Genome Sequence of *Stenotrophomonas maltophilia* PML168, Which Displays Baeyer-Villiger Monooxygenase Activity


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*Stenotrophomonas maltophilia* PML168 was isolated from Wembury Beach on the English Coast from a rock pool following growth and selection on agar plates. Here we present the permanent draft genome sequence, which has allowed prediction of function for several genes encoding enzymes relevant to industrial biotechnology, including a novel flavoprotein monooxygenase.

The biotechnological application of this enzyme is currently being evaluated.

**Nucleotide sequence accession numbers.** The nucleotide sequence for the draft genome sequence was deposited in EMBL under accession numbers CAJH01000001 to CAJH01000097.

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**REFERENCES**
