

HLA TYPING REPORT

Donor(s) Results

Last Name/URD#	: Bector	Date Of Birth	: 25-Mar-1962
First Name	: Rajeev	Histo ID	: H86523
Relation	: NA	Received On	: 24-Feb-2020
Specimen Type	: Buccal Swab	Collected On	: 22-Feb-2020

A*	B*	C*	DRB1*	DRB345*	DQB1*	DQA1*	DPB1*	DPA1*
01:01:01G	48:01:01G	06:02:01G	07:01:01G	NA	03:03:02G	NA	NA	NA
11:01:01G	57:01:01G	08:01:01G	15:06:01	NA	05:02:01G	NA	NA	NA

Typing Status : Complete

Null Allele Resolution Status : A*01:04:01:01N,A*01:04:01:02N,A*01:22N,A*01:56N,A*01:87N,A*11:21N,A*11:69N,A*11:210N,B*57:79N,C*06:46N,C*06:211:01:01N,C*06:211:01:02N,DQB1*03:375N,DQB1*05:90N excluded

Note:

1. Match ratio to Patient: Based on protein level in ARS (Exon 2 and 3 for Class I, Exon 2 for Class II).
2. Allele Database Version : 3.37.0 (Jul 2019)
3. Allele bearing suffix G : The first allele of the Included Alleles. Included alleles in the appendix
4. NMDP code is reported if G code is not available after analyzing non-ARS exons
5. Sequencing Technology : Illumina Sequencing
6. Reagent Information : Lot Number available upon request.

Report History

Reported Date	History	Comments
28-Feb-2020	Donor: Rajeev	

DIRECTOR / ASSOCIATE DIRECTOR



HLA TYPING REPORT - APPENDIX

Locus	G Code	NMDP Allele Code	Segment Sequenced	Included Alleles
Sample ID : H86523				
A	01:01:01G	01:CAVHH	Exon 1,2,3,4	01:01:01:01/01:01:01:02N/01:01:01:03/01:01:01:04/01:01:01:05/ 01:01:01:06/01:01:01:07/01:01:01:08/01:01:01:09/01:01:01:10/ 01:01:01:11/01:01:01:12/01:01:01:13/01:01:01:14/01:01:01:15/ 01:01:01:16/01:01:01:17/01:01:01:18/01:01:01:19/01:01:01:20/ 01:01:01:21/01:01:01:22/01:01:01:23/01:01:01:24/01:01:01:25/ 01:01:83/01:01:84/01:01:93/01:01:94/01:01:104/01:01:105/ 01:248Q/01:251/01:252/01:276/01:277/01:280/01:300/01:309/01:317
A	11:01:01G	11:CAVJD	Exon 1,2,3,4	11:01:01:01/11:01:01:02/11:01:01:03/11:01:01:04/11:01:01:05/ 11:01:01:06/11:01:01:07/11:01:01:08/11:01:01:09/11:01:01:10/ 11:01:01:11/11:01:01:12/11:01:01:13/11:01:01:14/11:01:01:15/ 11:01:79/11:01:87/11:263/11:270/11:278/11:280/11:295/11:303/ 11:324/11:328
B	48:01:01G	48:BBSD	Exon 2,3,4	48:01:01:01/48:01:01:02/48:01:07/48:43/48:47
B	57:01:01G	57:JJRT	Exon 2,3,4	57:01:01:01/57:01:01:02/57:01:01:03/57:01:01:04/57:01:01:05/ 57:01:01:06/57:01:28/57:01:37/57:103
C	06:02:01G	06:CAVPA	Exon 2,3,4,7	06:02:01:01/06:02:01:02/06:02:01:03/06:02:01:04/06:02:01:05/ 06:02:01:06/06:02:01:07/06:02:01:08/06:02:01:09/06:02:01:10/ 06:02:01:11/06:02:01:12/06:02:01:13/06:02:01:14/06:02:01:15/ 06:02:01:16/06:02:01:17/06:02:60/06:02:61/06:02:64/06:02:76/ 06:02:80/06:83/06:213/06:214/06:227/06:228/06:236/06:239/ 06:244/06:255/06:261/06:266/06:273/06:274/06:277/06:278
C	08:01:01G	08:CAVPJ	Exon 2,3,4,7	08:01:01:01/08:01:01:02/08:01:01:03/08:01:01:04/08:01:27/ 08:22:01:01/08:22:01:02/08:99/08:102/08:178/08:197
DQB1	03:03:02G	03:CAVTC	Exon 2,3	03:03:02:01/03:03:02:02/03:03:02:03/03:03:02:04/03:03:02:05/ 03:398
DQB1	05:02:01G	05:CAVTE	Exon 2,3	05:02:01:01/05:02:01:02/05:02:01:03/05:02:01:04/05:02:01:05/ 05:02:01:06/05:165/05:198/05:222
DRB1	07:01:01G	07:BMSUC	Exon 2,3	07:01:01:01/07:01:01:02/07:01:01:03/07:01:01:04/07:79/07:93
DRB1	15:06:01		Exon 2,3	