

Supplementary Material to: Evaluation of the occurrence of multiple paternity in *Squalus acanthias* in the South Atlantic region using nuclear markers

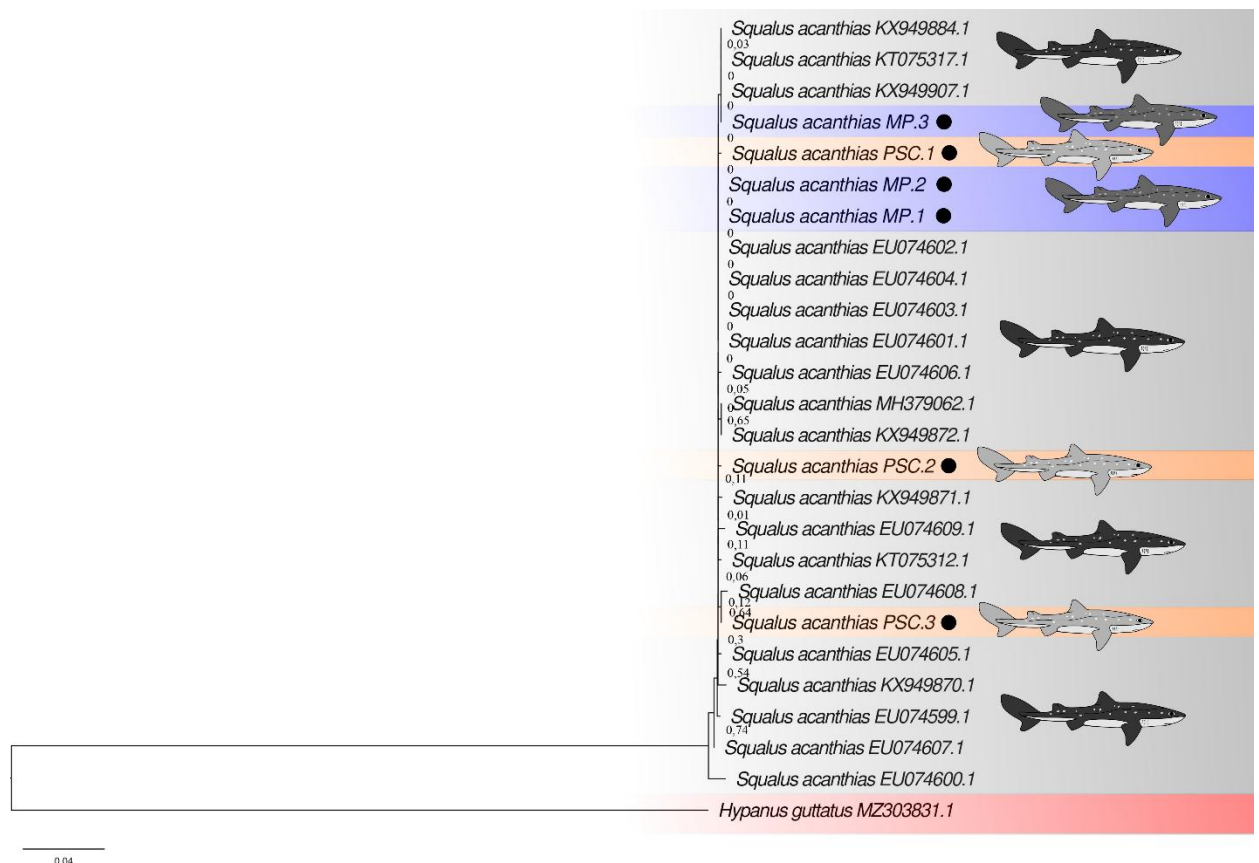


Figure S1 - Maximum Likelihood tree of *Squalus acanthias* specimens based on mitochondrial cytochrome c oxidase subunit I (COI) gene sequences under the K2P model. The asterisks (*) indicate the six samples sequenced in this study. Sequences highlighted in gray correspond to reference sequences obtained from public databases. Sequences in blue represent *S. acanthias* samples from Mar del Plata (MP) generated in this study, while those in orange represent *S. acanthias* samples from Puerto de Santa Cruz (PSC) generated in this study. The sequence highlighted in red corresponds to *Hypanus guttatus*, used as outgroup.