

Supporting Information

Antennal RNA-sequencing analysis reveals evolutionary aspects of chemosensory proteins in the carpenter ant *Camponotus japonicus*

Masaru K. Hojo^{1*}, Kenichi Ishii^{2*}, Midori Sakura¹, Katsushi Yamaguchi³, Shuji Shigenobu^{3,4}, Mamiko Ozaki¹

¹Department of Biology, Graduate School of Science, Kobe University, Kobe, Japan

²Department of Biological Sciences, Graduate School of Science, The University of Tokyo, Tokyo, Japan

³NIBB Core Research Facilities, National Institute for Basic Biology, National Institutes of Natural Sciences, Okazaki, Japan

⁴Department of Basic Biology, School of Life Science, Graduate University for Advanced Studies, Okazaki, Japan

*Equal contribution

Supplemental Tables

Supplemental Table S1. Nucleotide sequences of primers used for cloning of *CjapCSPs*.

Lower-case letters indicate additional sequences for restriction enzyme recognition.

| Gene | Forward primer (5'-3') | Reverse primer (5'-3') |
|------------------|--------------------------------------|--|
| <i>CjapCSP1</i> | aaggatccGAGGAAATGTACTCGG ACATG | aactcgagTCAACGGGTAAAGGGGATG |
| <i>CjapCSP2</i> | aaggatccGCCGAATCGGACAATA GCG | aactcgagTTATGAAACTCCAGCGTATT GTTG |
| <i>CjapCSP3</i> | aaggatccCGGCCACAAGACCATT ATAC | aactcgagTTAATTCTTCTTCAAGG CTCGATC |
| <i>CjapCSP4</i> | aaggatccGAGGACGTGCAGTATA CTAC | aactcgagTCAATCCAATCTGCCTCCTT C |
| <i>CjapCSP7</i> | aaggatccGACGATAAGTACACGA CCAAG | aactcgagTCACGCTTTATGTTGCGTT TCTC |
| <i>CjapCSP12</i> | aaggatccGAGGAAGAGAAATACG AGGATAA | aactcgagTTATTCTTAGCTTTCTT CATATCTTCTAAA |
| <i>CjapCSP13</i> | aaggatccGAAGAACTTATAGCG ATCAGTACG | aactcgagTTATTGACCAGCATTCTTCT TTTCATATCTTC |

Supplemental Table S2. Nucleotide sequences of primers used for RT-PCR.

| Gene | Forward primer (5'-3') | Reverse primer (5'-3') |
|---|-----------------------------|-----------------------------|
| Elongation factor 1 (<i>efl</i>) | CCCTTCGTCTTCCCCTCA | ACCATACCGGGTTTCAGCA C |
| 60S ribosomal protein L32 (<i>rpl32</i>) | CCAACAGGCTTCCGTAAAG TTC | TCTCTGCACAAAACCTGCAG TT |
| 60S ribosomal protein L18 (<i>rpl18</i>) | AAGAACCTGGACGGGAG AA | CGAGCTTTCGGTTACACG |
| Glyceraldehyde-3-phos- phate dehydrogenase (<i>gapdh</i>) | CATGACGACTGTACATGCG ATT | GCAGCGGAATAATGTTT G |
| <i>CjapCSP2</i> | ATTCGCGACGAAATGTCGA A | CTGGGGACGATTCTCGGTGT |
| <i>CjapCSP2</i> | GCGACAAGCGTTACTTGAC C | CAGGAGCCTCTCAAAACCA A |
| <i>CjapCSP3</i> | CAAATGCCCTCTGAAGCT C | CGCCTTCTCGCAATTCTT |
| <i>CjapCSP4</i> | AGATGCGGCAGAACTCAA | CGCTGTTCGCAATTAAACG A |
| <i>CjapCSP5</i> | CAGGGGATTGCGATGGATT T | CGGGGTACAACGACGACAT TT |
| <i>CjapCSP6</i> | GGTGTGTTGGACGAAGG A | TTCTGCCGTAGCCTTTGTT T |
| <i>CjapCSP7</i> | TGCCTGCTGGATAAGGGAA A | GCATTGCTGCACTCGGTCT |
| <i>CjapCSP8</i> | CCGATGCACGCAGTCTAA A | TGCCTATTTCCTTGCTCCTT G |
| <i>CjapCSP9</i> | TACGGGATCTGCACGCTTC T | TCGGTAATCGCATTGAGAA TGT |
| <i>CjapCSP10</i> | GTGCAAGACAGCTGATCAA AGA | ACCAATCGAGTACGATTCC AA |
| <i>CjapCSP12</i> | GTTCATGGCAACAGGACC A | TGCACTTAGTTGAAAAGCC TCAC |
| <i>CjapCSP13</i> | CCATGTCTAACAGCAGAAG CAAA | TCAACGATCTTCTGCCATT GT |

Supplemental Table S3. Nucleotide sequences of primers used for *in situ* hybridization

Lower-case letters indicate additional sequences for restriction enzyme recognition.

Note that the same primers listed in gene cloning (Supplemental Table S1) were used for *CjapCSP12* and *CjapCSP13*.

| Gene | Forward primer (5'-3') | Reverse primer (5'-3') |
|------------------|--------------------------------------|--|
| <i>CjapCSP1</i> | GGACATGTTCGATCACATCAA | CCTCGAAAACTACACATGATGC |
| <i>CjapCSP12</i> | aaggatccGAGGAAGAGAAATAC GAGGATAA | aactcgagTTATTCTTAGCTTTTC TTCATATCTTCTAAA |
| <i>CjapCSP13</i> | aaggatccGAAGAACTTATAGC GATCAGTACG | aactcgagTTATTGACCAGCATTCTT CTTTTCATATCTTC |

Supplemental Dataset

Supplemental dataset S1. Nucleotide sequences of twelve annotated CjapCSPs

Supplemental dataset S2. Amino acid sequences of twelve annotated CjapCSPs