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**Pathway Compartmentalization in Peroxisome of *Saccharomyces cerevisiae* to
Produce Versatile Medium Chain Fatty Alcohols**

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Figure S1

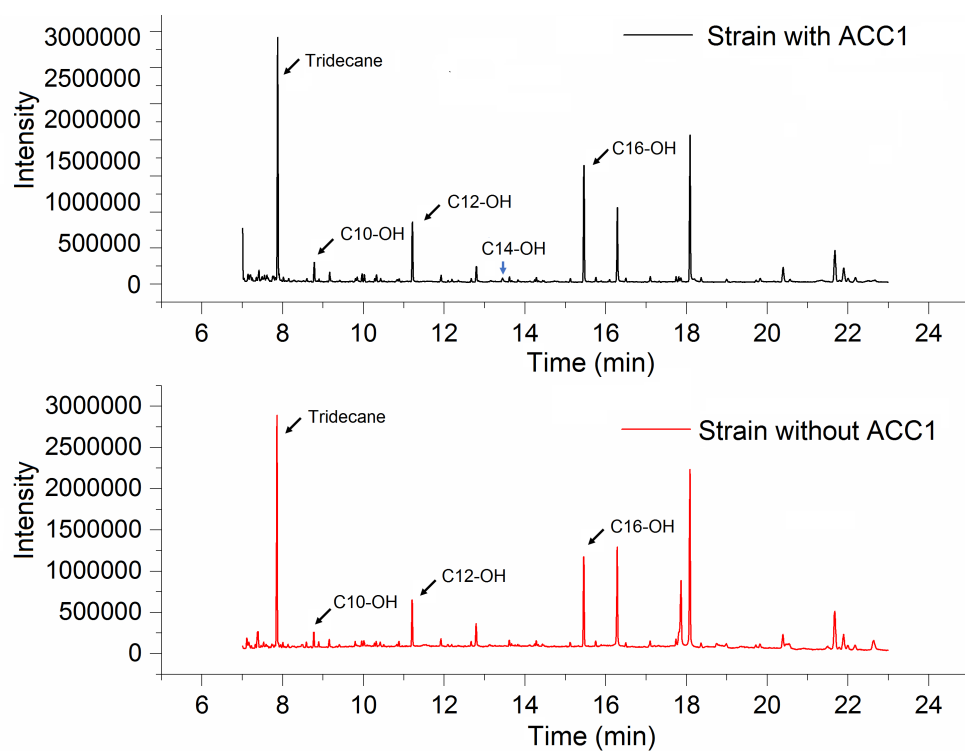


Figure S1: The GC spectrum of fatty alcohols produced by the strains with or without ACC1 overexpressed. Both of the strains were constructed in *S. cerevisiae* BY4741 with TaFAR expressed under constitutive TEF1 promoter and tagged with the PTS2 peptide (KL-QL) and the PEX7 was expressed under constitutive TPI1 promoter.

Figure S2

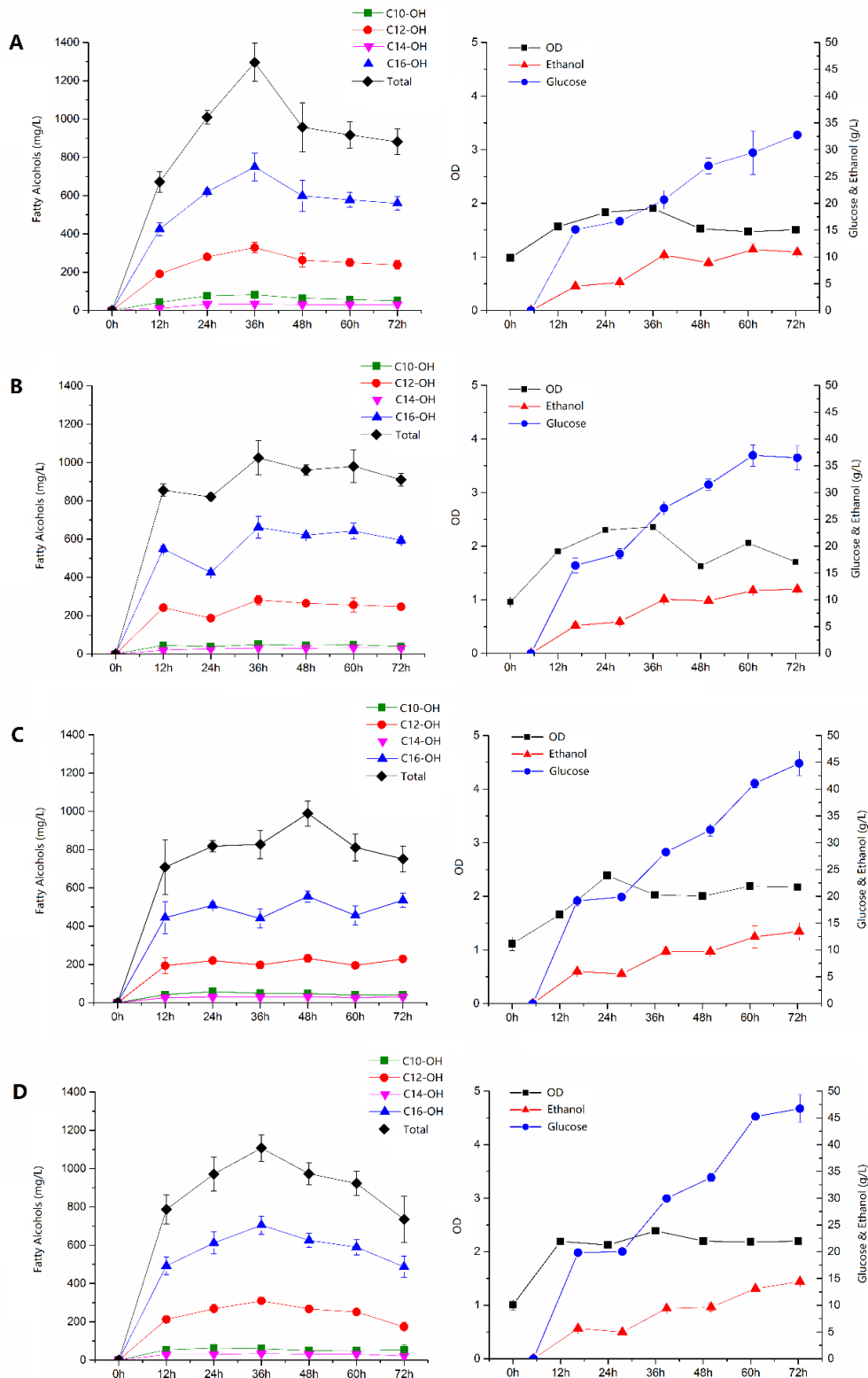


Figure S2. Medium optimization and fed-batch fermentation. (A) Nitrogen limited SC medium with C:N ratio at 50:1; (B) Nitrogen limited SC medium with C:N ratio at 50:1, supplement of 40 mM KCl and 10 mM KOH.; (C) Normal SC medium (control condition with C:N ratio at 3:1); and (D) Normal SC medium with supplement of 40 mM KCl and 10 mM KOH.

Figure S3

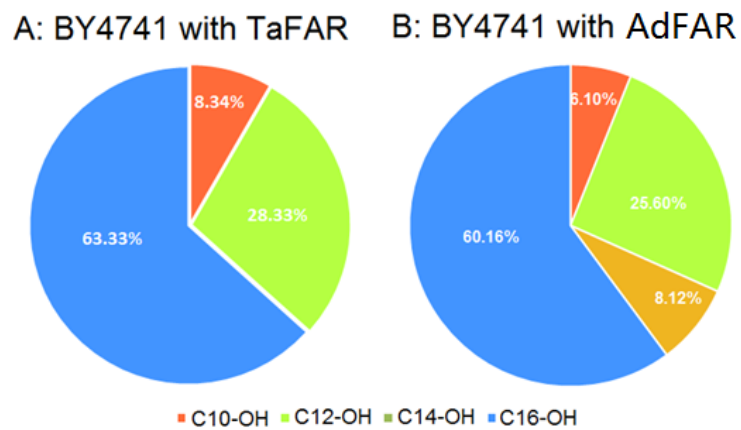


Figure S3: Profiles of fatty alcohols produced by expressing different FAR enzymes in peroxisome. (A) Strain MFAOH31 with TaFAR expressed in peroxisome (8.34% 1-decanol, 28.33% 1-dodecanol and 63.33% 1-hexadecanol); (B) Strain MFAOH38 with AdFAR1 expressed in peroxisome (6.10% 1-decanol, 26.60% 1-dodecanol, 8.12% 1-tetradecanol and 60.16% 1-hexadecanol).

Table S1 The primers used in this study.

Name	Sequence (5'-3')
TEF1p F	AAAGAAAGCATAGCAATCTAATCTAAGTTTAAATTACAAAATGGTTTCCATACCTGAATA
TEF1t R	GCGCAATTAACCCCTCACTAAAGGGAACAAAAGCTGGAGCTCGATAGCGCCGATCAAAGTA
TaFAR-SKL R	ATATAAAAGATATGCAACTAGAAAAGTCTTATCAATCTCCTCATAGCTTGGAGTATCTCA
TaFAR-SKF R	ATATAAAAGATATGCAACTAGAAAAGTCTTATCAATCTCCTCAGAAGCTTGGAGTATCTCA
TaFAR-SFL R	ATATAAAAGATATGCAACTAGAAAAGTCTTATCAATCTCCTCATAGGAAGGAGTATCTCA
TaFAR-SKV R	ATATAAAAGATATGCAACTAGAAAAGTCTTATCAATCTCCTCAAACCTTGGAGTATCTCA
TaFAR-LKL R	ATATAAAAGATATGCAACTAGAAAAGTCTTATCAATCTCCTCATAGCTTAGGTATCTCA
TaFAR-FKL R	ATATAAAAGATATGCAACTAGAAAAGTCTTATCAATCTCCTCATAGCTTGAAGTATCTCA
TaFAR-SHL R	ATATAAAAGATATGCAACTAGAAAAGTCTTATCAATCTCCTCATAGATGGGAGTATCTCA
TEF1t-SKL F	CTTCAGAGCCTCCAGTACTATGAGATACTCCAAGCTATGAGGAGATTGATAAGACTTTTC
TEF1t-SKF F	CTTCAGAGCCTCCAGTACTATGAGATACTCCAAGTTCTGAGGAGATTGATAAGACTTTTC
TEF1t-SFL F	CTTCAGAGCCTCCAGTACTATGAGATACTCCTCCTATGAGGAGATTGATAAGACTTTTC
TEF1t-SKV F	CTTCAGAGCCTCCAGTACTATGAGATACTCCAAGTTTGAGGAGATTGATAAGACTTTTC
TEF1t-LKL F	CTTCAGAGCCTCCAGTACTATGAGATACTCAAAGCTATGAGGAGATTGATAAGACTTTTC
TEF1t-FKL F	CTTCAGAGCCTCCAGTACTATGAGATACTTCAAGCTATGAGGAGATTGATAAGACTTTTC
TEF1t-SHL F	CTTCAGAGCCTCCAGTACTATGAGATACTCCATCTATGAGGAGATTGATAAGACTTTTC
RV-QL F	AAAGAAAGCATAGCAATCTAATCTAAGTTTAAATTACAAAATGAGTCAACGTGTTCAAAGTATTA AAGATCAACTTGTTTCCATACCTGAATATTAT
KL-QL F	AAAGAAAGCATAGCAATCTAATCTAAGTTTAAATTACAAAATGAGTCAAAAACCTCAAAGTATTA AAGATCAACTTGTTTCCATACCTGAATATTAT
KI-QL F	AAAGAAAGCATAGCAATCTAATCTAAGTTTAAATTACAAAATGAGTCAAAAAATCAAAGTATTA AAGATCAACTTGTTTCCATACCTGAATATTAT
KV-HL F	AAAGAAAGCATAGCAATCTAATCTAAGTTTAAATTACAAAATGAGTCAAAAAGTTCAAAGTATTA AAGATCATCTTGTTTCCATACCTGAATATTAT
TEF1p RV-QL R	AAAGAAAGCATAGCAATCTAATCTAAGTTTAAATTACAAAATGAGTCAACGTGTTCAAAG
TEF1p KL-QL R	AAAGAAAGCATAGCAATCTAATCTAAGTTTAAATTACAAAATGAGTCAAAAACCTCAAAG
TEF1p KI-QL R	AAAGAAAGCATAGCAATCTAATCTAAGTTTAAATTACAAAATGAGTCAAAAAATCAAAG
TEF1p KV-HL R	AAAGAAAGCATAGCAATCTAATCTAAGTTTAAATTACAAAATGAGTCAAAAAGTTCAAAG
PEX5 F	AAAGAAAGCATAGCAATCTAATCTAAGTTTAAATTACAAAATGGACGTAGGAAGTTGCTC
PEX5 R	ATATAAAAGATATGCAACTAGAAAAGTCTTATCAATCTCCTTAAAACGAAAATTCCTCTT
TEP1p PEX5 R	CAAGCGGATTATTTCCCACTGAGCAACTCCTACGTCCATTTTGTAAATAAAACCTTAGAT
TEP1t PEX5 F	CATGGACCTGAAAAGATTTAAAGGAGAATTTTCGTTTTAAGGAGATTGATAAGACTTTTC
PEX7 F1	AAAGAAAGCATAGCAATCTAATCTAAGTTTAAATTACAAAATGCTCAGATATCATATGCA
PEX7 R1	ATATAAAAGATATGCAACTAGAAAAGTCTTATCAATCTCCTCAACCTAAGCCGTTCCATA
TEF1p PEX7 R	CACCGTACCCACTAAAACCTTGATATGATATCTGAGCATTGTAATAAAACCTTAGAT
TEF1t PEX7 F	ATGGGATGGAAATTTATTTGTATGGAACGGCTTAGGTTGAGGAGATTGATAAGACTTTTC
TPI1p F	ATTCGATATTGTCGTAACAAATACTTTGATCGGCGTATCTATATCTAGGAACCCATCAG
TPI1p PEX7 R	CACCGTACCCACTAAAACCTTGATATGATATCTGAGCATTGTTAGTTTATGTATGTGTT
PEX7 F2	TTAAATCTATAACTACAAAAACACATACATAAACTAAAATGCTCAGATATCATATGCA
PEX7 R2	TAACATAATTACATGATATCGACAAAGGAAAAGGGCCTGTTCAACCTAAGCCGTTCCATA

CYC1t PEX7 F	ATGGGATGGAAATTTATTTGTATGGAACGGCTTAGGTTGAACAGGCCCTTTTCCTTTGT
CYC1t R	CCACCGCGGTGGCGGCCGCTCTAGAACTAGTGGATCCCCGCAAATTAAGCCTTCGAGC
TEF1p PEX3 R	GTCTCTGCAGAAGCGAACGTGATCTTTGATTTGGGGCCATTTTGTAAATAAAACCTAGAT
PEX3 F	AAAGAAAGCATAGCAATCTAATCTAAGTTTAAATTACAAAATGGCCCCAAATCAAAGATC
PEX3 R	ATATAAAAGATATGCAACTAGAAAAGTCTTATCAATCTCCTTAAGGCTTGAAGGAAAACG
TEF1t PEX3 F	CAACTTTGGCGTCTCCAGCTCGTTTTCTTCAAGCCTTAAGGAGATTGATAAGACTTTTC
TPI1p PEX19 R	ACTCGTTTTTCATTACTTTCGTTGTATGTTTGGCATTTTTAGTTTATGTATGTGTT
PEX19 F	TTAAATCTATAACTACAAAAACACATACATAAACTAAAAATGCCAAACATACAACACGA
PEX19 R	TAACATAATTACATGATATCGACAAAGGAAAAGGGCCTGTTTATTGTTGTTTGAACCGT
CYC1t PEX19 F	CCTTGATAAGGAATTAACCGACGGTTGCAAAACAATAAACAGGCCCTTTTCCTTTGT
TEF1p PEX14 R	CGAACAAATGCCTTACGATCTTTACTGACCACGTCACCTCATTTTGTAAATAAAACCTAGAT
PEX14F	AAAGAAAGCATAGCAATCTAATCTAAGTTTAAATTACAAAATGAGTGACGTGGTCAGTAA
PEX14R	CTTTAGACATACCACCAGATTGAAAATACAAATTTTACCTGGGATGGAGTCTTCGACCT
GFP F	CTCCATCCCAGGTGAAAAATTTGTATTTTCAATCTGGTGGTATGCTAAAGGTGAAGAATT
GFP R	ATATAAAAGATATGCAACTAGAAAAGTCTTATCAATCTCCTTATTTGTACAATTCATCCA
TEF1t GFP F	TGCTGGTATTACCCATGGTATGGATGAATTGTACAAATAAGGAGATTGATAAGACTTTTC
TaFAR mcherry R	TGCTAACCATAACCACCAGATTGAAAATACAAATTTTACCGTATCTCATAGTACTGGAGG
Cherry F	TATGAGATACGGTGAAAAATTTGTATTTTCAATCTGGTGGTATGGTTAGCAAAGGCGAGGA
Cherry R	ATATAAAAGATATGCAACTAGAAAAGTCTTATCAATCTCCTTATAGCTTGAACCACCAG
TEF1t cherry F	TGAAAAATTTGTATTTTCAATCTGGTGGTCCAAAGCTATAAGGAGATTGATAAGACTTTTC
AdFAR F	AAAGAAAGCATAGCAATCTAATCTAAGTTTAAATTACAAAATGTCTCAAAGACTACAAAG
AdFAR R	ATAAAAGATATGCAACTAGAAAAGTCTTATCAATCTCCTTATAGCTTGGAGTAACGCATG
TEF1P_AdFAR R	CCAAATGATCCTTGATACTTTGTAGTCTTTGAGACATTTTGTAAATAAAACCTAGATTAG
TEF1T_AdFAR F	CTTCAGGGCAAGCTCCACCATGCGTACTCCAAGCTATAAGGAGATTGATAAGACTTTTC