



## Genomic analysis of DNA from archived shark jaws

**Manuzzi, Alice**

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Page 18, line 9: “Manuzzi et al. 2018” to “Manuzzi et al. 2019”

Page 24, line 9: “were drawn” to “were drawn”

Page 61, Table 2: Some of the (\*) symbol indicating significance are wrong. Here is reported the correct table.

	<b>NWAtlantic</b>	<b>SWAtlantic</b>	<b>SWIndian</b>	<b>EIndian</b>	<b>SWPacific</b>	<b>Hawaii</b>
<b>NWAtlantic</b>		0.1628	0.0075*	0.0000*	0.0000*	0.0005*
<b>SWAtlantic</b>	0.0088		0.0257*	0.0124*	0.0115*	0.0115*
<b>SWIndian</b>	0.1065*	0.1551*		0.4204	0.3852	0.3880
<b>EIndian</b>	0.1091*	0.1452*	0.0051		0.6930	1.0000
<b>SWPacific</b>	0.1021*	0.1448*	0.0060	-0.0004		0.4950
<b>Hawaii</b>	0.0968*	0.1280*	0.0027	-0.0032	0.0006	

Page 83, line 17: Change from “A test of selection was performed using pcadapt (Luu et al. 2017), with qvalue of 0.1 as cut-off to identify putative selective outliers and bayescan () with” to “A test of selection was performed using pcadapt (Luu et al. 2017), with qvalue of 0.1 as cut-off to identify putative selective outliers and bayescan (Foll & Gaggiotti, 2008) with parameters -n 5000 -thin 10 -nbp 20 -pilot 5000 -burn 50000 -pr\_odds 100”

Page 105, Table S2: Wrong caption for the table, the correct version would be: “Pairwise FST estimates (lower diagonal) and p-values (upper diagonal) between all pairwise samples. Sample size per collection: GCA\_1970-1980 (6), GCA\_1990 (18), CRS\_2000 (16), TAS\_1910-1960 (20), TAS\_1970-1980 (13), TAS\_1990 (18), TAS\_2000 (7). The star symbol (\*) identifies significant values.”

Page 121, line 24: “Central Pacific (COP)” to “Central Pacific (CPO)”

Page 147, caption Figure 2: “(Weir & Cockerham 1996)” to “(Weir & Cockerham 1984)”

Page 173, caption Figure S8: “(Weir & Cockerham 1996)” to “(Weir & Cockerham 1984)”

Page 174, caption Figure S9: “(Weir & Cockerham 1996)” to “(Weir & Cockerham 1984)”