

Genomic data of two *Bacillus* and two *Pseudomonas* strains isolated from the acid mine drainage site at Mamut Copper Mine, Ranau, Malaysia

ABSTRACT

The genomic data of four bacteria strains isolated from the abandoned Mamut Copper Mine, an Acid Mine Drainage (AMD) site is presented in this report. Two of these strains belong to the genus *Bacillus*, while the other two belong to the genus *Pseudomonas*. The draft genome size of *Pseudomonas* sp. strain MCMY3 was 6,396,595 bp (GC: 63.3%), *Bacillus* sp. strain MCMY6 was 6,815,573 bp (GC: 35.2%), *Bacillus* sp. strain MCMY13 was 5,559,059 bp (GC: 35.5%) and *Pseudomonas* sp. strain MCMY15 was 7,381,777 bp (GC: 64.8%). These four genomes contained 493, 495, 495 and 579 annotated subsystems, respectively. The sequence data are available at GenBank sequence read archive with accession numbers SRX7859406, SRX7859404, SRX7859405 and SRX7293032 for strains MCMY3, MCMY6, MCMY13 and MCMY15, respectively.