Supplementary material S1
Fasta file of *S. nonagrioides* predicted protein sequences and corresponding transcript sequences

>SnonGOBP1
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>Snon-Fatty-acyl CoA reductase 1

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>Snon-Fatty-acyl CoA reductase 2

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>Snon-Fatty-acyl CoA reductase 3

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>Snon-Fatty-acyl CoA reductase 4

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>Snon-Fatty-acyl CoA reductase 5

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>Snon-Fatty-acyl CoA reductase 6

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>Snon-Fatty-acyl CoA reductase b

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>Snon-Fatty-acyl CoA reductase II

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>SnonGOBP1

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TTTTATACAATTTAATAACCGTCACCCTTCAGATTATACCTGCGTTCTTGATTGATTCATGGTGTATTTTAACTGGAAAGAAAGCAATACAAATTA
AAATACAGAACCGGCTGAAGAAAGTCGTAGATGCAGTGAAGTTCTTTTTACTCAACGAATGGAGGTTTTCTGATAACAATGTAAGAAATCTTTTA
AAAACTATGTCTTATGTGGACAAAGAAATGTTTAACTTTGACGTGCATTCGATTAACTGGGAAACCTGTATTAGAGATTACGTTCTAGGAGCTAG
GAAATACTTGTTGAAACATGAGAAA

Supplementary material S2

Maximum likelihood tree of candidate chemosensory proteins (CSPs) from *S. nonagrioides* and other Lepidoptera. Sequences used were from *B. mori* [63], *S. littoralis* [17, 18, 56], *H. melpomene* [58], *H. virescens* [47, 60, 61] and *P. xuthus* [65]. Signal peptide sequences were removed from the data set. Branch support was estimated by approximate likelihood-ratio test (aLRT) (circles: >0.95) [73]. Images were created using the iTOL web server [87]. The SnonCSPs identified in this study are in red.