

Supplemental Table S6. Fst values for RISP from arlequin based on pairwise comparisons of populations (pairwise sequence divergence). Values that were not significant based on permutation analyses are indicated by (NS). Significance is based on $\alpha=0.05$ in a Sequential Bonferroni procedure.

		1	2	3	4	5	6	7	8	9	10	11
		RP1	AB	IP	PVL	LH	SCN	LJS	LJP	BR	PES	SD
1	RP1	0										
2	AB	0.75429	0									
3	IP	0.27619	0.56562	0								
4	PVL	0.53356	0.41714	0.35616	0							
5	LH	0.99709	0.9952	0.99394	0.98755	0						
6	SCN	0.99522	0.99361	0.99252	0.98706	-0.02967 (NS)	0					
7	LJS	0.95093	0.9485	0.94659	0.93843	0.9532	0.95671	0				
8	LJP	0.97223	0.96981	0.96811	0.9601	0.97347	0.97416	0.06357	0			
9	BR	0.97566	0.97347	0.97203	0.96491	0.97707	0.97721	0.12096	0.23269	0		
10	PES	0.945	0.94243	0.94051	0.93206	0.77278	0.80061	0.87616	0.90152	0.91641	0	
11	SD	0.9911	0.98823	0.98617	0.9763	0.99404	0.9913	0.64197	0.76682	0.79607	0.90822	0