

6-Phosphofructokinase Genes				
Organism	Locus Tag	AA #104	AA #124	ATP or PPi Dependence
<i>Bacillus cereus</i> ATCC 14579	BC4600	G	G	ATP
<i>Caldanaerobacter subterraneus</i> subsp. <i>tengcongensis</i> MB4	TTE1816	G	G	ATP
<i>Ethanoligenens harbinense</i> YUAN-3T, DSM 18485	Ethha_1347	G	G	ATP
<i>Geobacillus thermoglucosidasius</i> C56-YS93	Geoth_0897	G	G	ATP
<i>P. furiosus</i> DSM 3638	PF0312	S	E	ATP
	PF1784	L	F	ATP
	TK0376	L	F	ATP
<i>Thermococcus kodakaraensis</i> KOD1	TK1110	S	E	ATP
<i>Thermotoga maritima</i> MSB8	TM0209	G	A	ATP
	TM0289	D	K	PPi
<i>Thermotoga neapolitana</i> DSM 4359	CTN_0395	D	K	PPi
	CTN_0476	G	A	ATP
<i>Thermotoga petrophila</i> RKU-1	Tpet_0623	D	K	PPi
	Tpet_0715	G	A	ATP
				ATP
<i>Caldicellulosiruptor bescii</i> DSM 6725	Athe_1265	G	G	
	Athe_1824	D	K	PPi
				ATP
<i>Caldicellulosiruptor saccharolyticus</i> DSM 8903	Csac_1830	G	G	
	Csac_2366	D	K	PPi
<i>Clostridium cellulolyticum</i> H10	Ccel_2223	D	K	PPi
	Ccel_2612	G	G	ATP
<i>Clostridium phytofermentans</i> ISDg	Cphy_0336	G	G	ATP
	Cphy_3345	D	K	PPi
<i>Clostridium thermocellum</i> ATCC 27405	Cthe_0347	D	K	PPi
	Cthe_1261	G	G	ATP
<i>Clostridium thermocellum</i> DSM 4150	CtherDRAFT_1670	G	G	ATP
<i>Thermoanaerobacter pseudethanolicus</i> 39E	Teth39_0494	D	K	PPi
	Teth39_0683	G	G	ATP