These alternative primers were designed by Emily Jackson based on a more comprehensive range of *Cronobacter* species strains which had been whole genome sequenced, than the original set by Baldwin.

They are particularly useful for *glnS* sequencing, which can be problematic due to the polymerase enzyme skipping a 'G' in one direction. This can cause confusion.

For PCR amplification For sequencing

glnS glnS

glnS-F GCATCTACCCGATGTACG glnS-F-S GGGTGCTGGATAACATCA

glnS-R-NEW TGTCACCGTTTGCATGGC glnS-R-S-NEW CAGACGCTTGTACTGCTT

gltB

gltB-F-NEW CGTGATGCTGAACTACCG

All sequences are 5' --> 3'

gltB-R CAGCACTTCCACCAGCTC

gyrB

gyrB-F TGCACCACATGGTATTCG

gyrB-R-NEW TGATCATCTTTCTGCGCGC

ppsA

ppsA-F GTCCAACAATGGCTCGTC

ppsA-R-NEW CATCATCTCGCGGATCTG