

S2 Table. Summary of mutations in thermoadapted mutants from *Zymomonas mobilis* CP4.

Locus tag	Gene	Classification	Product	Position	Reference	Alteration	Mutation	Amino acid change	Z4-80a	Z4-80b	Z4-80c	Z4-80d	Mutant
ZCP4_0028		Transcriptional regulation	signal transduction histidine kinase	32323	C	T	Transition	Met99Ile	+				
-			-	98768	G	GATGCCAGA	InDel						+
ZCP4_0125		Transporter	K+ transporter	142649	A	G	Transition	Thr403Ala		+			
ZCP4_0125		Transporter	K+ transporter	143231	C	T	Transition	Arg597Cys			+		
ZCP4_0567		Transcriptional regulation	DNA-directed RNA polymerase subunit beta	648575	G	A	Transition	Arg164Cys		+	+		
ZCP4_0588		General metabolism	formyltetrahydrofolate-dependent phosphoribosylglycinamide formyltransferase	670742	C	T	Transition	Leu118Phe				+	
ZCP4_0707		Transporter	ABC-type nitrate/sulfonate/bicarbonate transport system, ATPase component	799915	G	A	Transition	Pro438Leu		+			
-			-	1205885	T	TCAGGTCGC	InDel						+
-			-	1205931	T	TCGCAGGTG	InDel						+
-			-	1714424	TTATT	TT	InDel						+
-			-	1714426	A	AGGCT	InDel						+
-			-	1714428	TC	TCTC	InDel						+
ZCP4_1646		General metabolism	1-(5-phosphoribosyl)-5-((5-phosphoribosylamino)methylideneamino) imidazole-4-carboxamide isomerase	1856048	A	ATTATGACG	InDel	IYDGSL insertion					+
ZCP4_1702		Transporter	RND family efflux transporter, MFP1926213 subunit		G	A	Transition	Glu136Lys		+			
ZCP4_1702		Transporter	RND family efflux transporter, MFP1926238 subunit		C	T	Transition	Thr144Met			+		
ZCP4_1703		Transporter	efflux transporter, outer membrane factor lipoprotein, NodT family	1928285	A	G	Transition	Lys462Glu				+	
ZCP4_1739		Membrane stabilization	sphingosine/diacylglycerol kinase-like1966395 enzyme		C	T	Transition	Ala89Val		+			
ZCP4_1739		Membrane stabilization	sphingosine/diacylglycerol kinase-like1966601 enzyme		C	T	Transition	Leu158Phe			+		
ZCP4_1739		Membrane stabilization	sphingosine/diacylglycerol kinase-like1966796 enzyme		G	A	Transition	Val223Met				+	