfirmed
B*07:02:39	Unconfirmed	B*07:33:02	Unconfirmed	B*07:79	Unconfirmed	B*07:128	Unconfirmed
B*07:02:40	Unconfirmed	B*07:33:03	Unconfirmed	B*07:80	Confirmed	B*07:129	Unconfirmed
B*07:02:41	Confirmed	B*07:34	Unconfirmed	B*07:81	Unconfirmed	B*07:130	Unconfirmed
B*07:02:42	Unconfirmed	B*07:35	Unconfirmed	B*07:82	Confirmed	B*07:131	Confirmed
B*07:02:43	Unconfirmed	B*07:36	Confirmed	B*07:83	Unconfirmed	B*07:132	Confirmed
B*07:02:44	Unconfirmed	B*07:37	Confirmed	B*07:84	Unconfirmed	B*07:133	Unconfirmed
B*07:02:45	Unconfirmed	B*07:38	Confirmed	B*07:85	Unconfirmed	B*07:134	Unconfirmed
B*07:02:46	Unconfirmed	B*07:39	Unconfirmed	B*07:86	Confirmed	B*07:135N	Unconfirmed
B*07:02:47	Unconfirmed	B*07:40	Unconfirmed	B*07:87	Confirmed	B*07:136:01	Confirmed
B*07:02:48	Unconfirmed	B*07:41	Unconfirmed	B*07:88	Confirmed	B*07:136:02	Unconfirmed
B*07:03	Confirmed	B*07:42	Confirmed	B*07:89	Unconfirmed	B*07:137	Unconfirmed
B*07:04	Confirmed	B*07:43	Confirmed	B*07:90	Unconfirmed	B*07:138	Unconfirmed

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Allele	Status ¹	Allele	Status ¹	Allele	Status ¹
B*07:139	Unconfirmed	B*07:178	Unconfirmed	B*07:218	Unconfirmed
B*07:140	Unconfirmed	B*07:179	Unconfirmed	B*07:219	Confirmed
B*07:141	Unconfirmed	B*07:180	Unconfirmed	B*07:220	Unconfirmed
B*07:142	Unconfirmed	B*07:181N	Unconfirmed	B*07:221	Unconfirmed
B*07:143	Confirmed	B*07:182N	Unconfirmed	B*07:222	Confirmed
B*07:144	Unconfirmed	B*07:183	Unconfirmed	B*07:223	Unconfirmed
B*07:145	Unconfirmed	B*07:184	Unconfirmed	B*07:224	Unconfirmed
B*07:146	Unconfirmed	B*07:185	Unconfirmed	B*07:225	Unconfirmed
B*07:147	Unconfirmed	B*07:186	Unconfirmed	B*07:226	Unconfirmed
B*07:148	Confirmed	B*07:187	Unconfirmed	B*07:227	Unconfirmed
B*07:149	Confirmed	B*07:188	Unconfirmed	B*07:228	Unconfirmed
B*07:150	Unconfirmed	B*07:189	Unconfirmed	B*07:229	Unconfirmed
B*07:151:01	Confirmed	B*07:190	Unconfirmed	B*07:230	Unconfirmed
B*07:151:02	Unconfirmed	B*07:191	Unconfirmed	B*07:231N	Unconfirmed
B*07:152	Unconfirmed	B*07:192	Unconfirmed	B*07:232	Unconfirmed
B*07:153	Unconfirmed	B*07:193	Unconfirmed	B*07:233	Unconfirmed
B*07:154	Unconfirmed	B*07:194	Unconfirmed	B*07:234	Unconfirmed
B*07:155	Unconfirmed	B*07:195	Unconfirmed	B*07:235	Unconfirmed
B*07:156	Unconfirmed	B*07:196	Unconfirmed	B*07:236	Unconfirmed
B*07:157	Unconfirmed	B*07:197	Unconfirmed	B*07:237	Confirmed
B*07:158	Unconfirmed	B*07:198	Unconfirmed	B*07:238	Confirmed
B*07:159	Unconfirmed	B*07:199	Unconfirmed	B*07:239	Unconfirmed
B*07:160	Confirmed	B*07:200	Unconfirmed	B*07:240	Unconfirmed
B*07:161N	Unconfirmed	B*07:201N	Unconfirmed	B*07:241	Unconfirmed
B*07:162	Unconfirmed	B*07:202	Unconfirmed	B*07:242	Unconfirmed
B*07:163	Confirmed	B*07:203	Unconfirmed	B*07:243	Unconfirmed
B*07:164	Unconfirmed	B*07:204	Unconfirmed	B*07:244	Unconfirmed
B*07:165	Unconfirmed	B*07:205	Unconfirmed	B*07:245	Unconfirmed
B*07:166	Unconfirmed	B*07:206	Confirmed	B*07:246	Unconfirmed
B*07:167N	Unconfirmed	B*07:207	Unconfirmed	B*07:247	Unconfirmed
B*07:168	Unconfirmed	B*07:208	Confirmed	B*07:248	Confirmed
B*07:169	Unconfirmed	B*07:209	Unconfirmed	B*07:249	Unconfirmed
B*07:170	Unconfirmed	B*07:210	Unconfirmed	B*07:250	Unconfirmed
B*07:171	Unconfirmed	B*07:211	Unconfirmed	B*07:251N	Unconfirmed
B*07:172	Unconfirmed	B*07:212	Unconfirmed	B*07:252	Unconfirmed
B*07:173	Unconfirmed	B*07:213	Unconfirmed	B*07:253	Unconfirmed
B*07:174	Confirmed	B*07:214	Unconfirmed	B*07:254	Unconfirmed
B*07:175	Confirmed	B*07:215	Unconfirmed	B*07:255	Unconfirmed
B*07:176	Unconfirmed	B*07:216	Unconfirmed	B*07:256	Unconfirmed
B*07:177	Unconfirmed	B*07:217	Unconfirmed		

¹Allele status “confirmed” or “unconfirmed” as listed on the IMGT/HLA web page 2015-October-10, release 3.22.0, www.ebi.ac.uk/imgt/hla.

RESOLUTION IN HOMO- AND HETEROZYGOTES

Results file with resolution in HLA-B*07 homo- and heterozygotes is available upon request.

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SPECIFICITY TABLE

HLA-B*07 SSP subtyping

Specificities and sizes of the PCR products of the 87+1 primer mixes used for HLA-B*07 SSP subtyping

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	Amplified HLA-B*07 alleles ³	Other amplified HLA Class I alleles ⁴
1	395 bp	800 bp	*07:02:01-07:02:27, 07:02:29-07:02:48, 07:04- 07:07, 07:09, 07:10 ^w , 07:11- 07:12, 07:14-07:15, 07:17- 07:26, 07:28-07:31, 07:33:01-07:36, 07:39- 07:49N, 07:51-07:64, 07:66- 07:68:03, 07:70-07:82, 07:84, 07:86-07:124, 07:126- 07:179, 07:181N-07:182N, 07:184-07:206, 07:208- 07:218, 07:220-07:227, 07:229-07:256	*08:123, 08:132, 14:21, 38:26, 40:166 [?] , 42:01:01-42:02:01:02, 42:04-42:06, 42:08- 42:10, 42:12-42:21, 45:06, 67:01:01- 67:01:03, 67:03-67:05, 73:01 ^w -73:02 ^w , 81:02
2	285 bp	1070 bp	*07:02:01-07:06, 07:08- 07:11, 07:13, 07:15-07:17, 07:20-07:32, 07:34-07:39, 07:41-07:52, 07:54-07:59, 07:61-07:76, 07:79-07:83, 07:85-07:99, 07:101-07:121, 07:123-07:136:02, 07:138, 07:140-07:161N, 07:164, 07:166-07:192, 07:194- 07:195, 07:199-07:213, 07:215-07:224, 07:226, 07:228-07:234, 07:236- 07:238, 07:240-07:247, 07:249-07:254, 07:256	*08:79, 08:156, 35:66, 37:07, 40:15-40:16, 40:31-40:32, 40:45, 40:80, 40:98, 42:05:01-42:05:02, 48:01:01-48:01:06, 48:05-48:12, 48:14-48:16, 48:18-48:20, 48:22, 48:27-48:37, 81:01-81:08
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25⁹	305 bp 425 bp	1070 bp	*07:117 *07:27, 07:236, 07:240	*40:250 *08:126, 08:131, 08:139, 38:17, 40:188 [?]
26⁵	95 bp 205 bp 235 bp	800 bp	*07:28, 07:118 *07:51, 07:220, 07:226 *07:20, 07:24, 07:60, 07:100, 07:131, 07:220, 07:223	*08:14, 08:87, 27:32, 40:180, 44:54, 44:106, 44:135 *08:143, C*07:294, C*08:08:01-08:08:02 *08:21, 08:25, 15:07:01-15:07:03, 15:45, 15:55, 15:68, 15:126, 15:207, 15:324, 15:329, 15:331, 15:360, 35:05:01- 35:05:03, 35:22, 35:51, 35:58, 35:72, 35:89, 35:97, 35:114, 35:199, 35:232, 40:05, 40:71, 40:174, 44:54, 44:106, 44:135, 44:158, 44:184, 44:213, 46:12, 51:64, 51:148, 53:14, 55:51, 58:18, C*03:05, C*03:25, C*03:27, C*03:167, C*03:296
	290 bp		*07:42	
27	165 bp 215 bp	800 bp	*07:41 *07:29, 07:186	*08:123, 08:132, C*07:335
28^{5,8}	110 bp	1070 bp	*07:30	
29^{5,6,8}	85 bp	800 bp	*07:31, 07:34, 07:43	*18:110, 40:199, 41:08, 42:05:01-42:05:02, 48:37, C*07:38:01-07:38:02
	200 bp		*07:44	
30	190 bp	800 bp	*07:05:01-07:06, 07:32, 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, 07:209-07:210, 07:213-07:214, 07:222, 07:249	*08:40, 08:79, 13:01:01-13:01:02, 13:01:04-13:02:16, 13:02:18, 13:07N- 13:09, 13:11, 13:13:01-13:23, 13:25, 13:27-13:33, 13:36-13:45, 13:47, 13:49N- 13:50, 13:52, 13:54-13:61, 13:63N-13:71, 13:73-13:78, 13:80-13:85, 27:07:01- 27:07:04, 27:11, 27:20, 27:24, 27:27, 27:33, 27:43, 27:125, 27:130, 35:31, 35:88, 35:230, 39:47, 39:50, 40:01:01-40:01:20, 40:01:22-40:01:25, 40:01:27, 40:01:29- 40:01:31, 40:01:33-40:02:09, 40:02:11- 40:02:12, 40:02:14-40:02:17, 40:02:19- 40:02:24, 40:04, 40:06:01:01-40:06:09, 40:06:11-40:08, 40:10:01:01-40:16, 40:19, 40:21-40:23, 40:25, 40:27:01-40:27:02, 40:29-40:30, 40:34-40:38, 40:40, 40:43, 40:45-40:50, 40:53-40:58, 40:61-40:62, 40:64-40:70:02, 40:72:01-40:88, 40:90- 40:91, 40:93-40:97, 40:99-40:119, 40:121- 40:123, 40:125-40:130:02, 40:132, 40:134- 40:158, 40:160:01-40:162, 40:164-40:171, 40:173, 40:175-40:179, 40:182-40:183, 40:185-40:186, 40:188-40:198, 40:200- 40:266, 40:268-40:279, 40:281-40:283, 40:285-40:304, 40:306, 40:308-40:317, 41:19, 44:31, 48:01:01-48:01:06, 48:03:01- 48:07, 48:09-48:16, 48:18-48:22, 48:24, 48:28-48:36, 51:10, 51:16, 51:31, 51:34, 51:81-51:82, 51:93, 52:08, 54:26, 55:09, 55:22, 55:24, 58:27, 59:09, 78:08, 81:01- 81:08, C*02:68
31	215 bp	800 bp	*07:55, 07:100	*08:70, 15:07:01-15:07:03, 15:45, 15:68, 15:126, 15:207, 15:324, 15:331, 46:12, 48:19, C*02:60

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	280 bp		*07:19, 07:33:01-07:33:03, 07:53, 07:60, 07:100	*08:103, 08:134, 13:71, 14:05, 14:13, 15:07:01-15:07:03, 15:45, 15:55, 15:68, 15:126, 15:207, 15:324, 15:329, 15:331, 15:360, 18:14, 27:07:01-27:07:04, 27:11, 27:24, 27:32-27:34, 27:43, 27:138, 35:05:01-35:05:03, 35:22, 35:31, 35:51, 35:58, 35:72, 35:89, 35:97, 35:114, 35:199, 35:232, 37:09, 38:19, 39:03, 39:14, 39:24:01-39:24:02, 39:29, 39:37, 39:76, 40:02:01-40:03, 40:05, 40:08-40:09, 40:13, 40:18-40:19, 40:24, 40:27:01-40:27:02, 40:29, 40:35:01-40:35:02, 40:37, 40:39- 40:40, 40:50, 40:56-40:58, 40:71, 40:78, 40:82, 40:85, 40:89-40:91, 40:94, 40:97, 40:104-40:105, 40:107, 40:111, 40:115, 40:119, 40:122, 40:133Q, 40:142N-40:145, 40:157, 40:164, 40:169, 40:173-40:174, 40:176, 40:180-40:181, 40:189, 40:200- 40:203, 40:205-40:206, 40:209, 40:211, 40:214, 40:219-40:220, 40:224, 40:226, 40:229, 40:232, 40:246, 40:248, 40:254- 40:255, 40:266-40:267, 40:271, 40:274, 40:276, 40:283-40:284, 40:287, 40:289- 40:293, 40:296-40:297, 40:302-40:305, 40:309, 40:314, 42:08, 44:54, 44:106, 44:135, 44:158, 44:184, 44:213, 46:12, 48:04:01-48:04:02, 48:13, 48:24, 51:64, 51:81, 51:148, 53:14, 55:04, 55:49, 55:51, 58:18, 58:27, C*02:60, C*03:05, C*03:25, C*03:27, C*03:167, C*03:296, C*08:08:01-08:08:02
32¹⁰	190 bp	1070 bp	*07:34	*08:01:01-08:01:22, 08:01:24-08:05, 08:08N-08:10, 08:12:01-08:13, 08:15- 08:20, 08:22-08:24, 08:26:01-08:27, 08:29- 08:36, 08:38-08:39, 08:41-08:48, 08:50- 08:73, 08:75-08:76, 08:78, 08:80-08:86N, 08:88, 08:90-08:106, 08:109-08:110, 08:112-08:124, 08:126-08:155, 13:06, 13:53, 14:13, 15:42, 15:83, 15:86, 15:188, 15:224, 15:252, 18:26, 18:61, 18:97, 35:60, 35:87, 35:276, 37:12, 38:01:01, 38:01:03- 38:01:05, 38:01:07-38:02:05, 38:02:07- 38:07, 38:09-38:21, 38:23-38:28, 38:31- 38:51, 38:53-38:58, 39:01:01:01- 39:01:01:04, 39:01:03-39:01:18, 39:01:20- 39:06:05, 39:08-39:14, 39:16, 39:18-39:20, 39:22-39:24:02, 39:26-39:28, 39:30-39:31, 39:34-39:37, 39:39:01-39:39:02, 39:41- 39:42, 39:44-39:46, 39:48, 39:51-39:54, 39:56-39:66, 39:68-39:73, 39:75-39:88, 39:90, 39:92-39:95N, 39:97N, 39:99- 39:101, 39:103-39:107, 40:39, 40:51, 40:163, 40:184, 40:199, 41:01:01- 41:03:02, 41:05-41:09, 41:11-41:16, 41:20, 41:22-41:41, 42:01:01-42:02:01:02, 42:04- 42:08, 42:10-42:15, 42:17-42:19, 42:21, 44:166, 46:18, 48:37, 51:21, 51:36, 51:101,

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				51:143, 51:153, 51:157, 52:37, 54:01:01-54:02, 54:04-54:05N, 54:07- 54:08N, 54:10-54:13, 54:17-54:25, 54:27- 54:32, 54:34, 55:01:01-55:05, 55:07, 55:10-55:13, 55:15-55:21, 55:25-55:26, 55:28-55:40, 55:42-55:45, 55:47-55:50, 55:52-55:57, 55:59-55:67, 55:69-55:76, 56:10, 56:12, 56:23, 58:20, 59:01:01:01- 59:03, 59:05-59:08, 67:01:01-67:05, 78:07, C*01:90, C*15:24
33^{5,6}	90 bp	1070 bp	*07:149	*08:02, 08:117, 13:36, 15:89, 15:115, 18:09, 38:02:01-38:02:06, 38:08, 38:15, 38:18, 38:23, 38:29, 38:35, 38:43-38:50, 44:12, 44:232, 51:54, 51:78:01-51:78:02, 53:09, 53:11-53:13, 53:31, 53:36, 56:07, 59:08
	350 bp		*07:61	
34	155 bp	1070 bp	*07:62, 07:90	
35^{5,8}	105 bp	1070 bp	*07:63	
	215 bp		*07:143	*40:166, 42:02:01:01-42:02:01:02, 42:09, 42:17-42:18, 45:06, 54:02, 55:16, 56:35
36	180 bp	1070 bp	*07:167N, 07:201N	
	250 bp		*07:64	
37	210 bp	1070 bp	*07:65, 07:206	*08:26:03, 15:32:02, 18:04, 35:08:05, 35:42:02, 35:185, 39:63, 40:01:11, 40:58, 46:40:01, 56:05:01, 56:06, 78:01:01- 78:01:02, 78:02:02-78:03, 78:07-78:09
38⁸	190 bp	1070 bp	*07:103, 07:113	*08:64
	215 bp		*07:66	
39⁷	425 bp	1070 bp	*07:05:01-07:06, 07:34, 07:40, 07:53, 07:80, 07:90, 07:97, 07:105, 07:112, 07:123 ^w , 07:137-07:138, 07:140, 07:176, 07:182N, 07:201N, 07:206, 07:209- 07:210, 07:213, 07:222, 07:249	*08:123, 08:132, 08:146, 40:61 ^w , 40:166, 42:01:01-42:02:01:02, 42:04-42:05:02, 42:07-42:08, 42:10, 42:12-42:15, 42:17- 42:19, 42:21, 54:04, 54:11, 55:04, 55:08, 55:17, 55:20, 55:28, 55:49, 55:51, 56:05:01-56:06, 56:12, 56:15, 56:21, 81:01-81:04N, 81:06-81:08
40⁸	265 bp	1070 bp	*07:56:01-07:56:02, 07:72, 07:206	*08:26:01-08:26:03, 08:56:01-08:56:02, 38:28, 39:63, 40:21 [?] , 40:58 [?] , 40:137 [?] , 40:248 [?] , 40:282 [?] , 41:12, 41:36, 42:21, 49:03
	395 bp		*07:83	
41	400 bp	1070 bp	*07:58	
	425 bp		*07:156	
42	195 bp	1070 bp	*07:67N	
	225 bp		*07:71	
43	185 bp	1070 bp	*07:68:01-07:68:03, 07:214, 07:237	*15:138, 15:230, 15:241, 15:297, 18:35, 18:110, 35:66, 40:77, 40:87:01-40:87:02, 40:121, 40:158, 40:222, 40:237, 44:150, 48:12, 48:14, 48:26, 48:29
44⁸	150 bp	1070 bp	*07:10	*27:75 ^w , 73:01 ^w -73:02 ^w
45⁵	105 bp	1070 bp	*07:27, 07:50	*08:04, 08:17, 08:54, 08:110, 15:03:01:01- 15:03:05, 15:47:01-15:47:02, 15:54, 15:61- 15:62, 15:64:01-15:64:02, 15:68-15:69,

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46	425 bp	1070 bp	*07:13, 07:15, 07:92, 07:253 ^w	*08:15, 39:27, 40:73, 67:02, 73:01 ^w -73:02 ^w
47	215 bp	800 bp	*07:04, 07:19, 07:25, 07:146	*08:01:01-08:05, 08:07-08:08N, 08:10- 08:11, 08:14-08:15, 08:17-08:19N, 08:21- 08:24, 08:26:01-08:39, 08:41-08:48, 08:50- 08:54, 08:56:01-08:59:02, 08:61-08:69, 08:71-08:78, 08:80-08:83, 08:85-08:88, 08:90-08:106, 08:108-08:128, 08:130- 08:142, 08:144-08:155, 35:87, 37:09, 41:02:01-41:02:06, 41:04, 41:10-41:11, 41:13, 41:15, 41:18-41:19, 41:23-41:24, 41:27, 41:31, 41:36, 41:38-41:41, 42:01:01-42:02:01:02, 42:05:01-42:12, 42:14-42:18, 42:20-42:21, 44:106, 44:158, 44:166, 44:213
48	200 bp 280 bp	1070 bp	*07:76 *07:77, 07:193	*40:60, 40:245, C*07:38:01-07:38:02
49	200 bp	1070 bp	*07:82	*44:129
50⁵	75 bp 105 bp 165 bp	1070 bp	*07:74 *07:175 *07:87	*08:77, 48:27 *15:126
51	345 bp 470 bp	1070 bp	*07:97 *07:93	*08:114, 39:100 *08:39, 38:27
52⁸	290 bp	800 bp	*07:95	*40:233
53⁸	235 bp	800 bp	*07:14, 07:17, 07:28, 07:96:01-07:96:02, 07:115, 07:197, 07:225	*15:230, 15:297, 27:36, 40:42, 40:60, 40:124:01-40:124:02, 40:172, 40:280, 44:150, 48:17, 48:26, C*03:260, C*04:77, C*07:38:01-07:38:02
54⁶	160 bp 185 bp	1070 bp	*07:238 *07:98, 07:167N, 07:201N	
55	260 bp	1070 bp	*07:99	
56^{6,8}	220 bp 235 bp	1070 bp	*07:45 *07:20, 07:24, 07:60, 07:100, 07:131, 07:220, 07:223	*08:21, 08:25, 15:07:01-15:07:03, 15:45, 15:55, 15:68, 15:126, 15:207, 15:324, 15:329, 15:331, 15:360, 35:05:01- 35:05:03, 35:22, 35:51, 35:58, 35:72, 35:89, 35:97, 35:114, 35:199, 35:232, 40:05, 40:71, 40:174, 44:54, 44:106, 44:135, 44:158, 44:184, 44:213, 46:12, 51:64, 51:148, 53:14, 55:51, 58:18, C*03:05, C*03:25, C*03:27, C*03:167, C*03:296
57	150 bp 215 bp	1070 bp	*07:109 *07:70	

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58⁶	245 bp 440 bp	1070 bp	*07:112 *07:79	
59⁵	100 bp	800 bp	*07:115, 07:182N, 07:246	*08:69, 14:05, 38:19, 39:03, 39:24:01-39:24:02, 39:37, 39:76, 40:71, 40:314, 41:18, 42:06, 42:20, 48:07, C*08:08:01-08:08:02
	190 bp		*07:89, 07:113	*08:64
60	145 bp	1070 bp	*07:73	*40:166, 44:90, 45:06
61	205 bp 420 bp	1070 bp	*07:88 *07:105	
62	415 bp	800 bp	*07:16, 07:27, 07:37-07:38, 07:50, 07:75, 07:91, 07:180, 07:207, 07:219, 07:228	*08:03, 08:23, 08:54, 08:110, 08:133, 08:139, 14:14, 14:18, 15:68, 38:01:01-38:02:07, 38:04-38:12, 38:15-38:25, 38:27-38:40, 38:42-38:58, 39:05:01-39:05:02, 39:07-39:08, 39:11, 39:13:01-39:13:02, 39:20, 39:37, 39:49, 39:55-39:56, 40:01:01-40:01:43, 40:07, 40:09 [?] , 40:11:01 [?] -40:11:02 [?] , 40:14:01 [?] -40:15 [?] , 40:16, 40:18 [?] -40:19 [?] , 40:21 [?] -40:22N [?] , 40:23, 40:24 [?] -40:38 [?] , 40:42 [?] , 40:44 [?] -40:49 [?] , 40:51, 40:52 [?] -40:58 [?] , 40:59-40:60, 40:61 [?] -40:65 [?] , 40:67 [?] -40:69 [?] , 40:71 [?] , 40:72:01-40:73, 40:74 [?] -40:76 [?] , 40:77, 40:78 [?] , 40:79, 40:80 [?] -40:88 [?] , 40:90 [?] -40:93 [?] , 40:95 [?] -40:96 [?] , 40:98 [?] -40:121 [?] , 40:123 [?] , 40:124:01, 40:124:02 [?] -40:137 [?] , 40:138-40:140, 40:141 [?] , 40:145 [?] , 40:146, 40:148 [?] -40:149 [?] , 40:150, 40:151 [?] , 40:152-40:153, 40:154 [?] , 40:155:01N-40:155:02N, 40:156 [?] -40:165 [?] , 40:167 [?] -40:171 [?] , 40:172, 40:173 [?] -40:185 [?] , 40:186, 40:187 [?] -40:213 [?] , 40:215 [?] -40:228 [?] , 40:231 [?] -40:249 [?] , 40:250-40:253, 40:257 [?] , 40:258-40:265N, 40:272, 40:273 [?] -40:276 [?] , 40:277-40:278, 40:279 [?] -40:283 [?] , 40:285-40:286N, 40:287 [?] -40:300 [?] , 40:301, 40:306 [?] -40:313 [?] , 40:315 [?] , 40:316-40:317, 41:01:01-41:26, 41:28-41:41, 42:11, 44:18, 45:01:01-45:05, 45:07-45:17, 48:01:01-48:37, 49:01:01-49:14, 49:16-49:40, 50:01:01, 50:01:02 [?] , 50:01:03-50:01:04, 50:01:06-50:02, 50:04-50:20, 50:31-50:43
63⁵	65 bp	1070 bp	*07:94	*67:04, C*03:124, C*04:49, C*05:15, C*07:262, C*07:417, C*12:95
	230 bp		*07:135N	
64	210 bp	1070 bp	*07:69, 07:85, 07:180	*14:01:01-14:01:05, 14:07N-14:08:02, 14:10, 14:12, 14:14, 14:19, 14:26, 14:32, 14:40, 14:46-14:47, 18:33, 27:12, 27:16, 27:92, 27:119, 37:57, 38:01:01-38:02:07, 38:04, 38:08-38:25, 38:27-38:32, 38:34N-38:53, 38:55Q-38:58, 39:01:01:01-39:01:01:04, 39:01:03-39:01:22, 39:03, 39:05:01-39:07, 39:09:01-39:09:03, 39:11, 39:14-39:15, 39:18-39:19:02, 39:22, 39:24:01-39:38Q, 39:40:01N-39:48, 39:50-39:51, 39:53-39:57, 39:59-39:62,

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65	385 bp 450 bp	1070 bp	*07:54, 07:123 *07:111N	*45:06
66	435 bp	1070 bp	*07:36, 07:38, 07:81, 07:180, 07:219	*08:03, 08:52, 08:78, 38:01:01-38:01:10, 38:05-38:07, 38:09-38:14, 38:16, 38:19- 38:22, 38:24-38:28, 38:30-38:34N, 38:36- 38:42, 38:51-38:58, 40:19 [?] , 40:109 [?] , 40:117 [?] , 40:292 [?] , 44:18, 48:18, 49:01:01- 49:01:08, 49:03-49:14, 49:16-49:40
67⁷	230 bp	800 bp	*07:29, 07:72, 07:86, 07:114	*08:01:01-08:05, 08:07-08:56:02, 08:58- 08:93, 08:95-08:123, 08:125-08:127, 08:129-08:146, 08:148N-08:150, 08:152- 08:156, 14:11, 14:28, 38:33, 39:04, 39:63, 40:01:01-40:01:33, 40:01:35-40:01:43, 40:07, 40:14:01 [?] -40:15 [?] , 40:16, 40:21 [?] - 40:22N [?] , 40:23, 40:25 [?] , 40:30 [?] -40:34 [?] , 40:36 [?] , 40:38 [?] , 40:42 [?] -40:43 [?] , 40:45 [?] - 40:49 [?] , 40:51, 40:52 [?] -40:55 [?] , 40:58 [?] , 40:59-40:60, 40:61 [?] -40:63 [?] , 40:65 [?] , 40:66, 40:67 [?] , 40:69 [?] , 40:72:01-40:73, 40:76 [?] , 40:77, 40:79, 40:80 [?] -40:81 [?] , 40:84 [?] , 40:87:01 [?] -40:88 [?] , 40:92 [?] , 40:100 [?] -40:102 [?] , 40:106 [?] , 40:108 [?] , 40:110 [?] , 40:112 [?] - 40:114:02 [?] , 40:116 [?] -40:118N [?] , 40:121 [?] , 40:123 [?] , 40:124:01, 40:124:02 [?] -40:126 [?] , 40:128 [?] -40:130:02 [?] , 40:132 [?] , 40:134 [?] - 40:137 [?] , 40:138-40:140, 40:141 [?] , 40:146- 40:147, 40:150, 40:151 [?] , 40:152-40:153, 40:154 [?] , 40:155:01N-40:155:02N, 40:156 [?] , 40:158 [?] , 40:160:01 [?] -40:160:02 [?] , 40:163 [?] , 40:166 [?] , 40:168 [?] , 40:170 [?] -40:171 [?] , 40:172, 40:175 [?] , 40:178 [?] -40:179 [?] , 40:182 [?] -40:183 [?] , 40:185 [?] , 40:186, 40:187 [?] -40:188 [?] , 40:191 [?] - 40:194 [?] , 40:196 [?] -40:199 [?] , 40:204 [?] , 40:207 [?] -40:208 [?] , 40:212 [?] -40:213 [?] , 40:215 [?] - 40:219 [?] , 40:221 [?] -40:223 [?] , 40:227 [?] -40:228 [?] , 40:231 [?] , 40:233 [?] , 40:235 [?] -40:242 [?] , 40:245 [?] , 40:247 [?] , 40:249 [?] , 40:250-40:253, 40:257 [?] , 40:258-40:265N, 40:272, 40:273 [?] , 40:277-40:278, 40:279 [?] -40:282 [?] , 40:285- 40:286N, 40:288 [?] , 40:298 [?] -40:300 [?] , 40:301, 40:308 [?] , 40:310 [?] , 40:312 [?] , 40:315 [?] , 40:316, 41:01:01-41:41, 44:18, 45:01:01-45:17, 48:22, 49:01:01-49:14, 49:16-49:23, 49:25-49:30, 49:32-49:40, 50:01:01, 50:01:02 [?] , 50:01:03-50:02, 50:04-50:20, 50:31-50:42
68	335 bp 365 bp 415 bp	1070 bp	*07:148 *07:101 *07:13, 07:15, 07:160	*40:193 [?] *08:16, 08:123, 08:145, 42:18, 67:02
69^{6,8}	130 bp 190 bp	1070 bp	*07:136:01-07:136:02 *07:106	

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Lot-specific information

	245 bp		*07:108	
	465 bp		*07:102	*08:18, 73:01-73:02
70⁵	125 bp	800 bp	*07:104	
	205 bp		*07:114	
71¹¹	245 bp	1070 bp	*07:108	
	280 bp		*07:107	
	390 bp		*07:181N	
72	165 bp	1070 bp	*07:13, 07:110	*37:52, 38:11, 48:36, 54:02, 67:02, 81:06, C*03:137, C*03:182, C*07:102^w, C*07:351, C*12:82
73⁵	85 bp	1070 bp	*07:119, 07:169	*48:34
	240 bp		*07:49N	
74	345 bp	1070 bp	*07:120	*40:150, 57:29
	425 bp		*07:156	
75⁵	105 bp	1070 bp	*07:126, 07:175	*15:126
	250 bp		*07:121	
76⁸	130 bp	1070 bp	*07:136:01-07:136:02	
	170 bp		*07:128	
	285 bp		*07:122	
77	335 bp	800 bp	*07:148	*40:193 [?]
	405 bp		*07:116	
78⁵	110 bp	1070 bp	*07:125	
	160 bp		*07:231N	
	380 bp		*07:129	
79⁵	65 bp	1070 bp	*07:130	
	95 bp		*07:119	
80	405 bp	1070 bp	*07:161N	
81	140 bp	1070 bp	*07:147, 07:151:01- 07:151:02, 07:202	*15:116, 15:124, 18:88, 40:31, 40:33, 40:52, 40:59, 46:06, 46:43, 48:23, 57:59, 58:05, C*07:123, C*07:173, C*07:294
82	280 bp	1070 bp	*07:77, 07:163, 07:193, 07:248	*40:60, 40:245, C*07:38:01-07:38:02
83	175 bp	1070 bp	*07:174, 07:202, 07:222	*27:24, 40:31, 40:45, 40:80, 48:01:01- 48:01:06, 48:04:01-48:04:02, 48:06-48:07, 48:09, 48:11, 48:15-48:16, 48:18-48:20, 48:22, 48:24, 48:27-48:37, 81:01-81:08
84⁷	365 bp	1070 bp	*07:12, 07:18:01, 07:137, 07:162, 07:235	*35:76, 42:13, 56:09, 56:11
85⁵	110 bp	800 bp	*07:208	
	160 bp		*07:238	
86	195 bp	1070 bp	*07:219	*57:10, A*02:05:01-02:06:04, A*02:06:06- 02:06:21, A*02:08, A*02:10, A*02:14, A*02:21, A*02:28, A*02:41, A*02:44, A*02:51, A*02:54, A*02:57, A*02:61, A*02:72, A*02:79:01-02:79:02, A*02:84, A*02:91, A*02:99, A*02:106, A*02:108, A*02:122, A*02:126-02:127, A*02:137, A*02:142, A*02:144, A*02:154, A*02:170, A*02:172, A*02:178-02:180, A*02:229, A*02:232, A*02:244, A*02:248, A*02:259, A*02:271, A*02:278, A*02:286, A*02:290, A*02:295, A*02:300, A*02:310, A*02:324, A*02:328, A*02:330-02:331, A*02:333, A*02:337, A*02:344, A*02:355, A*02:359,

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Lot-specific information

A*02:373N, A*02:376, A*02:382, A*02:387, A*02:398, A*02:404-02:405, A*02:409, A*02:413, A*02:415, A*02:420- 02:421, A*02:428, A*02:433, A*02:438, A*02:453-02:454, A*02:465, A*02:470, A*02:472-02:476N, A*02:484, A*02:489, A*02:493, A*02:495-02:496, A*02:506N- 02:507, A*02:527, A*02:532, A*02:546, A*02:550, A*02:558, A*02:572, A*02:577, A*02:591-02:593, A*11:70:01-11:70:02, A*11:121, A*26:07:01-26:07:02, A*26:92			
87	210 bp	1070 bp	*07:251N
88 ¹²	-	-	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*07 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.

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.

⁴Due to the sharing of sequence motifs between HLA-B alleles, non-HLA-B*07 alleles will be amplified by primer mixes 1 to 20, 22 to 27, 29 to 33, 35, 37 to 40, 43 to 53, 56, 59, 60, 62 to 69, 72 to 75, 77, 81 to 84 and 86. In addition, a few HLA-A and HLA-C alleles will be amplified by primer mixes 3 to 5, 7, 9, 10, 12 to 14, 18, 20, 23, 26, 27, 29 to 32, 48, 53, 56, 59, 63, 72, 81, 82 and 86.

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

⁶Primer mixes 21, 24, 29, 33, 54, 56, 58 and 69 have a tendency to giving rise to primer oligomer formation.

⁷Primer mixes 9, 11, 39, 67 and 84 may give rise to a lower yield of HLA-specific PCR product than the other B*07 primer mixes.

⁸Primer mixes 14, 28, 29, 35, 38, 40, 44, 52, 53, 56, 69 and 76 may have tendencies of unspecific amplifications, most pronounced in primer mix 29.

⁹Primer mix 25 may give rise to unspecific amplification in B*73 alleles.

¹⁰Primer mix 32 may give rise to a faint band in many B*14, B*37, C*04, C*05, C*08 and C*18 alleles.

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¹¹Primer mix 71 may give rise to an unspecific amplification product of approx. 900 bp in some HLA-B alleles. This band should be disregarded in the interpretation of HLA-B*07 typings.

¹²Primer mix 88 contains a negative control, which will amplify more than 95% of 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 430 base pairs.

‘w’, might be weakly amplified.

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

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Lot No.: **3D9**

Lot-specific information

PRIMER SPECIFICATION

Well No.	1	2	3	4	5	6	7	8	9	10	11	12
Length of spec.	395	285	235	105	220	215	100	160	355	285	130	225
PCR product							215	235		420	210	
Length of int.	800	1070	1070	800	1070	1070	1070	1070	1070	800	1070	1070
pos. control ¹												
5'-primer(s) ²	44	363	412	540	103	103	97	412	693	44	103	419
	5'-ggC 3'	5'-AgC 3'	5'-ATg 3'	5'-Cgg 3'	5'-CCT 3'	5'-CCT 3'	5'-TCg 3'	5'-ATA 3'	5'-ggT 3'	5'-ggC 3'	5'-CCT 3'	5'-gTC 3'
			412		103	103	540	636		363	103	
			5'-ATg 3'		5'-CCT 3'	5'-CCT 3'	5'-gAC 3'	5'-CAC 3'		5'-Agg 3'	5'-CCT 3'	
							545			363		
							5'-AgT 3'			5'-Agg 3'		
3'-primer(s) ³	272	605	605	605	282	277	272	605	916	287	191	605
	5'-TgT 3'	5'-gCT 3'	5'-gCT 3'	5'-gCT 3'	5'-gCC 3'	5'-ggT 3'	5'-TgT 3'	5'-gCT 3'	5'-gAT 3'	5'-TCC 3'	5'-CgA 3'	5'-gCT 3'
							605	754		302	268	
							5'-gCT 3'	5'-AgA 3'		5'-ggC 3'	5'-gAg 3'	
										605	272	
										5'-gCT 3'	5'-TgC 3'	
											272	
											5'-TgA 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.	85	215	385	205	345	265	205	160	100	385	85	125
PCR product	290		420	230	425		250	205	165			
							295	275				
Length of int.	1070	800	1070	1070	1070	1070	800	1070	1070	1070	1070	800
pos. control ¹												
5'-primer(s) ²	355	363	44	103	44	379	363	25	481	322	560	103
	5'-TCA 3'	5'-AgC 3'	5'-ggC 3'	5'-CCT 3'	5'-ggC 3'	5'-ACC 3'	5'-AgC 3'	5'-CCg 3'	5'-CgA 3'	5'-gCC 3'	5'-CCT 3'	5'-CCT 3'
	560	363		103			409	363	545			103
	5'-CAC 3'	5'-AgC 3'		5'-CCT 3'			5'-ggC 3'	5'-AgC 3'	5'-AgT 3'			5'-CCT 3'
								636				527
								5'-CAC 3'				5'-TgT 3'
3'-primer(s) ³	605	539	259	268	218	605	572	131	605	419	605	191
	5'-gCT 3'	5'-TCA 3'	5'-CTC 3'	5'-gAg 3'	5'-gCA 3'	5'-gCT 3'	5'-gCg 3'	5'-ggA 3'	5'-gCT 3'	5'-CgT 3'	5'-gCT 3'	5'-CgA 3'
			287	292	301		616	527				605
			5'-TCC 3'	5'-gTA 3'	5'-gTC 3'		5'-AgA 3'	5'-CCg 3'				5'-gCT 3'
			302	292	302			754				
			5'-ggT 3'	5'-gTA 3'	5'-ggT 3'			5'-AgA 3'				
				292								
				5'-gTA 3'								
Well No.	13	14	15	16	17	18	19	20	21	22	23	24

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Lot No.: **3D9**

Lot-specific information

Well No.	25	26	27	28	29	30	31	32	33	34	35	36
Length of spec.	305	95	165	110	85	190	215	190	90	155	105	180
PCR product	425	205	215		200		280		350		215	250
		235										
		290										
Length of int. pos. control ¹	1070	800	800	1070	800	800	800	1070	1070	1070	1070	1070
5'-primer(s) ²	44 5'-ggC 3'	363 5'-AgC 3'	97 5'-TCg 3'	215 5'-gCC 3'	560 5'-CAC 3'	412 5'-ATA 3'	363 5'-AgC 3'	412 5'-ATA 3'	261 5'-AAC 3'	483 5'-gAA 3'	97 5'-TCC 3'	363 5'-AgC 3'
			148 5'-TgC 3'		852 5'-CAg 3'				756 5'-ACT 3'	503 5'-CCg 3'	207 5'-gAC 3'	
3'-primer(s) ³	181 5'-gTT 3'	418 5'-gTC 3'	272 5'-TgT 3'	282 5'-gCC 3'	605 5'-gCT 3'	559 5'-CTC 3'	538 5'-CCA 3'	559 5'-CgT 3'	311 5'-Cgg 3'	605 5'-gCT 3'	272 5'-TgT 3'	502 5'-CTA 3'
	301 5'-gTC 3'	527 5'-CCg 3'			916 5'-gAC 3'		603 5'-gTC 3'		970 5'-TCT 3'			570 5'-CCg 3'
		559 5'-CAg 3'										
		613 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.	210	190	425	265	400	195	185	150	105	425	215	200
PCR product		215		395	425	225						280
Length of int. pos. control ¹	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	800	1070
5'-primer(s) ²	141 5'-ATT 3'	363 5'-AgC 3'	272 5'-CTA 3'	44 5'-ggC 3'	625 5'-CCg 3'	89 5'-gAC 3'	463 5'-TgA 3'	165 5'-ACC 3'	206 5'-AgA 3'	44 5'-ggC 3'	363 5'-AgC 3'	113 5'-CCA 3'
					650 5'-CCT 3'	118 5'-C.g 3'						368 5'-gTC 3'
3'-primer(s) ³	311 5'-ggT 3'	506 5'-Tgg 3'	412 5'-gTT 3'	142 5'-TgC 3'	916 5'-gAC 3'	272 5'-TgT 3'	605 5'-gCT 3'	272 5'-TgC 3'	272 5'-Tgg 3'	299 5'-TCA 3'	538 5'-gTC 3'	272 5'-TgT 3'
		516 5'-gCg 3'		272 5'-TgA 3'								605 5'-gCT 3'
		539 5'-TCT 3'										
Well No.	37	38	39	40	41	42	43	44	45	46	47	48

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Lot-specific information

Well No.	49	50	51	52	53	54	55	56	57	58	59	60
Length of spec.	200	75	345	290	235	160	260	220	150	245	100	145
PCR product		105	470			185		235	215	440	190	
		165										
Length of int.	1070	1070	1070	800	800	1070	1070	1070	1070	1070	800	1070
pos. control ¹												
5'-primer(s) ²	341 5'-ggg 3'	363 5'-AgC 3'	44 5'-ggC 3'	355 5'-CCT 3'	409 5'-ggT 3'	363 5'-AgC 3'	386 5'-gCA 3'	363 5'-AgC 3'	98 5'-CTg 3'	44 5'-ggC 3'	363 5'-AgC 3'	167 5'-gCT 3'
									161 5'-Cgg 3'			
3'-primer(s) ³	2 nd I 5'-TCg 3'	398 5'-ggC 3'	221 5'-ACC 3'	605 5'-gCT 3'	605 5'-gCT 3'	484 5'-CgA 3'	605 5'-gCT 3'	544 5'-ggT 3'	272 5'-TgT 3'	121 5'-gCC 3'	419 5'-CgA 3'	272 5'-TgT 3'
		425 5'-CgC 3'	343 5'-T 3'			502 5'-CTA 3'		559 5'-CAg 3'		317 5'-ggC 3'	420 5'-gCT 3'	
		488 5'-CCA 3'				513 5'-TCA 3'					506 5'-Tgg 3'	
											514 5'-CTT 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.	205	415	65	210	385	435	230	335	130	125	245	165
PCR product	420		230		450			365	190	205	280	
								415	245		390	
									465			
Length of int.	1070	800	1070	1070	1070	1070	800	1070	1070	800	1070	1070
pos. control ¹												
5'-primer(s) ²	44 5'-ggC 3'	44 5'-ggC 3'	256 5'-ACT 3'	103 5'-CCT 3'	44 5'-ggC 3'	44 5'-ggC 3'	44 5'-ggC 3'	44 5'-ggC 3'	44 5'-ggC 3'	106 5'-CCA 3'	44 5'-ggC 3'	103 5'-CCT 3'
			363 5'-AgC 3'	103 5'-CCT 3'					363 5'-AgC 3'	188 5'-Cgg 3'	363 5'-AgC 3'	103 5'-CCT 3'
3'-primer(s) ³	82 5'-ggC 3'	292 5'-gTA 3'	282 5'-gCC 3'	272 5'-TgC 3'	259 5'-CTC 3'	309 5'-ATC 3'	106 5'-CAT 3'	212 5'-gCA 3'	119 5'-ggT 3'	272 5'-TgT 3'	119 5'-ggT 3'	226 5'-CAC 3'
	293 5'-ggA 3'	292 5'-gTA 3'	553 5'-CTA 3'		327 5'-TTT 3'			241 5'-CgA 3'	341 5'-CgT 3'		265 5'-CTA 3'	
		292 5'-gTA 3'						289 5'-AgC 3'	453 5'-TCT 3'		603 5'-TTC 3'	
									453 5'-TCC 3'			
									511 5'-CCg 3'			
Well No.	61	62	63	64	65	66	67	68	69	70	71	72

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Lot No.: **3D9**

Lot-specific information

Well No.	73	74	75	76	77	78	79	80	81	82	83	84
Length of spec.	85	345	105	130	335	110	65	405	140	280	175	365
PCR product	240	425	250	170	405	160	95					
				285		380						
Length of int.	1070	1070	1070	1070	800	1070	1070	1070	1070	1070	1070	1070
pos. control ¹												
5'-primer(s) ²	244	625	363	363	44	206	244	693	412	368	363	272
	5'-CgC 3'	5'-CCg 3'	5'-AgC 3'	5'-AgC 3'	5'-ggC 3'	5'-AgA 3'	5'-CgC 3'	5'-ggT 3'	5'-ATg 3'	5'-gTC 3'	5'-AgC 3'	5'-CTA 3'
	693	704		880		671	693					
	5'-ggT 3'	5'-TgT 3'		5'-CCT 3'		5'-CAg 3'	5'-ggT 3'					
3'-primer(s) ³	299	916	425	453	212	277	299	962	512	605	499	353
	5'-TCT 3'	5'-gAC 3'	5'-CgC 3'	5'-TCT 3'	5'-gCA 3'	5'-ggT 3'	5'-TCT 3'	5'-CAA 3'	5'-CCA 3'	5'-gCT 3'	5'-ggA 3'	5'-TgA 3'
	728		427	453	278	324	719					
	5'-CCT 3'		5'-gTT 3'	5'-TCC 3'	5'-ggA 3'	5'-TAC 3'	5'-AgA 3'					
	893		571	607		916						
	5'-CCT 3'		5'-CCg 3'	5'-CAC 3'		5'-gAC 3'						
				916								
				5'-gAC 3'								
Well No.	73	74	75	76	77	78	79	80	81	82	83	84

Well No.	85	86	87
Length of spec.	110	195	210
PCR product	160		
Length of int.	800	1070	1070
pos. control ¹			
5'-primer(s) ²	363	103	363
	5'-AgC 3'	5'-CCT 3'	5'-AgC 3'
3'-primer(s) ³	430	256	533
	5'-gCT 3'	5'-CCC 3'	5'-CT 3'
	484		
	5'-CgA 3'		
Well No.	85	86	87

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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Lot-specific information

CE
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101.512-24/03 – including *Taq* pol., IFU-01
101.512-24u/03u – without *Taq* pol., IFU-02

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Lot No.: **3D9**

Lot-specific information

CELL LINE VALIDATION SHEET									
HLA-B*07 SSP subtyping kit ²									
					Well				
					81	82	83	84	85
					86	87			
					Prod. No.:	201561681	201561682	201436483	201561684
						201561685	201561686	201561687	
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	9025	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		-	-	-	-

101.512-24/03 – including *Taq* pol., IFU-01
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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 7, 12, 13, 15, 17, 18, 20 to 23, 25, 27 to 29, 34, 36, 38, 40 to 43, 48 to 55, 57 to 61, 63, 65, 68, 70 to 82 and 84 to 87 were available.

The specificities of the primers in primer solutions 7, 12, 13, 15, 17, 18, 20, 21, 23, 25, 27, 29, 36, 38, 40, 43, 48, 50 to 53, 57, 59, 60, 63, 68, 70, 72, 73, 75 to 79, 81 to 82, 84 and 86 were tested by separately adding additional 5'-primers, respectively additional 3'-primers. In primer solutions 54, 58, 61, 65, 71, 80, 85 and 87 it was only possible to test the 5'-primer, the 3'-primer were not possible to test. In primer solutions 22, 28, 34, 41, 42, 49, 55 and 74 it was only possible to test the 3'-primers, the 5'-primers were not possible to test. In primer solutions 2, 3, 21, 27, 29, 35, 48, 57, 63, 70, 73, 76, 78 and 79 one or two 5'-primers were not possible to test, and in primer solutions 8, 10, 11, 15 to 17, 19, 20, 24 to 26, 33, 36, 38, 40, 50, 51, 56, 59, 63, 68, 69, 73 and 75 to 79 one or two 3'-primers were not possible to test. Additional primers in primer solutions 8, 10, 11, 19, 24, 26, 31, 33 and 69 were tested by separately adding one additional 5'-primer, or one additional 3'-primer.

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Lot-specific information

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