

Supplementary Materials

Table S1. Primer list

| Name | Sequence (5'-3') | Target gene | pRW412a region* |
|-----------|------------------------|--|-----------------|
| REV | CAGGAAACAGCTATGAC | Sequencing primer | - |
| UNIV | GTTTCCCAGTCACGAC | Sequencing primer | - |
| GM3F | AGAGTTGATCATGGC | 16S rRNA gene | - |
| GM4R | TACCTTGTACGACTT | 16S rRNA gene | - |
| IpfAfor | GTCTTCGTTGGTCACGACTC | <i>ipfA</i> | III |
| IpfBrev | TTCACTCGCGTGACCATG | <i>ipfB</i> | III |
| IpfDfor | ACCATGGTCACCGGAGTGA | <i>ipfD</i> | III |
| IpfDrev | GGTGCTATCCATCAATGGCTCC | <i>ipfD</i> | III |
| IpfFfor | GTTATGTTGGCAAGAGACCTGG | <i>ipfF</i> | III |
| IpfFrev4 | CTTTCTTGCCTCAGGATGGC | <i>ipfF</i> | III |
| HMSDFor- | TCCGGCATTTCATCGATG | <i>ipfM</i> | I |
| HMSDRev | ATGATGGCGCTGTTCTTGC | <i>ipfM</i> | I |
| IpfIfor- | GAGCGTTTGAAGTCTGGGC | <i>ipfI</i> | II |
| IpfIrev | CAATCACGGTGCCTGTGAAG | <i>ipfI</i> | II |
| DDFor | GACGGCGGAAGTTTGAA | Dienoate dehydrogenase gene (DD) | IV |
| DDRev | CTTGACGACGACGGTGT | Dienoate dehydrogenase gene (DD) | IV |
| DsbC2For- | CACAATGCATCCGCTACG | Thiol:disulfide gene involved in conjugative transfer (DsbC) | V+I |
| DsbC2Rev | CGGATCCTTCGCACAATAG | Thiol:disulfide gene involved in conjugative transfer (DsbC) | V+I |

* Roman numerals denote regions delimited by IS6 elements as indicated for pIBU218 of *R. wittichii* MPO218 (Aulestia et al., 2021).

Table S2. PCR amplification

| Target gene | pRW412a region* | <i>R. wittichii</i> RW1 | <i>S. granuli</i> RW412 | <i>R. wittichii</i> RW421 | <i>P. citronellolis</i> RW422 | <i>P. citronellolis</i> RW423 | <i>P. citronellolis</i> RW424 |
|--------------|-----------------|-------------------------|-------------------------|---------------------------|-------------------------------|-------------------------------|-------------------------------|
| <i>ipfAB</i> | III | ND | + | + | + | + | - |
| <i>ipfD</i> | III | ND | + | + | + | + | - |
| <i>ipfF</i> | III | ND | + | + | + | + | - |
| <i>ipfM</i> | I | - | + | - | + | + | - |
| <i>ipfI</i> | II | - | + | + | + | + | - |
| DD | IV | - | + | + | + | + | - |
| DsbC | V+I | - | + | - | + | + | - |

* Roman numerals in analogy to that used for pIBU218 of *R. wittichii* MPO218 (Aulestia et al., 2021).

ND: not determined

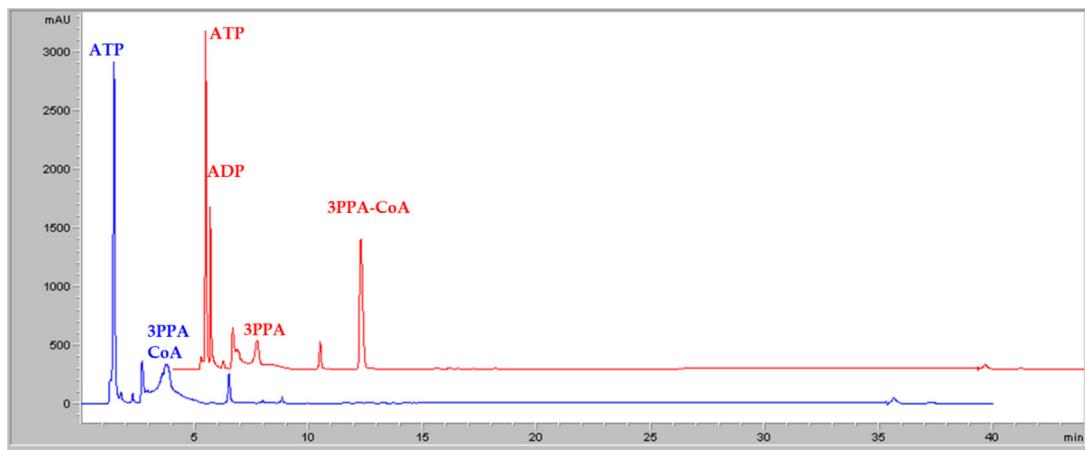


Figure S1. Overlay of HPLC chromatograms at 210 nm of the reaction components of 3-phenylpropanoic acid (3PPA) with IpFF initially (in blue) and after 35 minutes (in red).

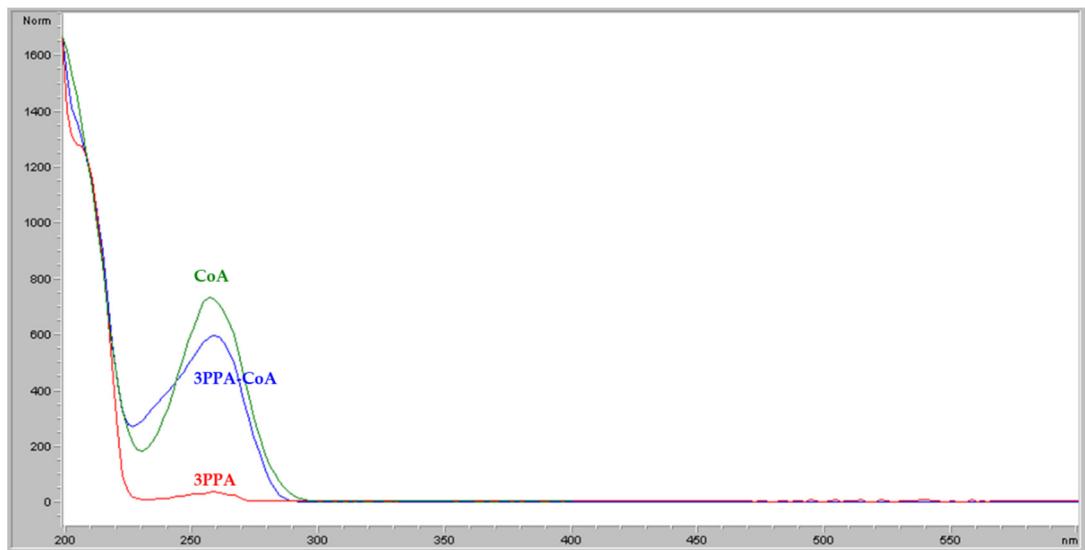


Figure S2. Overlay of spectra (200-600 nm) of the 3-phenylpropanoyl-CoA product (3PPA-CoA, in blue) compared to those of 3-phenylpropanoic acid (3PPA, in red) and coenzyme A (CoA, in green).

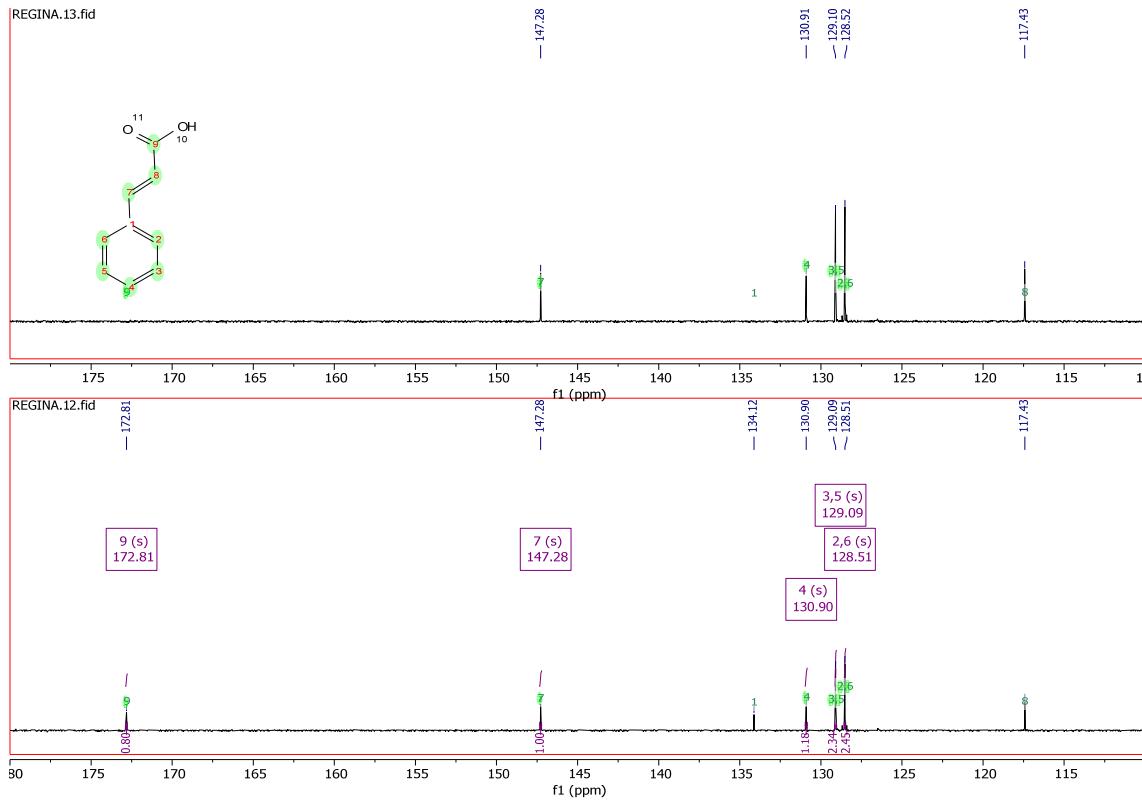


Figure S3. ^{13}C NMR (BB and DEPT135) spectrum of cinnamic acid, recorded at 151 MHz in deuteriochloroform.

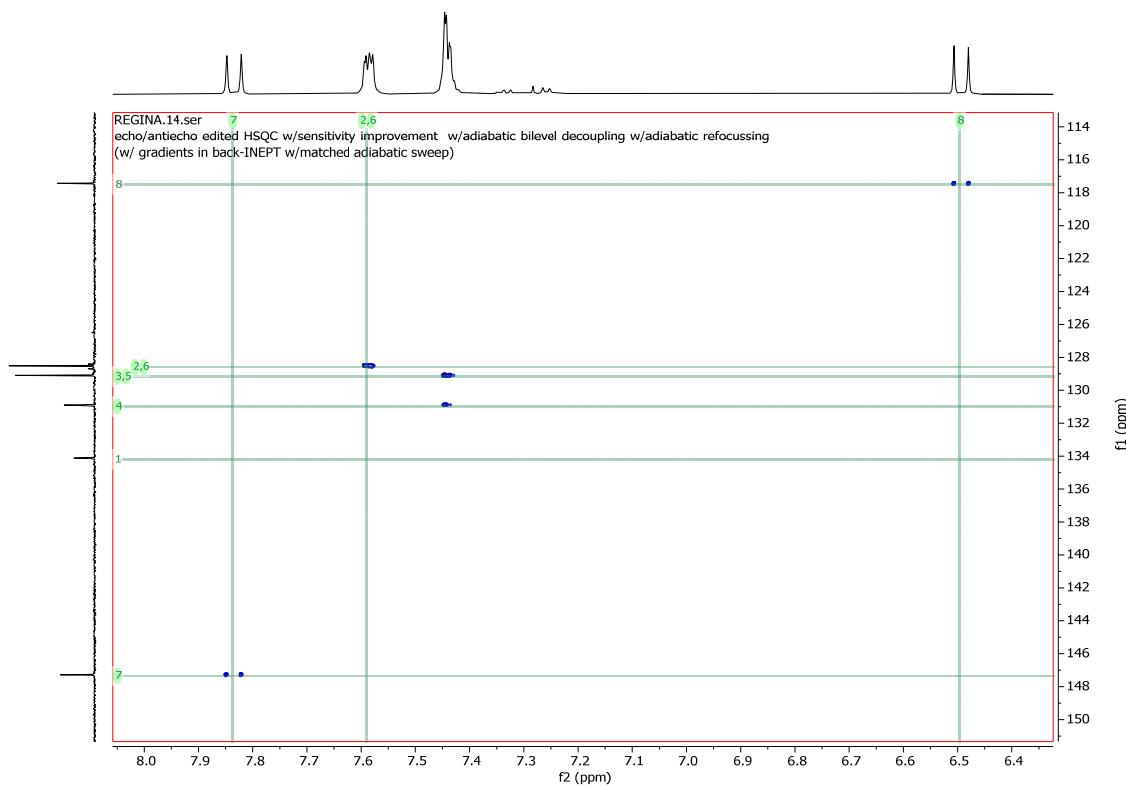


Figure S4. ^1H - ^{13}C HSQC 2D-NMR spectra of cinnamic acid.

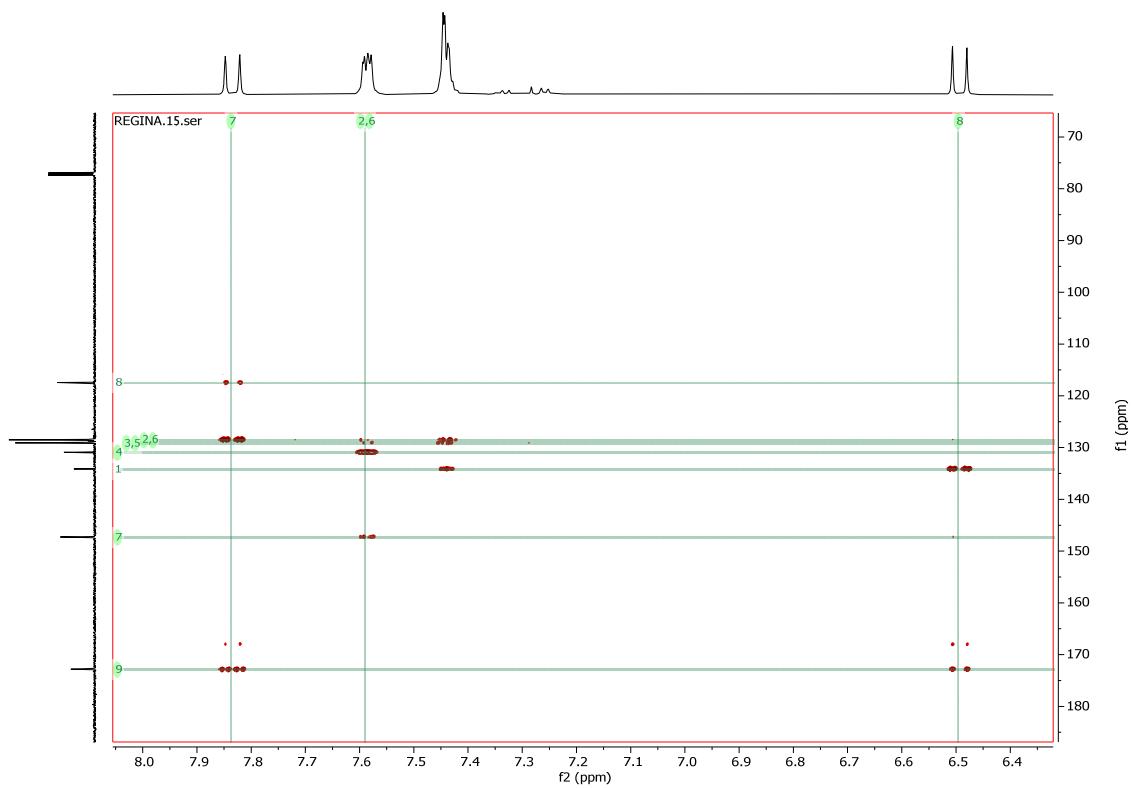


Figure S5. ^1H - ^{13}C HMBC 2D-NMR spectra of cinnamic acid to assign the quaternary carbons.