

## Supplementary Materials for:

# **$\beta$ -Farnesene Production from Low-cost Glucose in Lignocellulosic Hydrolysate by Engineered *Yarrowia lipolytica***

**Table S1** Plasmids used in this work.

Plasmids	Descriptions	Source
pRSF-HUH	hisG-P <sub>ura3</sub> -Ura3-T <sub>Ura3</sub> -hisG (HUH), Kana <sup>r</sup>	Lab storage
pRSF-ku70-HUH	HR donor to disrupt <i>ku70</i> locus	This work
pRSF-ku80-Leu2-HUH	HR donor to integrate P <sub>TEFin</sub> -Leu2-T <sub>XPR2</sub> into ku80 locus	This work
pRSF-inA1-aaBFS-HUH	HR donor to integrate P <sub>TEFin</sub> -aaBFS-T <sub>XPR2</sub> into intA1 locus	This work
pRSF-intC1-ERG10-tHMGR-HUH	HR donor to integrate P <sub>TEFin</sub> -ERG10-T <sub>ERG10</sub> and P <sub>TEFin</sub> -tHMGR-T <sub>tHMGR</sub> into intC1 locus	This work
pRSF-intC2-ERG12-ERG13-HUH	HR donor to integrate P <sub>EXP1</sub> -ERG12-T <sub>ERG12</sub> and P <sub>EXP1</sub> -ERG13-T <sub>ERG13</sub> into intC2 locus	This work
pRSF-intC3-IDI-ERG20-HUH	HR donor to integrate P <sub>TEFin</sub> -IDI-T <sub>IDI</sub> and P <sub>TEFin</sub> -ERG20-T <sub>ERG20</sub> into intC3 locus	This work
pRSF-intE1-ERG8-ERG19-HUH	HR donor to integrate P <sub>EXP1</sub> -ERG8-T <sub>EXP1</sub> and P <sub>EXP1</sub> -ERG19-T <sub>ERG19</sub> into intE1 locus	This work
pRSF-aaBFS-cjBFS-HUH	HR donor to integrate P <sub>TEFin</sub> -cjBFS-T <sub>XPR2</sub> into aaBFS locus	This work
pRSF-aaBFS-mcBFS-HUH	HR donor to integrate P <sub>TEFin</sub> -mcBFS-T <sub>XPR2</sub> into aaBFS locus	This work
pRSF-intE2-ERG10-tHMGR-HUH	HR donor to integrate P <sub>TEFin</sub> -ERG10-T <sub>ERG10</sub> and P <sub>TEFin</sub> -tHMGR-T <sub>tHMGR</sub> into intE2 locus	This work
pRSF-intE3-ERG12-ERG13-HUH	HR donor to integrate P <sub>EXP1</sub> -ERG12-T <sub>ERG12</sub> and P <sub>EXP1</sub> -ERG13-T <sub>ERG13</sub> into intE3 locus	This work
pRSF-intE4-IDI-ERG20-HUH	HR donor to integrate P <sub>TEFin</sub> -IDI-T <sub>IDI</sub> and P <sub>TEFin</sub> -ERG20-T <sub>ERG20</sub> into intE4 locus	This work
pRSF-intF1-ERG8-ERG19-HUH	HR donor to integrate P <sub>EXP1</sub> -ERG8-T <sub>EXP1</sub> and P <sub>EXP1</sub> -ERG19-T <sub>ERG19</sub> into intF1 locus	This work

pRSF-intF2-tHMGR-	HR donor to integrate $P_{TEFin}$ - $tHMGR$ - $T_{HMGR}$ and $P_{TEFin}$ - $tHMGR$ - $T_{HMGR}$ into intF2 locus	This work
pRSF-intF2- nadhHMGR-	HR donor to integrate $P_{TEFin}$ - $nadhHMGR$ - $T_{XPR2}$ and $P_{TEFin}$ - $nadhHMGR$ - $T_{XPR2}$ into intF2 locus	This work
nadhHMGR-HUH		
pRSF-intF2- nadhHMGR-tHMGR-	HR donor to integrate $P_{TEFin}$ - $nadhHMGR$ - $T_{XPR2}$ and $P_{TEFin}$ - $tHMGR$ - $T_{HMGR}$ into intF2 locus	This work
HUH		
pRSF-intF3-aaBFS- HUH	HR donor to integrate $P_{TEFin}$ - $aaBFS$ - $T_{XPR2}$ into intF3 locus	This work
pRSF-intF3-aaBFS- aaBFS-HUH	HR donor to integrate $P_{TEFin}$ - $aaBFS$ - $T_{XPR2}$ and $P_{TEFin}$ - $aaBFS$ - $T_{XPR2}$ into intF3 locus	This work

**Table S2** Primers used in this work.

ku70up-F	CGCACTTAATTAACACTACACTACACTACTGTACCATTCTACCC
ku70up-R	ttttacaacTGGTTCGTGTTCGTGGTGTGTCG
ku70-HUH-F	ACACGAACCAAGtgtaaaacgacggccagtcgaac
ku70-HUH-R	GTACTTTGGTAGCCGTAGGTCTCGTACTGCTTGAC
ku70dn-F	CTACCAAAAAGTACTATGGGAAGTGACTAGGGAGGC
ku70dn-R	agacacAGTGAACGACCAAGACTAAAGGGTGG
ku70-line-F	CACTACACTACACTACTTGTACCATTCTACCC
ku70-line-R	AGTGAACGACCAAGACTAAAGGGTGGC
ku80up-F	CGTTGCGCAGCAACCCACTGCCGCTAACCC
ku80up-R	ccggctctGCTCAAAGTCACTGGTATGGCAGTGTG
ku80-Leu2-tef-F	CTTGAGCagagaccgggttggcgccgtatttg
ku80-Leu2-tef-R	gttcgggttcctgcggtagtactgcaaaaagtgtcg
ku80-Leu2-F	ctaaccgcaggaaccgcgaaactaagaagaccaagactgac
ku80-Leu2-R	gtttacaacGACACGGGCATCTCACTTGCATATG
ku80-Leu2-HUH-F	CCGTGTCgttgtaaaacgacggccagtcgaac
ku80-Leu2-HUH-R	TGTATCCGTAGCCGTAGGTCTCGTACTGCTTGAC
ku80down-F	GGCTACGGATACATATTGGTGACAGTTACCGCAATC
ku80down-R	TCTCAAATGCCTGGATTCTGGATCCATTCCCTTGGTGTGCTTC
ku80-line-F	AGCAACCCACTGCCGCTCAACC
ku80-line-R	GATTCTGGATCCATTCCCTTGGTGTGCTTC

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intA1-aabfs-up-F	CGTTGCGCgaatgcgtcgatttttatccaaaaatac
intA1-aabfs-up-R	CCCGGTCTCTtgtgcactcgccgatatggttgatc
intA1-aabfs-tef-F	gegacaAGAGACCGGGTTGGCGGGCGTATT
intA1-aabfs-tef-R	GTGGACATTCTAGAGAATGATTCTTACTCAGAAGGAAATG
intA1-aabfs-F	TTCTCTAGAACATGTCCACCCTGCCCATCTCCTCTGTC
intA1-aabfs-R	AGGCTCGAGTTACACGACCATGGGGTGCACGAAGAAAG
intA1-aabfs-xpr-F	CTCGAGCCTGTCCCCACGTTGCCGGTCTTGC
intA1-aabfs-xpr-R	cgtttacaacACGGGCATCTCACTTGCATGTATG
intA1-aabfs-HUH-F	AGATGCCCGTgtgtaaaacgacggccagtcgaac
intA1-aabfs-HUH-R	ctagtggatGTAGCCGTAGGTCTCGTACTGCTTGAC
intA1-aabfs-dn-F	ACGGCTACatccactagtggctgccatagcactattg
intA1-aabfs-dn-R	TCTCAAATGCCTGgatttaaatgcggccgcaatgcacgc
intA1-aabfs-line-F	gaatgcgtgcgattttatccaaaaataccc
intA1-aabfs-line-R	gatttaaatgcggccgcaatgcac
intA1-cjbfs-tef-R	ATGTCCTTCTAGAGAATGATTCTTACTCAGAAGGAAATG
intA1-cjbfs-F	AATCATTCTCTAGAAAGGACATGAGCATTCCACTTGGCG
intA1-cjbfs-R	GGGGACAGGCTCGAGTCAAATGGCAAGAGGCTCTGTCAACAGC AGAG
intA1-cjbfs-xpr-F	TGCCATTGACTCGAGCCTGTCCCCACGTTGCCGGTCTTGC
intA1-mc-bfs-tef-R	TATGGTGCTTCTAGAGAATGATTCTTACTCAGAAGGAAATG
intA1-mcbfs-F	ATCATTCTCTAGAACGCACCATACCGTCTCCTCTGTAAGTTTTC
intA1-mcbfs-R	GGACAGGCTCGAGTCAAATAACCATGGGGTGCACAAAG
intA1-mcbfs-xpr-F	GTTATTGACTCGAGCCTGTCCCCACGTTGCCGGTCTTGCCTC
intC1-up-F	CGTTGCGCttgctggcccttgctcacatgttcttcctg
intC1-up-R	TGCAGTACAaaaatcataaataatagatgaatag
intC1-ERG10-tef-F	gattttGTACTGCAAAAAGTGTGGTCGGATG
intC1-ERG10-tef-R	GACACAAATGCGCCGCCAACCCGGTCTCTGAGCCCG
intC1-ERG10-F	GTCTCTGAGCCCGTCTACATTGTTCTACTG
intC1-ERG10-R	CGTGT CCTGCAGGCCGTGACGAATACGACTCTCG
intC1-ERG10-xpr-F	CTGCAGGACACGGGCATCTCACTTGCAT
intC1-ERG10-xpr-R	tcatagTAGGATCCA ACTACGGA ACTTGTGTTG
intC1-tHMGR-F	GGATCCTACctatgaccgtatgcaaatttcgaaccg
intC1-tHMGR-R	CTTTTGCA GTACaccaggctgtgaagggtgggtgagaagc
intC1-tHMGR-tef-F	gactgggtGTACTGCAAAAAGTGTGGTCGGATGACGTGG

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intC1-tHMGR-tef-R	tacaacAGAGACCGGGTTGGCGGCCGCATTG
intC1-tHMGR-HUH-F	CGGTCTCTgttgtaaaacgcacggccagtcgaac
intC1-tHMGR-HUH-F	ggatctgaGTAGCCGTAGGTCTCGTACTGCTTGAC
intC1-dn-F	TACGGCTACTcagatccactagtggcctagtcgtg
intC1-dn-R	TCTCAAATGCCTGgaagtggggatttaaatcgccgc
intC1-line-F	ttgtggccctttgctcacatgttc
intC1-line-R	gaagtggggatttaaatcgccgc
intC3-up-F	GCGTTGCGCccttattcgactcactatagaagttcctattc
intC3-up-R	CTTTTGCACTACCGtatacgggcatttcctctgac
intC1-IDI-tef-F	gtatacgGTACTGCAAAAAGTGTGGTCGGATG
intC1-IDI-tef-R	GTAAGACGTCGTAGAGACCGGGTTGGCGGCCGCATTG
intC1-IDI-F	CGGTCTCTACGACGTCTTACAGCGACAAAATCAAGAG
intC1-IDI-R	ATGAATAACAGTTGTATCTGGAGTCGAGACATGATG
intC1-ERG20-F	CAGATACAACGTATTATTGTGGGCCGAAGTGAAGAGG
intC1-ERG20-R	CTTTTGCACTCCAAAGGCAGAAATCGAAAGCGTGTCC
intC1- ERG20-tef-F	GCCTTGGAGTACTGCAAAAAGTGTGGTCGGATGACGTG
intC1- ERG20-HUH-R	cgacacgaacaatGTAGCCGTAGGTCTCGTACTGCTTGAC
intC3-dn-F	CGGCTACattgtcgatcgac
intC3-dn-R	TCTCAAATGCCTGaaatcgccgcgaatgcacgcgataatta
intC3-line-F	cgactcaatagaagtccatatcttagaaatgtataggacttc
intC3-line-R	aaatcgccgcgaatgcacgcgataatta
intC2-up-F	CATTAATTGCGTTGCGCggttctgataacgagaatcgtaatcc
intC2-up-R	gggccaaactcccttagagaataggaacttctatgtgagtc
intC2-ERG12-exp-F	cctattctctagaaggagttggcgccccgttttcg
intC2-ERG12-exp-R	AATGATGTAGTCCATgcttagatatgtctgtgttaaggggttgg
intC2-ERG12-F	atatctacagcATGGACTACATCATTGCGGCCAGG
intC2-ERG12-R	GAGCTCGGAAGAATGGAGATGGCCTATGACTCC
intC2-ERG13-F	ATTCTCCCCGAGCTCCACTTCTCGTGTGACAGATG
intC2-ERG13-R	ctacagcATGTCGCAACCCCAGAACGTTGGAATCAAAG
intC2-ERG13-exp-F	GGGGTTGCGACATgcttagatatgtctgtgttaaggggttgg
intC2-ERG13-exp-R	cgtttacaacaaggagttggcgccccgttttcg
intC2-HUH-F	ccaaactcccttgttaaaacgcacggccagtcgaacc
intC2-HUH-R	gtatccgaggagttcGTAGCCGTAGGTCTCGTACTGCTTGAC
intC2-dn-F	CCTACGGCTACgaactcctcgatcacccgagtgccgc

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intC2-dn-R	TCTCAAATGCCTGgcgaatgcacgcgatgattgaaacgcctg
intC2-line-F	ggttctgataacgagtaatcgtaatccgcaaataac
intC2-line-R	gccaatgcacgcgatgattgaaacgcctg
intE1-up-F	TTAATTGCGTTGCGCtaacaatgataaaccaggcccattgattgagac
intE1-up-R	aaactcctgcattgtccaagtacaatactaaacatactgtac
intE1-ERG8-exp-F	tggacaatgcaaggagttggcgccgttttcgagccc
intE1-ERG8-exp-R	CGAATAGGTGGTCATgctgttagatatgtcttgtgttaagggggttgg
intE1-ERG8-F	atctacagcATGACCACCTATTGGCTCCGGAAAGGC
intE1-ERG8-R	GCACTTGGACTCGAGCCATTACAGACGCGGAAAAG
intE1-ERG19-F	CTCGAGTCCAAGTGCAGAGATCCAGGACCAGGAGCGTTG
intE1-ERG19-R	tacagcATGATCCACCAGGCCTCCACCACCGCTCCGGTGAAC
intE1-ERG19-exp-F	CCTGGTGGATCATgctgttagatatgtcttgtgttaagggggtttg
intE1-ERG19-exp-R	ttttacaacccccacacgttgcgtgatgagcggcggcagattcgagcgttcc
intE1-HUH-F	acgtgtgggggtgtaaaacgcacggccagtcac
intE1-HUH-R	tatgttgtcgacgctGTAGCCGTAGGTCTCGTACTGCTTGAC
intE1-dn-F	CCTACGGCTACagcgtcgacaagcatacagccctcg
intE1-dn-R	TCTCAAATGCCTGaatgcacgcgatgttagaagcaattgga
intE1-line-F	taacaatgataaaccaggcccattgattgagac
intE1-line-R	gaatgcacgcgatgttagaagcaattggag
intE2-up-F	CGTTGCGCttcagtcgataccatgccgtcaactttcttg
intE2-up-R	ACTTTTGCACTACcaggatatacgagctacgtgggtgc
intE2-dn-F	CCTACGGCTACgtgtggattgcgatattgaagtgtgactcg
intE2-dn-R	TCTCAAATGCCTGgcgatgaaatacgcgcaggtttagactatacc
intE2-line-F	cagtcgataccatggccgtcaactttc
intE2-line-R	gcgatgaaatacgcgeaggtttagactatacc
intE3-up-F	CTTACATTAATTGCGTTGCGCtaagcggctctatccatggttactg
intE3-up-R	ggcccaaactcctgattccgaacagaaggaaatgcacgcgatc
intE3-dn-F	TACGAGACCTACGGCTACctgaactgtataccaatgtacttgtacgccttgaatc
intE3-dn-R	TCTCAAATGCCTGtgaaggaaatgcctaaaacctgaattgaacgaagcg
intE3-line-F	cggcttatccatggttactgaggagaaatgttc
intE3-line-R	tgaaggaaatgcctaaaacctgaattgaacgaagcg
intE4-up-F	AATTGCGTTGCGCgccccctggcggtctgataacgagtaat
intE4-up-R	GCACTTTTGCACTACatgaagtgaagttctatacttttagagaataggaacttct
intE4-dn-F	AGACCTACGGCTACgtagggaaacaccgcggAACAGGAAACATC

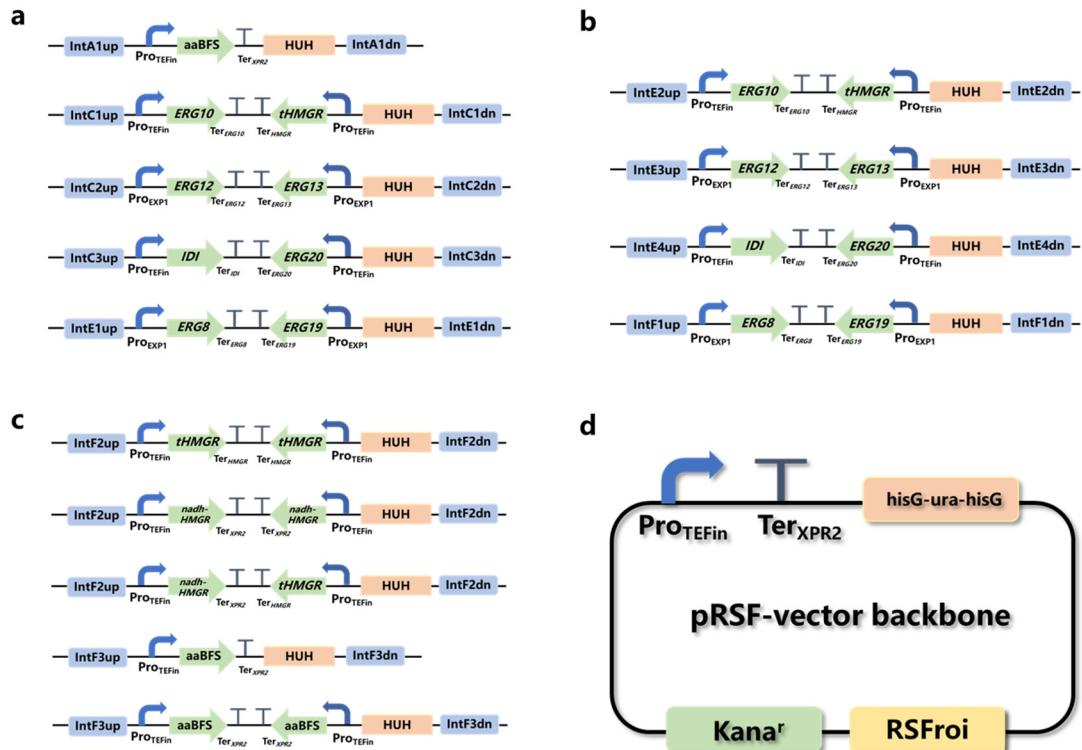
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intE4-dn-R	TCTCAAATGCCTGcacgcatttaacactggaccgtactgcc
intE4-line-F	ctggcggttctgataacgagtaatcg
intE4-line-R	cacgcgattaaacactggaccgtactgcc
intF1-up-F	CATTAATTGCGTTGCGCACACACAAGTTCAAGACTGTCCTTTG
intF1-up-R	ccaaactcctttatgccttactcgtagtgcataactgc
intF1-dn-F	GAGACCTACGGCTACgtggccttctgggaccgagaattagtgt
intF1-dn-R	TCTCAAATGCCTGggatttaatgcggccgcgaatgcac
IntF1-line-F	caagttcagacttgctcttgagtcttc
intF1-line-R	ggatttaaatgcggccgcgaatgcac
intF2-up-F	CATTAATTGCGTTGCGCgagaaaagtgggtacagtatgtacatac
intF2-up-R	TGCAGTACtcctcggtgggtattcgtagagcttcggctttgg
intF2-1-tef-F	ccgaggaGTACTGCAAAAGTGCTGGTCGGATGACGT
intF2-1-tef-R	cagactgggtAGAGACC GGTTGGCGGCGCATTG
intF2-1-thmgr-F	CCCGGTCTCTaccaggctgtgaagggtgggtgagaagcacgttc
intF2-1-thmgr-R	catagGTAGctatgaccgtatgcaaataattcgaaccgtttt
intF2-2-thmgr-F	catagCTACctatgaccgtatgcaaataattcgaaccgtttt
intF2-2-HUH-F	actggccgtcggttacaacAGAGACC GGTTGGCGGCGCATTG
intF2-2-HUH-R	taattgagaaggccactGTAGCCGTAGGTCTCGTACTGCTTGAC
intF2-dn-F	ACCTACGGCTACgtggcctctcaattaaataaac
intF2-dn-R	TCTCAAATGCCTGgcacgcattccctgattactatgatataatttc
IntF2-line-F	ggtacagtatgtacatacttgtaagggtgcaca
intF2-line-R	geacgcgattccctgattactatgatataatttc
intF2-nadh-tef-R	GCCAGTCTGCCGGTAGAGACC GGTTGGCGGCGCATTG
intF2-nadh-thmgr-F	ACCCGGTCTCTACCGGCAAGACTGGCCACATCGACGGACTG
intF2-nadh-thmgr-R	CGTAGTTGGATCTTAAGTGTCTCCAGCACTGTAGCTCTA
intF2-nadh-xpr-F	GAGAACACTTAAGATCCA ACTACGGAAC TTGTGTTGATG
intF2-nadh-xpr-R	cggtagGTAGGACACGGGCATCTCACTGCAT
intF2-2nsdh-xpr-R	AAGTGAGATGCCGTGTCAGGACACGGGCATCTCACTGCAT
intF2-2nsdh-xpr-F	AGATGCCGTGTCCTGACACGGGCATCTCACTGCATAT
intF3-up-F	TAATTGCGTTGCGCtgaagacaaggccgacgtatgagc
intF3-up-R	GAGTGAATTGgagggttagatcaggttatggat
intF3-1-tef-F	cgtcaaccctcCAATTCACTCACTCTCCGACTATCCAAACaa
intF3-2-tef-R	aagacactgcggAGAGACC GGTTGGCGGCGTATTG
intF3-2-HUH-F	cgcagtgtttcgctctggatctccagttgcgtatgaac

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intF3-2-HUH-R	cagaatgtcagcGTAGCCGTAGGTCTCGTACTGCTTGAC
intF3-dn-F	GGCTACgctgacattctcgctctggatcatctgc
intF3-dn-R	TCTCAAATGCCTGaaaggcacagagatcccttagagggatcag
IntF3-line-F	ctgaagacaaggccgacgatgatgac
intF3-line-R	gaaaggcacagagatcccttagagggatcag



**Figure S1 (a-c)** Schematic representation of gene chromosome integration in this study and (b) schematic representation of pRSF vector backbone. Blue rectangular boxes represent homologous arms, orange rectangular box represents the his-ura-his box, green arrow boxes represent genes, blue curved arrows represent promoters and the dark blue T represents terminators.

**Table S3** The exogenous DNA used in this study.

Genes	Sequence (codon optimized)
aaBFS	ATGTCCACCCTGCCCATTCCTCTGTCTCCTCTTCCACCTCCCC ATTGTCGTGGACGACAAGGACTCTACCAAGCCGACGTGATCCGACACA CCACCACCTTCAACGCCTCTATTGGGGTGACCAGTTCTCACCTACGA CGAGCCGAGGACCTGGTCATGAAGAACGAGCTGGAGGAGCTCAA GGAAGAGGTCAAGAACGGAGCTGATCACCATTAGGGCTCCAACGAGCC CATGCAGCACGTCAAGCTGATCGAGCTCATTGACGCCGTGAGCGACTC

	GGTATCGCTTACCACTTCGAGGAAGAGAGATCGAGGAAGCCCTGCAGCAC ATTCACGTCACCTACGGAGAGCAGTGGTGGACAAGGAGAACCTCCAG TCTATTCCCTGTGGTCCGACTGCTCCGACAGCAGGGTTCAACGTCTC TTCCGGAGTGTCAAGGACTCATGGACGAGAAGGGAAAGTTCAAGGA GTCTCTGTGCAACGACGCCAGGGCATCTGGCTCTACGAGGCCGCT TTCATGCGAGTCGAGGACGAGACCATTCTGGACAACGCCCTGGAGTTCT CCAAGGTGCACCTCGACATCATTGCTAAGGACCCCTTGTGACTCTCC CTGCGAACCCAGATTCAACCAGGCTCTGAAGCAGCCTCTCCGACGACGA CTGGCTCGAACATCGAGGCTCTGCACTACATGCCCATCTACCAGCAGGAGA CCTCCCACGACGAGGTCTGCTCAAGCTGGCTAAGCTCGACTTCTCCGT GCTCCAGTCTATGCACAAGAAGGAGCTGTCTCACATCTGCAAGTGGTGG AAGGACCTGGACCTCCAGAACAAAGCTCCCTCGAGGACCGAGTC GTGGAGGGATACTTCTGGATCCTGTCCATCTACTACGAGCCCCAGCACG CCCGAACCCGAATGTTCCATGAAGTCTGCATGGCTGGCTGGCGTGC GACGACACCTCGACAACTACGGCACCTACGAGGAGCTGGAGATCTC ACCCAGGCTGTCGAGAAGTGGTCTATTCCCTGTCTGGACATGCTCCCCG AGTACATGAAGCTGATCTACCAGGAGCTGGTCAACCTCACGTGGAGAT GGAGGAGTCCCTGGAGAACAGGAAGGCACAGGCCTACCAGATTCACTACGT GAAGGAGATGGCTAAGGAGCTCGTCCGAAACTACCTGGTGGAGGCCCG ATGGCTCAAGGAAGGCTACATGCCAACCTGGAGGAGTACATGTCTGC TCCATGGTACCGAACCTACGGCCTGATGACCGCTCGATCTACGTCG GCCGAGGTGACATCGTAACGAGGACACCTCAAGTGGGTGCTTCC CCCTCCCATCGTAAGGCCTCTCGTGATCATTGACTCATGGACGACA TTGTCTCTCACAAGGAAGAGCAGGAGCGAGGTACGTGGCTTCC TGAGTGCTACTCTAAGGAGTCCGGAGCCTTGAGGAAGAGGCC GTACATCTCCGAAAGGTCGAGGACGCCTGGAGGTACATCAACCGAGA GTCTCTCGACCTACCCTGTCCTTGCTCATGCC ACCTCGCTCGAACATGTGTGAGGTCTGTACTCCGTGAACGACGGCTTC CCACGCCAGGGTGACATGAAGTCCTACATGAAGTCTTCTCGTGC CCCCATGGTCGTGAA
<i>cjBFS</i>	ATGAAGGACATGAGCATTCACTTTGGCGGCGGTGTCTTCATCCACGG AGGAGACTGTGCGACCCATTGCTGACTCCATCCGACACTGTGGGGCAA CCACTCTTGAAATCTGCTGCCATGTGGAGACTATTGACGCCGCGACT CAGGAACAGCACGCCCTCTCAAGCAGGAGGTGCGTCGATGATCACC ACAACGGCCAACAAGCTGCCAAAAGCTCCACATGATTGATGCCGTT AACGGTTAGGAGTGGCCTACCACTTGAGAAGGAGATTGAGGACGAGC TGGGAAAGGTTCCATGATCTGGACTCTGACGATCTACGTGGTCTCC CTTCGGTTCAGACTATTCCGACAGCAGGGCTTAAGATCAGTTGTGACG TCTTGACAAGTTCAAGGATGATGAGGTAAGTTCAAGGAAAGTCTGAT CAACGATATCAGAGGCATGCTGTCCTTATGAAGCAGCCTATTGGCAA TCCGAGGAGAGGATCCTGGACGAGGCTATTGTGTTACAACCACCA CCTGAAGTCTGTCATCTCCATTCCGACCACTCGCATGCCAATTCCAACC TGGCTGAGCAGATCAGACACAGCTTGAGATTCCCTGCGAAAGGCC AGCCCGTCTGGAGGCTCGATACTTTGGACATCTACTCGCGAGACGAC

	CTGCATGACGAAACTCTGCTCAAATTGCCAAGCTGATTCAACATTCT GCAGGGCTGCTACCAGAAGGAGGCGTCATCATGACCCGATGGTGGAA CGACCTCGGCTCCCCAAGAAGGTTCCATATGCCAGAGACCGAACATC GAGACCTACATCTGGATGCTTCTGGTGTACGAGGCCAACCTCGC GTTGGCGAATCTCGCTCCAAGGTGGTGTATGATCACAAACCATAG ATGACACTTTGATGCATACGGTACCTCGAGGAGCTCACTGTTACT GAGGCCGTCACTAGGTGGACATTGGTCTCATCGACACCCTGCCGGAGT ACATGAAATTCAATTGTCAAGGCCTGTTGGATATCTACCGAGAGGCCGA GGAAGAGCTCGCAAAGGAGGGCCGCTCGTACGGAATCCCCTACGCTAA ACAGATGATGCAGGAGCTGATTATCCTACTTCACCGAAGCCAAGTGG CTGTACAAGGGCTACGTTCTACTTTGACGAGTATAAAAGCGTCGCACT ACGTTGATTGGCCTTCGAACCCCTGCCGCTCGTAGTTTGTGATCTAG GAGACTTATTGCCACCAAGGACAACCTCGAATGCATTCTCAAGAATGC TAAGTCGCTCAAGGCTACAGAAACCATTGGACGGCTCATGGACGACATT GCAGGTTACAAGTTGAGCAGAAACGGGCCACAACCCCTGCTGTC GAGTGCTACAAGAACCGAGCATGGAGTATCTGAAGAAGAAGCCGTAAA GAACCTTACTGGAAGTCGCCATTGAGGACATTAACGAGGAGC TGCTCAACCTACGACGGTACCTCTCCCCATGCTCCAACGACTCTGTAT TTGCTCGGTCAAGGTCAATTCTACGACGATGGTCACGACAGATAACAC CCATTCTCTAATGATGAAGCGACAAGTGGCTCTGCTGTTGACAGAGCCT CTTGCCATTG
<i>mcBFS</i>	ATGAGCACCATAACCGTCTCCTCTGTAAGTTTCTCGTCAAGCTCTCC GCTGGCCTTGATGACAAATTGAGTACCAAGCAGGACGTGGTGCACAC ACCATGAACCTCTCGGCAAGCATTGGGGAGACCAGTTTTAACTTATCA TGAGCCAGAGGATCTGGTGTCAAGAAGCAACTCGTCGAGCAGCTCAA GGAGGAGGTTAAGAAGGAACGTGATGACCATCAAGGGCTCCAACGAACC TATGCAGCATATCAAGCTTATGGAGCTGATTGATTGGTCCAGAGACTGG GTATTGCCTACCAACTTGAGGAGGAAATCGAGGAGGCCCTCCAACACAT CCACGTCACGTACGGAGAGCAGTGGATCGACAAGGAGAACCTGCAATC TGTGCTCTGTGGTCCGACTGCTCAGACAGCAGGGCTCAACGTGTCT TCTGGTGTGTTCAAGGACTACACCGATGAGAAGGGGGACTTCAAAGAG TCGTTGTGCAACGACGCACACGGCATCCTGGCTCTATGAAGCTGCTT ACATGCGAGTTGAGGGCGAGACCACTTGGACAAGGCCTCGAGTTCA CCAAGGTGCATCTCGACATCATCAGTAAGGACCCCTCTGTGACTTTAT CTGCGAACCGCAGTCCACCAGGCTCTAAACAACCTCTCCGCCGTCGG CTGGCAAGAACATCGAAGCGCTGCACTACATGCCGTCTACCAAGCAAGACT CCTCGCATAATGAGGTTCTGTTAAAGCTGCCAAACTGATTCTCCGTC CTACAGTCCATGCACAAGACGGAGCTCAGGCCACATTGCAAGTGGTGG AGGATCTCGATCTGCAGAACACAAGCTCCCTACGTGCGAGACAGAGTGGT CGAGGGCTACTCTGGATTCTGTCGATCTACTACGAGCCCCAGCGGGCC CGAACTCGAATGTTCTGATGAAGACTTGTATGTGGCTGGTGTCTGG ATGATACCTTGACAACATATGGAACATATGAGGAATTGGAGATTTTGCC CAGGCCGTTGAGCGGTGGTCCATAAGTGCATGGATATGCTGCCGAGT ACATGAAGCTGATTACCAAGGAGCTTGTGAATCTGCATGTGGACATGG

	AGAGTCCCTGAGAAAGGTGGCAAGACTTATCAGATCCAGTACGTCAAG GAAATGGCCAAGGAACTCGAAGGAACACTTGGTGGAGGCTCGATGG CTCAAGGAGGGATATGCCACACTAGAGGAGTACATGTCGGTTCTAT GGTACGGGAACTTACGGCTTGATGATTGCGCGCTGTACGTCGGTCTG GACGACATTGTGACAGAGGACACCTCAAGTGGGTTAGTTCTTACCTC CCATCATTAAGGCTTATGTGTCATTGTGAGATTGATGGACGATATTGTT CGCATCGTGAGGAACAGGAGCGAGGACACGTGGCTCGTATTGAGT GCTACTCCAAGAAAAGTGGTGCACCAGAAGAAGAACGCTGTGAATACA TCTCTCCAAGGTGAAAGACGCCTGGAAAGGTAATCAACCGAGAGTCACT TCGCCCACAGCCGTTCTTCTGCTTATGCCAGCTATTAAATTAGC ACGAATGTGTGAGGTGCTGTACTCTGTCAACGATGGACTCACTCATGCC GAGGGTACATGAAGTCCTACATGAAAAGCTACTTGTGCACCCCATGG TTATTGAA
<i>nadh-HMHR</i>	ATGACCGGCAAGACTGCCACATGACGGACTGAATTCTGAATCGAGA AGATGCGAGACCTGGACCCCCGCCCAAAGACTGGTGAGAGTGGCTGAAG CTGCTGGCCTGGAACCTGAAGCTATCTCTGCTCTGGCTGGCAATGGAGC TCTGCCCTGTCTCTGCTAATGGCATGATTGAAAATGTGATCGGCAAGT TCGAGCTGCCCTGGCGTTGCTACTAACCTTACTGTGAATGGCAGAGA TTACCTGATCCCCATGGCGTGGAAAGAGGCCCTCTGTGGTTGCTGCTGCTT CTTACATGGCCAGAACCGCCAGAGAGAACGGCGGCTTACTGCCCATGG AACTGCCCTCTGATGAGAGCTCAGATTCAAGTGGTGGACTGGCGAC CCCGAAGGAGCTAGACAAAGACTGCTGGCTCACAAAGCCGCTTATG GAAGCCGCGATGCTGTGGATCCTGTGCTGGTTGACTGGAGGAGGAT GTAGAGATATTGAGGTGCACGTGTTGAGACACCCCCGTGGAGCTAT GGTGGTGCTGCATCTTATTGTGGACGTGCGAGACGCCATGGCGCTAAT ACTGTGAATACCATGGCCGAGCGACTGGCCCCCGAACGTGGAACGAATTG CTGGCGGAACGTGCGACTGCGAATTCTGCTAATCTGCCGACCTGCG ACTGGTGCAGGCCAGACTGGATTAGCTCTGAAACCCCTGACTACTCAG GGCTATGACGGAGCTGACGTGGCTCGAGGAATGGTGGAGCTTGTGCC CTGGCTATTGTGGACCCCTATAGAGCCGCCACTCATAACAAGGGAATCAT GAACGGCATCGACCCCGTGGTGGCCACTGGAAACGATTGGAGAGC TATTGAGGCCGGAGCTCACGCCATGCCGCTAGAACTGGACACTATACTT CTCTGACCCGATGGAGCTGGCAACGACGGCAGACTGGTGGAACT TTGAGCTGCCCTGGCTTGGACTGGTGGAGGAGCTACTAAACTCA CCCTACCGCCCGAGCCGCCCTGGCTCTGATGCAAGTGGAAACCGCTACT GAGCTGGCCAAGTGACTGCCGCTGTGGACTGGCTAAATATGGCCG CTATCCGAGCCCTGGCTACCGAGGGAATTCAAGAGAGGCCATGACTCT GCACGCCGAAATATTGCCATCATGCCGGCGCTACTGGCGCCGATATTG ATAGAGTGACCAAGAGTGATCGTGGAGGCCGGAGACGTTCTGTGGCTAG AGCTAAACAAGTGCTGGAGAACACTTAA