Appendix 1: The Primer pairs from Gene bank database

Locus	Chromosomal Location	Repeat Length of PCR product	Annealing temperature (^{0} C)	Repeat motif	Primer sequences 5' to 3'	EMBI Accession number
UNH 207	U	138	58	(CA)n	F:ACA CAA CAA GCA GAT GGA GAC R:CAG GTG TGC AAG CAG AAG C	G12358
UNH 145	lg 8	162	55	(CA)n	F:CAT GCT GAA AGC TGA TTT R:ACC CAC ACC TAA AAT TAG AGA TA	G12297
UNH 146	lg 4	122	55	(CA)n	F:CCA CTC TGC CCC TCT AT R:AGC TGC GTC AAA CTC AAA G	G12298
UNH 172	lg 4	180	55	(CA)n	F:AAT GCC TTT AAA TGC CTT CA R:CTT TTA TAG TCG CCC TTT GTT A	G12324
UNH 166	lg 9	190	55	(CA)n	F:CCC TCA CAC ACA CTC TT R:GAT AAC GAC ACG ACA GTA C	G12318
UNH 190	lg 21	167	55	(CA)n	F:CGC GAT CGA GCA TTC TAA R:TGT CTG CAC GCG CTT TTG T	G12342
UNH 214	lg 10	165	50	(CA)n	F:TTC CAT AAT TGC TTT CTG GCA CGT TTT CCA TCA CTT CAA	G12365
UNH 216	lg 23	124	55	(CA)n	F:GGG AAA CTA AAG CTG AAA TA R:TGC AAG GAA TAT CAG CA	G12367
UNH 231	lg 6	176	50	(CA)n	F:GCC TAT TAG TCA AAG GCG T R:ATT TCT GCA AAA GTT TTC C	G12382
UNH 194	lg 22	195	56	(CA)n	F:ACT TAA TTT TTC AGC ATG ACA R:ACA CAG CCT GAA CTC TG	G12345
UNH 173	lg 13	174	56	(CA)n	F:CGT GAG AAA ACA ATG GT R:TAT TGA TTT TAT AGC TGT CGT G	G12325
UNH 149	lg 5	156	56	(CA)n	F:TTA AAA CCA GGC CTA CC R:GTT CTG AGC TCA TGC AT	G12364

Locus	Chromosomal Location	Repeat Length of PCR product	Annealing temperature (^{0} C)	Repeat motif	Primer sequences 5' to 3'	EMBI Accession number
UNH 213	lg 1	194	56	(CA)n	F:ACT GCT CCT CTT GTT TT R:TGT GAT AAG GTT AAT TAA AGT TAG G	G12364
UNH 211	lg 30	112	55	(CA)n	F:GGG AGG TGC TAG TCA TA R:CAA GGA AAA CAA TGG TGA TA	G12362
UNH 162	lg 19	195	55	(CA)n	F:CAG ACA CAG CAG AGG AT R:TGA TAA GTA ATT CAT CTG TTT G	G12314
UNH 120	lg 18	160	55	(CA)n	F:TAA GGC TCT ATG TGG TC R:TTA AAG GGG AAG AAA GA	G12273
UNH 104	lg 1	138	55	(CA)n	F:GCA GTT ATT TGT GGT CAC TA R:GGT ATA TGT CTA ACT GAA ATC C	G12257
UNH 111	lg 7	192	56	(CA)n	F:TGC TGT TCT TAT TTT CGC R:ATA AGA GTG TAT GCA TTA CTG G	G12264
UNH 172	lg 8	180	55	(CA)n	F:AAT GCC TTT AAA TGC CTT CA R:CTT TTA TAG TCG CCC TTT GTT A	G12324
UNH 233	U	173	55	(CA)n	F:TGC CGC CAT CTA TCT AT R:AAA CAC AAA GTG TGA GAC AGA TA	G12384

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations

UNH 104				
GT	PS	BN	EG	NS
202/146(1)	210/200(5)	175/172(1)	200/180(8)	200/180(4)
212/135(4)	210/210(2)	180/177(1)	180/180(2)	200/160(1)
209/185(2)	209/177(1)	200/200(1)	178/156(1)	200/159(1)
188/135(3)	160/162(2)	198/180(2)	180/155(1)	199/160(1)
180/180(3)	158/155(1)	200/177(1)	160/160(4)	198/157(1)
135/140(1)	160/158(1)	199/175(1)	200/158(1)	200/157(1)
140/140(1)	159/159(1)	200/180(3)	200/160(4)	197/171(1)
210/180(2)	197/139(2)	180/180(2)	180/160(5)	198/171(1)
206/158(1)	175/134(2)	176/174(1)	198/157(1)	200/179(1)
199/158(1)	178/140(2)	180/175(1)	199/159(1)	191/170(1)
200/161(6)	160/140(1)	177/176(1)	200/159(1)	206/181(1)
177/158(2)	180/140(3)	179/175(2)		200/162(1)
200/200(2)	175/140(1)	180/179(1)		191/152(1)
160/140(1)	180/180(2)	176/176(2)		200/200(2)
	171/132(1)	180/160(1)		202/181(1)
	188/140(1)	180/140(3)		200/157(1)
	179/141(2)	140/140(2)		192/185(1)
		138/149(1)		190/180(2)
		140/137(1)		201/160(1)
		136/134(1)		195/155(1)
		140/125(1)		200/165(1)
				209/180(1)
				180/180(3)

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations (Continued)

UNH 172				
GT	PS	BN	EG	NS
200/180(5)	210/200(11)	140/140(4)	210/180(4)	210/180(10)
191/172(1)	210/180(5)	181/139(1)	200/180(12)	208/171(1)
190/180(2)	200/200(3)	180/142(1)	209/180(1)	200/182(1)
192/184(1)	180/180(3)	208/171(1)	210/200(10)	197/175(1)
190/181(1)	200/180(2)	180/180(3)	200/200(3)	200/200(3)
206/171(1)	200/181(1)	175/169(1)		200/180(9)
180/180(2)	209/172(1)	212/175(1)		210/200(5)
200/200(6)	161/140(2)	210/180(1)		
178/170(1)	182/180(1)	200/160(1)		
195/175(1)	212/185(1)	140/140(2)		
190/170(3)		162/145(1)		
175/171(1)		160/142(1)		
179/180(1)		158/139(1)		
210/180(1)		162/140(1)		
205/176(1)		171/152(1)		
209/191(1)		139/140(1)		
200/195(1)		163/150(1)		
		182/142(1)		
		206/175(1)		
		149/143(1)		
		160/160(4)		

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations (Continued)

UNH 146				
GT	PS	BN	EG	NS
210/180(3)	140/120(4)	140/120(5)	200/180(5)	200/180(8)
181/163(1)	150/126(2)	160/140(3)	180/180(3)	180/180(3)
180/160(1)	120/120(3)	158/140(1)	210/200(2)	216/200(1)
160/160(3)	180/140(2)	158/136(1)	212/175(1)	200/200(2)
140/140(4)	177/140(1)	180/160(3)	180/160(2)	210/160(2)
161/132(1)	180/142(1)	160/160(2)	176/162(1)	209/161(1)
160/140(2)	176/139(1)	176/160(1)	172/160(1)	194/158(1)
158/136(1)	160/140(3)	192/178(1)	182/160(1)	192/164(1)
200/180(4)	200/180(2)	196/180(1)	191/161(1)	191/152(1)
162/141(1)	190/180(1)	165/177(1)	178/159(1)	199/158(1)
160/135(1)	188/165(1)	200/180(3)	171/132(1)	200/157(1)
159/157(1)	200/160(4)	178/165(1)	180/166(1)	206/182(1)
210/160(2)	212/136(1)	162/145(1)	182/180(1)	208/176(1)
177/158(1)	210/145(1)	177/159(1)	200/177(1)	198/157(1)
206/161(1)	141/121(1)	140/140(5)	160/160(4)	210/180(5)
200/140(3)	140/140(2)		200/200(4)	

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations (Continued)

UNH 207				
GT 180/160(4) 180/140(2) 160/160(3) 140/138(1) 140/140(6) 200/160(1) 209/161(1) 180/180(5) 210/180(3) 200/180(2) 200/200(2)	PS 210/200(2) 140/140(5) 200/200(5) 210/140(1) 200/180(5) 180/140(4) 210/172(1) 190/161(1) 180/180(2) 207/158(1) 181/140(1)	BN 200/140(4) 140/140(2) 180/142(1) 200/152(1) 195/160(1) 182/175(1) 200/160(2) 207/158(1) 210/180(2) 171/142(1) 160/160(2) 191/160(1) 210/192(1) 210/200(2) 179/180(1) 181/132(1) 200/200(3) 160/140(3)	EG 180/160(4) 210/180(3) 210/182(1) 180/180(2) 200/180(3) 195/171(1) 206/180(1) 180/163(1) 191/180(1) 210/195(1) 200/162(1) 197/160(1) 200/160(6) 200/200(4)	NS 200/160(10) 180/160(6) 180/140(7) 210/180(7)

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations (Continued)

UNH 166				
	PS	BN	EG	NS
200/180(8)	210/180(6)	200/180(5)	200/180(4)	180/160(3)
200/161(1)	200/180(4)	180/180(4)	200/200(3)	200/180(5)
180/160(7)	180/180(4)	210/190(1)	200/190(2)	211/180(1)
180/180(2)	197/171(1)	210/185(1)	210/180(2)	182/160(1)
210/160(3)	210/191(1)	190/162(1)	210/189(1)	190/190(2)
200/160(3)	186/164(1)	181/153(1)	191/160(1)	200/160(3)
210/171(1)	200/200(3)	181/159(1)	180/180(8)	192/167(1)
192/165(1)	187/160(1)	207/159(1)	190/180(2)	200/200(4)
207/182(1)	192/170(1)	180/140(3)	200/162(1)	171/160(1)
	210/190(2)	190/160(2)	210/199(1)	195/170(1)
	200/160(3)	187/162(1)	198/157(1)	190/170(2)
	210/200(1)	190/180(1)	190/190(4)	210/190(3)
	213/180(1)	200/191(1)		
	190/170(1)	179/160(1)		
		210/200(6)		

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations (Continued)

UNH 190				
GT	PS	BN	EG	NS
210/160(3)	180/180(5)	200/200(4)	180/160(12)	190/160(9)
200/160(4)	180/160(5)	212/200(1)	200/163(1)	200/180(4)
210/181(1)	210/182(1)	210/200(3)	200/160(5)	200/190(1)
210/180(3)	210/180(3)	200/160(7)	200/180(3)	180/180(3)
160/160(4)	210/193(1)	190/160(2)	210/160(1)	200/160(2)
210/200(6)	200/184(1)	160/160(2)	210/170(2)	213/190(1)
200/200(2)	208/187(1)	160/140(1)	215/180(2)	210/180(2)
210/210(1)	198/187(1)	200/140(2)	210/190(2)	190/176(1)
180/160(3)	210/190(1)	210/180(4)	200/175(1)	215/180(1)
	206/182(1)	180/180(2)	190/176(1)	187/161(1)
	195/170(1)	211/187(1)		210/170(1)
	200/160(4)	200/170(1)		200/170(3)
	160/160(3)			
	190/170(2)			

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations (Continued)

UNH 111				
GT 212/200(5) 200/200(5) 210/190(2) 200/180(3) 200/170(4) 210/191(1) 215/191(1) 190/170(2) 212/187(1) 220/200(3) 210/210(1) 190/170(2)	PS 180/160(1) 200/160(5) 180/180(1) 210/180(4) 210/200(3) 210/190(3) 185/160(1) 190/170(2) 215/172(1) 200/180(2) 160/150(2) 170/160(1) 200/170(1) 210/160(3)	BN 210/200(6) 200/180(11) 180/180(1) 210/180(4) 210/190(4) 204/170(1) 200/170(1) 195/170(2)	EG 200/160(13) 200/150(3) 200/200(3) 220/200(5) 210/190(1) 190/170(2) 215/170(3)	NS 200/160(4) 210/180(5) 215/180(1) 220/200(3) 210/190(1) 200/200(6) 210/160(7) 200/170(3)

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations (Continued)

UNH 173				
GT 210/200(5) 220/200(3) 200/200(7) 210/190(12) 210/180(1) 200/180(1) 190/190(1)	PS 200/160(10) 190/160(3) 180/180(4) 210/180(5) 200/120(5) 160/120(3)	BN 180/180(4) 200/180(2) 210/120(9) 200/200(3) 180/120(4) 200/120(3) 210/180(4) 210/190(1)	EG 210/180(3) 180/180(8) 200/180(11) 210/190(4) 200/171(1) 182/160(1) 190/190(2)	NS 200/180(6) 200/160(7) 180/180(3) 215/180(2) 200/200(3) 190/180(1) 190/170(4) 220/180(4)
UNH 189				
GT 210/200(5) 210/210(2) 200/160(7) 210/180(8) 190/170(8)	PS 200/160(7) 200/180(8) 180/180(2) 190/170(4) 180/140(3) 180/150(3) 160/140(3)	BN 210/180(3) 180/180(8) 200/180(11) 200/160(7)	EG 200/160(1) 200/159(1) 199/160(1) 200/180(2) 160/150(2) 170/160(1) 200/170(1) 210/160(3) 170/150(5) 210/180(7) 210/190(4) 180/180(2)	NS 190/170(11) 210/190(4) 190/150(2) 210/190(3) 200/160(8) 210/200(1) 180/160(1)

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations (Continued)

UNH 214				
GT 180/160(3) 160/160(2) 176/160(1) 160/140(8) 210/180(8) 190/170(8)	PS 170/150(5) 210/180(7) 200/160(8) 210/172(1) 190/161(1) 191/159(1) 204/170(1) 210/190(4) 220/200(1)	BN 220/200(6) 191/160(1) 210/192(1) 210/200(2) 160/160(2) 195/170(2) 190/170(5) 180/160(10) 191/150(1)	EG 180/160(4) 180/140(2) 160/160(3) 190/170(15) 210/210(1) 210/190(5)	NS 210/180(4) 190/160(12) 190/190(1) 220/200(3) 200/170(5) 193/170(1) 190/180(4)
UNH 216				
GT 223/200(1) 210/190(5) 200/160(16) 210/200(3) 182/160(1) 191/160(1) 190/170(3)	PS 210/180(6) 180/140(8) 190/140(4) 193/150(1) 200/160(5) 220/210(1) 220/185(1) 190/170(4)	BN 210/180(13) 190/180(5) 200/170(8) 192/170(1) 211/194(1) 205/180(1) 210/190(1)	EG 180/160(8) 200/180(15) 210/190(4) 200/171(1) 190/170(2)	NS 210/190(4) 214/170(1) 220/190(6) 195/170(2) 170/150(5) 180/160(4) 187/180(1) 190/160(4) 220/200(3)

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations (Continued)

UNH 149				
GT	PS	BN	EG	NS
210/200(5)	200/170(1)	212/175(1)	200/180(11)	200/180(6)
200/200(3)	210/160(3)	210/180(1)	210/190(4)	200/160(7)
200/160(7)	210/200(11)	200/160(1)	215/180(2)	180/180(3)
215/180(1)	210/180(5)	200/180(2)	200/160(13)	213/190(1)
210/180(4)	200/200(3)	210/120(9)		210/180(2)
190/160(9)	200/160(7)	200/200(3)		190/170(11)
190/190(1)		210/190(4)		
		204/170(1)		
		200/170(1)		
		195/170(2)		
		170/150(5)		

UNH 231				
GT	PS	BN	EG	NS
200/180(12)	200/140(9)	190/170(5)	160/140(6)	170/150(5)
200/161(1)	140/140(2)	180/160(10)	210/160(10)	180/160(4)
180/160(7)	180/142(1)	220/200(4)	190/190(2)	210/190(4)
180/180(2)	200/152(1)	200/200(3)	221/200(1)	220/200(3)
210/160(3)	195/160(1)	220/190(2)	190/180(5)	190/180(5)
190/170(2)	200/160(3)	200/191(1)	200/170(2)	193/150(1)
210/200(2)	200/170(3)	190/190(1)	212/180(1)	210/180(5)
	170/140(5)	210/200(4)	190/170(1)	
			210/210(2)	

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations (Continued)

UNH 233				
GT 210/160(13) 170/150(5) 210/180(7) 210/190(5)	PS 210/190(1) 206/182(1) 195/170(1) 200/160(4) 160/160(3) 190/170(2) 200/200(4) 171/160(1) 195/170(1) 220/200(8) 190/160(3) 200/200(1)	BN 160/160(4) 200/170(3) 200/192(1) 210/190(3) 190/170(9) 192/150(1) 205/180(1) 210/180(8)	EG 200/159(1) 199/160(1) 193/157(1) 200/157(1) 197/171(1) 196/171(1) 200/179(1) 191/170(1) 200/181(1) 210/200(6) 190/160(5) 200/200(3) 198/170(1) 170/170(6)	NS 190/150(10) 193/160(1) 200/180(6) 190/170(3) 170/150(5) 180/160(4) 197/150(1)

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations (Continued)

UNH 213				
GT 200/200(5) 210/190(2) 200/180(3) 200/170(4) 210/191(1) 190/170(9) 180/160(6)	PS 210/180(3) 210/193(1) 200/184(1) 208/187(1) 190/177(1) 210/190(1) 190/150(5) 200/170(8) 180/160(3) 172/150(1) 202/190(1) 210/200(7)	BN 180/180(5) 180/160(5) 210/182(1) 210/180(3) 210/193(1) 200/184(1) 190/180(4) 160/140(9)	EG 192/170(1) 211/194(1) 205/180(1) 210/190(1) 160/160(4) 200/170(3) 220/200(8) 190/160(3) 200/200(1) 210/180(7)	NS 180/160(1) 200/160(5) 180/180(1) 210/180(4) 210/200(3) 210/190(3) 190/150(10) 200/160(3)
UNH 145				
GT 210/180(6) 200/180(4) 180/180(4) 190/180(4) 160/140(9)	PS 210/190(5) 200/160(16) 210/200(3) 180/160(6)	BN 200/160(2) 209/150(1) 210/180(2) 171/152(1) 160/160(2) 191/160(1) 210/192(1) 210/200(2) 211/194(1) 200/180(1) 210/190(1) 190/150(10)	EG 200/180(5) 180/180(3) 210/200(2) 200/160(8) 190/180(2) 190/170(4) 212/180(1) 220/210(5)	NS 200/200(2) 160/140(1) 210/180(4) 210/200(3) 210/190(3) 180/140(10) 220/180(7)

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations (Continued)

UNH 162				
GT	PS	BN	EG	NS
200/180(2)	210/180(12)	196/160(1)	220/200(15)	170/140(6)
210/120(9)	190/170(5)	193/158(1)	200/200(4)	200/170(5)
200/200(3)	192/170(1)	200/157(1)	180/180(2)	180/160(10)
210/190(4)	200/200(3)	190/171(1)	210/190(7)	210/210(3)
200/170(1)	180/160(9)	196/171(1)	200/180(2)	172/146(1)
190/180(1)		200/170(10)		190/170(5)
170/150(6)		200/180(5)		
200/160(4)		209/180(1)		
		210/190(4)		
		180/160(5)		

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations (Continued)

UNH 194				
GT	PS	BN	EG	NS
210/180(3)	140/120(4)	140/120(5)	200/180(5)	200/180(8)
180/163(1)	150/126(2)	160/140(3)	180/180(3)	180/180(3)
180/160(1)	120/120(3)	158/140(1)	210/200(2)	216/200(1)
160/160(3)	180/140(2)	160/136(1)	212/170(1)	200/200(2)
140/140(4)	170/140(1)	180/160(3)	180/160(2)	210/160(2)
161/140(1)	180/142(1)	160/160(2)	176/150(1)	209/161(1)
160/140(2)	176/140(1)	176/160(1)	172/160(1)	194/160(1)
150/136(1)	160/140(3)	190/170(1)	182/160(1)	190/170(1)
200/180(4)	200/180(2)	196/180(1)	191/161(1)	190/160(1)
165/140(1)	190/180(1)	165/140(1)	188/160(1)	200/160(1)
160/135(1)	180/165(1)	200/180(3)	171/132(1)	200/157(1)
160/157(1)	200/160(4)	178/165(1)	180/166(1)	206/182(1)
210/160(2)	212/136(1)	166/140(1)	182/180(1)	208/176(1)
177/158(1)	210/145(1)	177/159(1)	200/177(1)	210/180(1)
206/161(1)	141/121(1)	140/140(5)	160/160(4)	210/180(5)
200/140(3)	140/140(2)		200/200(4)	
UNH 211				
GT	PS	BN	EG	NS
212/200(5)	180/160(1)	210/200(6)	200/160(13)	200/160(4)
200/200(5)	200/160(5)	200/180(11)	200/150(3)	210/180(5)
210/190(2)	180/180(1)	180/180(1)	200/200(3)	215/180(1)
200/180(3)	210/180(4)	210/180(4)	220/200(5)	220/200(3)
194/170(4)	210/200(3)	210/190(4)	210/190(1)	210/190(1)
210/190(1)	210/190(3)	204/170(1)	190/170(2)	200/200(6)
215/190(1)	180/164(1)	200/170(1)	210/170(3)	210/160(7)
190/160(2)	190/170(2)	190/170(2)		200/170(3)
212/187(1)	215/170(1)			
220/200(3)	200/180(2)			
210/210(1)	160/150(2)			
190/180(2)	170/160(1)			
	200/170(1)			
	210/160(3)			

Appendix 2: The observed genotypes for the 20 polymorphic loci, tested in the 5 populations (Continued)

UNH 120				
GT 210/160(6) 200/160(4) 210/181(1) 210/180(3) 160/160(4) 210/200(6) 200/200(2) 210/210(1)	PS 180/180(5) 180/160(5) 210/182(1) 215/180(3) 210/193(1) 201/184(1) 208/187(1) 198/187(1) 210/190(1) 206/180(1) 195/170(1) 200/160(4) 160/160(3)	BN 200/200(4) 212/200(1) 210/200(3) 200/160(7) 190/160(2) 160/160(2) 160/140(1) 200/140(2) 210/180(4) 180/180(2) 211/180(1) 200/170(1)	EG 180/160(12) 200/163(1) 200/160(5) 200/180(3) 210/160(1) 210/170(2) 215/180(2) 210/190(2) 200/175(1) 190/176(1)	NS 190/160(9) 200/180(4) 200/190(1) 180/180(3) 200/160(2) 213/190(1) 210/180(2) 190/170(1) 215/180(1) 180/160(1) 210/170(1) 200/170(3)

Appendix 3: Genetic distance measures of five populations.

pop ID	GT	PS	BN	EG	NS
GT	****				
PS	0.2187	****			
BN	0.4990	0.3709	****		
EG	0.3401	0.2940	0.4210	****	
NS	0.3036	0.3304	0.4115	0.0734	****

Nei's genetic distance (1972).

Genetic distance based on 20 microsatellite loci

pop ID	GT	PS	BN	EG	NS
GT	****				
PS	0.2065	****			
BN	0.4873	0.3577	****		
EG	0.3287	0.2813	0.4087	****	
NS	0.2918	0.3173	0.3988	0.0611	****

Nei's genetic distance (1978).

Genetic distance based on 20 microsatellite loci

Appendix 4: Overall Allele Frequency for 20 microsatellite loci among 5 populations

Allele \ Locus	UNH104	UNH172	UNH146	UNH207	UNH166	UNH190	UNH111	UNH173
Allele A	0.0376	0.1769	0.0214	0.0148	0.0269	0.0560	0.1836	0.1326
Allele B	0.2932	0.3269	0.1429	0.1556	0.4038	0.2910	0.4844	0.4205
Allele C	0.3045	0.3423	0.2893	0.4444	0.4385	0.3134	0.2227	0.3220
Allele D	0.2030	0.0154	0.1714	0.2185	0.1231	0.3321	0.1016	0.0682
Allele E	0.1617	0.0923	0.2750	0.1667	0.0077	0.0075	0.0078	0.0265
Allele F		0.0462	0.1000			0.0303		

Allele \ Locus	UNH149	UNH189	UNH214	UNH216	UNH231	UNH233	UNH213	UNH145
Allele A		0.0599	0.1224	0.0652	0.0931	0.1241	0.1678	0.0034
Allele B	0.3188	0.1901	0.4965	0.1993	0.4103	0.4793	0.6154	0.3699
Allele C	0.4819	0.4190	0.3042	0.5471	0.3966	0.2966	0.1888	0.3596
Allele D	0.1993	0.3310	0.0769	0.1884	0.1000	0.1000	0.0280	0.2671
Allele E								
Allele F								

Allele \ Locus	UNH162	UNH194	UNH211	UNH120
Allele A	0.3830	0.2148		0.1232
Allele B	0.4539	0.2958		0.3592
Allele C	0.1596	0.4225	0.5833	0.3275
Allele D	0.0035	0.0669	0.4167	0.1901
Allele E				
Allele F				

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