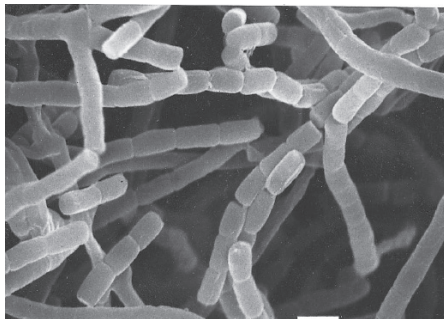


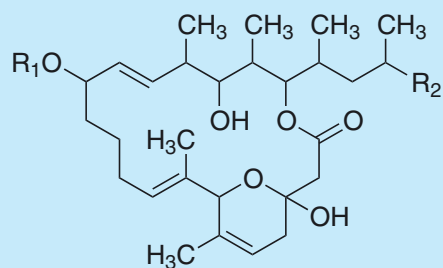
# Irumamycin

## 1. Discovery, producing organism and structures<sup>1-4)</sup>

Antifungal screening led to the discovery of the irumamycins from a culture broth of *Streptomyces subflavus*. Subsequently, a blocked mutant of the strain AM-3603<sup>T</sup> was found to accumulate two lactones, named irumanolides I and II.



*Streptomyces subflavus* subsp.  
*irumaensis* AM-3603<sup>T</sup>



	Irumamycin	3'-O-Decarbamoyl- irumamycin	Irumanolide I	Irumanolide II
R <sub>1</sub>			H	H
R <sub>2</sub>				

## 2. Physical data (Irumamycin)

White amorphous powder. C<sub>41</sub>H<sub>65</sub>NO<sub>12</sub>; mol wt 763.45. Sol. in MeOH, EtOH, acetone. Insol. in H<sub>2</sub>O, hexane, petroleum ether.

## 3. Biological activity<sup>5)</sup>

### 1) Antifungal activity

Test organism		MIC (μg/ml)* <sup>1</sup>					Prumycin
		Iruma- mycin	Venturi- cidin A <sup>6)</sup>	Botry- cidin	Aabo- mycin A	Blasti- cidin S	
<i>Pyricularia oryzae</i>	A* <sup>2</sup>	0.05	>0.03	0.05	<0.2	>100	>100
	B	>0.03	>0.03	0.03	<0.2	25	100
<i>Sclerotinia cinerea</i>	A	0.05	0.78	0.4	0.2	50	1.56
	B	0.1	0.78	0.78	0.2	25	6.25
<i>Rhizoctonia solani</i>	A	0.4	1.56	12.5	>100	>100	>100
	B	25	>25	>25	>100	>100	>100
<i>Botrytis cinerea</i>	A	—* <sup>3</sup>	—	—	>100	50	>100
	B	6.25	>25	12.5	>100	100	50
<i>Alternaria kikuchiana</i>	A	50	—	—	>100	50	>100
	B	12.5	5	12.5	100	12.5	50

\*1 Glucose-potato agar, 2–3 days.

\*2 A, isolated from an infected plant; B, laboratory stock culture.

\*3 —, Not tested.

## 2) Protective effect against plant-pathogenic fungi

Protection of plants from fungal infection by irumamycin in pot tests

Pathogenic organism	Plant infected	Irumamycin (ppm)	Protection value (%)
<i>Botrytis cinerea</i>	Cucumber	200	98
		100	97
		50	97
<i>Colletotrichum lagenarium</i>	Cucumber	200	100
		100	99
		50	92
<i>Cochliobolus miyabeanus</i>	Rice plant	200	92

Calculated from an equation:

$$\left( 1 - \frac{\text{No. of infectious lesions on treated plants}}{\text{No. of infectious lesions on control plants}} \right) \times 100$$

3) Antitrypanosomal activities<sup>8)</sup>

In the course of our screening program to discover new antitrypanosomal antibiotics, by screening metabolites from soil microorganisms, we have discovered 10 compounds that display potent antitrypanosomal activity *in vitro*. We reported the results of sensitivity tests on the GUTat 3.1 strain of *T. b. brucei* and the STIB900 strain of *T. b. rhodesiense* using the four commonly used antitrypanosomal drugs as well as detailing the novel antitrypanosomal activity and cytotoxicity of the ten antibiotics *in vitro*.

*In vitro* antitrypanosomal activity and cytotoxicity of 10 microbial metabolites

Compound	IC <sub>50</sub> (ng/mL)				
	Antitrypanosomal activity		Cytotoxicity MRC-5	Selective index	
	GUTat 3.1	STIB900		MRC-5/ GUTat 3.1	MRC-5/STIB900
Aureothin	1.4	1.1	>25,000	>17,857	>22,727
Cellocidin	150	30	5,910	39	179
Destomycin A	330	210	21,140	64	101
Echinomycin	20	14	6,310	316	451
Hedamycin	14	18	>25,000	>1,786	>1,389
Irumamycin	20	31	>12,500	>625	>403
LL-Z 1272b	49	59	13,620	278	231
<i>O</i> -methylnanaomycin A	210	16	4,890	23	306
Venturicidin A	120	540	>25,000	>208	>185
Virustomycin A	0.45	480	80	178	0.2

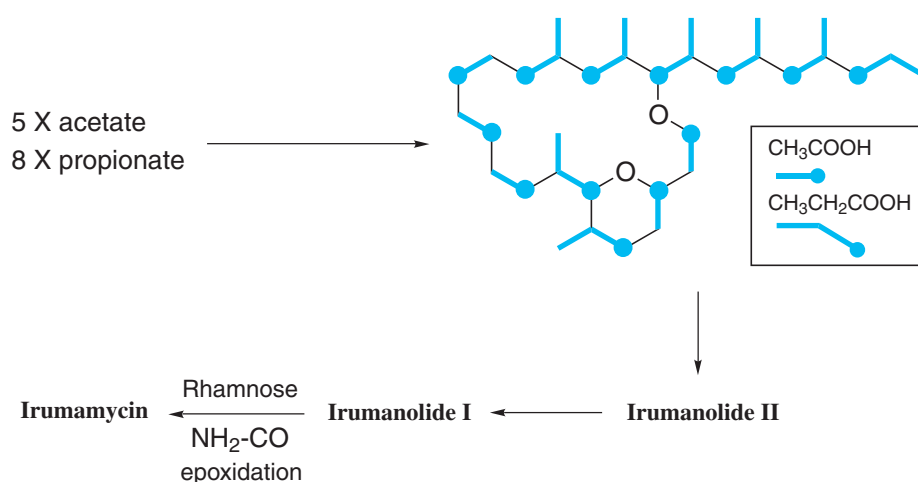
GUTat 3.1: *Trypanosoma brucei brucei* strain

STIB900: *Trypanosoma brucei rhodesiense* strain

MRC-5: Human lung fibroblast cell line

#### 4. Biosynthesis<sup>4,7)</sup>

The carbon skeleton of the aglycone moiety originated from 5 acetate and 8 propionate units. The biosynthetic pathway to irumamycin was proposed as shown below:



#### 5. References

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