Determining the Frequency of *RH* Blood Group System in the Maltese

Population

Antonella Giordmaina¹, Joseph Borg², Jesmond Debono¹, Neville Debattista¹

¹ Hospital Blood Bank, Pathology Department, Mater Dei Hospital, Malta, ² Faculty of Health Science, University of Malta, Malta

Introduction

The Rh blood group system does not only refer to the RhD antigen, but it is a complex protein based system.

The 5 most clinically significant antigens from this blood group system are the D,C,c,E and e since they are the cause of most alloimmunisations.

RHD and RHCE genes are distributed on chromosome 1p34.1-1p36. They are very similar, closely linked loci (figure 1). The 2 RH genes have opposite orientation, face each other with their 38 ends, and are separated by about 30 000 bp. A third gene, SMP1, has the same orientation as RHD and is positioned between RHD and RHCE. The RHD gene is flanked on both sides by the 2 highly homologous Rhesus boxes.

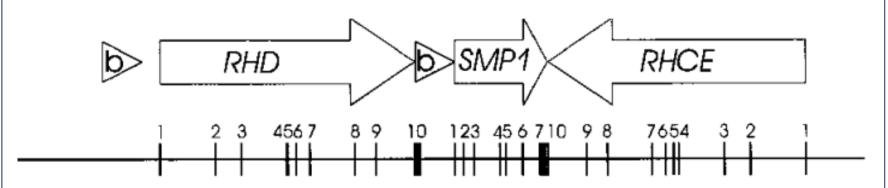


Figure 1: Schematic structure of the RH gene locus.

The positions and orientations of the genes and the Rhesus boxes are indicated by open arrows and triangles, respectively. The exons are shown as vertical bars, and their exon numbers are indicated., which are noted by (b)

Methods and Materials

400 blood donor samples and 397 neonatal blood samples were enrolled in this study. An allele-specific polymerase chain reaction (AS-PCR) method was used to determine the presence of *RHD*, *RHCE*E* and *RHCE*e*, while multiplex PCR was used to test for *RHCE*C/c*. The PCR products were analysed by Agarose-Gel electrophoresis (Figure 2,3,4,5).

81 from these 400 blood donor samples were tested by serology for the RhD, C,c,E and e antigens, and the results were used for comparison with the results obtained by genotyping.



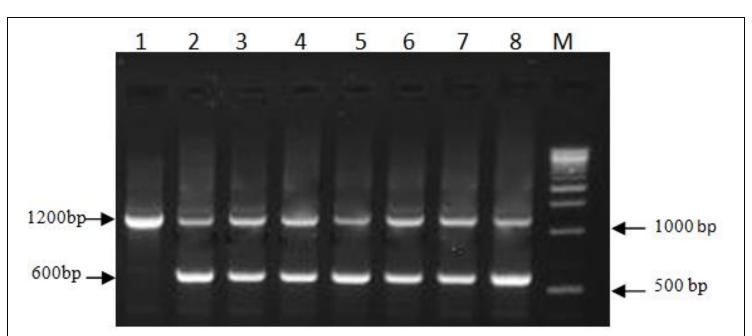


Figure 2. *RHD* products on a 1% agarose gel after electrophoresis

Samples from lane 2 to lane 8 are *RHD* positive samples,. Sample 1 is an *RHD* negative sample since only the control fragment was obtained.

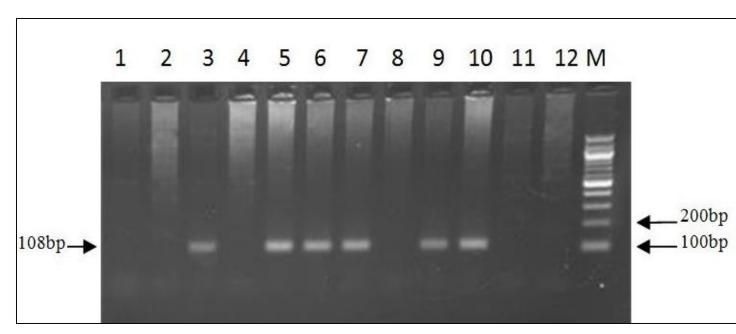


Figure 4: RHCE*E product on a 2.7% agarose gel after electrophoresis

Samples 1, 2, 4,8,11 and 12 are RHCE*E negative, while the rest are RHCE*E neg ative.

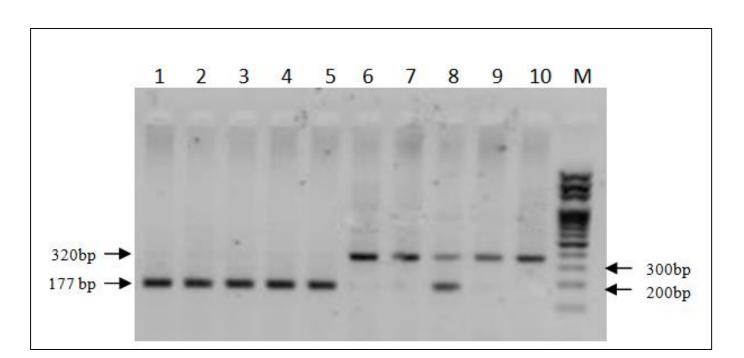


Figure 3: RHC products on a 2% agarose gel after electrophoresis Samples 1 to 5 are recessive homozygotes (cc), samples 6, 7, 9 and 10 are dominant homozygous (CC) whilst sample 8 is heterozygote (Cc).

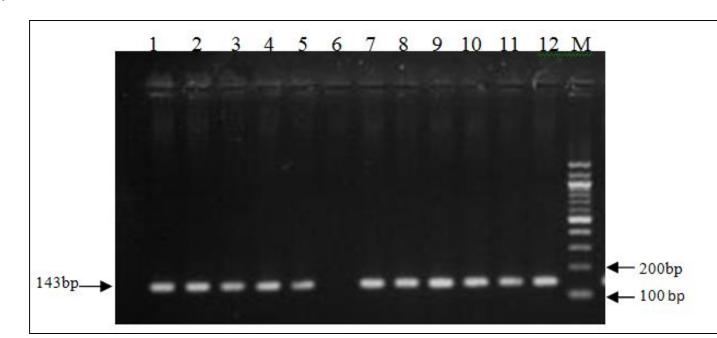


Figure 5: RHCE*e product on a 2.7% agarose gel after electrophoresis Sample 6 is an RHCE*e negative while the rest are RHCE*e positive.

Results

Rh D Positive					Rh D Negative			
Genotype	Cases	Percentage	Frequency	Genotype	Cases	Percentage	Frequency	
DCCEE	0	00.00	00.00	dCCEE	0	00.00	00.00	
DCcEE	2	00.25	0.003	dCcEE	0	00.00	00.00	
DccEE	12	01.50	0.015	dccEE	0	00.00	00.00	
DCCEe	11	01.38	0.014	dCCEe	0	00.00	00.00	
DCCee	195	24.46	0.245	dCCee	0	00.00	00.00	
DCcEe	93	11.67	0.117	dCcEe	0	00.00	00.00	
DccEe	53	06.64	0.064	dccEe	3	00.38	0.004	
DCcee	305	38.27	0.383	dCcee	5	00.63	0.006	
Dccee	62	07.78	0.078	dccee	55	06.90	0.069	

Table 1: The RHDCE genotype frequencies in the population

Discussion and Conclusion

Out of 797 samples, the most common genotype was Dccee (38.27%) followed by DCCee (24.46%). In RHD negative samples dccee was the most frequents (6.9%) (Table 1).

The molecular techniques used, offer a fast in-house testing system to obtain the *RHD* and *RHCE* genotype status.

However this may merit further development to be used in a clinical setting

Contact

Antonella Giordmaina,

Hospital Blood Bank,
Mater Dei Hospital,
Triq Id-Donaturi Tad-Demm,
Msida, Malta
MSD 2090

References

- 1. Faas, Simsek, Bleeker, Overbeeke, Cujipers, Borne, K. d., et al. (1995). Rh E/e Genotyping by Allele-Specific Primer Amplification. *Blood: Journal of the American Society of Hematology*, 85 (3), 829-832.
- 2. Flegel, W. A. (2011). Molecular genetics and clinical applications for RH. *Transfusion and Apherisis Science*, 44 (1), 81-91.
- 3. Johnsen, J. M. (2015). Using red blood cell genomics in transfusion medicine. American Society of Haematology.
- 4. Halima, A. B., Bahri, R., Esteban, E., Moral, P., & Chaabani, H. (2015). Variation of Rhesus Haplotype Frequencies in North Africans and in Worldwide Population Analyses. *International Journal of Human Genetics*, 15 (1), 21-31.
- Lejla Lasić, N. L., Silajdžić, E., Pojskić, L., Hadžiselimović, R., & Pojskić, N. (2013). Molecular genetic variance of RH blood group system within human population of Boznia and Hersegovina. Association of Basic Medical Sciences of FBIH, 13 (1), 10-13.
- 5. Wegner, F.F. & Flegel, W.A. (2000). RHD gene deletion occurred in the Rhesus box. Blood (95), 3662-3668