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**CTS** Collaborative Transplant Study

|             |                 |
|-------------|-----------------|
| Manual No.  | 27              |
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| Product No. | 127             |
| Lot No.     | DQA10-5         |

## WORKING INSTRUCTION

### HLA-DQA1\* CTS-PCR-SSP TRAY KIT

#### LOCUS- AND LOT-SPECIFIC MANUAL

To be applied to the following product:

| Product No. | Description                                                                                                                                                                                                             |
|-------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 127         | HLA-DQA1* low resolution CTS-PCR-SSP Tray Kit   |

#### 1. Main differences

- **Between Lot DQA10-5 (the current lot) and Lot DQA10-4:**  
The kit was updated to cover new alleles included in the IMGT/HLA Database of January 2025. Deleted and renamed alleles were taken into consideration.

#### 2. Introduction

- Intended use: This kit reveals a low resolution typing of HLA-DQA1\* by the PCR-SSP method.
- Allele coverage: IMGT/HLA Sequence Database Release 3.59.0, January 2025, except: DQA1\*02:39, DQA1\*05:05:21
- This manual is only valid for **Lot No. DQA10-5**.
- This manual should be used together with the Main Manual (General Information) which is the 'Working instruction for the CTS-PCR-SSP TRAY and MINITRAY KITS' (Manual No. 100A).

### 3. Content

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| Table 1: Allele specificities and sizes of the PCR products of each HLA-DQA1* CTS-PCR-SSP primer mix (Lot No. DQA10-5) based on IMGT/HLA Sequence Database Release 3.59.0, January 2025 ..... | 4 |
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|                                                                                                                                                                                                  |   |
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| Table 2: Amplification patterns of HLA-DQA1* alleles detected by the HLA-DQA1* CTS-PCR-SSP primer mixes (Lot No. DQA10-5) based on IMGT/HLA Sequence Database Release 3.59.0, January 2025 ..... | 6 |
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#### 4. Kit Composition

- Number of PCR primer mixes per test: 11:
  - 10 HLA-specific mixes
  - 1 negative control mix
- Please note: Well positions E2 - A2, E4 - A4, E6 - A6, E8 - A8, E10 - A10, E12 - A12 are empty!
- Number of tests per tray: 6
- Number of trays per kit: 10
- The primer mixes are aliquoted and dried in PCR-Trays.
- PCR buffer: 3.0 ml of Mastermix SSP (without Taq polymerase).

For storage condition, please refer to Section 1 of the 'Working instruction for the CTS-PCR-SSP TRAY and MINITRAY KITS' (Manual No. 100A) supplied along with this product.

#### 5. Materials, Reagents and Equipment not supplied

Please refer to Section 2 of the 'Working instruction for the CTS-PCR-SSP TRAY and MINITRAY KITS' (Manual No. 100A) supplied along with this product.

#### 6. Sample Requirements, PCR and Gel Electrophoresis

Please refer to Section 3 to 6 of the 'Working instruction for the CTS-PCR-SSP TRAY and MINITRAY KITS' (Manual No. 100A) supplied along with this product.

#### 7. Result Evaluation

- Check the approximate size of the PCR product against the Primer Mix Specificity Table (Appendix / Table 1) to confirm the correct product size.
- Use the Amplification Pattern Table (Appendix / Table 2) to make the allele assignments or use the SCORE Software for detailed result interpretation.

#### 8. Interpretation Hints

- Weak or false positive reactions can occur if you use a different Taq polymerase. Also the quality and quantity of DNA is a crucial factor and can affect the mix reactivities. Under suboptimal test conditions, some mixes could give rise to false positive reactions (if there are any potentially false positive reactions when using this specific lot, they are indicated in Section 9).
- Alleles that are known to amplify weakly are listed with "w" (= weak) in the tables (Appendix).
- Please also refer to Section 7 and 8 of the 'Working instruction for the CTS-PCR-SSP TRAY and MINITRAY KITS' (Manual No. 100A) supplied along with this product.

#### 9. Special notes

HLA-DQA1\*05:04 generates two PCR fragments with Mix 9.

Potentially false positive reactions and primer dimer formation: Mix 5

#### 10. Troubleshooting

Please refer to Section 8 of the 'Working instruction for the CTS-PCR-SSP TRAY and MINITRAY KITS' (Manual No. 100A) supplied along with this product.

#### 11. Precaution

Please refer to Material Safety Data Sheet for the CTS-PCR-SSP TRAY and MINITRAY KITS (Manual No. 100B) supplied along with this product.

#### 12. Contact

If you have any particular questions concerning this kit, which are not answered in this or the Main Manual, please do not hesitate to contact us at:

Phone: +49 6221 564013; Fax: +49 6221 564200

E-mail: dna.labor@med.uni-heidelberg.de

### 13. Appendix

**Table 1:** Allele specificities and sizes of the PCR products of each **HLA-DQA1\*** CTS-PCR-SSP primer mix (**Lot No. DQA10-5**) based on IMGT/HLA Sequence Database Release 3.59.0, January 2025

| Position |    |    |    |    |     | Mix   | Allele                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  | Serology | Size   |
|----------|----|----|----|----|-----|-------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------|--------|
| H1       | H3 | H5 | H7 | H9 | H11 | Mix 1 | DQA1*01:01:01:01-01:01:01:03/01:01:01:05-01:01:14/01:04:01:01-01:05:05/01:07/01:12/01:18/01:22/01:26-01:27/01:29w/01:34-01:35/01:37/01:43/01:49/01:53/01:55-01:56/01:59-01:61/01:64:01-01:64:02/01:66:01-01:67:02/01:74/01:77/01:80/01:83/01:86/01:88N-01:90/01:95-01:96/01:98-01:99/01:107/01:110/01:112/01:114/01:117-01:118/01:120/01:122Q/01:126/01:130-01:131/01:137/01:148-01:149/01:153?/01:154-01:157/01:160/01:161?/01:164-01:166/01:169/01:172/01:177                                                                                                                                                                                         | -, Null  | 145 bp |
| G1       | G3 | G5 | G7 | G9 | G11 | Mix 2 | DQA1*01:02:01:01-01:03:12/01:06/01:08-01:11/01:13-01:17/01:19:01-01:19:01:02/01:21/01:23-01:25/01:28/01:30-01:33/01:36/01:38:01-01:42/01:44-01:48/01:50-01:52/01:54/01:57-01:58/01:62-01:63:01:02/01:65/01:68-01:73/01:75-01:76/01:78-01:79/01:81-01:82/01:84-01:85/01:87:01-01:87:01:02/01:91/01:92?/01:93-01:94/01:97/01:100-01:106/01:108-01:109/01:111/01:113N/01:115-01:116/01:119/01:121/01:123-01:125/01:127-01:129/01:132:01-01:136N/01:138-01:147/01:150-01:152/01:158/01:159?/01:162-01:163/01:167-01:168/01:170-01:171/01:173-01:176/01:178-01:179                                                                                           | -, Null  | 145 bp |
| F1       | F3 | F5 | F7 | F9 | F11 | Mix 3 | DQA1*01:01:01:01-01:01:01:03/01:01:01:05-01:01:13/01:01:14?/01:02:01:01-01:02:22/01:02:23?/01:02:24-01:02:28/01:04:01:01-01:04:05/01:04:06?/01:04:07?/01:04:08-01:09/01:11-01:13/01:16N/01:18-01:19:01:02/01:21-01:23/01:25-01:28/01:31-01:32/01:34-01:43/01:46/01:48-01:49/01:51:01-01:56/01:58-01:59/01:60?/01:61-01:64:02/01:66:01-01:67:02/01:69/01:71-01:74/01:75?/01:77/01:80-01:81/01:83/01:85-01:86/01:88N-01:91/01:92?/01:93-01:96/01:98-01:100/01:103-01:107/01:109-01:112/01:114-01:116/01:117?/01:118/01:120/01:122Q/01:124-01:135/01:137/01:140-01:145/01:148-01:158/01:160-01:161/01:163-01:169/01:171-01:172/01:174-01:175/01:177-01:179 | -, Null  | 170 bp |
| E1       | E3 | E5 | E7 | E9 | E11 | Mix 4 | DQA1*01:03:01:01-01:03:12/01:10:01-01:10:02/01:14-01:15N/01:17/01:24/01:30/01:33/01:44-01:45/01:47/01:50/01:57/01:65/01:68/01:70/01:76/01:78-01:79/01:82/01:84?/01:87:01-01:87:01:02/01:97/01:102/01:108/01:113N/01:119/01:121/01:123/01:136N/01:138-01:139/01:146-01:147/01:159/01:162/01:170/01:173/01:176                                                                                                                                                                                                                                                                                                                                            | -, Null  | 170 bp |
| D1       | D3 | D5 | D7 | D9 | D11 | Mix 5 | DQA1*01:03:02?-01:03:03?/01:03:05?/01:04:01:01-01:04:02:02/01:04:03?-01:04:04?/01:04:05-01:05:04/01:05:05?/01:07/01:08?-01:09?/01:12?-01:13?/01:15N?/01:17?-01:18?/01:20?/01:22?/01:26?/01:28?-01:29?/01:31?-01:37?/01:40Q?/01:42?/01:46?-01:47?/01:50?/01:52?-01:53?/01:55?-01:56?/01:59?-01:62?/01:64:01-01:64:02/01:67:01?/01:68?-01:69?/01:71?-01:75?/01:77?-01:79?/01:80/01:81?/01:83?/01:85?/01:86/01:90?-01:98?/01:100/01:101?/01:103?/01:105?/01:109?-01:112?/01:114/01:115?-01:118?/01:131/01:132:01?/01:134?-01:135?/01:137/01:139?/01:141?/01:148?-01:149?/01:152?-01:154?/01:155/01:158/01:159?-01:162?/01:164-01:165/01:172?-              | -, Null  | 200 bp |

| Position |    |    |    |     |     | Mix    | Allele                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           | Serology | Size             |
|----------|----|----|----|-----|-----|--------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------|------------------|
|          |    |    |    |     |     |        | 01:173?/01:177/01:178?-01:179?, DQA1*02:01:02?/02:01:04?/02:01:12?-02:01:13?/<br>02:01:21?/02:04?-02:05?/02:07?-02:10?/02:12?-02:13?/02:16?/02:18?/02:21?-02:24?/<br>02:26?-02:27?/02:30?-02:31?/02:33?-02:34?/02:38N?/02:40?, DQA1*03:01:03?-<br>03:01:04?/03:05?-03:06?/03:08?/03:12?/03:16?/03:19?/03:23?-03:24?/03:26?-03:27N?/<br>03:29?-03:30?/03:32N?-03:39?/03:41?/03:43?/03:46?-03:49?/03:51?/03:61?-03:62?/<br>03:64?-03:65?, DQA1*04:01:03?/04:01:05?/04:01:09?/04:03N?/04:06?/04:10?-04:12N?/<br>04:15?/04:17?, DQA1*05:04?/05:10?/05:12?/05:15N?-05:17N?/05:22?/05:26?-05:28?/<br>05:30?/05:32?/05:34?-05:35:01?/05:37?/05:39?-05:46?/05:50?/05:52?/05:54:02Q?/<br>05:60?-05:64?/05:66?-05:69?/05:76?-05:77?/05:80?/05:82?/05:86?-05:88:01?/05:92?-<br>05:95?/05:97?-05:100?, DQA1*06:01:02?-06:01:03?/06:02?/06:04?-06:05N?/06:07? |          |                  |
| C1       | C3 | C5 | C7 | C9  | C11 | Mix 6  | DQA1*02:01:01:01-02:01:20/02:01:21?/02:01:22-02:05/02:07-02:37/02:38N?/02:40-<br>02:49                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           | -, Null  | 105 bp           |
| B1       | B3 | B5 | B7 | B9  | B11 | Mix 7  | DQA1*03:01:01:01/03:01:01:03-03:01:01:07/03:01:03-03:01:11/03:01:12?/03:01:13-<br>03:02:01:02/03:02:01:04-03:28/03:29?/03:30-03:45/03:46?/03:47-03:84                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            | -, Null  | 130 bp           |
| A1       | A3 | A5 | A7 | A9  | A11 | Mix 8  | DQA1*01:02:12, DQA1*04:01:01:01-04:01:08/04:01:09?/04:01:10-04:26                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | -, Null  | 215 bp           |
| H2       | H4 | H6 | H8 | H10 | H12 | Mix 9  | <b>DQA1*05:01:01:01-05:01:02/05:01:04-05:05:20/05:05:22-05:09:02/05:10w/05:11:01:01-<br/>05:33/05:34?/05:35:01-05:48/05:49?/05:50-05:86/05:87?/05:88:01-05:92/05:94/05:95?/<br/>05:96-05:118/05:119?/05:120</b>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  | -, Null  | <b>see below</b> |
|          |    |    |    |     |     |        | DQA1*05:04                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | -        | 205 bp           |
|          |    |    |    |     |     |        | DQA1*05:01:01:01-05:01:02/05:01:04-05:05:20/05:05:22-05:09:02/05:10w/05:11:01:01-<br>05:33/05:34?/05:35:01-05:48/05:49?/05:50-05:86/05:87?/05:88:01-05:92/05:94/05:95?/<br>05:96-05:118/05:119?/05:120                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           | -, Null  | 190 bp           |
| G2       | G4 | G6 | G8 | G10 | G12 | Mix 10 | DQA1*05:93, DQA1*06:01:01:01-06:05N/06:06?/06:07/06:08?/06:09-06:12                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              | -, Null  | 105 bp           |
| F2       | F4 | F6 | F8 | F10 | F12 | Mix 11 | Negative Control                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |          | 440 bp           |

**Well positions:** E2 - A2, E4 - A4, E6 - A6, E8 - A8, E10 - A10, E12 - A12 are empty!

**Amplification control (internal positive control):** 440 base pairs (bp)

**w** = weak

**?** = nucleotide sequence information not available for the primer matching sequence or alleles with unknown reactivities

**Bold:** mixes which result in PCR fragments of different sizes (the specificities are first indicated all in one row, then split into several groups in the subsequent rows depending on the fragment size)

**Table 2:** Amplification patterns of **HLA-DQA1\*** alleles detected by the HLA-DQA1\* CTS-PCR-SSP primer mixes (**Lot No. DQA10-5**) based on IMGT/HLA Sequence Database Release 3.59.0, January 2025

| Allele                                                                                                                                                                                                                                                                                                                                                           | Serology | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 |
|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------|---|---|---|---|---|---|---|---|---|----|
| DQA1*01:01:01:01-01:01:01:03/01:01:01:05-01:01:13/01:27/01:43/01:49/01:66:01:01-01:66:01:02/01:67:02/01:88N-01:89/01:99/01:107/01:120/01:122Q/01:126/01:130/01:156-01:157/01:166/01:169                                                                                                                                                                          | -, Null  | 1 |   | 3 |   |   |   |   |   |   |    |
| DQA1*01:01:14                                                                                                                                                                                                                                                                                                                                                    | -        | 1 |   | ? |   |   |   |   |   |   |    |
| DQA1*01:02:01:01-01:02:11/01:02:13:01-01:02:22/01:02:24-01:02:28/01:06/01:11/01:16N/01:19:01:01-01:19:01:02/01:21/01:23/01:25/01:38:01:01-01:39/01:41/01:48/01:51:01:01-01:51:01:02/01:54/01:58/01:63:01:01-01:63:01:02/01:104/01:106/01:124-01:125/01:127-01:129/01:132:02-01:133Q/01:140/01:142-01:145/01:150-01:151/01:163/01:167-01:168/01:171/01:174-01:175 | -, Null  |   | 2 | 3 |   |   |   |   |   |   |    |
| DQA1*01:02:12                                                                                                                                                                                                                                                                                                                                                    | -        |   | 2 | 3 |   |   |   |   | 8 |   |    |
| DQA1*01:02:23                                                                                                                                                                                                                                                                                                                                                    | -        |   | 2 | ? |   |   |   |   |   |   |    |
| DQA1*01:03:01:01-01:03:01:17/01:03:04/01:03:06-01:03:12/01:10:01-01:10:02/01:14/01:24/01:30/01:44-01:45/01:57/01:65/01:70/01:76/01:82/01:87:01:01-01:87:01:02/01:102/01:108/01:113N/01:119/01:121/01:123/01:136N/01:138/01:146-01:147/01:170/01:176                                                                                                              | -, Null  |   | 2 |   | 4 |   |   |   |   |   |    |
| DQA1*01:03:02-01:03:03/01:03:05/01:15N/01:17/01:33/01:47/01:50/01:68/01:78-01:79/01:97/01:139/01:162/01:173                                                                                                                                                                                                                                                      | -, Null  |   | 2 |   | 4 | ? |   |   |   |   |    |
| DQA1*01:04:01:01-01:04:02:02/01:04:05/01:04:08-01:05:04/01:07/01:64:01-01:64:02/01:80/01:86/01:114/01:131/01:137/01:155/01:164-01:165/01:177                                                                                                                                                                                                                     | -        | 1 |   | 3 |   | 5 |   |   |   |   |    |
| DQA1*01:04:03-01:04:04/01:05:05/01:12/01:18/01:22/01:26/01:34-01:35/01:37/01:53/01:55-01:56/01:59/01:61/01:67:01/01:74/01:77/01:83/01:90/01:95-01:96/01:98/01:110/01:112/01:118/01:148-01:149/01:154/01:160/01:172                                                                                                                                               | -        | 1 |   | 3 |   | ? |   |   |   |   |    |
| DQA1*01:04:06-01:04:07                                                                                                                                                                                                                                                                                                                                           | -        | 1 |   | ? |   | 5 |   |   |   |   |    |
| DQA1*01:08-01:09/01:13/01:28/01:31-01:32/01:36/01:40Q/01:42/01:46/01:52/01:62/01:69/01:71-01:73/01:81/01:85/01:91/01:93-01:94/01:103/01:105/01:109/01:111/01:115-01:116/01:132:01/01:134-01:135/01:141/01:152/01:178-01:179                                                                                                                                      | -        |   | 2 | 3 |   | ? |   |   |   |   |    |
| DQA1*01:20                                                                                                                                                                                                                                                                                                                                                       | -        |   |   |   |   | ? |   |   |   |   |    |
| DQA1*01:29                                                                                                                                                                                                                                                                                                                                                       | -        | w |   |   |   | ? |   |   |   |   |    |
| DQA1*01:60/01:117                                                                                                                                                                                                                                                                                                                                                | -        | 1 |   | ? |   | ? |   |   |   |   |    |
| DQA1*01:75                                                                                                                                                                                                                                                                                                                                                       | -        |   | 2 | ? |   | ? |   |   |   |   |    |
| DQA1*01:84                                                                                                                                                                                                                                                                                                                                                       | -        |   | 2 |   | ? |   |   |   |   |   |    |
| DQA1*01:92                                                                                                                                                                                                                                                                                                                                                       | -        |   | ? | ? |   | ? |   |   |   |   |    |

| Allele                                                                                                                                                                                                                                                                                                                                                             | Serology | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 |
|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------|---|---|---|---|---|---|---|---|---|----|
| DQA1*01:100/01:158                                                                                                                                                                                                                                                                                                                                                 | -        |   | 2 | 3 |   | 5 |   |   |   |   |    |
| DQA1*01:101                                                                                                                                                                                                                                                                                                                                                        | -        |   | 2 |   |   | ? |   |   |   |   |    |
| DQA1*01:153/01:161                                                                                                                                                                                                                                                                                                                                                 | -        | ? |   | 3 |   | ? |   |   |   |   |    |
| DQA1*01:159                                                                                                                                                                                                                                                                                                                                                        | -        |   | ? |   | 4 | ? |   |   |   |   |    |
| DQA1*02:01:01:01-02:01:01:12/02:01:03/02:01:05-02:01:11/02:01:14-02:01:20/<br>02:01:22-02:03/02:11/02:14-02:15/02:17/02:19-02:20/02:25/02:28-02:29/02:32/02:35-<br>02:37/02:41-02:49                                                                                                                                                                               | -, Null  |   |   |   |   |   | 6 |   |   |   |    |
| DQA1*02:01:02/02:01:04/02:01:12-02:01:13/02:04-02:05/02:07-02:10/02:12-02:13/<br>02:16/02:18/02:21-02:24/02:26-02:27/02:30-02:31/02:33-02:34/02:40                                                                                                                                                                                                                 | -        |   |   |   |   | ? | 6 |   |   |   |    |
| DQA1*02:01:21/02:38N                                                                                                                                                                                                                                                                                                                                               | -, Null  |   |   |   |   | ? | ? |   |   |   |    |
| DQA1*03:01:01:01/03:01:01:03-03:01:01:07/03:01:05-03:01:11/03:01:13-03:02:01:02/<br>03:02:01:04-03:04/03:07/03:09-03:11/03:13-03:15/03:17-03:18/03:20-03:22/03:25/<br>03:28/03:31/03:40/03:42/03:44-03:45/03:50Q/03:52-03:60/03:63/03:66-03:84                                                                                                                     | -, Null  |   |   |   |   |   |   | 7 |   |   |    |
| DQA1*03:01:03-03:01:04/03:05-03:06/03:08/03:12/03:16/03:19/03:23-03:24/03:26-<br>03:27N/03:30/03:32N-03:39/03:41/03:43/03:47-03:49/03:51/03:61-03:62/03:64-03:65                                                                                                                                                                                                   | -, Null  |   |   |   |   | ? |   | 7 |   |   |    |
| DQA1*03:01:12                                                                                                                                                                                                                                                                                                                                                      | -        |   |   |   |   |   |   | ? |   |   |    |
| DQA1*03:29/03:46                                                                                                                                                                                                                                                                                                                                                   | -        |   |   |   |   | ? |   | ? |   |   |    |
| DQA1*04:01:01:01-04:01:02:04/04:01:04/04:01:06-04:01:08/04:01:10-04:02/04:04-<br>04:05/04:07-04:09/04:13-04:14N/04:16/04:18-04:26                                                                                                                                                                                                                                  | -, Null  |   |   |   |   |   |   |   | 8 |   |    |
| DQA1*04:01:03/04:01:05/04:03N/04:06/04:10-04:12N/04:15/04:17                                                                                                                                                                                                                                                                                                       | -, Null  |   |   |   |   | ? |   |   | 8 |   |    |
| DQA1*04:01:09                                                                                                                                                                                                                                                                                                                                                      | -        |   |   |   |   | ? |   |   | ? |   |    |
| DQA1*05:01:01:01-05:01:02/05:01:04-05:03:03/05:05:01:01-05:05:20/05:05:22-<br>05:09:02/05:11:01:01-05:11:01:02/05:13-05:14/05:18-05:21/05:23-05:25/05:29Q/05:31/<br>05:33/05:35:02-05:36N/05:38/05:47-05:48/05:51/05:53-05:54:01Q/05:55-05:59/<br>05:65:01:01-05:65:01:03/05:70-05:75/05:78-05:79/05:81/05:83-05:85N/05:88:02-05:91/<br>05:96/05:101-05:118/05:120 | -, Null  |   |   |   |   |   |   |   |   | 9 |    |
| DQA1*05:04/05:12/05:15N-05:17N/05:22/05:26-05:28/05:30/05:32/05:35:01/05:37/<br>05:39-05:46/05:50/05:52/05:54:02Q/05:60-05:64/05:66-05:69/05:76-05:77/05:80/<br>05:82/05:86/05:88:01/05:92/05:94/05:97-05:100                                                                                                                                                      | -, Null  |   |   |   |   | ? |   |   |   | 9 |    |
| DQA1*05:10                                                                                                                                                                                                                                                                                                                                                         | -        |   |   |   |   | ? |   |   |   | w |    |
| DQA1*05:34/05:87/05:95                                                                                                                                                                                                                                                                                                                                             | -        |   |   |   |   | ? |   |   |   | ? |    |
| DQA1*05:49/05:119                                                                                                                                                                                                                                                                                                                                                  | -        |   |   |   |   |   |   |   |   | ? |    |
| DQA1*05:93, DQA1*06:01:02-06:01:03/06:02/06:04-06:05N/06:07                                                                                                                                                                                                                                                                                                        | -, Null  |   |   |   |   | ? |   |   |   |   | 10 |
| DQA1*06:01:01:01-06:01:01:05/06:01:04/06:03/06:09-06:12                                                                                                                                                                                                                                                                                                            | -        |   |   |   |   |   |   |   |   |   | 10 |

| Allele           | Serology | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 |
|------------------|----------|---|---|---|---|---|---|---|---|---|----|
| DQA1*06:06/06:08 | -        |   |   |   |   |   |   |   |   |   | ?  |

**w** = weak

**?** = nucleotide sequence information not available for the primer matching sequence or alleles with unknown reactivities



## 14. Certificate of Analysis



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**CTS** Collaborative Transplant Study

# Certificate of Analysis

## HLA-DQA1\* CTS-PCR-SSP Tray Kit

|                                              |         |
|----------------------------------------------|---------|
| Product number                               | 127     |
| Lot number                                   | DQA10-5 |
| Number of HLA-specific primer mixes per test | 10      |

### Mix specifications

The specificity of each primer pair has been tested against a panel of well characterized DNAs.

### Result

No false positive or false negative amplifications were obtained under our test conditions of the bulk reagents.

|                  |                                              |
|------------------|----------------------------------------------|
| Date of approval | 06.08.2025                                   |
| Approved by      | H. Tran, M.D.<br>Quality Control, Supervisor |