



**Strain identifier**

**BacDive ID:** 11142      **DOI:** 10.13145/bacdive11142.20190402.4  
**Type strain:** yes      **Designation:** MSL-09  
**Culture col. no.:** DSM 19266, KCTC 19272

**Sections**

- [Name and taxonomic classification](#)
- [Morphology and physiology](#)
- [Culture and growth conditions](#)
- [Isolation, sampling and environmental information](#)
- [Application and interaction](#)
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**Name and taxonomic classification**

<a href="#">Ref.: 7985</a>	<b>Domain</b>	Bacteria
<a href="#">Ref.: 7985</a>	<b>Phylum</b>	Actinobacteria
<a href="#">Ref.: 7985</a>	<b>Class</b>	Actinobacteria
<a href="#">Ref.: 7985</a>	<b>Order</b>	Actinomycetales
<a href="#">Ref.: 7985</a>	<b>Family</b>	Nocardioidaceae
<a href="#">Ref.: 7985</a>	<b>Genus</b>	Nocardioides
<a href="#">Ref.: 7985</a>	<b>Species</b>	Nocardioides koreensis
<a href="#">Ref.: 7985</a>	<b>Full Scientific Name</b>	Nocardioides koreensis Dastager et al. 2008
<a href="#">Ref.: 7985</a>	<b>Designation:</b>	MSL-09
<a href="#">Ref.: 7985</a>	<b>Type strain:</b>	yes

**Prokaryotic Nomenclature Up-to-date (PNU)**

<a href="#">Ref.: 20215</a>	<b>Domain</b>	Bacteria
<a href="#">Ref.: 20215</a>	<b>Phylum</b>	Actinobacteria
<a href="#">Ref.: 20215</a>	<b>Class</b>	Actinobacteria
<a href="#">Ref.: 20215</a>	Literature reference	Int. J. Syst. Bacteriol. 47:483*
<a href="#">Ref.: 20215</a>	<b>Family</b>	Nocardioidaceae
<a href="#">Ref.: 20215</a>	<b>Genus</b>	Nocardioides
<a href="#">Ref.: 20215</a>	Taxonomical status	genus (AL)



Ref.: 20215	Literature reference	Int. J. Syst. Bacteriol. 30:225
Ref.: 20215	<b>Species</b>	Nocardioides koreensis
Ref.: 20215	Taxonomical status	sp. nov. (VP)
Ref.: 20215	Literature reference	Int. J. Syst. Evol. Microbiol. 58:2292*
Ref.: 20215	<b>Full Scientific Name</b>	Nocardioides koreensis Dastager et al. 2008

**Morphology and physiology**

Ref.: 32484	<b>Gram stain</b>	positive
Ref.: 32484	<b>Cell length</b>	0.8-3.2 µm
Ref.: 32484	<b>Cell width</b>	0.2-0.7 µm
Ref.: 32484	<b>Cell shape</b>	rod-shaped
Ref.: 32484	<b>Motility</b>	yes
Ref.: 19779	<b>Cultivation medium used</b>	ISP 2
Ref.: 19779	<b>Colony color</b>	Light ivory (1015)
Ref.: 19779	<b>Incubation period</b>	10-14 days
Ref.: 19779	<b>Cultivation medium used</b>	ISP 3
Ref.: 19779	<b>Colony color</b>	Light ivory (1015)
Ref.: 19779	<b>Incubation period</b>	10-14 days
Ref.: 19779	<b>Cultivation medium used</b>	ISP 4
Ref.: 19779	<b>Colony color</b>	Light ivory (1015)
Ref.: 19779	<b>Incubation period</b>	10-14 days
Ref.: 19779	<b>Cultivation medium used</b>	ISP 5
Ref.: 19779	<b>Incubation period</b>	10-14 days
Ref.: 19779	<b>Cultivation medium used</b>	ISP 6
Ref.: 19779	<b>Colony color</b>	Light ivory (1015)
Ref.: 19779	<b>Incubation period</b>	10-14 days
Ref.: 19779	<b>Cultivation medium used</b>	ISP 7



Ref.: 19779

**Incubation period** 10-14 days

Ref.: 32484

Ref.: 32484

<b>Halophily</b>	<b>Salt</b>	<b>Tested relation</b>	<b>Salt conc.</b>
	NaCl	growth	0-5 %
	NaCl	optimum	0-5 %

Ref.: 19779

**API coryne**

API ID	317
NIT	+
PYZ	-
PYRA	-
PAL	+
betaGUR	-
betaGAL	+
alphaGLU	-
betaNAG	-
ESC	-
URE	+
GEL	-
Control	n.d.
GLU	+
RIB	-
XYL	-
MAN	-
MAL	+
LAC	+
SAC	+
GLYG	-
CAT	n.d.

Ref.: 19779

**API zym**

API ID	603
Control	n.d.
Alkaline phosphatase	+
Esterase (C 4)	+
Esterase Lipase (C 8)	+
Lipase (C 14)	-
Leucine arylamidase	-

Valine arylamidase	-
Cystine arylamidase	-
Trypsin	-
Alpha-chymotrypsin	-
Acid phosphatase	+
Naphthol-AS-BI-phosphohydrolase	+
Alpha-galactosidase	-
Beta-galactosidase	-
Beta-glucuronidase	-
Alpha-glucosidase	+
Beta-glucosidase	-
N-acetyl-beta-glucosaminidase	-
Alpha-mannosidase	-
Alpha-fucosidase	-

	<b>Metabolite utilization</b>	<b>Chebi ID</b>	<b>Metabolite</b>	<b>Utilization activity</b>	<b>Kind of utilization tested</b>
<a href="#">Ref.: 32484</a>		17716	Lactose	+	carbon source
<a href="#">Ref.: 32484</a>		37684	Mannose	+	carbon source
<a href="#">Ref.: 32484</a>		28053	Melibiose	+	carbon source
<a href="#">Ref.: 19779</a>	<b>Medium Name (multicellularity)</b>		ISP 2		
<a href="#">Ref.: 19779</a>	<b>Multicellular complex forming ability</b>		no		
<a href="#">Ref.: 19779</a>	<b>Medium Name (multicellularity)</b>		ISP 3		
<a href="#">Ref.: 19779</a>	<b>Multicellular complex forming ability</b>		no		
<a href="#">Ref.: 19779</a>	<b>Medium Name (multicellularity)</b>		ISP 4		
<a href="#">Ref.: 19779</a>	<b>Multicellular complex forming ability</b>		no		
<a href="#">Ref.: 19779</a>	<b>Medium Name (multicellularity)</b>		ISP 5		
<a href="#">Ref.: 19779</a>	<b>Multicellular complex forming ability</b>		no		



Ref.: 19779	<b>Medium Name (multicellularity)</b>	ISP 6
Ref.: 19779	<b>Multicellular complex forming ability</b>	no
Ref.: 19779	<b>Medium Name (multicellularity)</b>	ISP 7
Ref.: 19779	<b>Multicellular complex forming ability</b>	no
Ref.: 32484	<b>Decomposition/lysis</b>	aggregates in chains
Ref.: 32484	<b>Oxygen tolerance</b>	aerobe
Ref.: 32484	<b>Ability of spore formation</b>	no

**Culture and growth conditions**

Ref.: 7985	<b>Culture medium</b>	R2A MEDIUM (DSMZ Medium 830), 28°C
Ref.: 7985	<b>Culture medium growth</b>	yes
Ref.: 7985	<b>Culture medium link</b>	<a href="https://www.dsmz.de/microorganisms/medium/pdf/DSMZ_Medium830.pdf">https://www.dsmz.de/microorganisms/medium/pdf/DSMZ_Medium830.pdf</a>
Ref.: 19779	<b>Culture medium</b>	ISP 2
Ref.: 19779	<b>Culture medium growth</b>	yes
Ref.: 19779	<b>Culture medium composition</b>	Name: ISP 2 / Yeast Malt Agar (5265); 5265 Composition Malt extract 10.0 g/l Yeast extract 4.0 g/l Glucose 4.0 g/l Agar 15.0 g/l Preparation: Sterilisation: 20 minutes at 121°C pH before sterilisation: 7.0 Usage: Maintenance and Taxonomy Organisms: All Actinomycetes
Ref.: 19779	<b>Culture medium</b>	ISP 3
Ref.: 19779	<b>Culture medium growth</b>	yes

Ref.: 19779	<b>Culture medium composition</b>	Name: ISP 3; 5315 Composition Dog oat flakes 20.0 g/l Trace element solution (5314) 2.5 ml/l Agar 18.0 g/l Preparation: Oat flakes are cooked for 20 minutes, trace element solution and agar are added (in the case of non rolled oat flakes the suspension has to be filtered). Sterilisation: 20 minutes at 121°C pH before sterilisation: 7.8 Usage: Maintenance and taxonomy (e.g. SEM As liquid medium for metabolite production) Organisms: All Actinomycetes Trace element solution 5314 Name: Trace element solution 5314; 5314 Composition CaCl <sub>2</sub> x H <sub>2</sub> O 3.0 g/l Fe-III-citrate 1.0 g/l MnSO <sub>4</sub> 0.2 g/l ZnCl <sub>2</sub> 0.1 g/l CuSO <sub>4</sub> x 5 H <sub>2</sub> O 0.025 g/l Sodium tetra borate 0.2 g/l CoCl <sub>2</sub> x 6 H <sub>2</sub> O 0.004 g/l Sodium molybdate 0.01 g/l Preparation: Use double distilled water. Sterilisation: 20 minutes at 121°C pH before sterilisation: Usage: Trace element solution for different media Organisms:
Ref.: 19779	<b>Culture medium</b>	ISP 4
Ref.: 19779	<b>Culture medium growth</b>	yes
Ref.: 19779	<b>Culture medium composition</b>	Name: ISP 4; DSM 547 Solution I: Difco soluble starch, 10.0 g. Make a paste of the starch with a small amount of cold distilled water and bring to a volume of 500 ml. Solution II: CaCO <sub>3</sub> 2.0 g K <sub>2</sub> HPO <sub>4</sub> (anhydrous) 1.0 g MgSO <sub>4</sub> x 7 H <sub>2</sub> O 1.0 g NaCl 1.0 g (NH <sub>4</sub> ) <sub>2</sub> SO <sub>4</sub> 2.0 g Distilled water 500.0 ml Trace salt solution (see below) 1.0 ml The pH should be between 7.0 and 7.4. Do not adjust if it is within this range. Mix solutions I and II together. Add 20.0 g agar. Liquify agar by steaming at 100°C for 10 to 20 min. Trace element solution: FeSO <sub>4</sub> x 7 H <sub>2</sub> O 0.1 g MnCl <sub>2</sub> x 4 H <sub>2</sub> O 0.1 g ZnSO <sub>4</sub> x 7 H <sub>2</sub> O 0.1 g Distilled water 100.0 ml
Ref.: 19779	<b>Culture medium</b>	ISP 5
Ref.: 19779	<b>Culture medium growth</b>	yes
Ref.: 19779	<b>Culture medium composition</b>	Name: ISP 5 (5323) Composition L-Asparagine 1.0 g/l Glycerol 10.0 g/l K <sub>2</sub> HPO <sub>4</sub> 1.0 g/l Salt solution (see preparation) 1.0 ml/l Agar 20.0 g/l Preparation: Salt solution 1.0 g FeSO <sub>4</sub> x 7 H <sub>2</sub> O 1.0 g MnCl <sub>2</sub> x 4 H <sub>2</sub> O 1.0 g ZNSO <sub>4</sub> x 7 H <sub>2</sub> O in 100 ml water Sterilisation: 20 minutes at 121°C pH before sterilisation: 7.2 Usage: Maintenance and taxonomy Organisms: All Actinomycetes
Ref.: 19779	<b>Culture medium</b>	ISP 6
Ref.: 19779	<b>Culture medium growth</b>	yes
Ref.: 19779	<b>Culture medium composition</b>	Name: ISP 6 (5318) Composition Peptone 15.0 g/l Proteose peptose 5.0 g/l Ferric ammonium citrate 0.5 g/l Sodium glycerophosphate 1.0 g/l Sodium thiosulfate 0.08 g/l Yeast extract 1.0 g/l Agar 15.0 g/l Sterilisation: 20 minutes at 121°C pH before sterilisation: Usage: Production of melanoid pigments Organisms: All Actinomycetes
Ref.: 19779	<b>Culture medium</b>	ISP 7
Ref.: 19779	<b>Culture medium growth</b>	yes

Ref.: 19779 **Culture medium composition** Name: ISP 7 (5322) Composition Glycerol 15.0 g/l L-Tyrosine 0.5 g/l L-Asparagine 1.0 g/l K<sub>2</sub>HPO<sub>4</sub> 0.5 g/l NaCl 0.5 g/l FeSO<sub>4</sub> x 7 H<sub>2</sub>O 0.01 g/l Trace element solution 5343 1.0 ml/l Agar 20.0 Sterilisation: 20 minutes at 121°C pH before sterilisation: 7.3 Usage: Production of melanoid pigments Organisms: All Actinomycetes

		<b>Kind of temperature</b>	<b>Temperature</b>
Ref.: 7985	<b>Temperatures</b>	growth	28 °C
Ref.: 19779		optimum	28 °C
Ref.: 32484		growth	27-37 °C
Ref.: 32484		optimum	30 °C

Ref.: 7985 **Temperature range** mesophilic  
 Ref.: 19779 **Temperature range** mesophilic  
 Ref.: 32484 **Temperature range** mesophilic

		<b>Kind of pH</b>	<b>pH</b>
Ref.: 32484	<b>pH</b>	growth	07-08
Ref.: 32484		optimum	07-08

### Isolation, sampling and environmental information

Ref.: 7985 **Sample type/isolated from** soil, farming field  
 Ref.: 7985 **Geographic location (country and/or sea, region)** Bigeum Island  
 Ref.: 7985 **Country** Republic of Korea  
 Ref.: 7985 **Continent** Asia

<b>Isolation sources categories</b>	<b>Cat1</b>	<b>Cat2</b>	<b>Cat3</b>
	#Engineered	#Agriculture	#Field
	#Environmental	#Terrestrial	#Soil

### Application and interaction

Ref.: 19779 **Biosafety level** 1 Risk group (German classification)  
 Ref.: 7985 **Biosafety level** 1 Risk group (German classification)

### Molecular biology



Ref.: 7985      **GC-content**      69.9 mol% high performance liquid chromatography (HPLC)  
Ref.: 32484      **GC-content**      69.9 mol%

	Sequence database	Sequence accession description	Sequence accession number	Sequence length(bp)	Associated NCBI tax ID
Ref.: 7985	GenBank Direct submission	Nocardioides sp. MSL 09 16S ribosomal RNA gene, partial sequence	EF466115	1398	433651

**Strain availability**

Ref.: 7985      **Culture collection no.**      DSM 19266, KCTC 19272

Ref.: 7985      **Strain history**      <- Chang-Jin Kim; MSL-09

**Associated Passport(s) in StrainInfo**

Ref.: 20218      834855 - <http://www.straininfo.net/strains/834855>

Ref.: 20218      834856 - <http://www.straininfo.net/strains/834856>

**References**

Ref.: 7985      Leibniz Institut DSMZ-Deutsche Sammlung von Mikroorganismen und Zellkulturen GmbH; Curators of the DSMZ; DSM 19266

Ref.: 19779      Wink, J.: Compendium of Actinobacteria. HZI-Helmholtz-Centre for Infection Research, Braunschweig.

Ref.: 20215      D.Gleim, M.Kracht, N.Weiss et. al.: Prokaryotic Nomenclature Up-to-date - compilation of all names of Bacteria and Archaea, validly published according to the Bacteriological Code since 1. Jan. 1980, and validly published nomenclatural changes since.

Ref.: 20218      Verslyppe, B., De Smet, W., De Baets, B., De Vos, P., Dawyndt P. StrainInfo introduces electronic passports for microorganisms.. Syst Appl Microbiol. 37: 42-50 2014 (10.1016/j.syapm.2013.11.002, 24321274)

Ref.: 32484      Barberan A, Caceres Velazquez H, Jones S, Fierer N. Hiding in Plain Sight: Mining Bacterial Species Records for Phenotypic Trait Information. mSphere 2: None-None 2017 (10.1128/mSphere.00237-17, None) - **originally annotated from #28704**

Ref.: 28704      IJSEM 2292 2008 (10.1099/ijms.0.65566-0)

\* These References are textmined

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