

CHEMBIOCHEM

Supporting Information

A Retro-biosynthesis-Based Route to Generate Pinene-Derived Polyesters

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Sequences of CHMO_{Acineto_QM}

The gene and protein sequences of CHMO_{Acineto_QM} are given below:

5'-

```
atgggcagcagccatcatcatcatcacagcagcggcctggtgccgcggcagccatattggctagcatgactg
gtggacagcaaatgggtcgcggatccatgtcacaaaaatggatttgatgctatcgtgattggtggtggtttggcggg
ctttatgcagtcaaaaaattaagagacgagctcgaacttaaggtcaggcttttgataaagccacggatgtcgcaggta
cttggtactggaaccgftaccaggtgacgtgacggatacagaaacccacctctactgctattcttgggataaagaatta
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gcatgattaaagaagagctatcaattcaataccgcggtcaatcggctcattacaacgaagcagatgccttggggaa
gtcaccactgaatatggtgataagtacagggcgcggttcctcatcactgctttaggcttattgtctgcgcctaactgcca
acatcaaaggcattaatcagtttaagggtgagctgcatcataccagccgctggccagatgacgtaagtttgaaggta
aacgtgtcggcgtgattggtacgggtccaccggtgttcaggttattacggctgtggcacctctggctaaacacctcactg
tctccagcgttctgcacaatacagcgttccaattggcaatgatccactgtctgaagaagatgtaaaaagatcaaaga
caattatgacaaaattgggatggtgatggaattcagcccttgccttggcctgaatgaaagcacagtgccagcaatg
agcgtatcagctgaagaacgcaaggcagttttgaaaaggcatggcaaacaggtggcgggttccgttcatgttgaaa
ctttcggatgattgccaccaatattggaagccaatattcgaagcgaataattcattaagggtaaaattgctgaaatcgtc
aaagatccagccattgcacagaagcttatgccacaggatttgatgcaaaacgtccgttgtgtgacagtggttactaca
acaccttaaccgtgacaatgtccgtttagaagatgtgaaagccaatccgattgttgaattaccgaaaacgggtgtgaa
actcgaaaatggcgttccgttgaattagacatgctgatactggccacaggtttgatgccgtcgatggcaactatgtgcg
catggacattcaaggtaaaaacggcctggccattaaagactactggaaagaaggccgctcgagctatatgggtgtctg
cgtaataactatccaaacatgttcatggtgcttggaccgaatggcccgtttaccaacctgccgcatcaattgaatcac
aggtggaatggatcagtgataccattcaatacaggttgaaaacaatgttgaatccattgaatgcacaaaagaagcg
gaagaacaatggactcaaactgcccgaatattgcggaaatgacctattccctaaagcgaatcctggatgtttggtgc
gaatatcccgggcaagaaaaacacggttacttctatctcgggtggtttaaaagaatatcgcagtgcgctagccaactgc
aaaaacctgacctatgaagggtttgatattcaattacaacggtcagatatcaagcaacctgccaatgcctaa-3'
```

Translated sequence of CHMO_{Acineto_QM}:

```
MGSSHHHHHSSGLVPRGSHMASMTGGQQMGRGSMQKMDFDAIVIGGGFGGL
YAVKKLRDELELVQAFDKATDVAGTWYWNRYPGALTDTEHLYCYSWDKELLQS
LEIKKKYVQGPVDRKYLQQVAEKHDLKKSQYQFNTAVQSAHYNEADALWEVTTEYG
DKYTARFLITALGLLSAPNLPNIKGINQFKGELHHTSRWPDDVSFEGKRVGVIGTGST
GVQVITAVAPLAKHLTVFQRSAQYSVPIGNDPLSEEDVKKIKDNYDKIWDGVWNSAL
AFGLNESTVPAMSVSAEERKAVFEKAWQTGGGFRFMFETFGDIATNMEANIEAQN
IKGKIAEIVKDPAIAQKLMQDLYAKRPLCDSGYNTFNDRDNVRLDVKANPIVEITEN
GVKLENGDFVELDMLILATGFDAVDGNYVRMDIQGKNGLAIKDYWKEGPSSYMGV
CVNNYPNMFVLPNGPFTNLPPSIESQVEWISDTIQYTVENNVESIECTKEAEEQ
WTQTCANIAEMTLFPKAQSWIFGANIPGKKNVYFYLGGLKEYRSALANCKNHAYE
GFDIQLQRSDIKQPANA
```

For reference, the original protein sequence of wild type CHMO_{Acineto} (GenBank: BAA86293.1^[1]), is given below:

```
MSQKMDFDAIVIGGGFGGLYAVKKLRDELELVQAFDKATDVAGTWYWNRYPGAL
TDTEHLYCYSWDKELLQSLEIKKKYVQGPVDRKYLQQVAEKHDLKKSQYQFNTAVQ
SAHYNEADALWEVTTEYGDKYTARFLITALGLLSAPNLPNIKGINQFKGELHHTSRW
PDDVSFEGKRVGVIGTGSTGVQVITAVAPLAKHLTVFQRSAQYSVPIGNDPLSEEDV
```

KKIKDNYDKIWDGVVNSALAFGLNESTVPAMSVSAEERKAVFEKAWQTGGGFRFM
 FETFGDIATNMEANIEAQNFIKGKIAEIVKDPAIAQKLMQDLYAKRPLCDSGYNTF
 NRDNVRLLEDVKANPIVEITENGVKLENGDFVELDMLICATGFDAVDGNYVRMDIQGK
 NGLAMKDYWKEGPSSYMGVTVNNYPNMFMLGPNPFTNLPPSIESQVEWISDTI
 QYTVENNVESIEATKEAEEQWTQTCANIAEMTLFPAQSWIFGANIPGKKNNTVYFYL
 GGLKEYRSALANCKNHAYEGFDIQLQRSDIKQPANA

Construction of the F246A variant

Primers in the 5' to 3' direction are specified below:

F246A_f

ggtgatggaattcagcccttgccGCGggcctgaatg

F246A_r

ggcactgtgctttcattcaggccGCGggcaagg

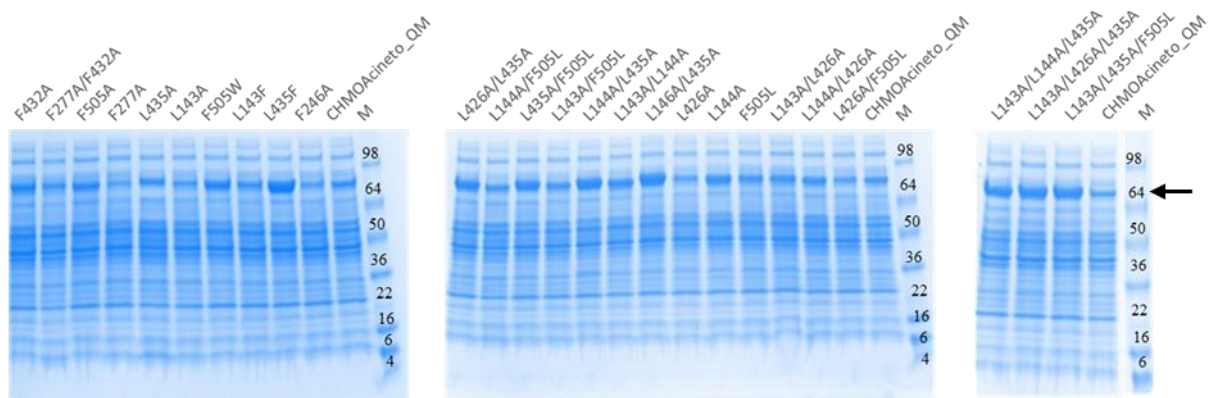


Figure S1. SDS-PAGE analysis (4-15%) of CHMO_{Acineto_QM} and its variants expressed in *E. coli* BL21(DE3). Samples were taken after 20 h of induction (with 0.05 mM IPTG) at 25°C. Lane M, pre-stained protein molecular ladder SeeBlue Plus 2. Relevant bands of CHMO_{Acineto_QM} and variants are indicated by the arrow.

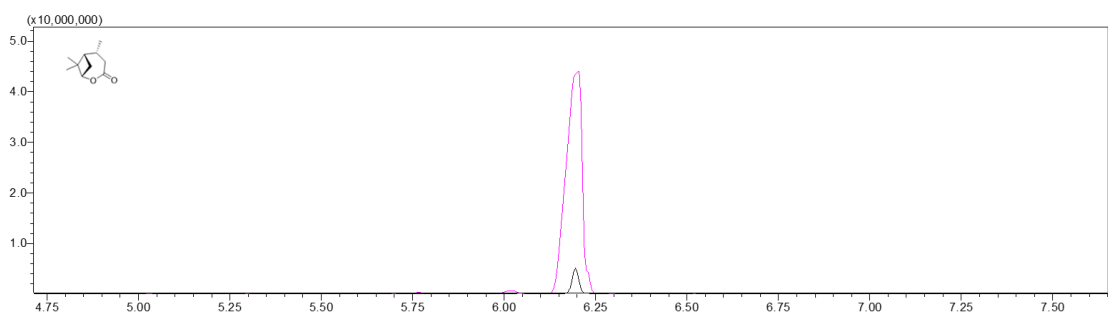


Figure S2. GC-spectra of the “normal lactone” **10** chemically (*purple*) and enzymatically (*black*) produced.

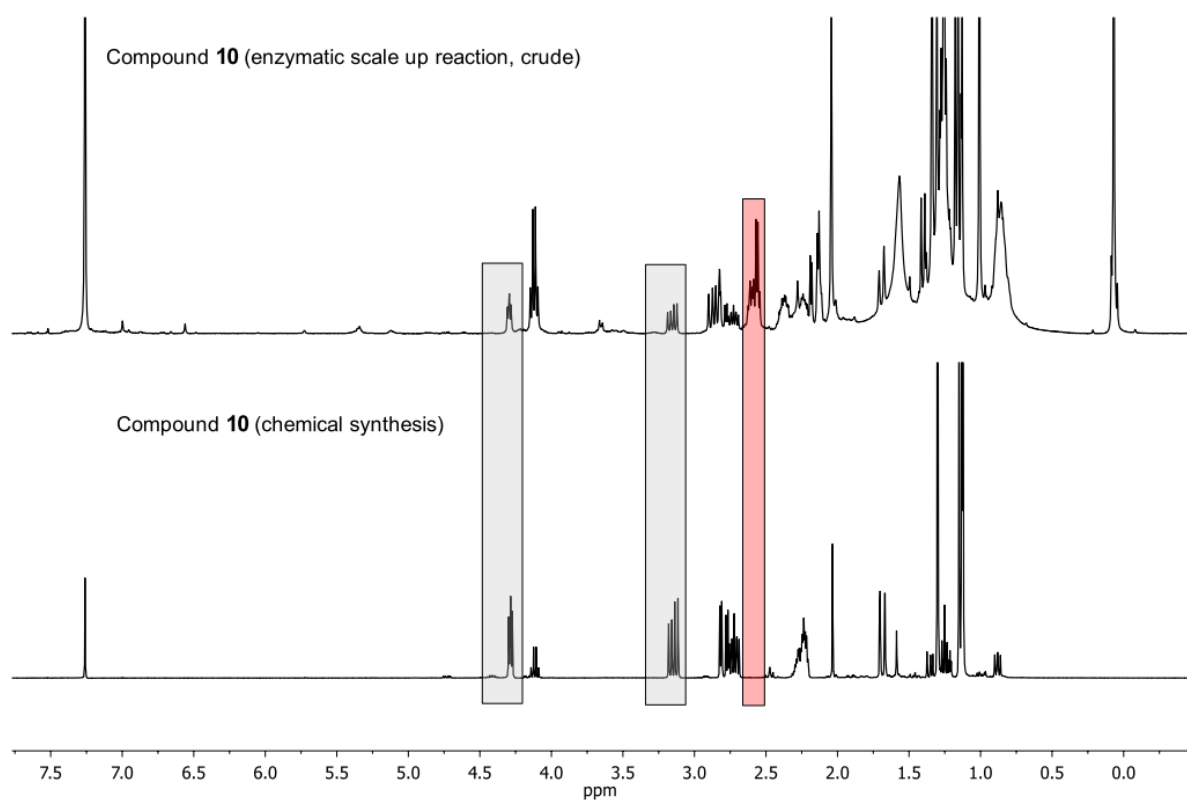


Figure S3. $^1\text{H-NMR}$ spectra of extracted reaction mixture from enzymatic upscaling (top) compared to $^1\text{H-NMR}$ of compound **10** (bottom). Rectangles used to highlight the characteristic peaks of compound **10** (grey) and **9** (red).

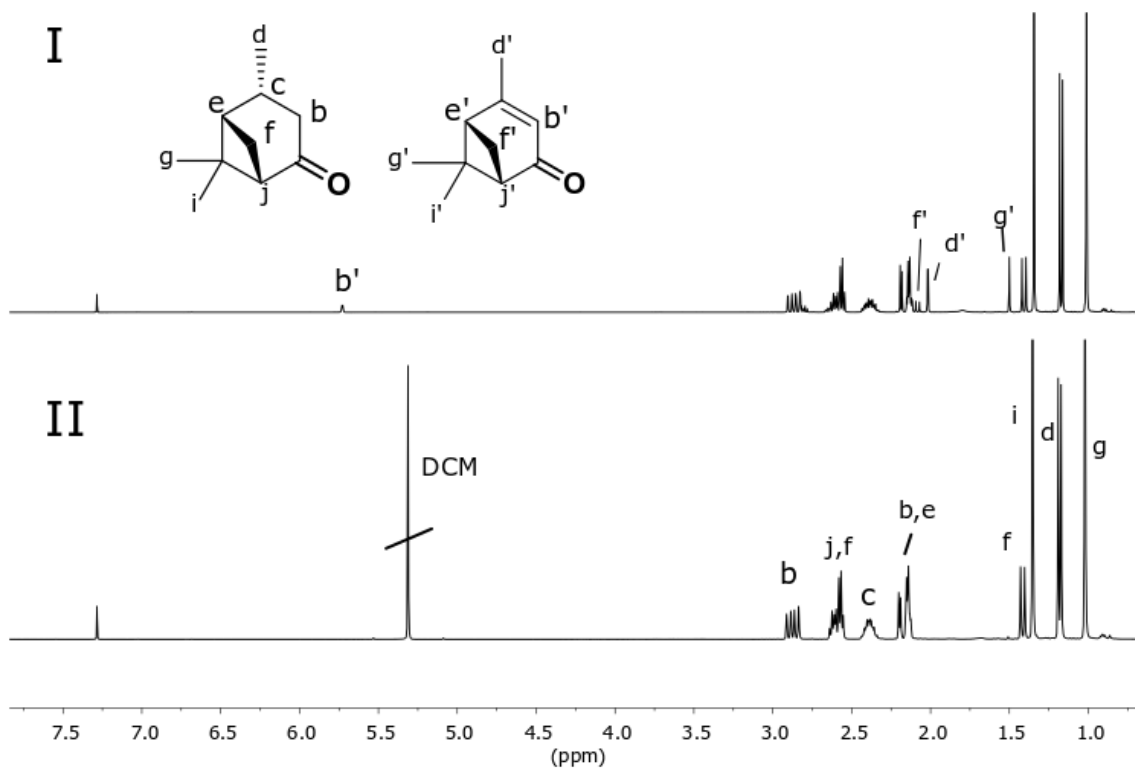


Figure S4. $^1\text{H-NMR}$ spectra of (I.) a mixture containing both **8** and **9** and (II.) the chemically reduced mixture of (I.) in order to contain only **9**.

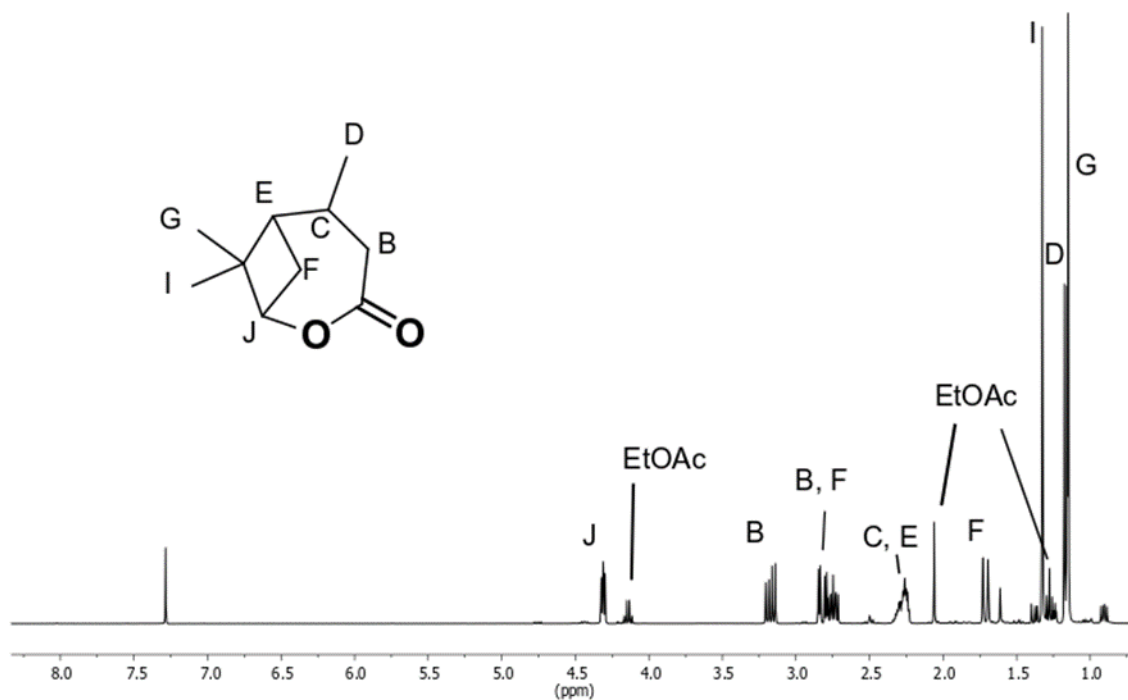


Figure S5 $^1\text{H-NMR}$ spectra of **10**.

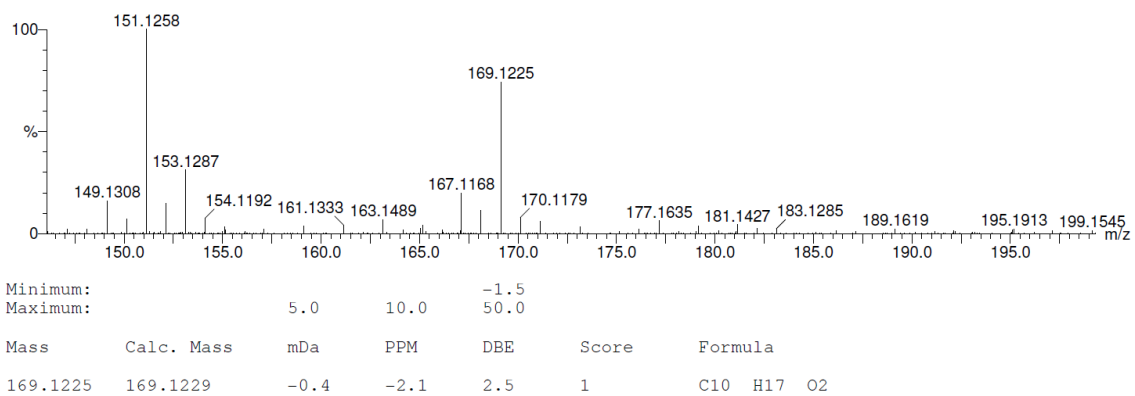


Figure S6. High-resolution GC/MS-ToF mass spectrum of **10**. Spectra was acquired with chemical ionization and methane as a reagent gas.

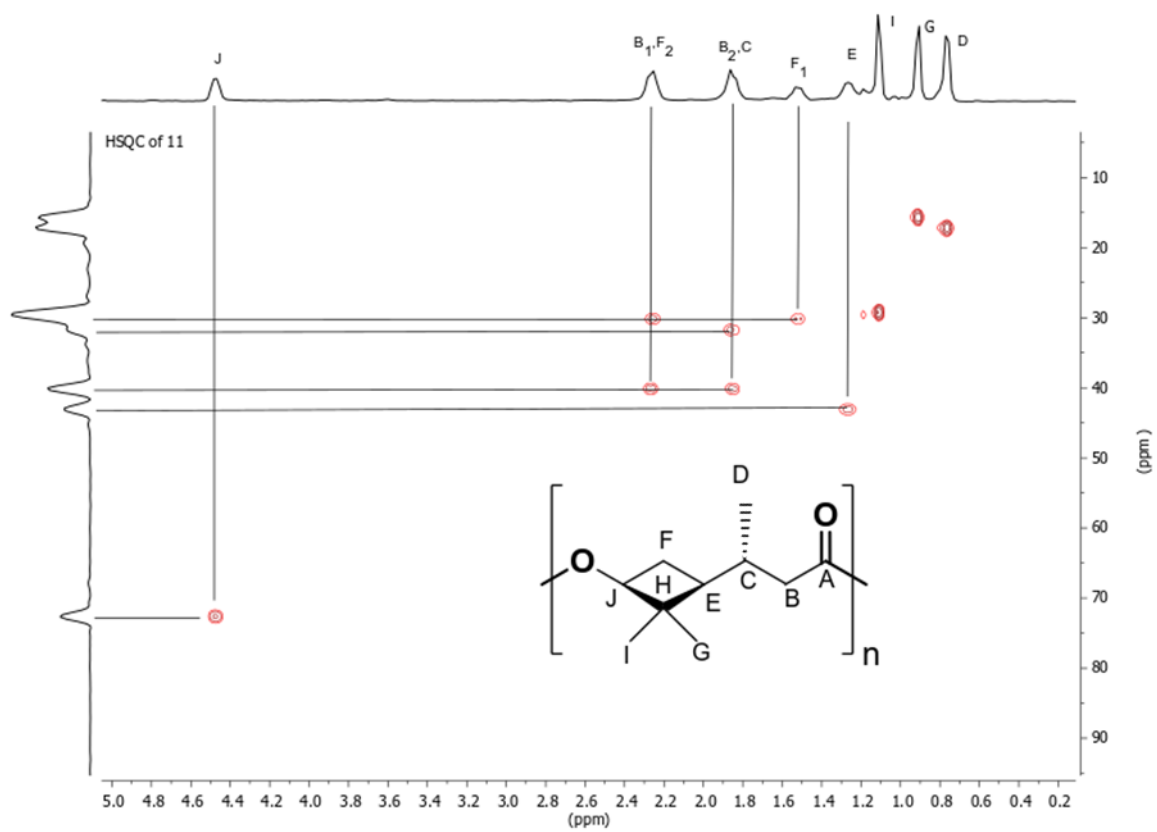


Figure S7. HSQC correlation plot of **11** polymerized using MSA/benzyl alcohol for 24 h at 70 °C.

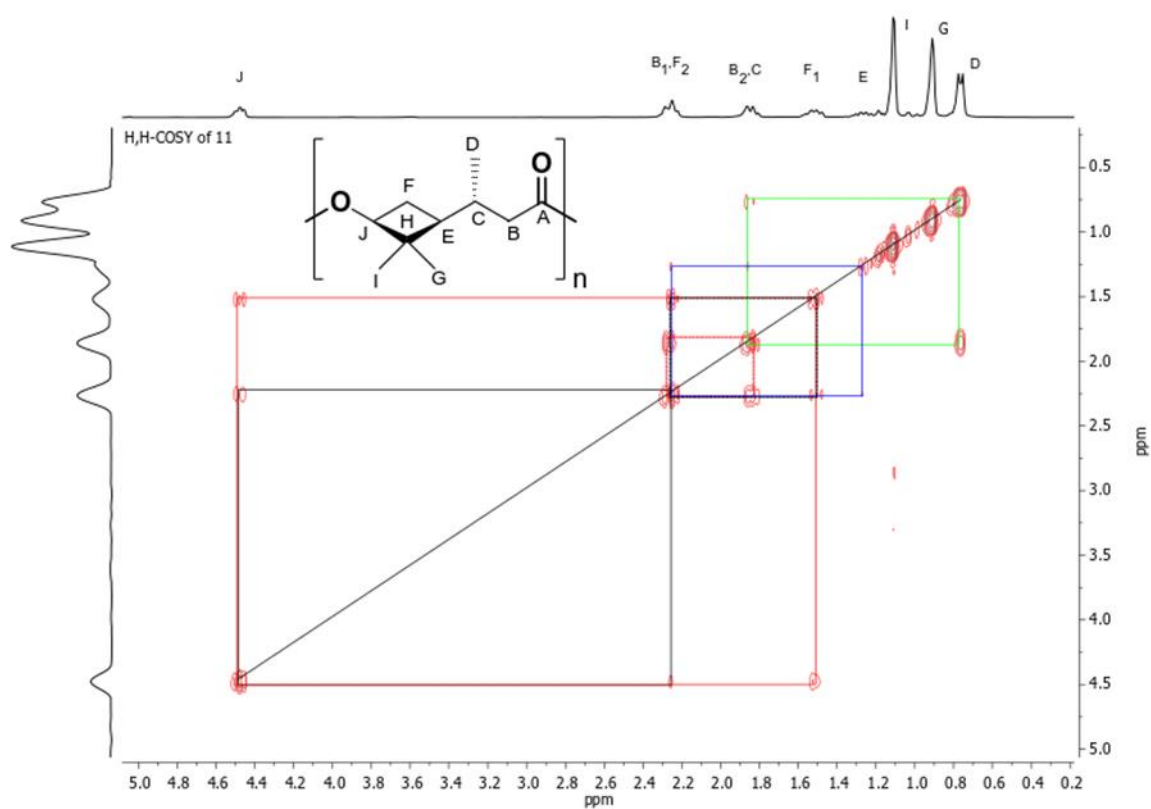


Figure S8. H,H-COSY of **11** polymerized with MSA/benzyl alcohol for 24 h at 70 °C. Rectangles indicate the coupling between the specific protons. Large black: J-F₂ coupling, large red: J-F₁ coupling, small black: F₁-F₂ coupling, small red: B₁-B₂ coupling, blue: E-F₂ coupling and green: C-D coupling.

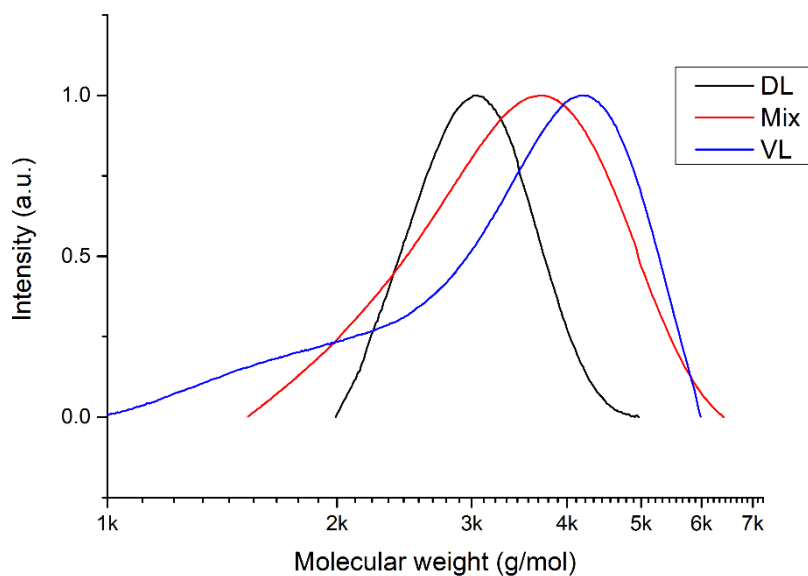


Figure S9. Molecular weight distributions for the synthesized polyesters. Black: poly- ϵ DL; Red: poly(ϵ DL- ϵ VaL); Blue: poly- ϵ VaL.

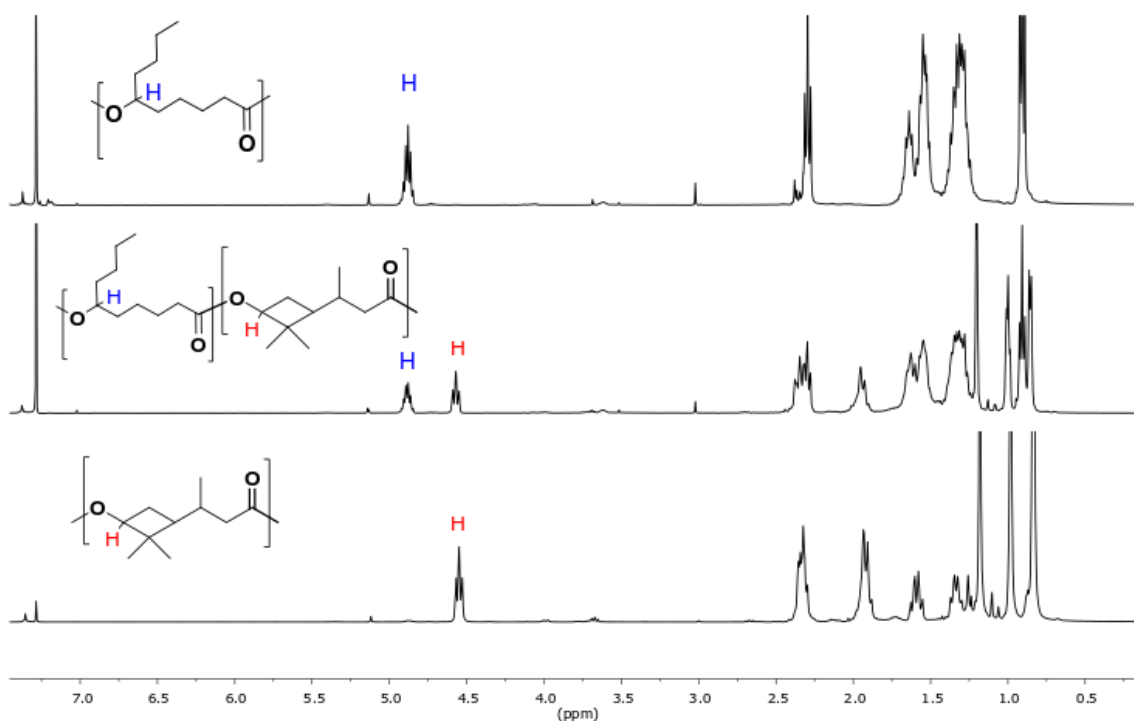


Figure S10. $^1\text{H-NMR}$ spectra of precipitated polymers (top: poly- ϵ DL; middle: poly(ϵ DL- ϵ VaL); bottom: poly- ϵ VaL) polymerized using MSA/benzyl alcohol for 24 h at 70 $^\circ\text{C}$.

Supplementary References

- [1] Y. C. Chen, O. P. Peoples, C. T. Walsh, *J. Bacteriol.* **1988**, *170*, 781-789.