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A prospective study on the biodegradation of petroleum hydrocarbons mediated by selected marine bacterial isolates

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Abstract

Petroleum hydrocarbons are an indispensable energy source and serve as a fundamental feedstock for petroleum oil refiners and petrochemical industries. The release of petroleum hydrocarbon pollutants in the environment is associated with anthropogenic activities during crude oil production, exploration, and transportation. The widespread petroleum hydrocarbons pollutants in the environment is associated with anthropogenic activities during crude oil production, exploration, and transportation. The widespread petroleum hydrocarbons pollutants in the environment is associated with anthropogenic activities during crude oil production, exploration, and transportation. The widespread petroleum hydrocarbons pollutants in the environment with petroleum hydrocarbons as a carbon source for their growth and metabolic activities. Therefore, this study focused on the isolation, molecular identification and characterization of bacterial isolates from seawater samples collected at the coastal area nearby oil Refinery Company in Oman. Six marine bacteria were isolated from the contaminated seawater and their taxonomical identification was performed based on 16S rRNAgene sequence. The degradation potential of bacterial isolates was assessed by growing them in F/2 medium supplemented with crude oil as a sole carbon source. Two bacterial isolates belonging to the genus Niallia showed the strongest growth (S1.5: 1.01 ± 0.27 and S2.1: 0.8 ± 0.23) and were subjected to further Gas Chromatography–Mass Spectrometry (GC-MS) analysis determining which n-alkanes in crude oil were degraded by the isolates. The G-C-MS chromatogram indicated that S2.1 exhibited greater efficiency in degrading a higher number of carbon n-alkanes (C13-C34) com-pared to S1.5 (C13-C18). The t-test analysis confirmed the efficiency of S2.1 (t = 3.182; p < 0.01) (93.61 ± 1.48 %) compared to S1.5 (86.08 ± 1.85 %). Molecular docking revealed robust binding affinities between selected n-alkanes and alkane hydroxylases, facilitating the subsequent degradation poten

Author Keywords

Biodegradation; GC-MS; Hydrocarbon-degrading bacteria; Marine ecosystem; N-alkanes; Niallia

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