

CcdB_F	MQFKVYTYK-RESRYRLFVDVQSDIIDTPGRRMVIPLASARLLSDKV-SRELYPVV-HIG	57
CcdB_pSLT	MQFKVYTCK-RESRYRLFVDVQSDIIDTPGRRMAVPLVSARLLSEKV-PRDLYPVM-HIG	57
CcdB_CFT073	MQFTVYRSRGRNAAFPFDVTSDIIGEINRRIVIPLTPIERFSRIRPPERLNPIILLVD	60
CcdB-1	MQFTVYRSRGRNAAFPFDVTSDIIGEINRRIVIPLTPIERFSRIRPPERLNPIILLVD	60
	***.** : *:: : :.* ***. .**.:** . :* . * *:: :.	
CcdB_F	DESWRMMTTQMVSVPVSVIGEEVADLSHRENDIKNAINLMF <sup>W</sup> GI	101
CcdB_pSLT	DEPYRLLTTDMTSVPATVIGEEVADLSLRENDIKNAINLMF <sup>R</sup> GI	101
CcdB_CFT073	GKEYVLMTHETATVPVNALGTKFCDA <sup>S</sup> AHRTLIK <sup>S</sup> ALDFML <sup>L</sup> GI	104
CcdB-1	GKEYVLMTHETATVPVNALGTKFCDA <sup>S</sup> AHRTLIK <sup>S</sup> ALDFML <sup>L</sup> GI	104
	.: : :.* : .:**...* :..* * :.. **.***:*. **	

**Figure S4.** Multiple amino acid sequence alignment of CcdB proteins. CcdB proteins encoded by F plasmid (GenBank AAA96062), pSLP plasmid (WP\_001159863), UPEC CFT073 (WP\_001160966) and NRG857c (YP\_006118417) were alignment by CLUSTAL O(1.2.4). Identical amino acid residues are highlighted in black. Residue W99, reported to be crucial for CcdB toxicity in *E. coli*, along with its variation, are highlighted in yellow.