

Table 3. Distribution of simulated TMRCA, in years, for autosomal loci under each scenario simulated on the basis of median estimates

	Model		
	AFREG	ASEG	MREBIG
Quantiles for the TMRCA distribution			
0.00	103,525	102,675	133,500
0.05	135,950	207,586	429,200
0.10	140,175	283,550	540,300
0.15	185,696	343,859	643,325
0.20	249,775	397,745	742,800
0.25	310,244	449,575	845,650
0.30	369,975	500,275	864,300
0.35	430,000	551,900	866,425
0.40	492,250	605,650	868,725
0.45	556,739	661,050	871,200
0.50	626,200	721,175	873,875
0.55	699,875	785,775	876,900
0.60	782,100	857,310	880,225
0.65	873,400	933,725	883,975
0.70	975,775	1,020,908	888,300
0.75	1,095,000	1,121,425	893,475
0.80	1,242,105	1,242,280	899,725
0.85	1,428,979	1,397,750	907,850
0.90	1,689,533	1,620,278	919,275
0.95	2,143,954	1,991,964	938,850
1.00	7,870,725	6,685,300	1,189,675
Average TMRCA over all loci	807,262	862,251	813,530
Loci where MRCA is located in Africa, %	100	100	82.1
Loci where MRCA is located in Asia, %	0	0	17.9
Loci where MRCA is located in America, %	0	0	0

Model labels are as in Fig. 1