CORRECTION



Correction to: Phylogenetics and antibacterial properties of exopolysaccharides from marine bacteria isolated from Mauritius seawater

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In the above-mentioned paper, the wrong figure was provided for Fig. 6. The correct one should be:

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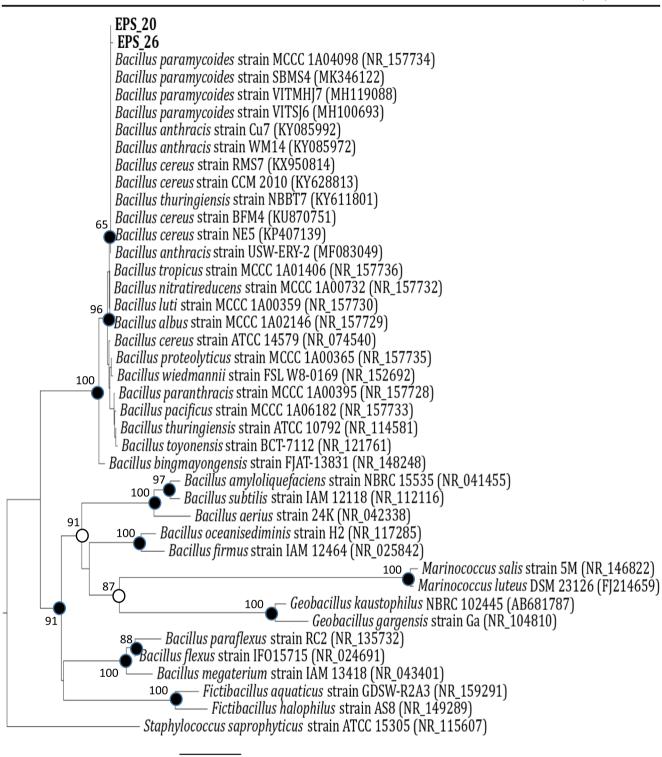


Fig. 6 Neighbour-joining tree based on 16S rRNA sequences showing the relationship of isolates EPS-20 and EPS-26 with other strains from the *Bacillacceae* family. Bootstrap values (expressed as percentages of 1000 replications) ≥50 are shown at branch nodes. \blacksquare represent nodes obtained for all phylogenetic trees while \blacksquare represent nodes obtained

0.02

only on two of the tree topologies. Number in parentheses indicate GenBank nucleotide accession numbers. *Staphylococcus saprophyticus* ATCC 15305 was chosen as outgroup. Bar represents 0.02 substitution per nucleotide position

