

Strain		DSM 106873
Genus		<i>Streptomyces</i>
Species		<i>cahuitamycinicus</i>
Status		
Risk group		1 (provisional classification by DSMZ)
Type strain		13K301, KCTC 49110
Genbank accession number		16S rRNA gene: MG770843
Reference		
Author		Saygin H, Ay H, Guven K, Cetin D, Sahin N.
Title		<i>Streptomyces cahuitamycinicus</i> sp. nov., isolated from desert soil and reclassification of <i>Streptomyces galilaeus</i> as a later heterotypic synonym of <i>Streptomyces bobii</i>
Journal		Int J Syst Evol Microbiol
Volume		70
Page		4050-4060
Year		2020
Morphology		
Agar	ISP 2 - growth/G	Good
Agar	ISP 2 - colony colour/R	1001 beige
Agar	ISP 2 - aerial mycelium/A	Sparse, 9003 signal white
Agar	ISP 2 - soluble pigment/S	None
Agar	ISP 3 - G	Good
Agar	ISP 3 - R	1013 oyster white
Agar	ISP 3 - A	Good, 9003 signal white
Agar	ISP 3 - S	3022 salmon pink
Agar	ISP 4 - G	Good
Agar	ISP 4 - R	1014 ivory
Agar	ISP 4 - A	None
Agar	ISP 4 - S	1011 brown beige
Agar	ISP 5 - G	Good
Agar	ISP 5 - R	1013 oyster white
Agar	ISP 5 - A	Good, 9003 signal white
Agar	ISP 5 - S	3012 beige red
Agar	ISP 6 - G	Sparse
Agar	ISP 6 - R	1015 light ivory, 1024 ochre yellow
Agar	ISP 6 - A	None
Agar	ISP 6 - S	None
Agar	ISP 7 - G	Good
Agar	ISP 7 - R	7003 moss grey

Agar	ISP 7 - A	Good, 9002 grey white
Agar	ISP 7 - S	8000 green brown
Agar	suter with tyrosine - G	Good
Agar	suter with tyrosine - R	7013 brown grey, 8014 sepia brown
Agar	suter with tyrosine - A	Sparse, 9002 grey white
Agar	suter with tyrosine - S	8001 ochre brown, 9005 jet black
Agar	suter without tyrosine - G	Good
Agar	suter without tyrosine - R	1001 beige
Agar	suter without tyrosine - A	None
Agar	suter without tyrosine - S	None
	Sporechains/Sporangia	
Physiology		
Melanin		1
pH	range	
pH	optimum	
temperature	range	
temperature	optimum	
sodium chloride tolerance		2,5%
lysozyme tolerance		
use of carbohydrates	glucose	+
use of carbohydrates	arabinose	+
use of carbohydrates	sucrose	+
use of carbohydrates	xylose	++
use of carbohydrates	inositol	++
use of carbohydrates	mannose	+
use of carbohydrates	fructose	+
use of carbohydrates	rhamnose	+
use of carbohydrates	raffinose	++
use of carbohydrates	cellulose	+
Api zym	Phosphatase alcaline	5
Api zym	Esterase (C4)	3
Api zym	Esterase Lipase (C8)	3
Api zym	Lipase (C14)	3
Api zym	Leucin arylamidase	5
Api zym	Valine arylamidase	2
Api zym	Cystine arylamidase	1
Api zym	Trypsin	3
Api zym	Chymotrypsin	5
Api zym	Phosphatase acid	5
Api zym	Naphtol-AS-BI-phosphohydrolase	5
Api zym	alpha galactosidase	0
Api zym	beta galactosidase	5
Api zym	beta glucuronidase	0
Api zym	alpha glucosidase	2

Api zym	beta glucosidase	0
Api zym	N-acetyl-beta-glucosaminidase	5
Api zym	alpha mannosidase	4
Api zym	alpha fucosidase	0
Api coryne	nitrate reduction	-
Api coryne	Pyraziamidase	-
Api coryne	Pyrrolidonyl arylamidase	+
Api coryne	Alkaline phosphatase	+
Api coryne	beta glucuronidase	-
Api coryne	beta galactosidase	+
Api coryne	alpha glucosidase	-
Api coryne	N-acetyl -beta glucosaminidase	+
Api coryne	Esculin (beta glucosidase)	-
Api coryne	Urease	-
Api coryne	Gelatine(hydrolysis)	+
Api coryne	Glucose fermentation	-
Api coryne	Ribose fermentation	-
Api coryne	Xylose fermentation	-
Api coryne	Mannitol fermentation	-
Api coryne	Maltose fermentation	-
Api coryne	Lactose fermentation	-
Api coryne	Sucrose fermentation	-
Api coryne	Glycogen fermentation	-

APICORYNE



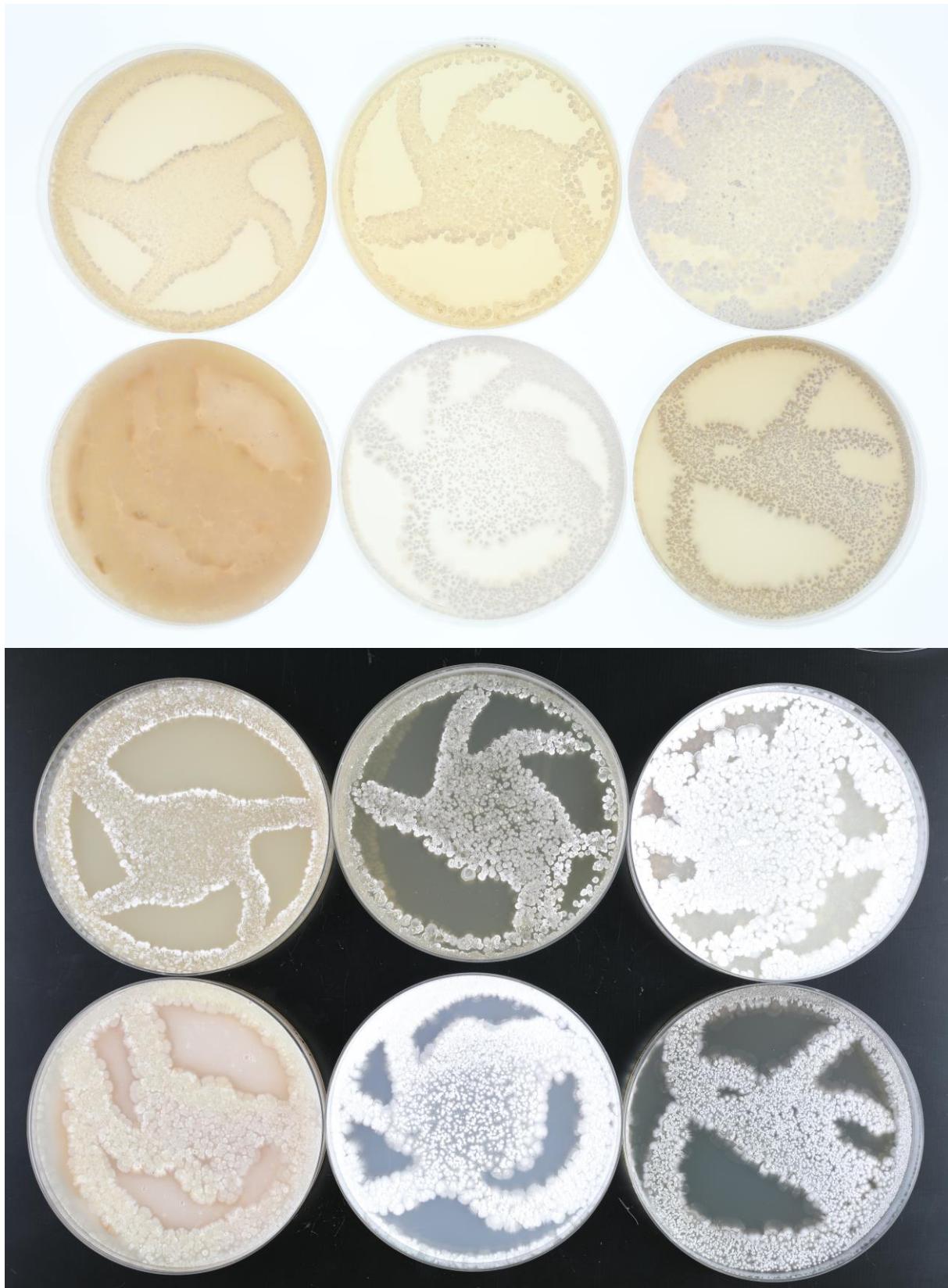
Abbildung 1: Apicoryne-Teststreifen mit Keim DSM.

APIZYM

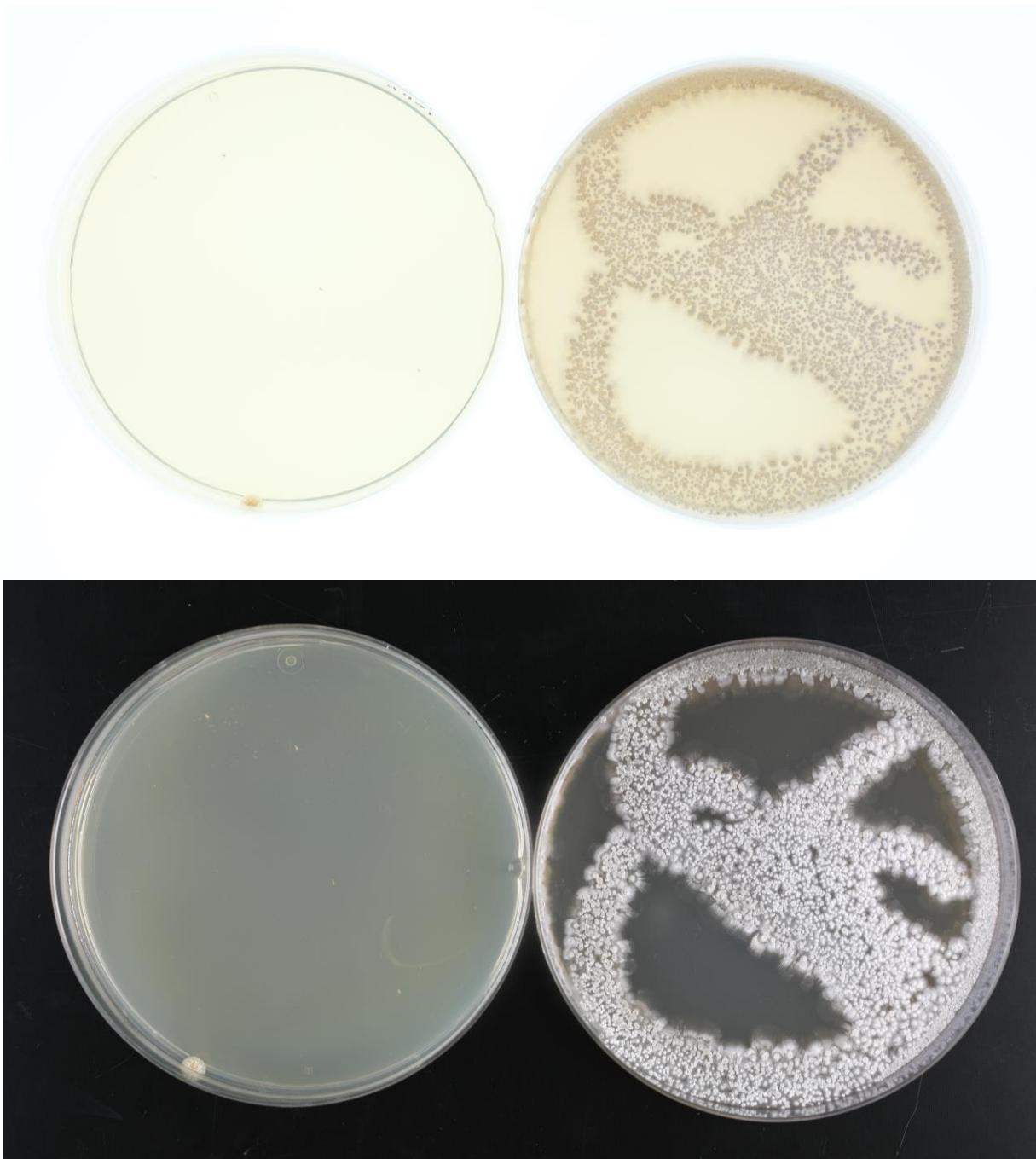


Abbildung 2: Apizym-Teststreifen mit Keim DSM.

Plates (65, ISP2, ISP3, ISP4, ISP5, ISP7)



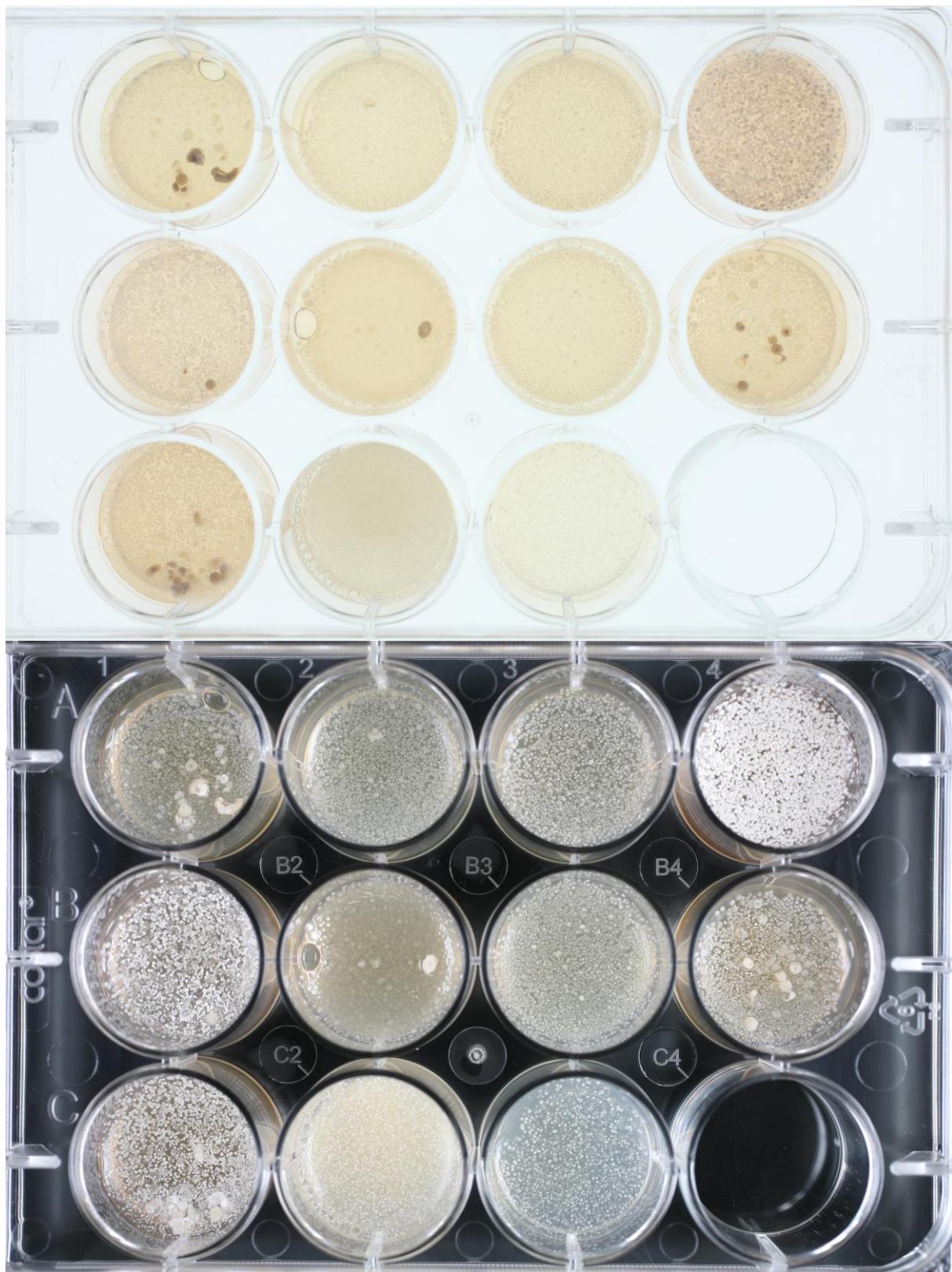
(ISP6, ISP7)



(SSM+T, SSM-T)



Carbon utilization test (from top left to bottom right: glucose, arabinose, sucrose, xylose, inositol, mannose, fructose, rhamnose, raffinose, cellulose)



Sodium chloride tolerance test (from top left to bottom right: 0%, 2,5%, 5%, 7,5%, 10%)

