

	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	W	X	Y	Z	AA	AB	AC	AD	AE	AF	AG	AH	AI			
1	Donor Code	Loci	Allele 1	Allele 2	Typing Date																																	
2	Sample ID 1	A	02:01:01G	33:01:01G	10/29/2021																																	
3	Sample ID 1	B	14:02:01G	15:01:01G	10/29/2021																																	
4	Sample ID 1	C	03:04:01G	08:02:01G	10/29/2021																																	
5	Sample ID 1	DRB1	01:02:01G	04:01:01G	10/29/2021																																	
6	Sample ID 1	DRB4	01:DJXS		10/29/2021																																	
7	Sample ID 1	DQB1	03:02:01G	05:01:01G	10/29/2021																																	
8	Sample ID 1	DQA1	01:01:01G	03:01:01G	10/29/2021																																	
9	Sample ID 1	DPB1	04:01:01G	04:01:01G	10/29/2021																																	
10	Sample ID 1	DPA1	01:03:01G	01:03:01G	10/29/2021																																	
101																																						
102																																						
103	ARS - Antigen Recognition Site																																					
104	Reference - Human Immunology 68, 392-417 (2007)																																					
105	@American Society for Histocompatibility and Immunogenetics, 2007																																					
106	<b>Note:</b>																																					
107	Alleles bearing suffix 'G' in A,B,C locus has identical sequences in exon 2 and exon 3 antigen recognition sites.																																					
108																																						
109	Alleles bearing suffix 'G' in DRB1,DRB3,DRB4,DRB5,DQB1,DQA1,DPB1,DPA1 locus has identical sequences in exon 2 antigen recognition sites.																																					
110																																						
111	Please find below the list of alleles included for each allele suffix with 'G'.																																					
112	Allele Database Version used in the report is 3.43.0(Jan 2021)																																					
113	Sequencing Technology : Illumina Sequencing																																					
114	NMDP code is reported if G code is not available after analyzing non-ARS exons																																					
115																																						
116																																						
117																																						



