

**Table S1: Nucleotide composition per species based on mitochondrial NADH2 gene sequence data of sharks landed from by-catch along the northern Pacific Coast of Colombia.**

<b>Species</b>	<b>C</b>	<b>T</b>	<b>A</b>	<b>G</b>
<i>Mustelus lunulatus</i>	29.08%	30.01%	30.41%	10.49%
<i>Sphyrna lewini</i>	32.46%	27.12%	31.52%	8.91%
<i>Mustelus henlei</i>	9.97%	30.66%	29.72%	29.65%
<i>Carcharhinus falciformis</i>	29.22%	29.98%	31.71%	9.10%
<i>Carcharhinus limbatus</i>	8.82%	31.89%	29.40%	29.89%