

Draft genome sequence of *Flavobacterium sp.* strain PL002, isolated from Antarctic *Porphyra* algae

ABSTRACT

Here, we report the draft genome sequence of *Flavobacterium sp.* strain PL002, isolated from Antarctic *Porphyra algae*. The 4,299,965-bp genome sequence is assembled into 170 contigs, has 32.92% GC content, and 3,734 predicted genes.