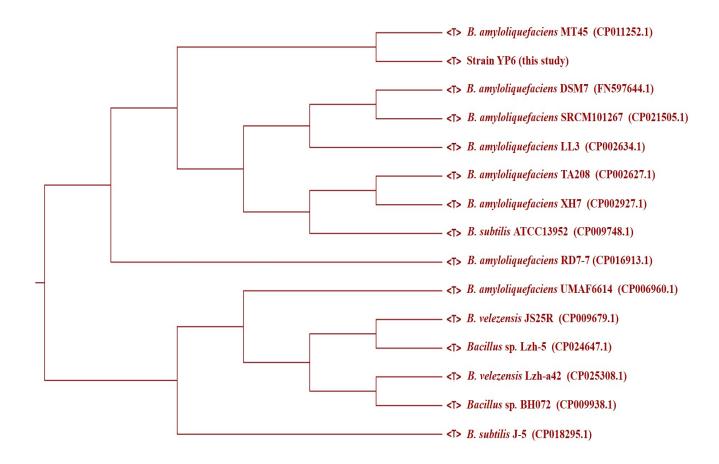


**Appendix A.** The plant-growth-promoting activity and OPs-degrading capability of strain YP6. (a) The solubilizing-phosphate effect of strains; (b) The contents of indole-3-acetic acid (IAA) and siderophores by strain YP6; (c) Degradation of various types of OPs by strain YP6. The average values and standard error were calculated from three independent replicates. The conditions of culture for (a) and (b): YP6 was incubated of 1-5 d at 30 °C, 200 r/min in LB medium (containing 10 g/L rock phosphate powder). The conditions of culture for (c): YP6 was incubated of 48 h at 30 °C, 200 r/min in LB medium with 6 types of OPs with initial concentration of 50 mg/L, respectively.



Appendix B. Phylogenetic tree from the genomes of strain YP6 and other Bacillus strains using CVTree web tool. (K=6)

Appendix C. The partial genes involved in solubilizing-phosphorus, OPs-degradation, IAA and siderophores synthesis in B. amyloliquefaciens YP6

Capabilities	Description	Gene	Strand	Start	End
		name			
Solubilizing-phosphorus/OPs	alkaline phosphatase	phoD	+	256,152	257,903
degradation		phoA	-	922,335	920,956
	phosphatase	phoE	-	1,007,871	1,007,296
		ycsE	-	1,073,123	1,072,311
		bcrC	-	2,069,264	2,068,653
	carboxylesterase	yvaK	-	3,284,370	3,283,624
Producing indole-3-acetic	tryptophan synthase subunit alpha	trpA	-	2,313,324	2,312,527
acid (IAA)	tryptophan synthase subunit beta	$trp\mathbf{B}$	-	2,314,519	2,313,317
	indole-3-glycerol-phosphate synthase	$trp\mathbb{C}$	-	2,315,910	2,315,158
	anthranilate phosphoribosyltransferase	$trp\mathbf{D}$	-	2,316,919	2,315,903
	anthranilate synthase component I	<i>trp</i> E	-	2,318,438	2,316,891
	acetyltransferase	cgeE	-	2,080,138	2,079,359
		epsM	-	3,345,785	3,345,138
Producing siderophores	isochorismatase	entB	-	3,110,569	3,109,649
	isochorismate synthase	menF	-	3,000,480	2,999,068
		entC	-	3,113,427	3,112,231
	2,3-dihydro-2,3-dihydroxybenzoate	entA	-	3,114,235	3,113,450
	dehydrogenase				