

## Olerup SSP<sup>®</sup> DQB1\*02

**Product number:** 101.213-24 – including *Taq* polymerase  
101.213-24u – without *Taq* polymerase  
**Lot number:** 08S  
**Expiry date:** 2015-October-01  
**Number of tests:** 24  
**Number of wells per test:** 12

### CHANGES COMPARED TO THE PREVIOUS DQB1\*02 LOT (63N):

Well	5'-primer	3'-primer	rationale
6	Exchanged	-	Decrease tendencies of primer oligomer formation.
8	New	New	New primer pair for the DQB1*02:10 allele.
9	New	New	New primer pair for the DQB1*02:07 allele.
10	New	New	New primer pair for the DQB1*02:08 allele.
11	New	New	New primer pair for the DQB1*02:09 allele.
12	New	New	New primer pair for the DQB1*02:11 allele.

**THE NUMBER OF WELLS** is increased from 8 to 12.

#### ALLELE COVERAGE:

DQB1\*02:01 to DQB1\*02:11, i.e. all the currently recognized DQB1\*02 alleles, give rise to unique amplification patterns; [www.ebi.ac.uk/imgt/hla](http://www.ebi.ac.uk/imgt/hla), 2013-January-11, release 3.11.0.

The DQB1\*02 kit cannot distinguish the silent mutation in the DQB1\*02:01:01-02:01:06, DQB1\*02:01:08 and DQB1\*02:02:01-02:02:02 alleles.

The DQB1\*02:01:07 and the DQB1\*03:01:01:01-03:23, 03:25-03:47, 04:01:01-04:09, 05:01:01:01-05:13, 05:15-05:18, 06:01:01-06:37 and 06:39-06:52 alleles give rise to identical amplification patterns with the DQB1\*02 primer set. These alleles can be distinguished by the DQ low resolution kit and/or respective DQB1 high resolution kit.

#### RESOLUTION IN DQB1\*02 HOMO- AND HETEROZYGOTES:

Very good.

#### INFLUENCE ON THE INTERPRETATION OF DQB1\*02 SUBTYPINGS BY NON-DQB1\*02 ALLELES:

None of importance.

#### MODIFICATIONS MADE DUE TO COMMENTS FROM CUSTOMERS:

No suggestions received.