



Erratum

Erratum to “Frequencies of polymorphisms of Rh, Kell, Kidd, Duffy and Diego systems of Santa Catarina, southern Brazil” [Rev Bras Hematol Hemoter. 2016;38(3):199–205]



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Please consider the following correction:

Results

Page 201 – “Statistical differences were observed in the comparison with Brazilian populations. The genotypes RHCE*Ee, KEL*01.1/KEL*02, FY*01/FY*01, JK*01/JK*01, and JK*02/02 were more prevalent in the population from the state of Santa Catarina in comparison with the state of São Paulo. In addition, RHCE*Cc, RHCE*cc, RHCE*ee, FY*01/FY*02, FY*01/FY*02.N01, FY*02/FY*02, FY*02/FY*02.N01, FY*02.N01/FY*02.N01, and JK*01/JK*02 genotypes were more common in the population from the state of São Paulo” **should read as** “Statistical differences were observed in the comparison with Brazilian populations. The genotypes RHCE*Ee, KEL*01.1/KEL*02, FY*01/FY*02, JK*01/JK*01, and JK*02/02 were more prevalent in the population from the state of Santa Catarina in comparison with the state of São Paulo. In addition, RHCE*Cc, RHCE*cc, RHCE*ee, FY*01/FY*02.N01, FY*02/FY*02, FY*02/FY*02.N01, FY*02.N01/FY*02.N01, and JK*01/JK*02 genotypes were more common in the population from the state of São Paulo.”

Page 201 – “When the data were compared with European populations, only the genotypes RHCE*Cc, RHCE*EE, RHCE*ee, KEL*01.1/KEL*01.1, KEL*01.1/KEL*02, KEL*02/KEL*02 JK*01/JK*01, and DI*01/DI*01 did not differ statistically (Table 4)” **should read as** “When the data were compared with European populations, only the genotypes RHCE*Cc, RHCE*EE, RHCE*ee, KEL*01.1/KEL*01.1, KEL*01.1/KEL*02, KEL*02/KEL*02, FY*01/FY*01, JK*01/JK*01, and DI*01/DI*01 did not differ statistically (Table 4).”

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Tables

Table 1 should read as:

System	Genotype	Frequency (n(%))	
		373 donors	31 patients
Duffy	FY*01/FY*01	57 (15.3)	3 (9.7)
	FY*01/FY*02	176 (47.2)	11 (35.5)

Table 2 should read as:

Genotype	JOA N33
<i>Rh System</i>	
RHCE*Ee	0.12 (4)
RHCE*ee	0.82 (27)
RHCE*E	0.12
RHCE*e	0.88

Table 3 should read as:

Genotype	SC N373	PR N400	SP-POP1 N948	SP-POP2 N250	SP-POP3 N308
<i>Duffy System</i>					
FY*01/FY*01	0.15 (57)	0.12 (50)	0.12 (114)	0.14 (34)	0.12 (36)
FY*01/FY*02	0.47 (176)	0.40 (157)*	0.45 (426)	0.27 (68)*	0.34 (105)*

Table 4 should read as:

Genotype	SC N373	Austria ²⁷ N200	Naples ²⁸ N225	Zurich ²⁹ N4000
<i>Duffy System</i>				
FY*01/FY*01	0.15 (57)	0.20 (40)	0.18 (40)	0.18 (703)
FY*01/FY*02	0.47 (176)	0.36 (72)*	0.43 (98)	0.47 (1862)