## SUPPLEMENTARY INFORMATION

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# Supplementary table 1. Strains and plasmids

| Plasmid or      | Relevant genotype or feature   | Reference  |
|-----------------|--|------------|
| Strain          |  |            |
| Plasmid         |  |            |
| pKAS32          | Suicide vector for allelic exchange in <i>V. cholerae</i> , Amp <sup>R</sup>             | [1]        |
| pLAFR2          | Broad-host-range cosmid; <i>mob</i> , Tet <sup>R</sup>                                   | [2]        |
| pBB1            | <i>V. harveyi luxCDABE</i> on pLAFR2, Tet <sup>R</sup>                                   | [3]        |
| pDH345          | <i>V. cholerae cqsS</i> on pSLS4, Kan <sup>R</sup>                                       | [4]        |
| pWN1365         | <i>V. cholerae cqsS</i> <sup>C170F</sup> on pDH345, Kan <sup>R</sup>                     | This study |
| pWN1960         | <i>V. cholerae cqsS</i> <sup>C170F</sup> on pKAS32, Amp <sup>R</sup>                     | This study |
| pJMH280         | <i>V. harveyi cqsS</i> on pGEM-T, Amp <sup>R</sup>                                       | [5]        |
| pWN1513         | <i>V. harveyi cqsS</i> <sup>F175C</sup> on pJMH280, Amp <sup>R</sup>                     | This study |
| pJMH282         | <i>V. harveyi cqsS</i> on pLAFR2, Tet <sup>R</sup>                                       | [5]        |
| pWN1515         | <i>V. harveyi cqsS</i> <sup>F175C</sup> on pLAFR2, Tet <sup>R</sup>                      | This study |
| pWN1327         | <i>V. harveyi cqsA</i> on pET-28B, untagged, Kan <sup>R</sup>                            | This study |
| WN1666          | <i>V. harveyi cqsA</i> on pET-28B, N-terminal His <sub>6</sub> -tagged, Kan <sup>R</sup> | This study |
| Vibrio cholerae |  |            |
| C6706str        | Wild type  | [6]        |
| BH1523          | $\Delta cqsA$  | This study |
| DH197           | $\Delta cqsA, \Delta luxQ$   | This study |
| WN1102          | $\Delta cqsA, \Delta luxQ, pBB1$   | This study |

| WN1981         | $\Delta cqsA, \Delta luxQ, cqsS^{C170Y}$                      | This study |
|----------------|---|------------|
|                |   | 2          |
|                | C170V   |            |
| WN1993         | $\Delta cqsA$ , $\Delta luxQ$ , $cqsS^{C1/01}$ /pBB1          | This study |
|                |   |            |
| Vibrio harvevi |   |            |
|                |   |            |
|                |   |            |
| BB120          | Wild type   | [7]        |
|                |   |            |
| DALL602        | Apart   | [5]        |
| JMIN003        | ΔcqsA   | [5]        |
|                |   |            |
| JMH626         | $\Delta cqsA, \Delta luxN, \Delta luxQ$                       | [5]        |
|                |   |            |
| WN1207         | A age A age S Alw N Alw BO                                    | This study |
| WIN1397        | $\Delta cqsA, \Delta cqsS, \Delta uuxN, \Delta uuxPQ$         | This study |
|                |   |            |
| WN1492         | $\Delta cqsA, \Delta cqsS, \Delta luxN, \Delta luxPQ/pJMH282$ | This study |
|                |   |            |
| 11024          |   |            |
| WN1834         | $\Delta cqsA, \Delta cqsS, \Delta luxN, \Delta luxPQ/pWN1515$ | This study |
|                |   |            |
| 1              |   |            |

#### 2. HPLC and SFC conditions

For NP-HPLC, four liters of the cell-free supernant from an M9 culture *E. coli* overexpressing *V. harveyi* CqsA (see materials and methods for details) was extracted into MTBE (4 x 500 mL), the combined organic extract was washed with a saturated brine solution (1 L) and was dried over Na<sub>2</sub>SO<sub>4</sub>. The dried organics were carefully concentrated *in vacuo* at room temperature to yield a concentrated organic extract in ~2-5 mL of organic solvent. Complete concentration of this sample leads to significant loss of biological activity. This concentrated organic sample was diluted with hexanes (5 mL) and was purified by preperative HPLC on a system composed 2 PrepStar SD-1 pumps, a Knauer K-2501 multi-wavelength detector set at 280nm, a Rainin FC-1 fraction collector and using a PrincetonSPHER Premier (2 x 25 cm) column. The mixture was fractionated by application of a gradient from 9:1 hexane:MTBE to 100% MTBE at 20 mL/min over 25 min, with fractions collected every minute. Ea-C8-CAI-1 (by HRMS and bioassay) has a retention time of 8 minutes, C8-CAI-1 and the  $\alpha$ -hydroxy ketone regioisomer (by HRMS, NMR and bioassay) elute as two smaller peaks between 9 and 10 minutes.



Due to the observed instability of purified or unpurified Ea-C8-CAI-1, we turned to an alternate strategy for purification. Accordingly, we developed a SFC (supercritical fluid chromatography) method using liquid  $CO_2$  along with a small amount of per-deuterated methanol (d4-MeOH) as a co-solvent. Importantly, this approach directly provides concentrated fractions in d4-MeOH, suitable for direct NMR analysis.

For the purification, four liters of the cell-free supernant from an M9 culture *E. coli* overexpressing *V. harveyi* CqsA (see materials and methods for details) was extracted into MTBE (4 x 500 mL), the combined organic extract was washed with a saturated brine solution (1 L) and was dried over Na<sub>2</sub>SO<sub>4</sub>. The dried organics were carefully concentrated *in vacuo* at room temperature to yield a concentrated organic extract in ~2-5 mL of organic solvent. Purification was achieved using a Berger Multigram II SFC system equipped with 2 Varian SD-1 pumps, a Knauer K-2501 multi-wavlength dectector set at 280nm, a Knauer K-1900 pump, a Vatran SGP-50-100 condenser, using a PrincetonSPHER Premier (2 x 25 cm) column. Fractionation of the crude mixture was achieved by application of an isocratic method using a mixture of 8% d4-MeOH/CO<sub>2</sub> (100 bar) at 50 mL/min. Fractions were collected manually and the product was found to have a retention time of 3.5 min. This fraction was subjected to direct <sup>1</sup>H-NMR analysis, and found to contain a mixture of Ea-C8-CAI-1 along with C8-CAI-1 and its  $\alpha$ -hydroxy ketone regioisomer.



## **3. HRMS** analyses (Corrective factor determination and cell-free fluids analyses)

| Molecule              | 1        | ion abundance | 9        | n        | nultiplicative | factor of mole | cular ion to d | <sub>2</sub> -CAI1 |
|-----------------------|----------|---------------|----------|----------|----------------|----------------|----------------|--------------------|
|                       | Sample 1 | Sample 2      | Sample 3 | Sample 1 | Sample 2       | Sample 3       | Average        | Standard           |
|                       |          |               |          |          |                |                |                | Deviation          |
| Ea-C8-CAI-1           | 49.4     | 183.7         | 263      | 1.30     | 0.81           | 0.74           | 0.95           | 0.31               |
| C8-CAI-1              | 57.6     | 174.1         | 192.2    | 1.12     | 0.85           | 1.02           | 1.00           | 0.14               |
| Ea-CAI-1              | 33       | 73.5          | 108.9    | 1.95     | 2.01           | 1.80           | 1.92           | 0.11               |
| d <sub>2</sub> -CAI-1 | 64.4     | 148.1         | 195.6    | 1.00     | 1.00           | 1.00           | 1.00           | 0.00               |

A. HRMS data for corrective factor determination:

## B. HRMS data for the *V. harveyi* cell-free fluids:

| Molecule              | Ion Abu  | indance from | HRMS     | Calculate | d Concentra | tion (nM) | Average | Standard  |
|-----------------------|----------|--------------|----------|-----------|-------------|-----------|---------|-----------|
| Wolecule              | Sample 1 | Sample 2     | Sample 3 | Sample 1  | Sample 2    | Sample 3  | (nM)    | Deviation |
| Ea-C8-CAI-1           | 203.5    | 329.6        | 135.7    | 62.3      | 49.5        | 36.6      | 49.5    | 12.8      |
| C8-CAI-1              | 62       | 104.6        | 34.2     | 198.6     | 165.0       | 97.0      | 153.5   | 51.8      |
| Ea-CAI-1              | 0        | 0            | 0        | 0.0       | 0.0         | 0.0       | 0.0     | 0.0       |
| CAI-1                 | 0        | 0            | 0        | 0.0       | 0.0         | 0.0       | 0.0     | 0.0       |
| d <sub>2</sub> -CAI-1 | 155.4    | 316.9        | 176.3    | 500.0     | 500.0       | 500.0     | 500.0   | 0.0       |

C. HRMS data for the V. cholerae cell-free fluids:

| Molecule              | Ion Abu  | indance from | HRMS     | Calculate | d Concentra | tion (nM) | Average | Standard  |
|-----------------------|----------|--------------|----------|-----------|-------------|-----------|---------|-----------|
| Wolceute              | Sample 1 | Sample 2     | Sample 3 | Sample 1  | Sample 2    | Sample 3  | (nM)    | Deviation |
| Ea-C8-CAI-1           | 141.3    | 83.9         | 92.1     | 23.1      | 15.0        | 19.8      | 19.3    | 4.1       |
| C8-CAI-1              | 0        | 0            | 0        | 0.0       | 0.0         | 0.0       | 0.0     | 0.0       |
| Ea-CAI-1              | 348.7    | 321.1        | 320.5    | 115.2     | 116.0       | 139.0     | 123.4   | 13.5      |
| CAI-1                 | 132.3    | 111.6        | 98.2     | 227.6     | 209.9       | 221.8     | 219.8   | 9.0       |
| d <sub>2</sub> -CAI-1 | 290.7    | 265.8        | 221.4    | 500.0     | 500.0       | 500.0     | 500.0   | 0.0       |

The detection limits for the above molecules were determined to be: Ea-C8-CAI-1, 6.25 nM; C8-CAI-1, 25 nM; Ea-CAI-1, 12.5 nM; CAI-1, 25 nM. This was determined by the analysis of a series of samples containing pure samples of each of the listed molecules at concentrations of 50 nM, 25 nM, 12.5 nM, 6.25 nM, 3.125 nM, and 1.56 nM. Detection limits are described as the last sample for which a HRMS peak was observed.

#### 4. Ea-C8-CAI-1 synthesis by V. harveyi CqsA in vitro (A) and in vivo (B)

(A) The substrate required for *V. harveyi* CqsA to produce Ea-C8-CAI-1 was tested in buffer (10 mM HEPES, 0.1 M NaCl) in the presence or absence of C8-CoA (100 μM), SAM (1 mM), and *V. harveyi* CqsA (500 nM). The reaction mixture was incubated at room temperature for 1 hr and terminated by addition of an equal volume of acetonitrile. The reaction mixture was assayed for stimulation of bioluminescence expression in the *V. harveyi* CAI-1 reporter strain JMH626. Only reactions containing CqsA, SAM, and C8 CoA resulted in induction of bioluminescence. HRMS analysis of the reaction products revealed the molecular ion of Ea-C8-CAI-1. RLU denotes relative light units.



(B) Heavy atom labeling was used to examine whether SAM could be used *in vivo* by CqsA as a substrate. *E. coli* overexpressing *V. harveyi* CqsA (WN1327, top) produced Ea-C8-CAI-1 (184.169 ion) and C8-CAI-1 (187.169 ion). A *metE E. coli* mutant (WN1590, middle) grown in the presence of unlabeled methionine carrying the CqsA overexpression plasmid produced the identical two molecules. Consistent with the idea that SAM is used *in vivo* for Ea-C8-CAI-1 synthesis and C8-CAI-1 is derived from Ea-C8-CAI-1, WN1590 grown in the presence of d8-L-methionine (bottom) produced triple-deuterium labeled Ea-C8-CAI-1 (187.189 ion) and C8-CAI-1 (190.188 ion) (See [8] for detailed explanations).



#### 5. Chemical Syntheses.

**Analytical methods.** NMR spectra were recorded using a Bruker Avance II spectrometer (500 MHz for <sup>1</sup>H; 125 MHz for <sup>13</sup>C) equipped with either a <sup>1</sup>H-optimized TCI (H/C/N) cryoprobe or a <sup>13</sup>C-optimized dual C/H cryoprobe. Chemical shifts are reported in parts per million (ppm) and were calibrated according to residual solvent. High-resolution mass spectral analyses were performed using an Agilent 1200-series electrospray ionization–time-of-flight (ESI-TOF) mass spectrometer in the positive ESI mode. Note, the NMR data for synthetic and natural Ea-CAI-1 also appear in the supplement to Wei et al. [8] as evidence for production of Ea-CAI-1 and conversion to CAI-1 in *V. cholerae*.

**Chemical Reactions.** Unless otherwise noted, all reactions were performed in flame-dried glassware under an atmosphere of nitrogen. All chemicals purchased from commercial vendors were used without further purification. Anhydrous Sure/Seal<sup>TM</sup> solvents were purchased from commercial vendors.

**Purification.** Flash chromatography was performed using standard grade silica gel 60 230-400 mesh from SORBENT Technologies. Analytical thin-layer chromatography was carried out using Silica G TLC plates, 200  $\mu$ m with UV<sub>254</sub> fluorescent indicator (SORBENT Technologies), and visualization was performed by staining (anisaldehyde, ceric ammonium molybdate, or ninhydrin) and/or by absorbance of UV light.

All synthetic molecules were determined to be pure by NMR and HRMS, Ea-CAI-1 and C8-Ea-CAI-1 were not purified and were characterized directly from the final deprotection step. NMR data of these products containing peaks consistent with the removed protecting group (TIPS-F) and internal standard (PhMe) are provided.



(Z)-*tert*-butyl 1-(methoxy(methyl)amino)-1-oxobut-2-en-2-ylcarbamate (S1). L-Vinylglycine hydrochloride [9,10](2.14 g, 15.6 mmol) was dissolved in dioxane/H<sub>2</sub>O (182 mL, 0.085 M, 1:1 dioxane:H<sub>2</sub>O) and was treated with NaHCO<sub>3</sub> (2.62 g, 31.1 mmol, 2.0 eq) and Boc<sub>2</sub>O (3.56 g,

16.3 mmol, 1.05 eq) sequentially at room temperature. The resulting mixture was heated to reflux for 2.5 hr, was cooled and concentrated *in vacuo* to a volume of ca. 80 mL. The resulting solution was acidified with 1 N HCl to ~pH 6.5, was extracted with DCM (3 x 50 mL), dried over Na<sub>2</sub>SO<sub>4</sub> and concentrated to dryness. The resulting oil (2.82 g) was dissolved in THF (156 mL, 0.1 M to VGly-HCl) at ambient temperature and was treated with HOBt (6.3 g, 46.7 mmol, 3.0 eq to VGly-HCl), HNMe(OMe)-HCl (1.75 g, 17.9 mmol, 1.15 eq to VGly-HCl), EDC (3.43 g, 17.9 mmol, 1.15 eq to VGly-HCl), and Et<sub>3</sub>N (10.8 mL, 77.8 mmol, 5.0 eq to VGly-HCl) sequentially and the mixture was allowed to stir for 8 hr at ambient temperature. The crude mixture was filtered through a plug of celite and concentrated in vacuo. The resulting oil was purified by silica gel chromatography eluting with a gradient from hexanes to 60% EtOAc/hexanes. Fractions containing the desired product were combined and concentrated in vacuo to provide S1 as a pale yellow oil (2.2 g, 58% from VGly-HCl). <sup>1</sup>H-NMR (500 MHz, CDCl<sub>3</sub>)  $\delta$  6.22 (bs, 1H), 5.67 (bs, 1H), 3.64 (s, 3H), 3.22 (s, 3H), 1.68 (d, J = 7.0 Hz, 3H), 1.41 (s, 9H); <sup>13</sup>C-NMR (125 MHz, CDCl<sub>3</sub>) δ 167.4, 152.9, 129.5, 120.9, 80.5, 61.1, 34.4, 28.3, 12.6; HRMS (ESI-TOF) calcd for  $C_{11}H_{20}N_2O_4Na$ , 267.1321 m/z [M+Na]; observed, 267.1323 m/z[M+Na]<sup>+</sup>. The configuration of this molecule was assigned on the basis of its <sup>1</sup>H-NMR spectra [10].



(*Z*)-*tert*-butyl 4-oxotridec-2-en-3-ylcarbamate (S2). To a solution of (*Z*)-*tert*-butyl 1-(methoxy(methyl)amino)-1-oxobut-2-en-2-ylcarbamate (S1) (172.5 mg, 0.71 mmol) in THF (7 mL, 0.1 M) at 0°C was added nonyl-MgBr (3.5 mL, 3.5 mmol, 5.0 eq, 1.0 M in diethyl ether) and the mixture was stirred at 0°C for 2 h. The resulting mixture was quenched with sat. NH<sub>4</sub>Cl (20 mL), extracted with Et<sub>2</sub>O (3 x 20 mL), combined organics were washed with sat. NaCl (20 mL), dried over Na<sub>2</sub>SO<sub>4</sub> and concentrated *in vacuo*. The resulting oil was purified by silica gel chromatography eluting with a gradient from hexanes to 20% EtOAc/hexanes to provide S2 as a clear colorless oil (166.3 mg, 76%). <sup>1</sup>H-NMR (500 MHz, CDCl<sub>3</sub>)  $\delta$  6.50 (q, *J* = 7.1 Hz, 1H), 6.42 (s, 1H), 2.64 (t, *J* = 7.5 Hz, 2H), 1.85 (d, *J* = 7.1 Hz, 3H), 1.65-1.55 (m, 2H), 1.44 (s, 9H), 1.32-

1.16 (m, 12H), 0.86 (t, J = 6.9 Hz, 3H); <sup>13</sup>C-NMR (125 MHz, CDCl<sub>3</sub>)  $\delta$  198.0, 153.1, 135.3, 130.4, 80.5, 36.9, 32.1, 29.7, 29.6, 29.5, 29.5, 28.4, 24.9, 22.9, 15.3, 14.3; HRMS (ESI-TOF) calcd for C<sub>18</sub>H<sub>33</sub>NO<sub>3</sub>Na 334.2358 *m*/*z* [M+Na]; observed, 334.2351 *m*/*z* [M+Na]<sup>+</sup>.



(*Z*)-*tert*-butyl 4-oxoundec-2-en-3-ylcarbamate (S3). Prepared in an analogous manner to S2 from (*Z*)-*tert*-butyl 1-(methoxy(methyl)amino)-1-oxobut-2-en-2-ylcarbamate (S1) and heptyl-MgBr in 71% yield. <sup>1</sup>H-NMR (500 MHz, CDCl<sub>3</sub>)  $\delta$  6.50 (q, *J* = 7.1 Hz, 1H), 6.42 (s, 1H), 2.64 (t, *J* = 7.5 Hz, 2H), 1.84 (d, *J* = 7.1 Hz, 3H), 1.64-1.51 (m, 2H), 1.44 (s, 9H), 1.33-1.18 (m, 8H), 0.85 (t, *J* = 6.9 Hz); <sup>13</sup>C-NMR (125 MHz, CDCl<sub>3</sub>)  $\delta$  198.0, 153.1, 135.3, 130.4, 80.5, 36.9, 31.9, 29.5, 29.3, 28.4, 24.9, 22.8, 15.3, 14.3; HRMS (ESI-TOF) calcd for C<sub>16</sub>H<sub>29</sub>NO<sub>3</sub>Na 306.2045 *m/z* [M+Na]; observed, 306.2041 *m/z* [M+Na]<sup>+</sup>.



(Z)-3-aminotridec-2-en-4-one (Ea-CAI-1). To a solution of (Z)-tert-butyl 4-oxotridec-2-en-3ylcarbamate (S2) (54.9 mg, 0.18 mmol) in CH<sub>2</sub>Cl<sub>2</sub> (1.2 mL, 0.15 M) at 0°C was added Et<sub>3</sub>N (74  $\mu$ L, 0.53 mmol, 3.0 eq) followed by TIPSOTf (142  $\mu$ L, 0.53 mmol, 3.0 eq) and the mixture was stirred at 0°C for 2 h. To this mixture was added additional Et<sub>3</sub>N (74  $\mu$ L, 0.53 mmol, 3.0 eq) and TIPSOTf (142  $\mu$ L, 0.53 mmol, 3.0 eq) and the mixture was allowed to slowly warm to ambient temperature and was stirred for an additional 18 h. The prolonged reaction time, temperature profile, and the portionwise addition of Et<sub>3</sub>N and TIPSOTf were found to be important for optimal conversion. At this time, TLC analysis showed complete consumption of the starting material (Rf=0.2, 10% EtOAc/hexanes, ninhydrin) to a single new product spot (Rf=0.4, 10% EtOAc/hexanes, ninhydrin) and the reaction was quenched with sat. NH<sub>4</sub>Cl (10 mL), extracted with CH<sub>2</sub>Cl<sub>2</sub> (3 x 10 mL), the combined organics were washed with sat. NaCl, dried over Na<sub>2</sub>SO<sub>4</sub> and concentrated *in vacuo*. The resulting oil was flashed through a plug of silica gel, eluting with 10% EtOAc/hexanes, was concentrated to dryness and used without further purification. The resulting oil was dissolved in d6-DMSO (2.1 mL with 3  $\mu$ L PhMe as an internal standard), was treated with TASF (145.4 mg, 0.53 mmol, 3 eq to **S2**) and was vortexed for 5 min before NMR analysis. <sup>1</sup>H-NMR analysis enabled determination of the concentration of Ea-CAI-1 by integration of the Ea-CAI-1 vinyl methyl doublet ( $\delta$  1.65) to the PhMe methyl singlet ( $\delta$  2.29). The concentration of Ea-CAI-1 determined in this manner was 13.6 $\mu$ M (0.029 mmol, 16% from **S2**). <sup>1</sup>H-NMR (500 MHz, d6-DMSO)  $\delta$  5.53 (q, *J* = 7.1 Hz, 1H), 4.31 (s, 2H), 2.60 (t, *J* = 7.3 Hz, 2H), 1.65 (d, *J* = 7.0 Hz, 3H), 1.53-1.41 (m, 2H), 1.30-1.17 (m, 12H), 0.85 (t, *J* = 6.4 Hz, 3H); <sup>13</sup>C-NMR (125 MHz, d6-DMSO)  $\delta$  197.1, 141.8, 106.9, 35.2, 31.4, 29.0, 29.0, 28.8, 25.1, 22.2, 14.0, 12.2; HRMS (ESI-TOF) calcd for C<sub>13</sub>H<sub>26</sub>NO 212.2014 *m*/*z* [M+H]; observed, 212.2010 *m*/*z* [M+H]<sup>+</sup>.



(*Z*)-3-aminoundec-2-en-4-one (C8-Ea-CAI-1). Prepared in an analogous manner to Ea-CAI-1 from (*Z*)-*tert*-butyl 4-oxoundec-2-en-3-ylcarbamate (S3) in 13% overall yield. <sup>1</sup>H-NMR (500 MHz, d6-DMSO/PhMe internal standard)  $\delta$  5.51 (q, *J* = 7.1 Hz, 1H), 4.30 (s, 2H), 2.59 (t, *J* = 7.4 Hz, 2H), 1.65 (d, *J* = 7.1 Hz, 3H), 1.51-1.43 (m, 2H), 1.25-1.19 (m, 8H), 0.84 (t, *J* = 6.8 Hz, 3H); HRMS (ESI-TOF) calcd for C<sub>11</sub>H<sub>22</sub>NO 184.1701 *m*/*z* [M+H]; observed, 184.1699 *m*/*z* [M+H]<sup>+</sup>.

Refer to [4,11,12] for syntheses of CAI-1, Am-CAI-1, C8-CAI-1, and Am-C8-CAI-1.



## 6. NMR analyses of natural and synthetic CAI-1 type compounds.













| Synthetic Ea-CAI-1 <sup>1</sup> H-NMR   | Natural Ea-CAI-1 <sup>1</sup> H-NMR   |
|---|---|
| 5.53 (q, <i>J</i> = 7.1 Hz, 1H)   | 5.53 (q, <i>J</i> = 7.1 Hz, 1H)   |
| 4.31 (s, 2H)  | 4.28 (s, 2H)  |
| 2.60 (t, $J = 7.3$ Hz, 2H)  | 2.60 (t, <i>J</i> = 7.4 Hz, 2H)   |
| 1.65 (d, <i>J</i> = 7.0 Hz, 3H)   | 1.65 (d, <i>J</i> = 7.1 Hz, 3H)   |
| 1.53-1.41 (m, 2H)   | 1.52-1.41 (m, 2H)   |
| 1.30-1.17 (m, 12H)  | 1.31-1.13 (m, 12H)  |
| 0.85 (t, J = 6.4 Hz, 3H)  | 0.85 (t, <i>J</i> = 6.9 Hz, 3H)   |
|   |   |
| Synthetic Ea-CAI-1 <sup>13</sup> C-NMR  | Natural Ea-CAI-1 <sup>13</sup> C-NMR  |
| Synthetic Ea-CAI-1 <sup>13</sup> C-NMR<br>197.1   | Natural Ea-CAI-1 <sup>13</sup> C-NMR<br>197.2   |
| Synthetic Ea-CAI-1 <sup>13</sup> C-NMR<br>197.1<br>141.8  | Natural Ea-CAI-1 <sup>13</sup> C-NMR<br>197.2<br>141.8  |
| Synthetic Ea-CAI-1 <sup>13</sup> C-NMR<br>197.1<br>141.8<br>106.9   | Natural Ea-CAI-1 <sup>13</sup> C-NMR<br>197.2<br>141.8<br>106.9   |
| Synthetic Ea-CAI-1 <sup>13</sup> C-NMR<br>197.1<br>141.8<br>106.9<br>35.2   | Natural Ea-CAI-1 <sup>13</sup> C-NMR<br>197.2<br>141.8<br>106.9<br>35.2   |
| Synthetic Ea-CAI-1 <sup>13</sup> C-NMR<br>197.1<br>141.8<br>106.9<br>35.2<br>31.4   | Natural Ea-CAI-1 <sup>13</sup> C-NMR<br>197.2<br>141.8<br>106.9<br>35.2<br>31.4                                 |
| Synthetic Ea-CAI-1 <sup>13</sup> C-NMR<br>197.1<br>141.8<br>106.9<br>35.2<br>31.4<br>29.0   | Natural Ea-CAI-1 <sup>13</sup> C-NMR<br>197.2<br>141.8<br>106.9<br>35.2<br>31.4<br>29.0                         |
| Synthetic Ea-CAI-1 <sup>13</sup> C-NMR<br>197.1<br>141.8<br>106.9<br>35.2<br>31.4<br>29.0<br>29.0                                 | Natural Ea-CAI-1 <sup>13</sup> C-NMR<br>197.2<br>141.8<br>106.9<br>35.2<br>31.4<br>29.0<br>29.0                 |
| Synthetic Ea-CAI-1 <sup>13</sup> C-NMR<br>197.1<br>141.8<br>106.9<br>35.2<br>31.4<br>29.0<br>29.0<br>28.8                         | Natural Ea-CAI-1 <sup>13</sup> C-NMR<br>197.2<br>141.8<br>106.9<br>35.2<br>31.4<br>29.0<br>29.0<br>28.8         |
| Synthetic Ea-CAI-1 <sup>13</sup> C-NMR<br>197.1<br>141.8<br>106.9<br>35.2<br>31.4<br>29.0<br>29.0<br>28.8<br>25.1                 | Natural Ea-CAI-1 <sup>13</sup> C-NMR<br>197.2<br>141.8<br>106.9<br>35.2<br>31.4<br>29.0<br>29.0<br>28.8<br>25.1 |
| Synthetic Ea-CAI-1 <sup>13</sup> C-NMR<br>197.1<br>141.8<br>106.9<br>35.2<br>31.4<br>29.0<br>29.0<br>28.8<br>25.1<br>22.2         | Natural Ea-CAI-1 <sup>13</sup> C-NMR   197.2   141.8   106.9   35.2   31.4   29.0   29.0   28.8   25.1   22.2   |
| Synthetic Ea-CAI-1 <sup>13</sup> C-NMR<br>197.1<br>141.8<br>106.9<br>35.2<br>31.4<br>29.0<br>29.0<br>28.8<br>25.1<br>22.2<br>14.0 | Natural Ea-CAI-1 <sup>13</sup> C-NMR   197.2 141.8   106.9 35.2   31.4 29.0   29.0 28.8   25.1 22.2   14.1 14.1 |



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