

Supplementary Data 1. Bowfin Scpp gene predictions.

The initiation codon, the termination codon, and the poly(A) signal are shown in red, and the nucleotides at the beginning of exons are highlighted in yellow. The nucleotide sequences for mepe, dmp1, enam, scpp1p20 are incomplete.

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>mepe: 5' end & 3' end missing
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>ibsp
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>dmp1: a unsequenced gap shown by NNNNNNNNNNNNN
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>odam

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>scpp3ba: borders between exons 3 and 4 not determined

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