Supplementary Information

VALIDATION OF NEW D-LOOP HAPLOTYPES OF THE LOGGERHEAD TURTLES FROM SOUTHWEST ATLANTIC

Figure S1. Polymorphic sites between the new D-loop haplotypes (indicated with asterisks) do *Caretta caretta* of the Southwest Atlantic compared with the known CCA1.1, CCA4.1, CCA4.2, CCA4.3, CCA24.1, and CCA25.1.

Base pair	s [10	20	30	40	50	60	70	80]
00444	10000		OTTA	0.7.0.7	17010		ATOO	0014	00
CCA1.1	ACCG	JAATT	GITA	UIGI	ATCAG	AGAAA	AICC	CGIA	GU
CCA4.1	T		A.	A		.T(3A	A	
CCA4.2	T		A.	A	<mark>-</mark>	.T(GC	AA	
CCA4.3	T		A.	A		.TA	GC.	.AA	
CCA4.4*	T		A	AT		G	CA	A	
CCA4.5*	T	C	A	A	Τ	T	.GC	AA	
CCA4.6*	T		A	A	T	T	.GC	AA	
CCA4.7*	T		A	CA.	T	T	G	CAA	
CCA4.8*	TC	,		AA.	T	T	G	CAA	
CCA24.1	T	G.		AA.		T	GC	AA	
CCA25.1	T		A	CA.		T	GC	AA	

Figure S2. ABGD results measure of distance between the Caretta caretta lineages compared with other worldwide.

