

**Table S2. Nested Procrustes ANOVA on shape for device comparison: Only specimens from the Baltic Sea population.** We applied the following nested hierarchical structure: Specimen > Device > Operator > Landmark replica. The R-squared values (Rsq) give an estimate of the relative contribution of each factor to the total shape variation.

<b>Variables</b>	<b>Df</b>	<b>SS</b>	<b>MS</b>	<b>Rsq</b>	<b>F</b>	<b>Z</b>	<b>Pr(&gt;F)</b>
<i>Device Comparison Dataset: Only specimens from the Baltic Sea population</i>							
Specimen	16	0.348	0.022	0.894	184.509	20.325	0.001
Specimen:Device	17	0.014	0.001	0.035	6.803	16.463	0.001
Specimen:Device:Operator	34	0.020	0.001	0.050	4.887	16.615	0.001
Residuals (Landmark replica)	68	0.008	0.000	0.021			
Total	135	0.389					