Kumar RR., Jadeja VJ., *BioImpacts*, 2018, 8(4), 253-261 doi: 10.15171/bi.2018.28 http://bi.tbzmed.ac.ir/

Characterization and partial purification of an antibacterial agent from halophilic actinomycetes *Kocuria* sp. strain rsk4

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Isolation of halophilic actinomycetes



Figure S1: Isolate rsk4

Antibacterial potential of strain rsk4



Figure S2: Antibiotic susceptibility test of Staphylococcus aureus



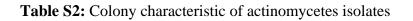
Figure S3: Antibacterial assay of rsk by agar well diffusion method

Taxonomic characterization of the organism

Table S1: A	ntibiotic	resistance	tests	of s	strain	rsk4
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Characteristic	Strain rsk4
Growth in the presence of antibiotics	
Ampicillin	+
Ciprofloxacin	-
Colistin	-
Co-timoxazole	-
Gentamicin	+
Nitrofurantoin	-
Streptomycin	-
Tetracycline	-
Cephalothin	-
Clindamycin	-
Erythromycin	-
Ofloxacin	+
Penicillin-G	+
Vancomycin	+

"+" growth, "-" no growth



Medium	Size	Shape	Elevation	Opacity	Texture	Margin	Colour
ISP2	Small	Circular	Raised	Opaque	Moist	Smooth	Orange
ISP4	Small	Circular	Raised	Opaque	Moist	Smooth	Orange
ISP5	Small	Circular	Raised	Opaque	Moist	Smooth	Orange
ISP7	Small	Circular	Raised	Opaque	Moist	Smooth	Orange
Nutrient agar	Small	Circular	Raised	Opaque	Moist	Smooth	Reddish orange
tryptone soya agar	Small	Circular	Raised	Opaque	Moist	Smooth	Reddish orange
starch casein	Small	Circular	Raised	Opaque	Moist	Smooth	Reddish orange

Table S3: Sequence alignment of strain rsk4

Select: <u>All None</u> Selected:0 X Alignments BOwnload <u>GenBank</u> Graphics Distance tree of results						0
Description	Max score	Total score	Query cover	E value	ldent	Accession
Kocuria palustris strain TAGA27 16S ribosomal RNA gene, partial sequence	2078	2078	99%	0.0	94%	<u>NR 026451.1</u>
Kocuria assamensis strain S9-65 16S ribosomal RNA, partial sequence	2061	2061	99%	0.0	94%	<u>NR 132604.1</u>
Kocuria atrinae strain P30 16S ribosomal RNA gene, partial seguence	1897	1897	99%	0.0	92%	<u>NR 116744.1</u>
Kocuria rosea strain DSM 20447 16S ribosomal RNA gene, partial sequence	1890	1890	99%	0.0	92%	<u>NR 044871.1</u>
Kocuria sediminis strain FCS-11 16S ribosomal RNA gene, partial sequence	1873	1873	99%	0.0	92%	<u>NR 118222.1</u>
Kocuria rhizophila strain TA68 16S ribosomal RNA gene, partial sequence	1871	1871	99%	0.0	92%	<u>NR 026452.1</u>
Kocuria flava strain HO-9041 16S ribosomal RNA gene, partial seguence	1869	1869	99%	0.0	92%	<u>NR 044308.1</u>
Kocuria turfanensis strain HO-9042 16S ribosomal RNA gene, partial seguence	1868	1868	99%	0.0	92%	<u>NR 043899.1</u>
Kocuria dechangensis strain NEAU-ST5-33 16S ribosomal RNA, partial sequence	1866	1866	99%	0.0	92%	<u>NR 137239.1</u>
Kocuria polaris strain CMS 76or 16S ribosomal RNA gene, partial sequence	1862	1862	99%	0.0	92%	<u>NR 028924.1</u>
Kocuria gwangalliensis strain SJ2 16S ribosomal RNA gene, partial sequence	1860	1860	99%	0.0	92%	<u>NR 116266.1</u>
Kocuria carniphila strain CCM 132 16S ribosomal RNA gene, partial sequence	1860	1860	99%	0.0	92%	<u>NR 027193.1</u>
Kocuria salsicia strain 104 16S ribosomal RNA gene, partial sequence	1855	1855	99%	0.0	91%	<u>NR 117299.1</u>
Kocuria varians strain G33 16S ribosomal RNA gene, partial sequence	1855	1855	99%	0.0	91%	<u>NR 029297.1</u>
🗌 Kocuria aegyptia strain YIM 70003 16S ribosomal RNA gene, partial seguence	1853	1853	99%	0.0	91%	<u>NR 043511.1</u>
Kocuria subflava strain YIM 13062 16S ribosomal RNA, partial sequence	1844	1844	99%	0.0	91%	<u>NR 144586.</u>
Kocuria arsenalis strain CM1E1 16S ribosomal RNA, partial sequence	1842	1842	98%	0.0	92%	<u>NR 148610.1</u>
Kocuria varians strain ATCC 15306 16S ribosomal RNA gene, partial sequence	1836	1836	96%	0		
Kocuria halotolerans strain YIM 90716 16S ribosomal RNA, complete sequence	1834	1834	99%	0	∃ Que	estions co