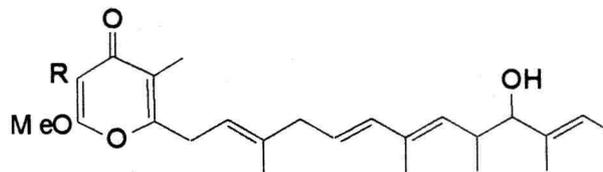


Name:	<i>Streptomyces pactum</i>
Authors:	Bhuyan et al. 1962
Status:	Approved Lists
Reference(s):	Int. J. Syst. Bacteriol. 30:395 (AL)
Risk group:	1 (German classification)
Type strain:	ATCC 12434, IMET 43357, NRRL 2939, DSM 40530

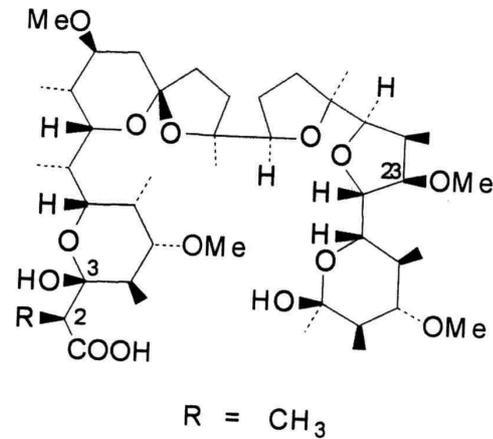
Secondary metabolites of *Streptomyces pactum*

Actinopyrones, exhibiting coronary vasodilating activities and are weakly active against gram-positive bacteria and dermatophytes

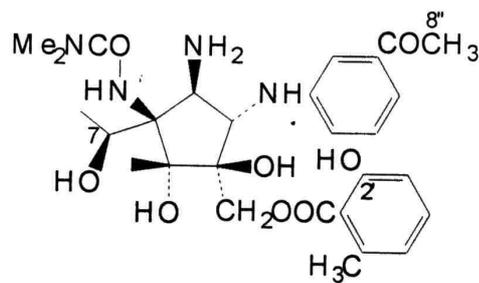


Actinopyrone A R = CH₃
Actinopyrone B R = H
Actinopyrone C R = CH₂CH₃

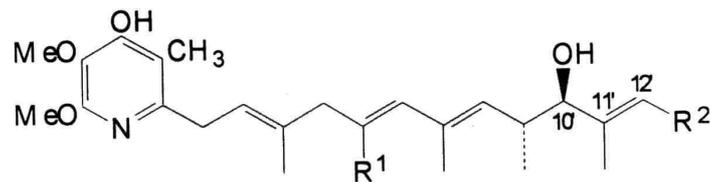
Lonomycin A, polyether antibiotic, ionophore and protozoacide



Pactamycin, shows activity against gram-positive and -negative bacteria and shows antineoplastic activity



Piericidin A, pyridine antibiotic family, inhibits NADH oxidase of ox-heart mitochondria in absence of bovine serum albumen, shows also insecticidal activity



- Piericidin A₁, R¹ = H, R² = CH₃
 A₂, R¹ = R² = CH₃
 A₃, R¹ = H, R² = CH(CH₃)₂
 A₄, R¹ = CH₃, R² = CH(CH₃)₂

Genus: Streptomyces

FH 2018

Species: pactum

Numbers in other collections: IMET 43357

Morphology:

	G	R
<u>ISP 2</u>	good	grey
	A	SP
	beige	beige
	G	R
<u>ISP 3</u>	good	colorless
	A	SP
	sparse	none
	G	R
<u>ISP 4</u>	good	beige
	A	SP
	grey	beige
	G	R
<u>ISP 5</u>	good	beige
	A	SP
	grey	none
	G	R
<u>ISP 6</u>	good	colorless
	A	SP
	none	none
	G	R
<u>ISP 7</u>	good	beige
	A	SP
	grey beige	none

Spore chains: RAb

Spore surface: smooth

Sporangia:

Fragmentation:

Melanoid pigment: - - - -

NaCl resistance:

Lysozyme resistance:

pH: Value- Optimum-

Temperature : Value- Optimum- 28 °C

Carbon utilization:

Glu	Ara	Suc	Xyl	Ino	Man	Fru	Rha	Raf	Cel
+	-	-	-	-	-	-	-	-	-

Enzymes:

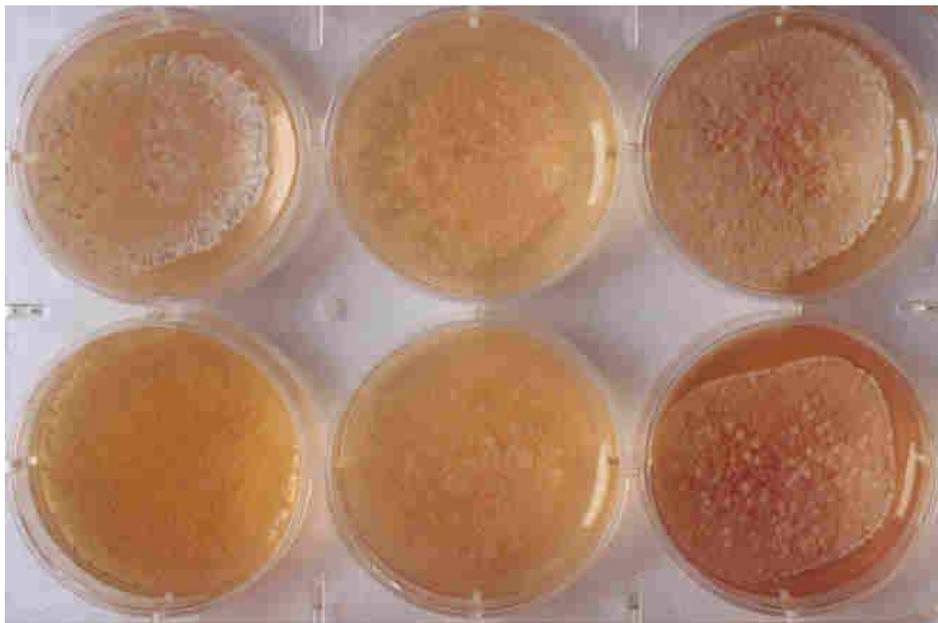
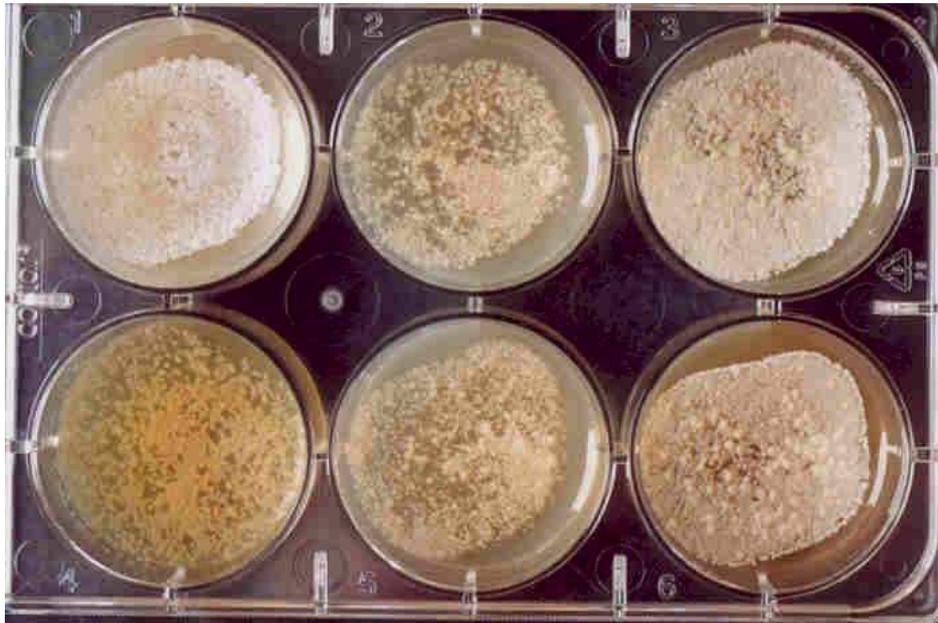
Gel	Cit	Ure	Arg	Onp	Trp	Lys	Odc	VP	Ind	H2S
+	-	+	+	+	-	+	+	-	-	-

Comments:



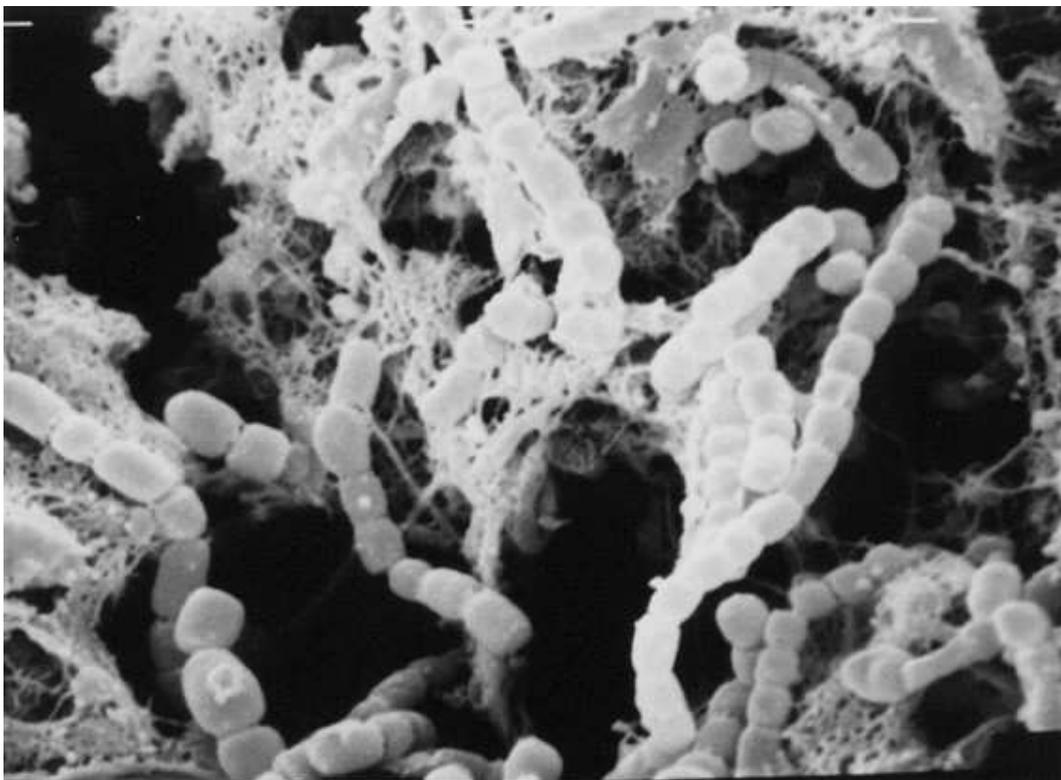
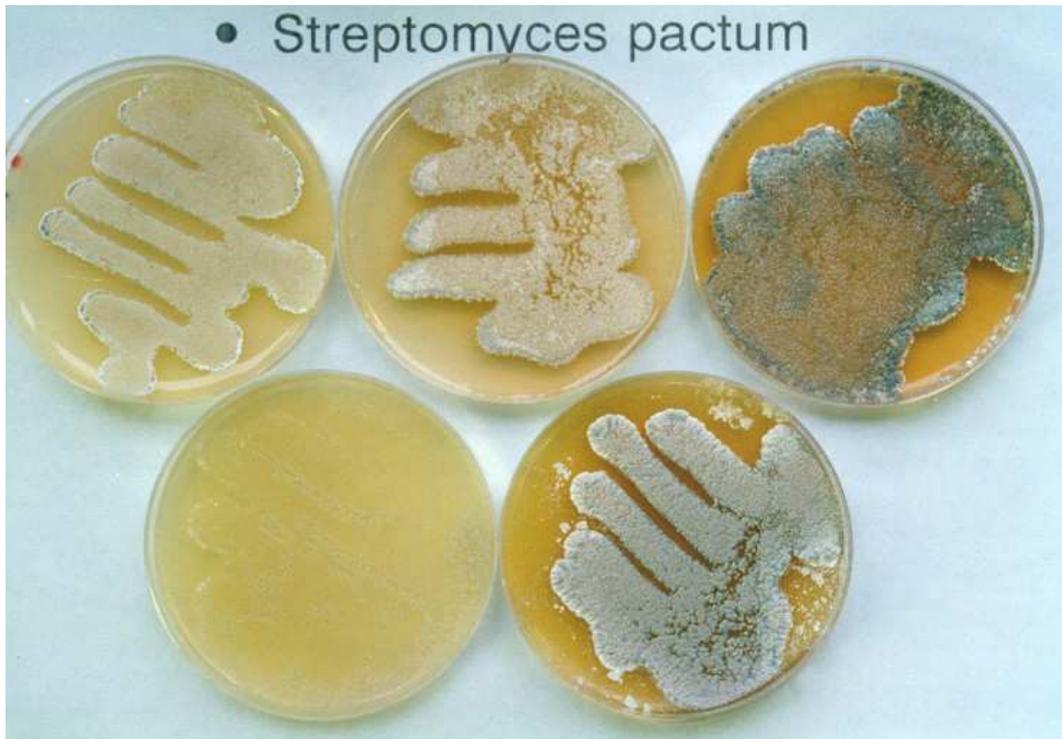
Streptomyces pactum

A and B – Agar plates medium 5006, 5265 and 5315



Streptomyces pactum

C and D – Microplate with ISP- and melanin media



Streptomyces pactum

E – Agar plates medium 5315, 5317, 5323, 5318 and 5322

F – Spore chain morphology and spore surface in SEM (x 10.000)

