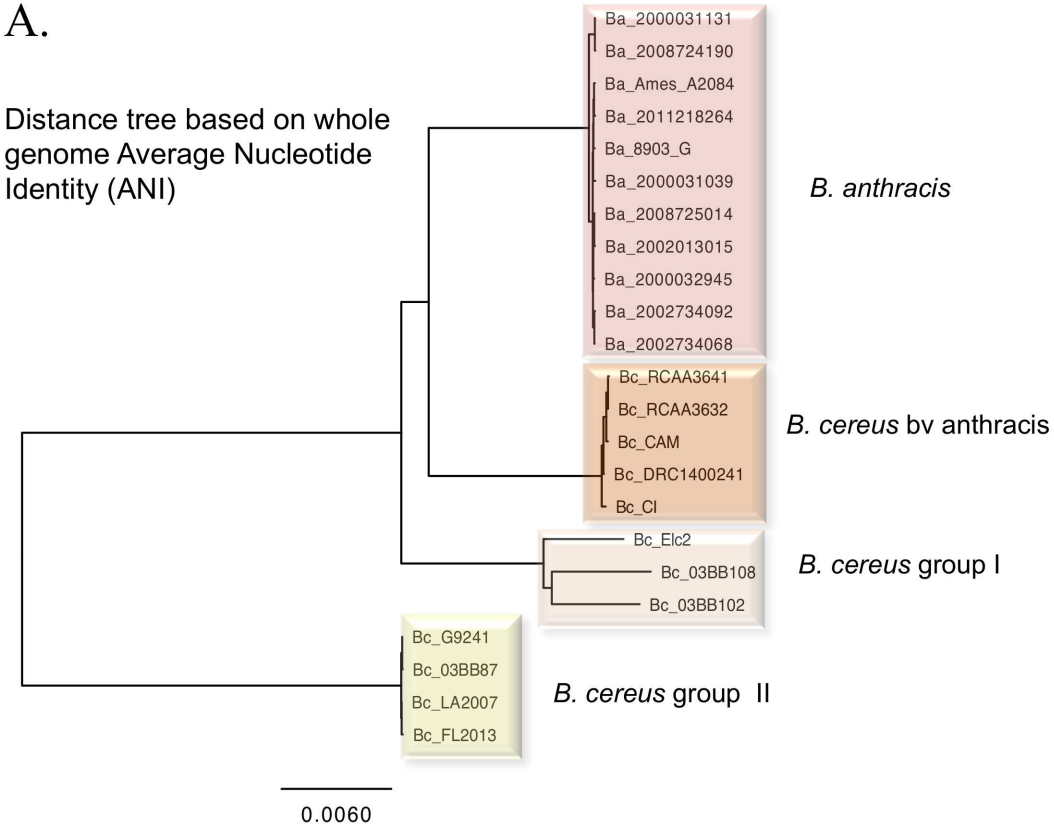
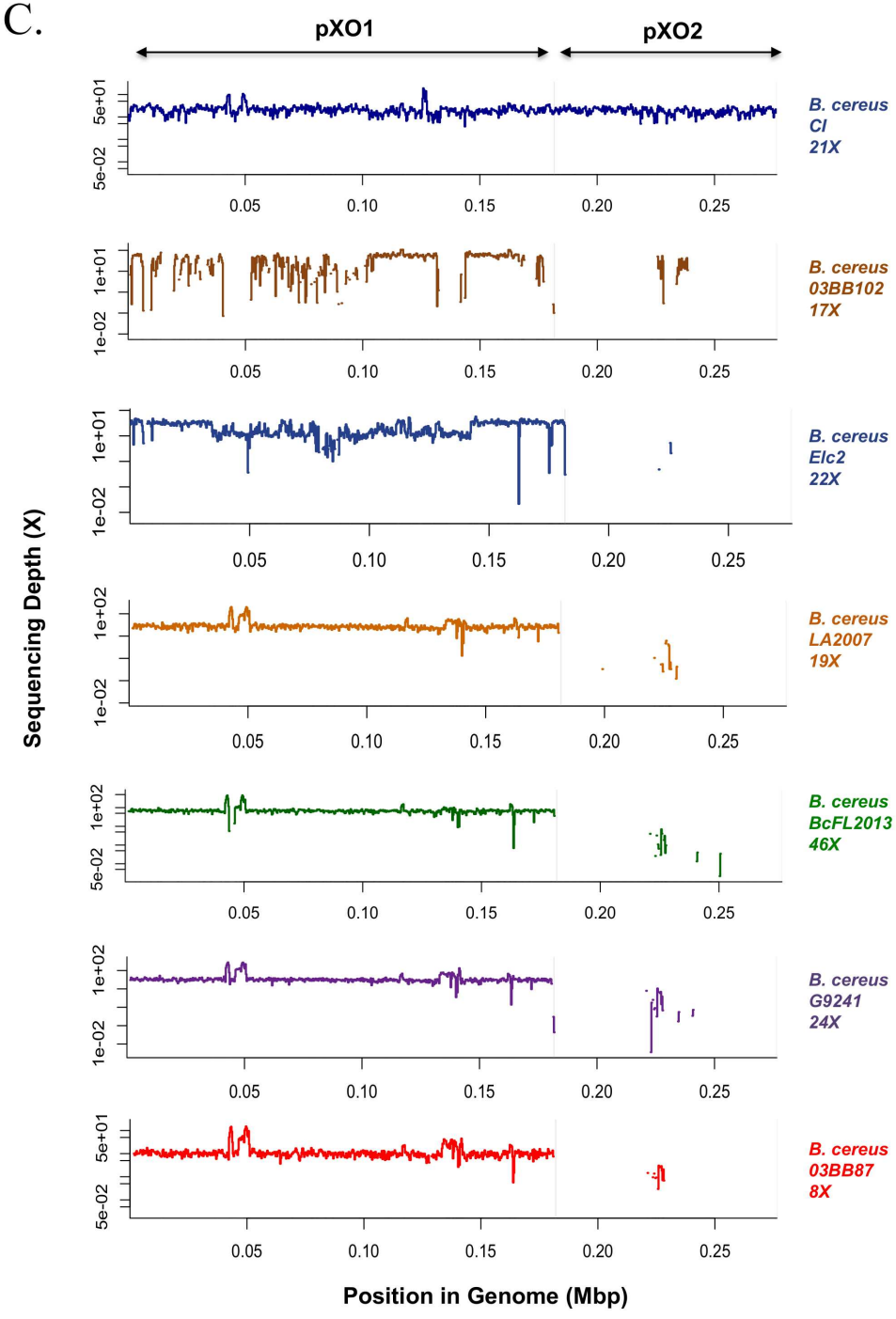
A.

Distance tree based on whole genome Average Nucleotide Identity (ANI)



B.

100-ANI	*B. anthracis*	*B. cereus* bv anthracis	*B. cereus* group I	*B. cereus* group II
B. anthracis	0.04	-	-	-
***B. cereus* bv anthracis**	1.89	0.03	-	-
***B. cereus* group I**	2.36	2.46	1.01	-
***B. cereus* group II**	5.24	5.09	5.48	0.01

C.