

Additional File 1

Table S1. Population information for the *Ae. aegypti* samples used in this study. Populations represented in both microsatellite and SNPs datasets are indicated by bold characters. In the analyses focused on the broader geographical region instead of the population level the populations were grouped as indicated in the “Broad Region” column.

Population	Country	Broad Region	Collected	N (msats/SNPs)
Francesville & Lope forest	Gabon	Africa	2014	34/0
Bijagos	Gouinea-Bissau	Africa	2009	30/0
Rabai-out	Kenya	Africa	2009-2012	34/0
Nairobi	Kenya	Africa	2012	34/8
Dakar	Senegal	Africa	2005	34/0
N'goye	Senegal	Africa	2007	34/0
Goudiry	Senegal	Africa	2012	34/12
Sedhiou	Senegal	Africa	2012	34/12
Kichwamba	Uganda	Africa	2009	34/0
Johannesburg	South Africa	Africa	2015	18/9
Luanda	Angola	Africa	2016	28/0
Townsville	Australia	Asia	2009	34/0
Cairns	Australia	Asia	2013	34/12
Tahiti	French Polynesia	Pacific	2010	34/12
Attock	Pakistan	Asia	2014	34/10
Cebu City	Philippines	Asia	2013	34/8
Riyadh	Saudi Arabia	Asia	2012	34/11
Rayong	Thailand	Asia	2009	34/0
Prachuabkhirikan	Thailand	Asia	2009	34/0
Bangkok	Thailand	Asia	2013	34/11
Ho Chih Minh	Vietnam	Asia	2013	34/12
Hanoi	Vietnam	Asia	2013	34/12
Zugdidi (coast)	Georgia	Black Sea	2015	22/0
Tbilisi (inland)	Georgia	Black Sea	2015	15/5
Hopa	Turkey	Black Sea	2015	20/10
Pazar	Turkey	Black Sea	2015	34/0
Cordoba	Argentina	New World	2006	26/0
Salta	Argentina	New World	2003, 2014	27/0
Cachoeiro	Brazil	New World	2008	34/0
Maraba	Brazil	New World	2010	34/12
Natal	Brazil	New World	2010	34/0
Jacobina	Brazil	New World	2013	34/12
Rio de Janeiro	Brazil	New World	2014	34/7
Cali	Colombia	New World	2013	34/6
Dominica	Commonwealth of Dominica	New World	2009	34/0
Siquirres	Costa Rica	New World	2014	34/6
Tijuana	Mexico	New World	2013	20/10
Las Palomas, GRO	Mexico	New World	2012	34/0
Mazatan, CHP	Mexico	New World	2012	34/0

Tapachula Norte, CHP	Mexico	New World	2012	34/12
Nogales, SON	Mexico	New World	2013	34/9
Chetumal, QRO	Mexico	New World	2013	34/0
Amacuzac, MOR	Mexico	New World	2014	34/12
Madeira	Portugal	Atlantic	2012	34/0
Patillas	Puerto Rico	New World	2014	34/12
	Republic of			
Trinidad	Trinidad and Tobago	New World	2014	34/12
Maricopa County, AZ	USA	New World	2013	34/0
Madera, CA	USA	New World	2013	34/10
Clovis, CA	USA	New World	2013	34/10
New Orleans, NO	USA	New World	2012	34/12
Miami, FL	USA	New World	2011	34/0
Columbus, GA	USA	New World	2011	34/8
North Key West, FL	USA	New World	2013	34/12
Hawaii	USA	Pacific	2010	25/0
Bolivar	Venezuela	New World	2004	34/0
Zulia	Venezuela	New World	2004	34/0
<i>Aedes mascarensis</i>	Mauritius	Indian Ocean	2014	0/2*
Total				1,795/308

*used as outgroup only for the phylogenetic analysis

Table S2. Priors and posteriors for the ABC analysis testing scenarios on the origin of Black Sea populations. Priors for splitting time and for population size are all uniform distributions. Posterior probabilities for each scenario depicted in Fig. 2 are shown with 95% confidence intervals. Error and posterior distributions of parameters (mean with 90% confidence interval) are reported for the best-fit scenarios. Divergence times are measured in generations.

Parameter	Details	Prior	Posterior
Colonization scenario	Scenario 1 – Black Sea from merge between Asia and New World Scenario 2 – Black Sea from New World Scenario 3 – Black Sea from Asia Scenario 4 – Black Sea from Africa Scenario 5 – Black Sea from New World	N/A	0.004 [0.000, 0.023] 0.411 [0.391, 0.430] 0.560 [0.545, 0.574] 0.026 [0.007, 0.044] 0.001 [0.0000, 0.019]
SCENARIO 2			
Effective population size	Africa New World Black Sea Asia	100 – 450,000 100 – 400,000 100 – 400,000 100 – 400,000	199,000 [82,900 – 370,000] 70,400 [14,400 – 207,000]* 80,800 [55,300 – 232,000]* 193,000 [40,800 – 350,000]*
¹ Split time	New World from Africa Asia from America Black Sea from America	4,500 – 6,000 500 – 1,800 100 – 1,800	5,340 [4,720 – 5,930] 1,550 [1,080 – 1,790] 876 [305 – 1,630]
Confidence	Type I error (Simulated under scenario 2) Type II error (Simulated under scenario 1) Type II error (Simulated under scenario 3) Type II error (Simulated under scenario 4) Type II error (Simulated under scenario 5)	N/A	0.120 0.000 0.180 0.012 0.032
SCENARIO 3			
Effective population size	Africa Asia New World Black Sea	100 – 450,000 100 – 400,000 100 – 400,000 100 – 400,000	207,000 [88,500 – 379,000] 196,000 [41,800 – 351,000]* 72,700 [14,200 – 205,000]* 79,900 [55,000 – 231,000]*
¹ Split time	New World from Africa Asia from America Black Sea from Asia	4,500 – 6,000 500 – 1,800 100 – 1,800	5,350 [4,730 – 5,930] 1,570 [981 – 1,750] 953 [431 – 1,490]
Confidence	Type I error (Simulated under scenario 3) Type II error	N/A	0.300

	(Simulated under scenario 1)	N/A	0.002
	Type II error		
	(Simulated under scenario 2)	N/A	0.100
	Type II error		
	(Simulated under scenario 4)	N/A	0.000
	Type II error		
	(Simulated under scenario 5)	N/A	0.044
Mutation rate	Microsatellite –Generalized Stepwise Mutation Model (GSM)	9.0×10^{-6} - 1×10^{-5}	9.5×10^{-6} [9.07×10^{-6} - 9.96×10^{-6}]

¹ Time in generations.

* Cases where the median of the absolute error/true value is > 0.3 that indicate low accuracy in the estimation of the specific parameter and scenario.

Table S3. Climatic data for Black Sea sampling localities (indicated by bold characters) and indicative worldwide sampling localities where *Ae. aegypti* is established as downloaded from www.worldclim.com.

Site	TempAnnual	maxTempWarmMonth	minTempColdMonth	meanTempWetQuar	meanTempDryQuar	meanTempWarmQuar	meanTempColdQuar
Nairobi	17.8	26.3	10.1	18.7	16.2	19.1	15.9
Johannesburg	16.2	26.5	2.2	19.9	10.8	20.2	10.7
Bangkok	28	34.6	20.3	28.1	26.2	29.5	26
Hanoi	23.9	33.1	13.4	28.5	17.5	29.1	17.5
Zugdidi	13.6	27.6	0.4	21.8	12.2	21.8	5.2
Tsereteli	11.8	29.2	-4.1	19.7	1.3	21.9	1.3
Hopa	12.4	25.4	0.5	10.1	10.5	19.9	4.9
Pazar	8.1	23.3	-5.5	11.8	0.3	16.8	-1.2
Jacobina	23	30.3	15.7	24.5	22.5	24.5	20.8
Rio de Janeiro	22.4	29.7	15.2	24.9	20	25.2	20
Nogales	16.5	34	-0.5	24.4	19.4	24.9	8.4
New Orleans	20.3	32.7	5.8	27.2	16.7	27.5	12.3
Miami	24.2	31.5	15.5	27	20	27.8	20

Temperature is estimated in ° C. Abbreviations: TempAnnual: mean annual temperature, maxTempWarmMonth: maximum temperature for the warmest month of the year, minTempColdMonth: minimum temperature for the coldest month of the year, meanTempWetQuar: mean temperature for the wettest quarter of the year, meanTempDryQuar: mean temperature for the driest quarter of the year, meanTempWarmQuar: mean temperature for the warmest quarter of the year, meanTempColdQuar: mean temperature for the coldest quarter of the year.

Table S4. Genetic diversity. Summary of the population genetic diversity statistics for the 56 populations of *Ae. aegypti* consisting the microsatellite dataset.

Population	Na	Ne	Ho	He	uHe	Fis	AR(30)
Luanda, Angola	6.17	3.40	0.62	0.67	0.68	0.07	5.36
Dakar, Senegal	4.92	2.62	0.61	0.57	0.57	-0.08	4.20
Bijagos, Gouinea-Bissau	6.42	3.59	0.60	0.65	0.66	0.09	5.70
Kichwamba, Uganda	6.08	3.77	0.62	0.69	0.70	0.10	5.58
N'goye, Senegal	6.50	3.84	0.64	0.69	0.70	0.07	5.53
Goudiry, Senegal	4.25	2.76	0.50	0.59	0.60	0.16	3.85
Sedhiou, Senegal	4.83	2.74	0.60	0.58	0.59	-0.03	4.34
Francesville, Gabon	7.17	4.02	0.62	0.70	0.71	0.10	6.15
Johannesburg, South Africa	5.92	3.33	0.62	0.63	0.64	<0.01	5.66
Maricopa, AZ, USA	4.33	2.52	0.54	0.56	0.57	0.05	3.95
Tijuana, Mexico	2.92	2.40	0.62	0.55	0.56	-0.13	2.91
Madera, CA, USA	3.92	2.62	0.53	0.54	0.55	0.02	3.60
Clovis, CA, USA	3.25	2.36	0.44	0.48	0.48	0.05	3.14
New Orleans, NO, USA	5.83	3.06	0.58	0.61	0.62	0.05	4.86
Cali, Colombia	4.17	2.25	0.46	0.52	0.53	0.15	3.56
Bolivar, Venezuela	4.25	2.61	0.55	0.56	0.57	<0.01	3.82
Cachoeiro, Brazil	4.00	2.24	0.46	0.49	0.50	0.09	3.50
Zulia, Venezuela	3.92	2.58	0.55	0.52	0.53	-0.09	3.65
Maraba, Brazil	4.08	2.47	0.53	0.56	0.57	0.04	3.64
Natal, Brazil	3.75	2.21	0.48	0.48	0.49	0.01	3.30
Miami, FLO, USA	5.42	3.45	0.65	0.63	0.63	-0.05	4.68
Las Palomas, Mexico	3.83	2.67	0.56	0.56	0.57	-0.02	3.57
Mazatan, Mexico	3.25	2.24	0.51	0.48	0.48	-0.05	3.14
Columbus, GA, USA	4.83	3.22	0.64	0.62	0.63	<0.01	4.32
Tapachula, Mexico	3.58	2.22	0.50	0.49	0.49	-0.02	3.27
Jacobina, Brazil	3.83	2.07	0.43	0.44	0.45	0.05	3.35
Nogales, Mexico	4.00	2.40	0.55	0.55	0.56	<0.01	3.70
Chetumal, Mexico	3.67	2.47	0.53	0.49	0.50	-0.06	3.48
Siquirres, Costa Rica	4.33	2.88	0.58	0.60	0.61	0.04	3.96
Rio de Janeiro, Brazil	4.42	2.15	0.42	0.48	0.49	0.11	3.89
Amacuzac, Mexico	4.00	2.87	0.55	0.55	0.56	<0.01	3.76
Cordoba, Argentina	4.08	2.56	0.56	0.56	0.58	0.03	3.75
Salta, Argentina	5.17	3.40	0.63	0.63	0.65	0.00	4.85
Key West, FLO, USA	4.92	3.18	0.62	0.62	0.63	0.01	4.37
Patillas, Puerto Rico	4.33	2.98	0.57	0.54	0.55	-0.07	4.03
Trinidad	4.17	2.66	0.47	0.53	0.53	0.15	3.87
Dominica	3.50	2.33	0.44	0.46	0.47	0.02	3.28
HCM, Vietnam	5.17	2.74	0.60	0.59	0.60	-0.02	4.46
Hanoi, Vietnam	3.42	2.09	0.46	0.47	0.48	0.09	3.20
Cebu City, Philippines	5.50	2.77	0.58	0.58	0.59	<0.01	4.66

Attock, Pakistan	4.75	2.94	0.52	0.59	0.59	0.12	4.31
Saudi Arabia	5.08	2.97	0.58	0.58	0.60	<0.01	4.53
Rayong, Thailand	4.33	2.48	0.56	0.55	0.56	<0.01	3.94
Prachuabkhirikan, Thailand	4.75	2.58	0.53	0.59	0.56	0.09	4.15
Townsville, Australia	3.83	2.63	0.51	0.55	0.56	0.07	3.63
Cairns, Australia	4.08	2.74	0.59	0.6	0.60	0.02	3.76
Bangkok, Thailand	4.67	2.84	0.59	0.59	0.60	-0.01	4.20
Madera, Portugal	3.58	2.43	0.48	0.54	0.55	0.12	3.30
Hawaii, USA	3.25	2.15	0.52	0.49	0.50	-0.07	3.12
Tahiti, French Polynesia	4.33	2.50	0.57	0.58	0.59	0.02	3.93
Rabai-out, Kenya	8.08	3.86	0.66	0.69	0.71	0.06	6.43
Nairobi, Kenya	7.75	3.61	0.60	0.68	0.70	0.11	6.20
Hopa, Turkey	3.17	2.32	0.50	0.52	0.54	0.03	3.10
Pazar, Turkey	3.08	2.00	0.53	0.47	0.48	-0.04	2.70
Coast, Georgia	3.00	2.16	0.50	0.51	0.52	<0.01	2.92
Inland, Georgia	2.67	2.00	0.42	0.45	0.46	0.05	2.67

Abbreviations: Na: total number of different alleles, Ho: observed Heterozygosity, He: expected Heterozygosity, uHe: Unbiased Expected Heterozygosity, Ne: Number of effective alleles, Fis: Fixation index, AR(30): allele richness estimated by rarefaction (N=30 genes).

Figure Legends

Figure S1. STRUCTURE bar plots based on the microsatellites dataset, including equal number of populations from each region. Population names are reported on their X axes. For each STRUCTURE run only the number of genetic clusters supported by the Evanno method is presented. For details see legend of Fig. 4.

Figure S2. STRUCTURE bar plots based on the microsatellites dataset, including equal number of *Ae. ae. aegypti* populations from each region. For each STRUCTURE run only the number of genetic clusters supported by the Evanno method is presented. For details see legend of Fig. 4.

Figure S3. Principal Components Analysis (PCA) on the global (a) and the *Ae. ae. aegypti* (b) microsatellite dataset as implemented and plotted using the ade4 package in R. Populations originated from different regions are presented with different colors. A barplot of the eigenvalues of the each PCA analysis is shown in the respective inset. *Abbreviations:* Aus, Australia; Pac, Pacific; Atl, Atlantic; BS, Black Sea.

Figure S4. Discriminant Analysis of Principal Components (DAPC) for the *Ae. ae. aegypti* populations based on the microsatellite dataset. The graph represents the individuals as dots and the groups as inertia ellipses. A barplot of eigenvalues for the discriminant analysis (DA eigenvalues) is displayed in the inset. The number of bars represent the number of discriminant functions that retained in the analysis and the eigenvalues correspond to the ratio of the variance between groups over the variance within groups for each discriminant function.

Figure S5. Discriminant Analysis of Principal Components (DAPC) for the *Ae. aegypti* populations collected from Turkey and Georgia based on the microsatellite dataset. For details see legend of Fig. S4.

Figure S6. Assignment test as implemented in Geneclass2 for the Black Sea populations and using the remaining worldwide populations as reference panel. The individual mosquitoes from Georgia and Turkey were assigned to the populations presented on the X axis of the barplots. The percentage of assignment for each one the Black Sea population to each one of the reference panel's population is presented on the Y axis.

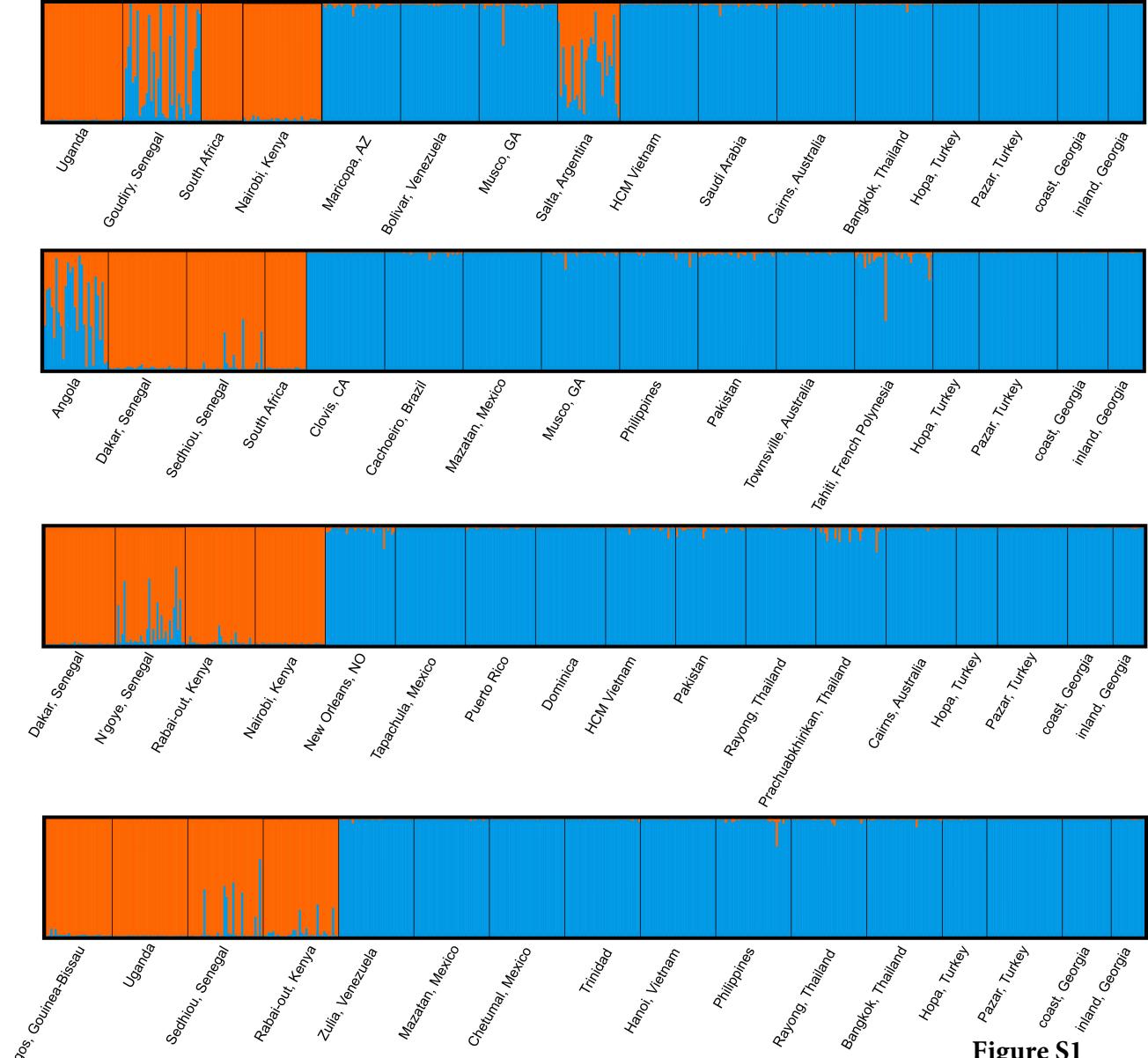


Figure S1

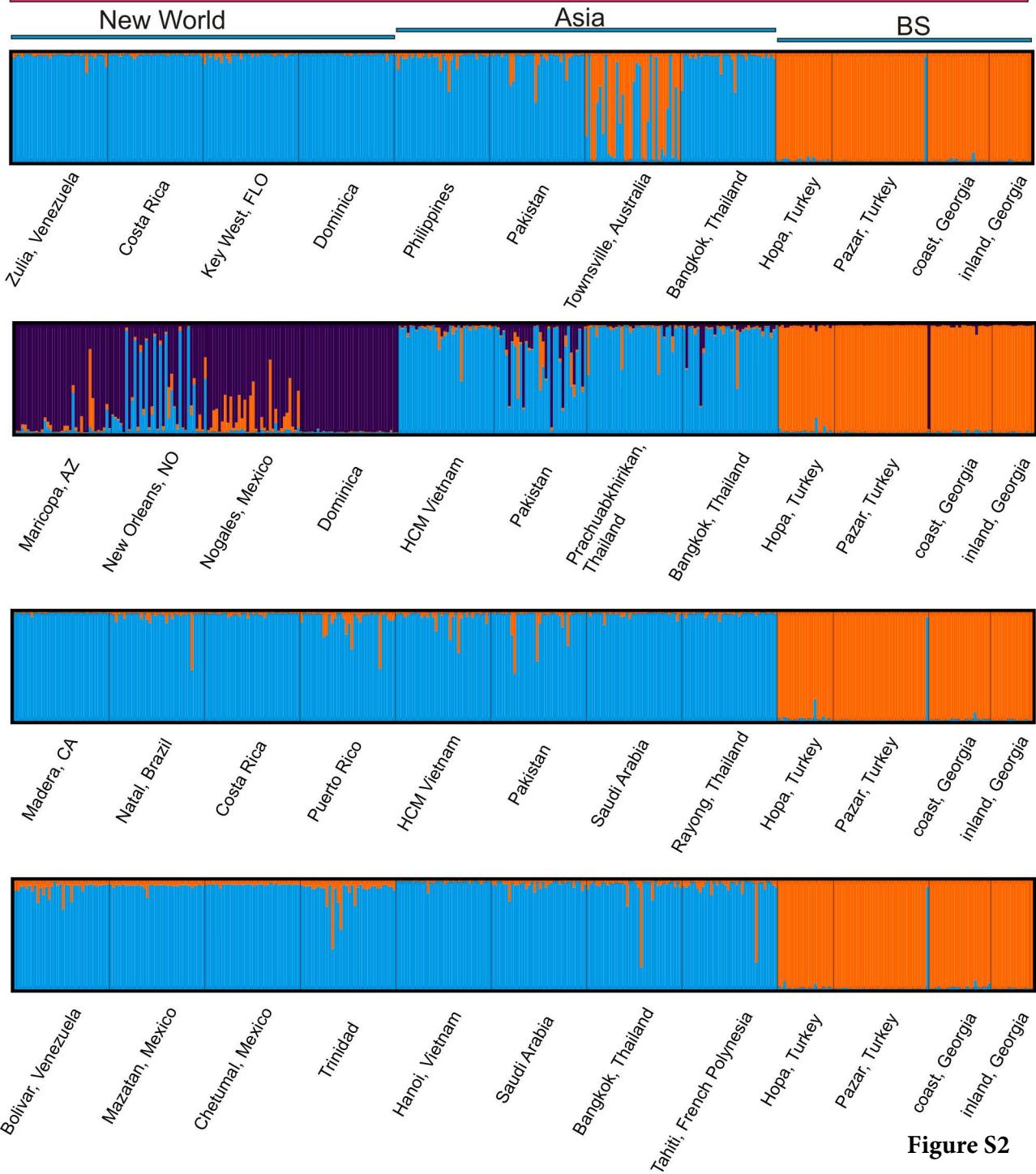
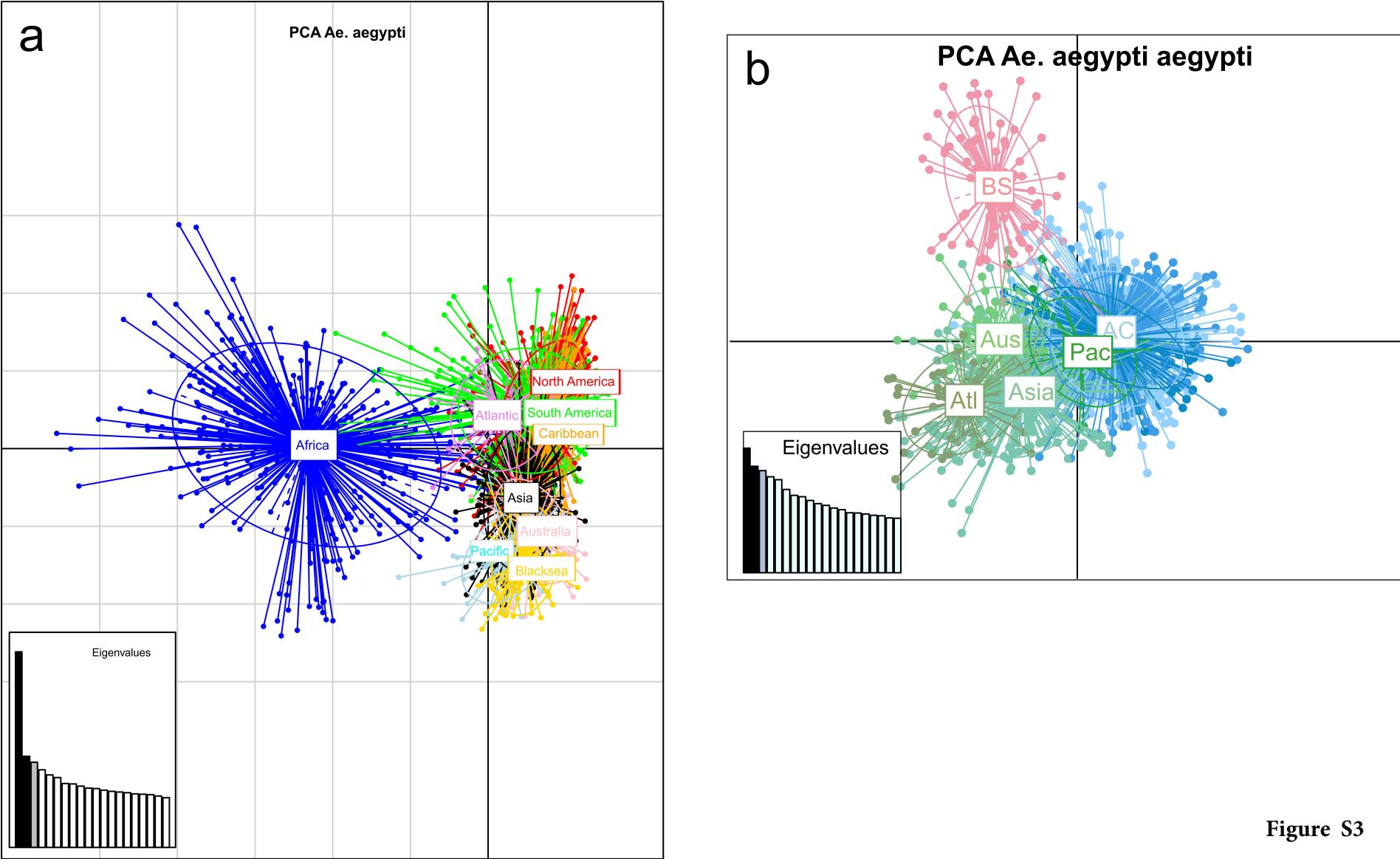


Figure S2



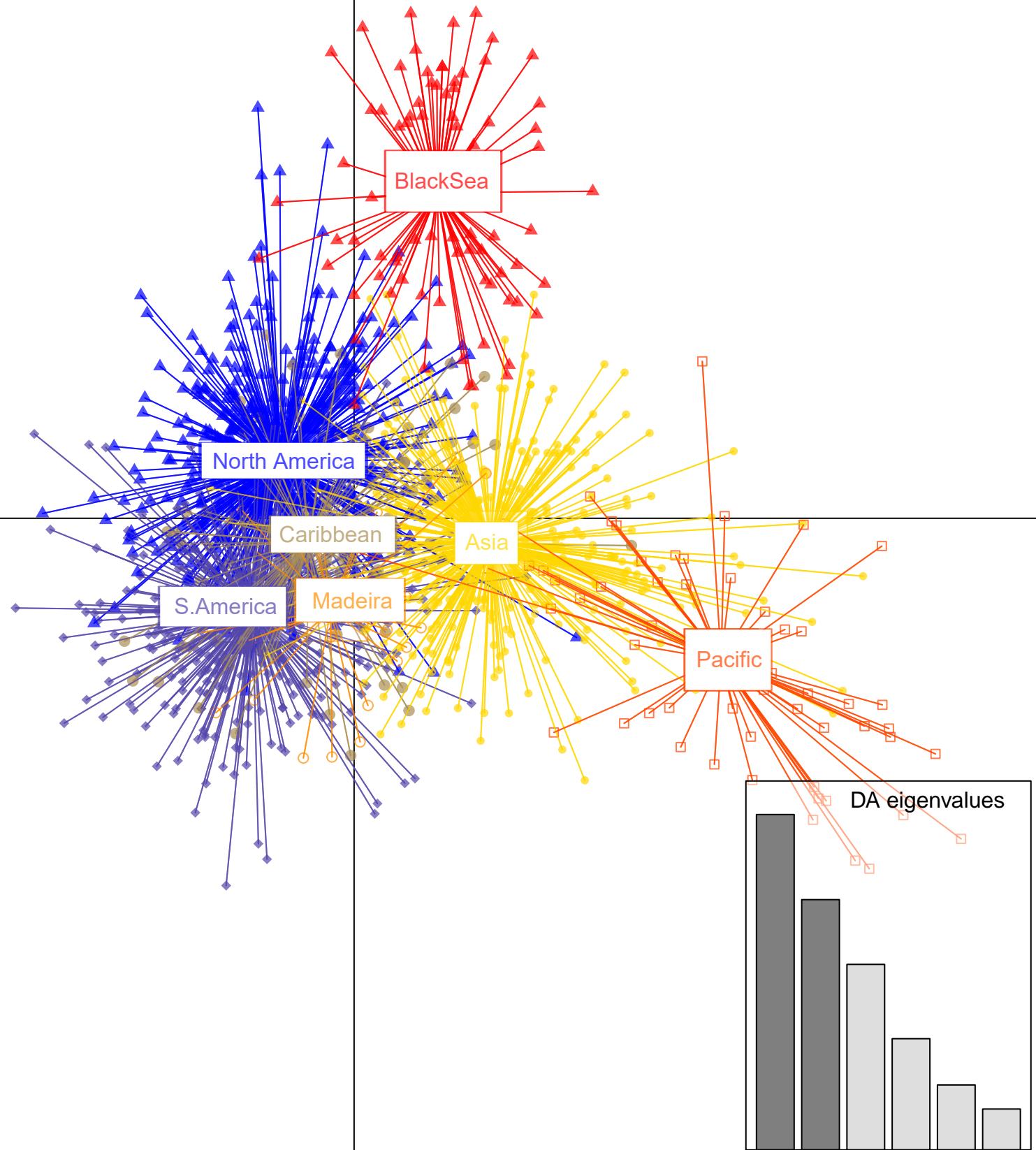


Figure S4

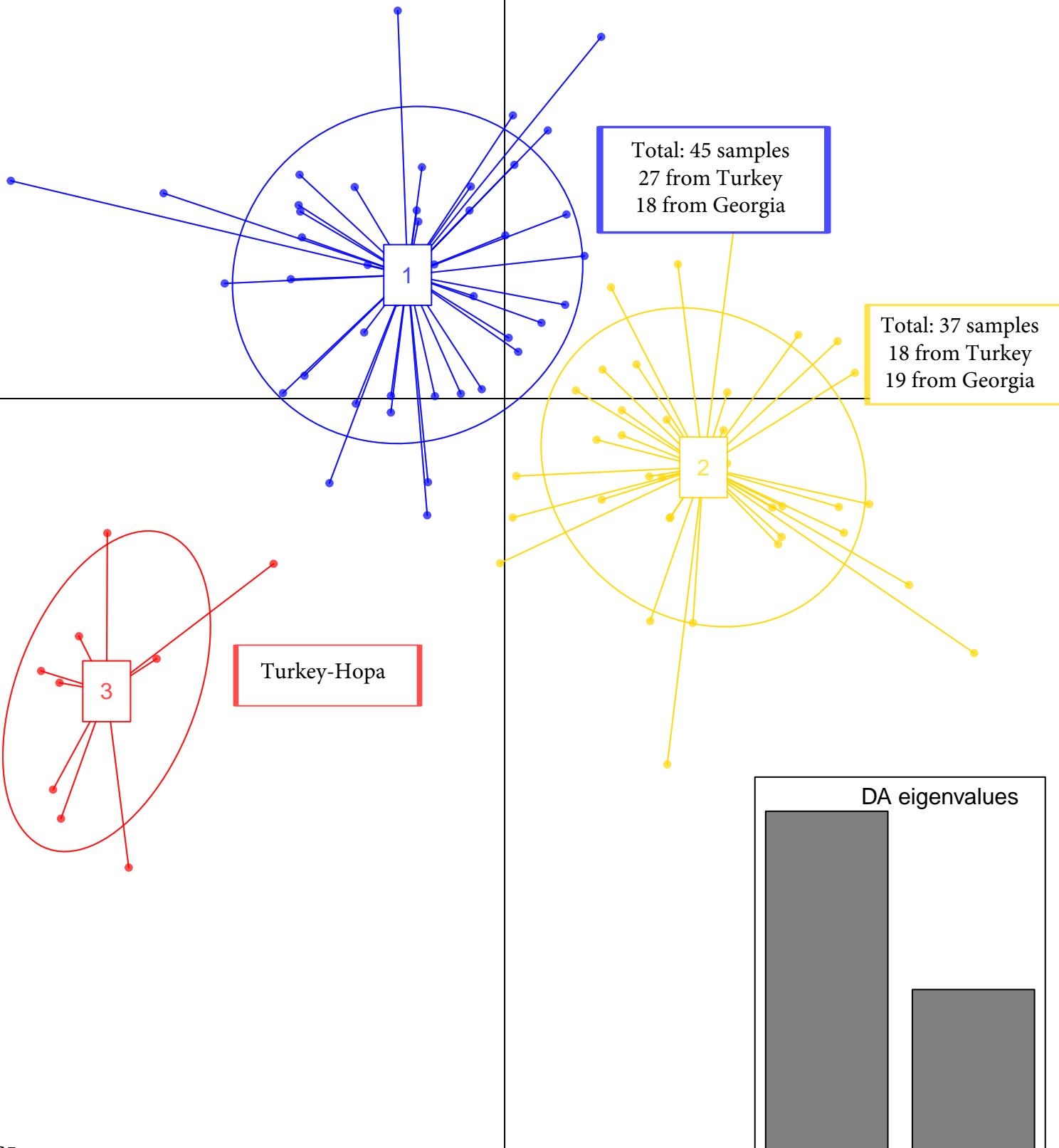
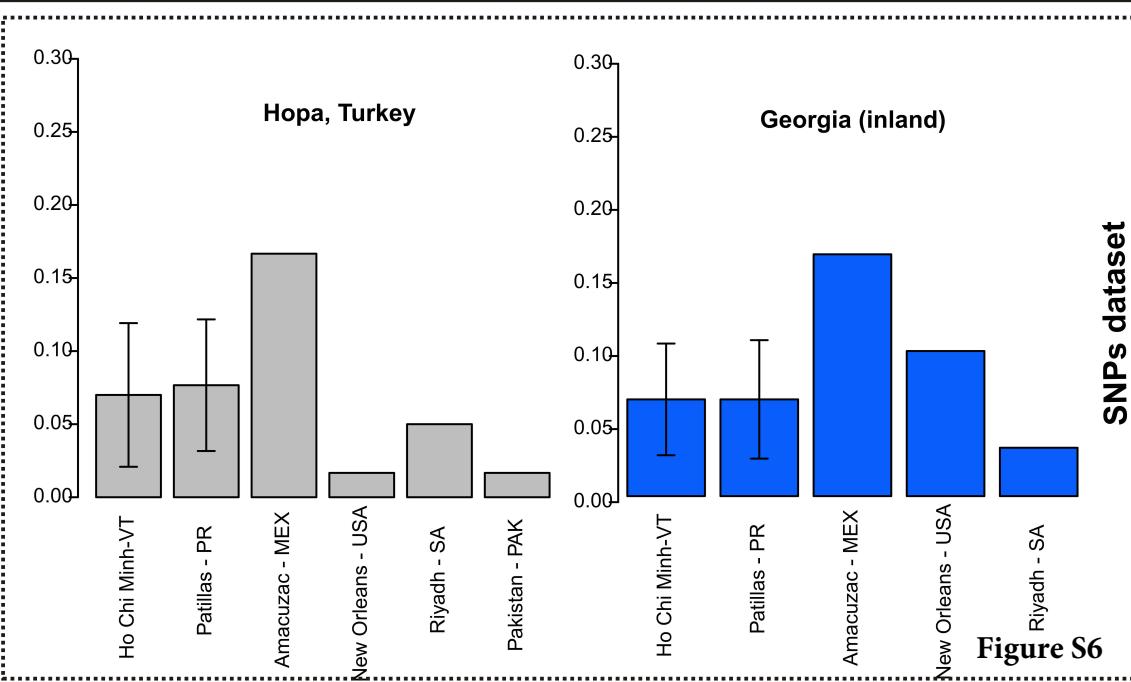
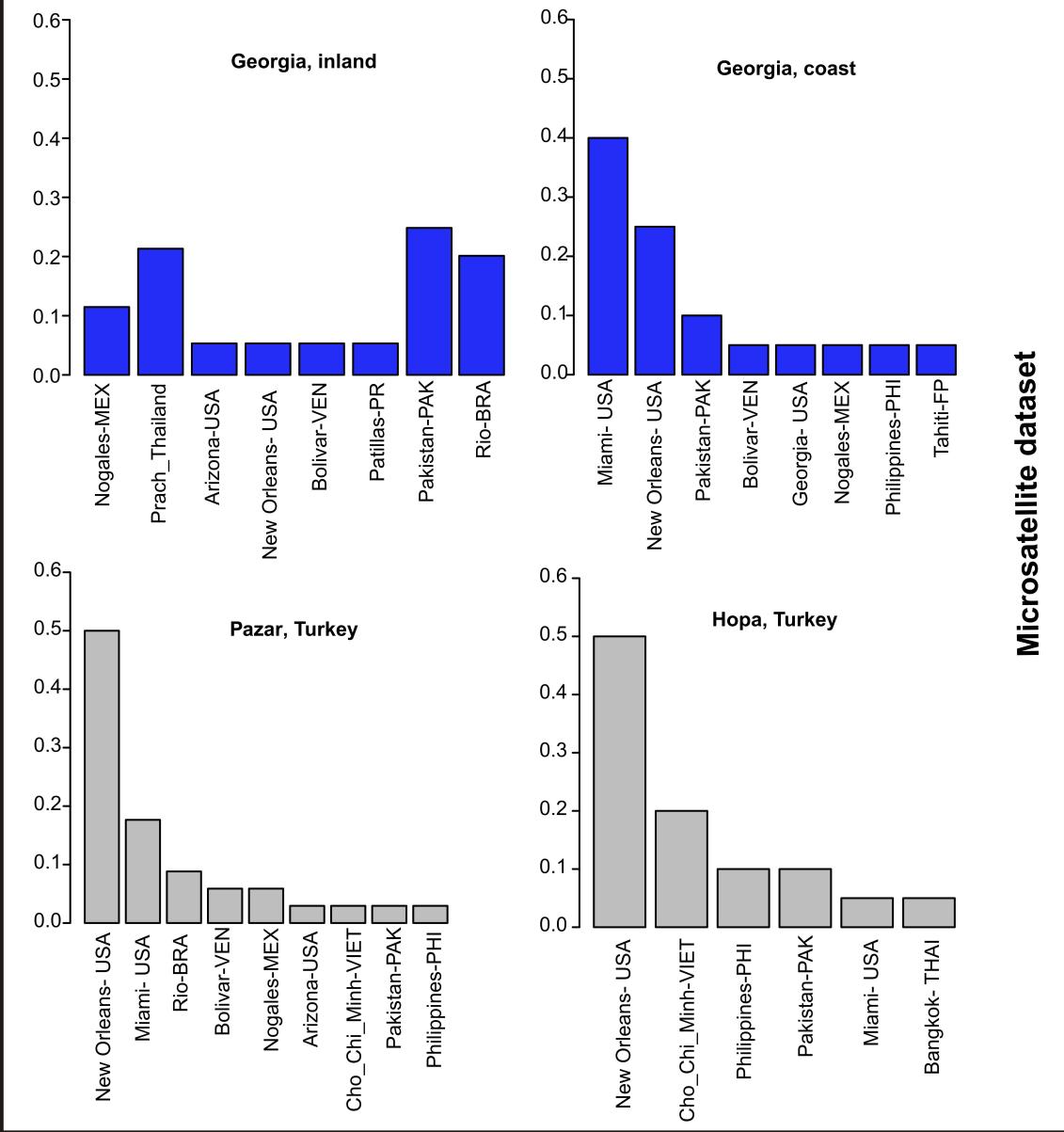


Figure S5

Microsatellite dataset



SNPs dataset

Figure S6