

Table 3. Database identities with ORFs in type 2 locus

ORF	Blastx hit	Accession no.	Organism	Expect value	Reference
<i>cpsA</i>	Undecaprenyl pyrophosphate synthetase	O31751	<i>B. subtilis</i>	2e-78	Unpublished data
<i>cpsB</i>	Phosphatidate cytidylyltransferase	O31752	<i>B. subtilis</i>	6e-46	Unpublished data
<i>cpsC</i>	Glycerophosphotransferase Teichoic acid synthesis (TagF)	P13485	<i>B. subtilis</i>	e-100	1
<i>cpsD</i>	Glycosyl transferase (Cap33fJ)	CAA07403	<i>S. pneumoniae</i>	3e-12	2
<i>cpsE</i>	Putative sugar transferase	CAB73862	<i>C. jejuni</i>	2e-34	3
<i>cpsF</i>	No good match				
<i>cpsG</i>	UDP- <i>N</i> -acetylenolpyruvoyl glucosamine reductase (MurB)	P18579	<i>B. subtilis</i>	2e-40	4
<i>cpsH</i>	No good match				
<i>cpsI</i>	UDP-galactopyranose mutase (cap33fN)	CAA07407	<i>S. pneumoniae</i>	e-137	2
<i>cpsJ</i>	Teichoic acid translocation ATP-binding protein (TagH)	P42954	<i>B. subtilis</i>	2e-80	5
<i>cpsK</i>	Teichoic acid translocation permease protein (TagG)	P42953	<i>B. subtilis</i>	3e-60	5

References:

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