Table.S1 The primers used in this study.

Primer name	Primer sequence (5'-3')
capA-up	GGGGTACCCTGACATTTTCGGTGCGAG
<i>capA</i> -down	GCTCTAGATTCATCCTCCCGGATAAGGC
capB-up	GGGGTACCTCGACTCCCAGCTGG
capB-down	GCTCTAGACGCAGCTCTGCTGTC
<i>capJ</i> -up	GGGGTACCAGTCGTGCGGCTTCG
<i>capJ</i> -down	GCTCTAGAGACCACGCTGATGCC
regl-up	GGGGTACCGATGGTCTGAAAGAAGT
<i>regl-</i> down	GCTCTAGAAACGATGCTGATTGC
<i>regII-</i> up	GGGGTACCTGACCAGACCAGC
regII-down	GCTCTAGAGCGGCGACGGTTA
regIII-up	GGGGTACCTGGATGCGTGGAT
regIII-down	GCTCTAGAGCCATACCGGACT
Fosmid-T7	TAATACGACTCACTATAGGG
Fosmid-RP	CTCGTATGTTGTGGAATTGTGAGC

Table.S2 The isoelectric points of all the overexpressed genes.

Gene name	Isoelectric point (pI)	
сарВ	5.72	
capC	4.76	
capD	5.52	
capE	5.29	
capFi	4.55	
capFii	4.77	
capV	4.98	
capG	5.05	
сарТ	4.35	

Fig.S1

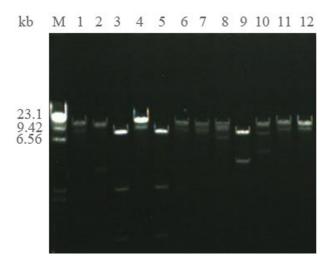


Fig.S1 Agarose electrophoresis of fosmid plasmid after BamHI digestion. Bands 1-12 were plasmids for randomly selected strains. All of the lanes 1-12 contains a vector strip at about 8kb location, the size of other strips are different.

Fig.S2

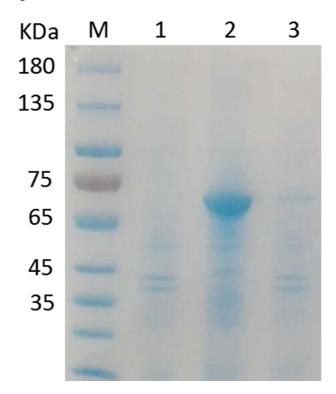


Fig.S2 The SDS-PAGE of the CapB protein overexpressed strains. 1: negative control strains BL21-pUC19-GFP(empty vector) 2: *capB* was constructed into the vector pET30a as a positive control (T7 as promoter) 3: *capB* was constructed into the vector pUC19 using its own promoter. In the position of 70KDa, compared with negative control, *capB* gene expression quantity increased significantly of at lane 2 and 3.