

Compendium of Actinobacteria from Dr. Joachim M. Wink
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Strain		DSM 45601
Genus		<i>Allonocardiopsis</i>
Species		<i>opalescens</i>
Status		
Risk group		L1
Type strain		CPCC 203428, KCTC 19844
Genbank accession numbers		16S rRNA gene: JQ309825
Reference		
Author		Du, H. J., Zhang, Y. Q., Liu, H. Y., Su, J., Wei, Y. Z., Ma, B. P., Guo, B. L., Yu, L. Y.
Title		<i>Allonocardiopsis opalescens</i> gen. nov., sp. nov., a new member of the suborder <i>Streptosporangineae</i> , from the surface-sterilized fruit of a medicinal plant
Journal		<i>Int J Syst Evol Microbiol</i>
Volume		63 (Pt3)
Page		900-4
Year		2013
Morphology		
Agar	ISP 2 - growth/G	good
Agar	ISP 2 - colony color/R	ivory (1014)
Agar	ISP 2 - aerial mycelium/A	none
Agar	ISP 2 - soluble pigment/S	none
Agar	ISP 3 - G	sparse
Agar	ISP 3 - R	ivory (1014)
Agar	ISP 3 - A	none
Agar	ISP 3 - S	none
Agar	ISP 4 - G	good
Agar	ISP 4 - R	ivory (1014) and sand yellow (1002)
Agar	ISP 4 - A	none
Agar	ISP 4 - S	ivory(1014)
Agar	ISP 5 - G	sparse
Agar	ISP 5 - R	light ivory (1015)
Agar	ISP 5 - A	none
Agar	ISP 5 - S	none
Agar	ISP 6 - G	sparse
Agar	ISP 6 - R	ivory (1014)
Agar	ISP 6 - A	none
Agar	ISP 6 - S	golden yellow (1004)
Agar	ISP 7 - G	sparse
Agar	ISP 7 - R	oyster white (1013)
Agar	ISP 7 - A	none
Agar	ISP 7 - S	none

Agar	suter with tyrosine - G	sparse
Agar	suter with tyrosine - R	light ivory (1015)
Agar	suter with tyrosine - A	none
Agar	suter with tyrosine - S	none
Agar	suter without tyrosine - G	good
Agar	suter without tyrosine - R	light ivory (1015)
Agar	suter without tyrosine - A	none
Agar	suter without tyrosine - S	none
	Sporechains/Sporangia	
Physiology		
Melanin		-
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)	2
Api zym	Esterase Lipase (C8)	3
Api zym	Lipase (C14)	1
Api zym	Leucin arylamidase	5
Api zym	Valine arylamidase	5
Api zym	Cystine arylamidase	2
Api zym	Trypsin	2
Api zym	Chymotrypsin	3
Api zym	Phosphatase acid	3
Api zym	Naphtol-AS-BI-phosphohydrolase	2
Api zym	alpha galactosidase	5
Api zym	beta galactosidase	4
Api zym	beta glucuronidase	5
Api zym	alpha glucosidase	5
Api zym	beta glucosidase	5
Api zym	N-acetyl-beta-glucosaminidase	4

Api zym	alpha mannosidase	5
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 45601.

Apizym

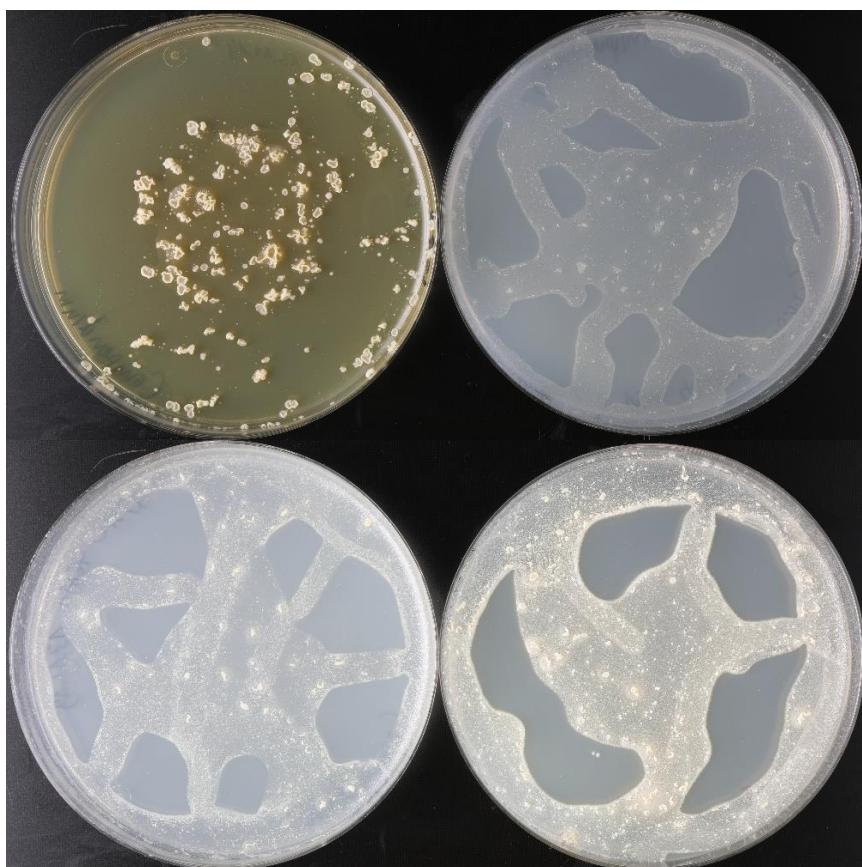
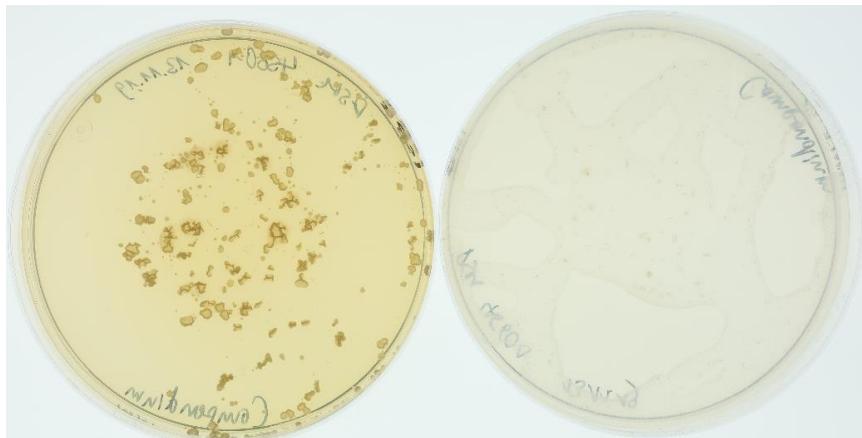


Abbildung 2: Apizym-Teststreifen mit Keim DSM 45601.

Plates (535, 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%)**

