

Table 4: The 10 most frequent *KIR3DL1/S1* and *KIR3DL2* haplotypes in European American, Ugandan, Papuan, Egyptian and Spanish populations

Population	Telomeric		Frequency
	<i>KIR3DL1/S1</i>	<i>KIR3DL2</i>	
European American ‡(Total observed 66)	<i>S1*013</i>	<i>*007</i>	0.189
	<i>L1*001</i>	<i>*001</i>	0.115
	<i>L1*002</i>	<i>*002</i>	0.112
	<i>L1*005</i>	<i>*001</i>	0.079
	<i>L1*015</i>	<i>*002</i>	0.071
	<i>L1*004</i>	<i>*003</i>	0.071
	<i>L1*004</i>	<i>*005</i>	0.063
	<i>L1*008</i>	<i>*009</i>	0.040
	<i>L1*001</i>	<i>*011</i>	0.028
	<i>L1*020</i>	<i>*009</i>	0.027
Ugandan ‡(Total observed 81)	<i>L1*015</i>	<i>*001</i>	0.092
	<i>L1*031</i>	<i>*001</i>	0.077
	<i>L1*001</i>	<i>*001</i>	0.072
	<i>L1*004</i>	<i>*003</i>	0.052
	<i>L1*059^a</i>	-	0.046
	<i>L1*018</i>	<i>*001</i>	0.041
	<i>L1 NEG</i>	<i>*019</i>	0.040
	<i>L1*022</i>	<i>*001</i>	0.036
	<i>L1*015</i>	<i>*013</i>	0.033
	<i>L1 NEG</i>	<i>*006</i>	0.032
Papuan ‡(Total observed 16)	<i>S1*013</i>	<i>*007</i>	0.481
	<i>L1*017</i>	<i>*002</i>	0.243
	<i>L1 NEG</i>	<i>*007</i>	0.124
	<i>L1*005</i>	<i>*010</i>	0.100
	<i>L1*015</i>	<i>*002</i>	0.014
	<i>L1*001</i>	<i>*001</i>	0.005
	<i>L1*005</i>	<i>*001</i>	0.005
	<i>S1*013</i>	<i>*010</i>	0.005
	<i>L1 NEG</i>	<i>*010</i>	0.005
	<i>L1 NEG</i>	<i>*070</i>	0.005
Egyptian ‡(Total observed 40)	<i>L1*001</i>	<i>*001</i>	0.174
	<i>S1*013</i>	<i>*007</i>	0.116
	<i>L1*002</i>	<i>*002</i>	0.093
	<i>L1*004</i>	<i>*005</i>	0.070
	<i>L1*004</i>	<i>*003</i>	0.052
	<i>L1*005</i>	<i>*010</i>	0.050
	<i>L1*008</i>	<i>*009</i>	0.047
	<i>S1*013</i>	<i>*006</i>	0.041

	<i>L1*014</i>	<i>*032</i>	0.035
	<i>L1*005</i>	<i>*001</i>	0.031
Spanish ‡(Total observed 24)	<i>L1*005</i>	<i>*001</i>	0.138
	<i>S1*013</i>	<i>*007</i>	0.138
	<i>L1*001</i>	<i>*001</i>	0.124
	<i>L1*002</i>	<i>*002</i>	0.100
	<i>L1*004</i>	<i>*005</i>	0.100
	<i>L1*004</i>	<i>*003</i>	0.095
	<i>L1*015</i>	<i>*002</i>	0.071
	<i>L1*007</i>	<i>*008</i>	0.048
	<i>L1*001</i>	<i>*011</i>	0.038
	<i>L1*009</i>	<i>*011</i>	0.024

‡The number of distinct haplotypes identified by analyzing two telomeric genes (*KIR3DL1/S1* and *KIR3DL2*);

^a*L1*059* is an allele of *KIR3DL1/2v*, a fusion gene derived from *KIR3DL1* and *KIR3DL2*.