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Isolation and characterization of highly active keratinolytic microorganisms with promising potential for waste sheep wool processing

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Authors:

Blaž Petek, Maja Vodušek, Tomaž Accetto, Maša Zorec, Polona Zalar, Jernej Oberčkal, Romana

Marinšek Logar*

Affiliation of corresponding author

University of Ljubljana, Biotechnical faculty, Department of Microbiology, Jamnikarjeva 101, 1000 Ljubljana, Slovenia, romana.marinsek@bf.uni-lj.si

Heatmap diagram of Average nucleotide identity metric of isolated strains versus strains of other species which appeared most related upon preliminary analysis using the 16SrRNA gene sequence. 95% (white and red colour) is considered to be the species boundary. A: RS3014; B: S3021; C: RS308; D: S3027; E: S3018.

