

Supplementary Tables

Table S1: Primers used in the study.

Name		
Construction of ΔyeaJ		
Name	#	DNA Sequence (5'-3') ^a
YeaJ-TU	319	TATGGTAAACTAACGCATCCGACTTTTCAGGGCTGTAGTTCATC TAGACAGCTGCATGCAT
YeaJ-TD	320	CCATTTCTCACTCCTGGTGCAATATGAACATTTACTGCGAAGCAGTGTAGGCTGGAGCTG
YeaJ-F1U	321	GCGACAATAGTCACGACTGAA
YeaJ-F1D	322	CCAGCGCAGTACCACGTATT
YeaJ-F2U	348	CAGTCTATTAGAGGAATTCTCATCG
YeaJ-F2D	349	CGCGTTCTCCTGGGTTAATA
YeaJ-MU	350	CCGACTTTCAGGGCTGTAGTTCATCTGCTTCGCAGTAAATGTTTCATATTG
YeaJ-MD	351	CAATATGAACATTTACTGCGAAGCAGATGAACTACAGCCCTGAAAGTCGG
ThyA-F	395	ATGATGGTTTGGGCATATCG
ThyA-R	396	ACGAGCGCCAAATACAGATT
Construction of pYeaJ		
Name	#	DNA Sequence (5'-3') ^a
YeaJ-F- <i>Xba</i> I	364	GCTCTAGACAGTCTATTAGAGGAATTCTCATCG
YeaJ-R- <i>Hind</i> III	365	CCCAAGCTT CGCGTTCTCCTGGGTTAATA
pBAD-F	405	ATGCCATAGCATTTTTATCC
pBAD-R	406	GATTTAATCTGTATCAGG
Construction of pWYeaJ		
YeaJ-F-<i>Sma</i>I	495	CGTACCCGGGGCGACAATAGTCACGACTGAA
YeaJ-F-<i>Hind</i>III	365	CCCAAGCTT CGCGTTCTCCTGGGTTAATA
M13-20	M13	GTAAAACGACGGCCAGT
M13 Reverse	M13R	CATGGTCATAGCTGTT

^a Underlining indicates restriction endonuclease recognition sites