

A Platform for the Discovery of New Macrolide Antibiotics

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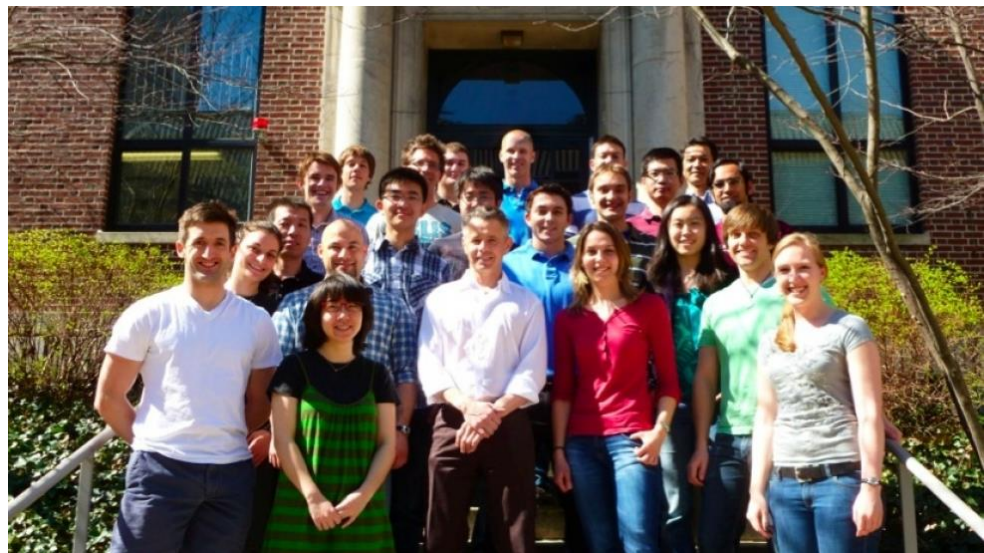
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Presented by Alexander Chatterley

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The Myers Group

- ❖ Andrew Myers graduated from MIT 1981 with Bachelors of science. Graduate student and a brief post doc with E. J. Corey. Began independent career at Caltech (1986) then moved to Harvard in 1998.
- ❖ Comprised of 17 members (6 grad students/8 post docs/3 staff).
- ❖ Research focuses include synthesis of complex natural antibiotics and the development of methodology to aid in this.

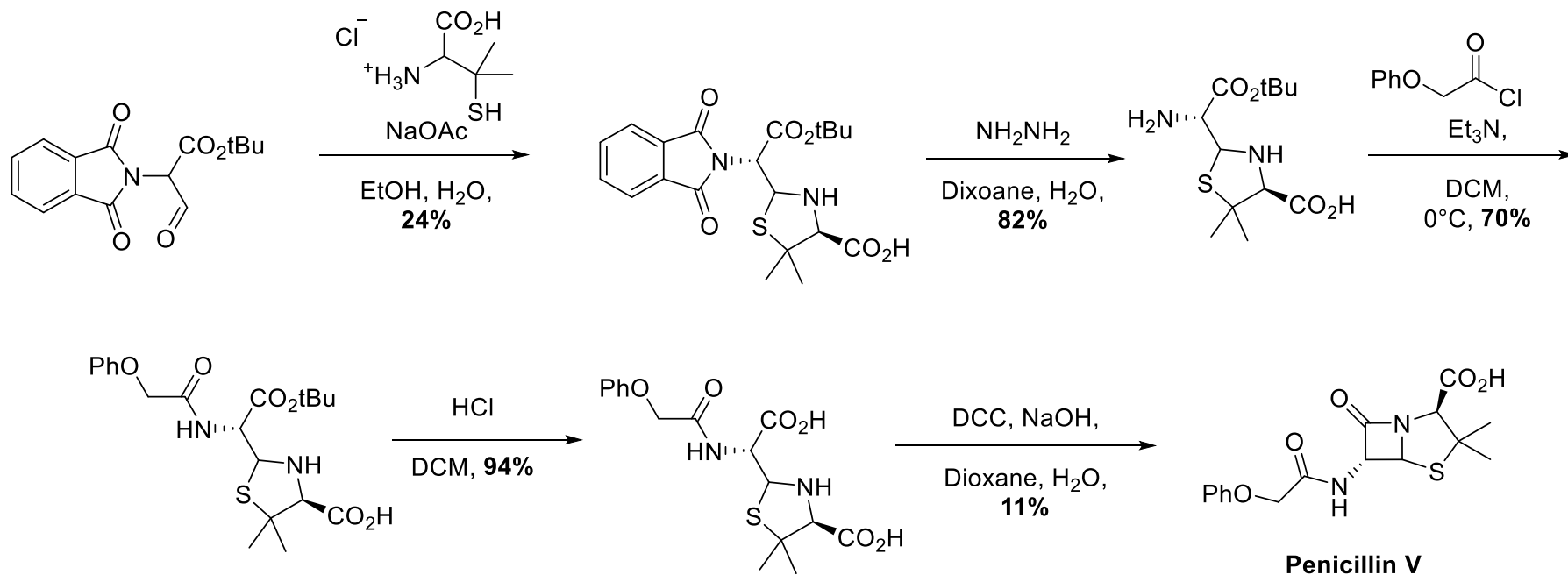


Antibiotics

Antibiotics are one of the major corner stones of modern medicine.

- ❖ Early reports of using mould as a poultice in pre-BC times to treat open wounds by applying moulds.
- ❖ Use became more refined throughout history, particularly in the late renaissance through to early industrial. For example the use of mould by apothecaries in England to treat injuries.
- ❖ First anti-biotic isolated by British biologist Sir Andrew Fleming in 1928, arguably began the modern anti-bacterial era.
- ❖ First chemical synthesis of penicillin in 1957 by Dr Edward Sheehan (supervisor to E. J. Corey).

First synthesis of Pencillin



Antibacterial Apocalypse

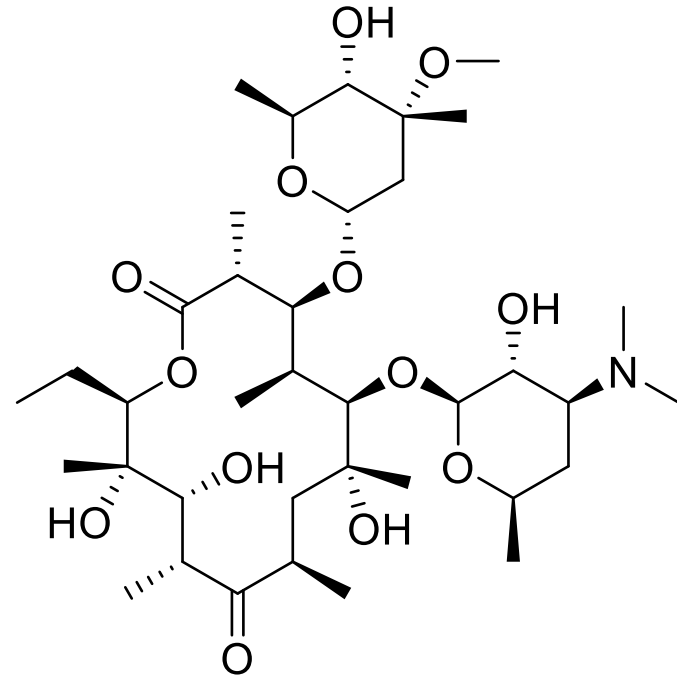
Antibiotic resistance is rapidly becoming a global health concern.

- ❖ Over use of antibiotics in people and animals has caused multiple strains of drug resistance bacteria to evolve.
- ❖ No new class of antibiotics have been discovered in the 1980's.
- ❖ Antibiotic research while crucial, is not attractive to the private sector due to the business model.

Classes of antibiotics

There are several classes of antibiotics:

- ❖ Penicillin's
- ❖ Aminoglycosides
- ❖ Carbapenems
- ❖ Cephalosporin's
- ❖ Fluoroquinolones
- ❖ Sulphonamides
- ❖ **Macrolides**



Erythromycin A

Macrolide Antibiotics

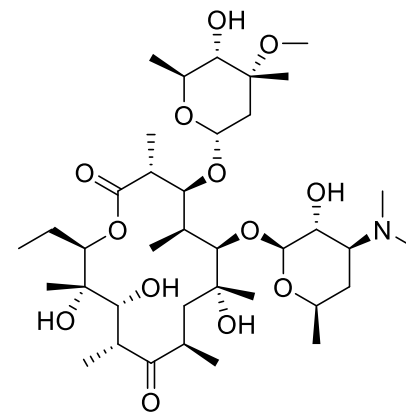
Erythromycin, one of the earliest macrolides was isolated from Philippine soil samples in 1949.

- ❖ Characterised by their large macrocyclic ring structure, Usually 14, 15 or 16 membered.
- ❖ Active against Gram-positive and to limited Gram-negative bacteria.
- ❖ Binds reversibly to the P site on the 50S subunit of the bacterial ribosome. This inhibits protein synthesis resulting in bacterial cell death.

Macrolide synthesis

First total synthesis of Erythromycin A was completed by Woodward in 1981.

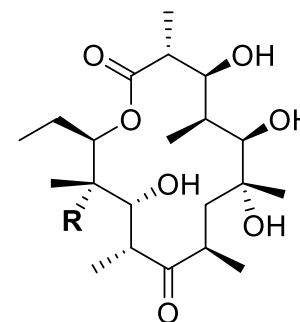
- ❖ 0.089% total yield,
- ❖ 52 steps total.
- ❖ 48 students worked on it.



Erythromycin A

Total synthesis of Erythronolide A and B carried out by Corey in 1978/9.

Again, these were very complex synthesis, involving multiple students (including K.C. Nicolaou).



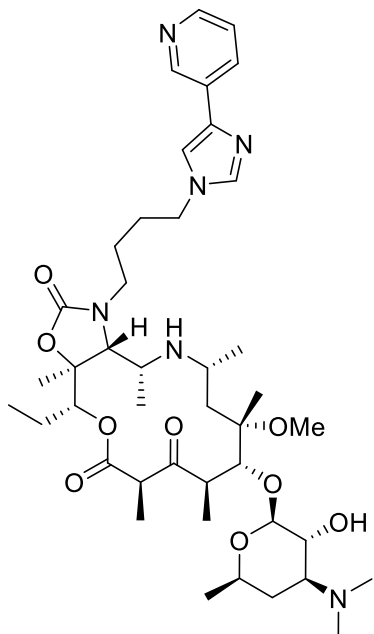
R = OH A
H B

Erythronolide

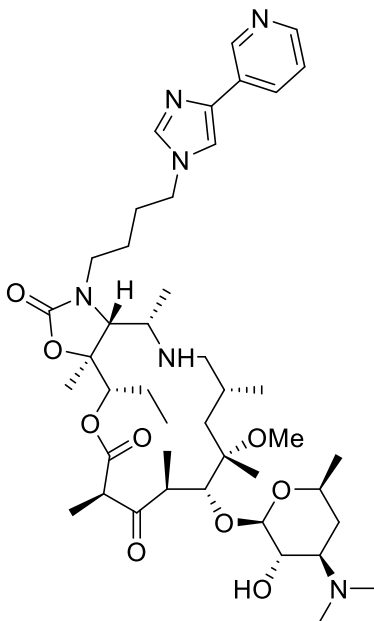
This Paper

Meyers group embarked on a quest to synthesise a library of macrolides as a platform for the discovery of new antibiotics.

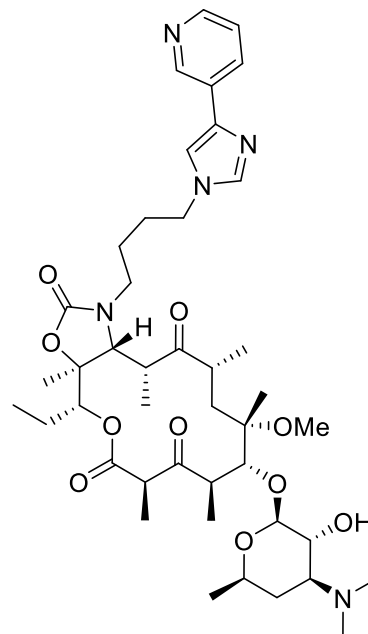
They divided this approach into three classes:



14-Membered azaketolides

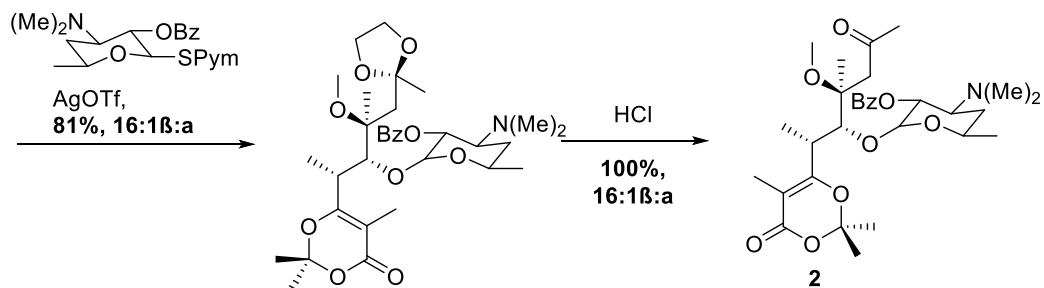
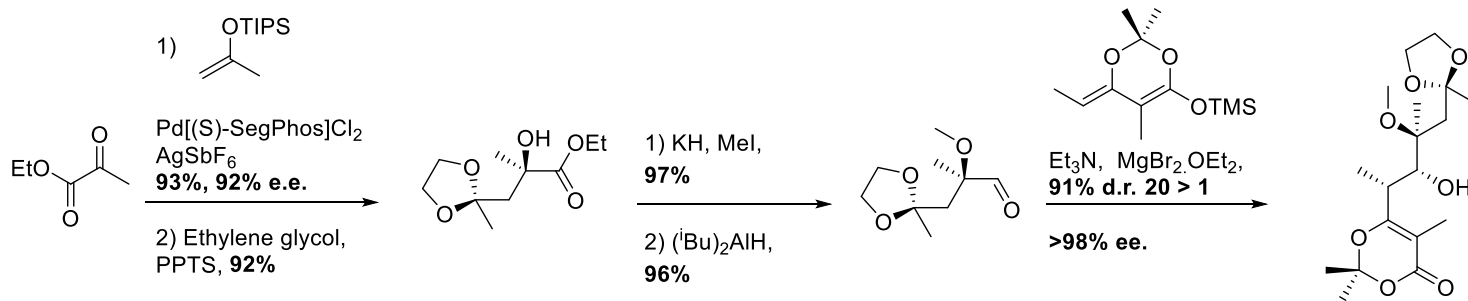
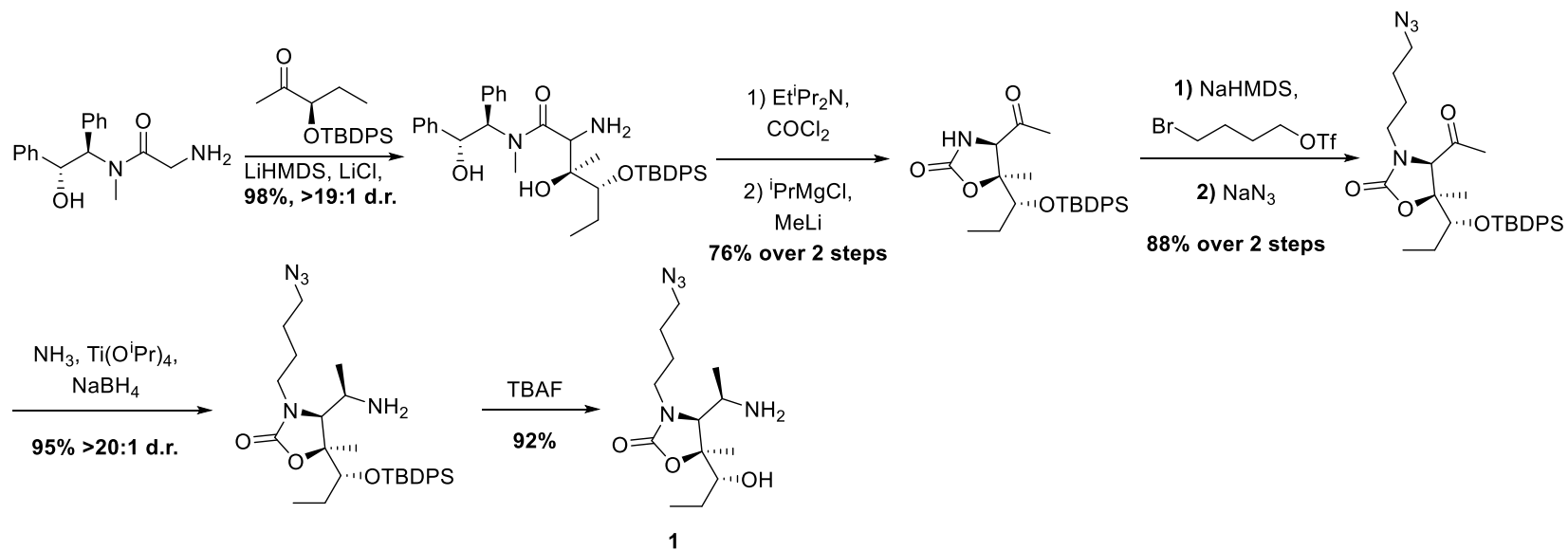


15-Membered azaketolides

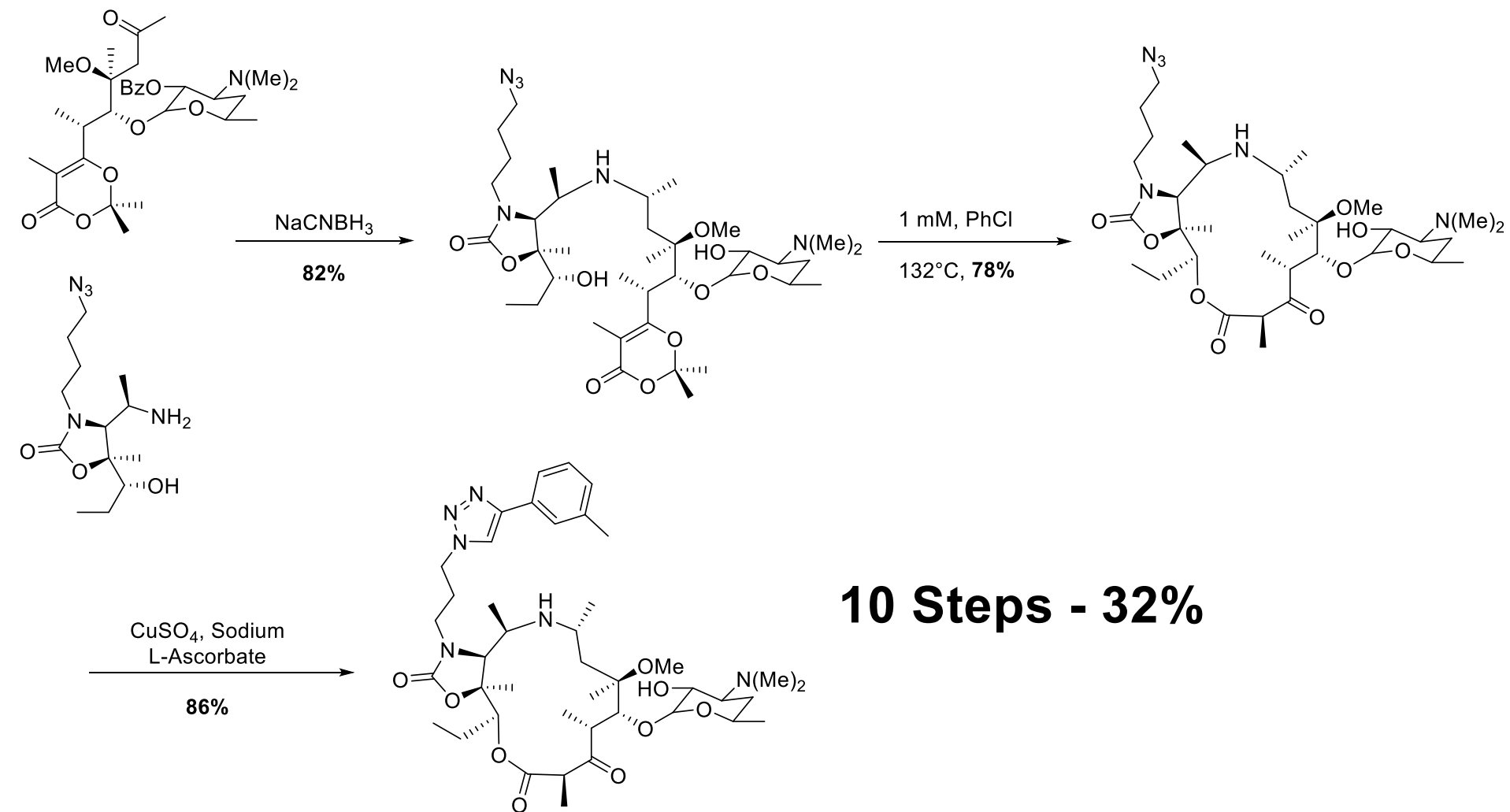


15-Membered ketolides

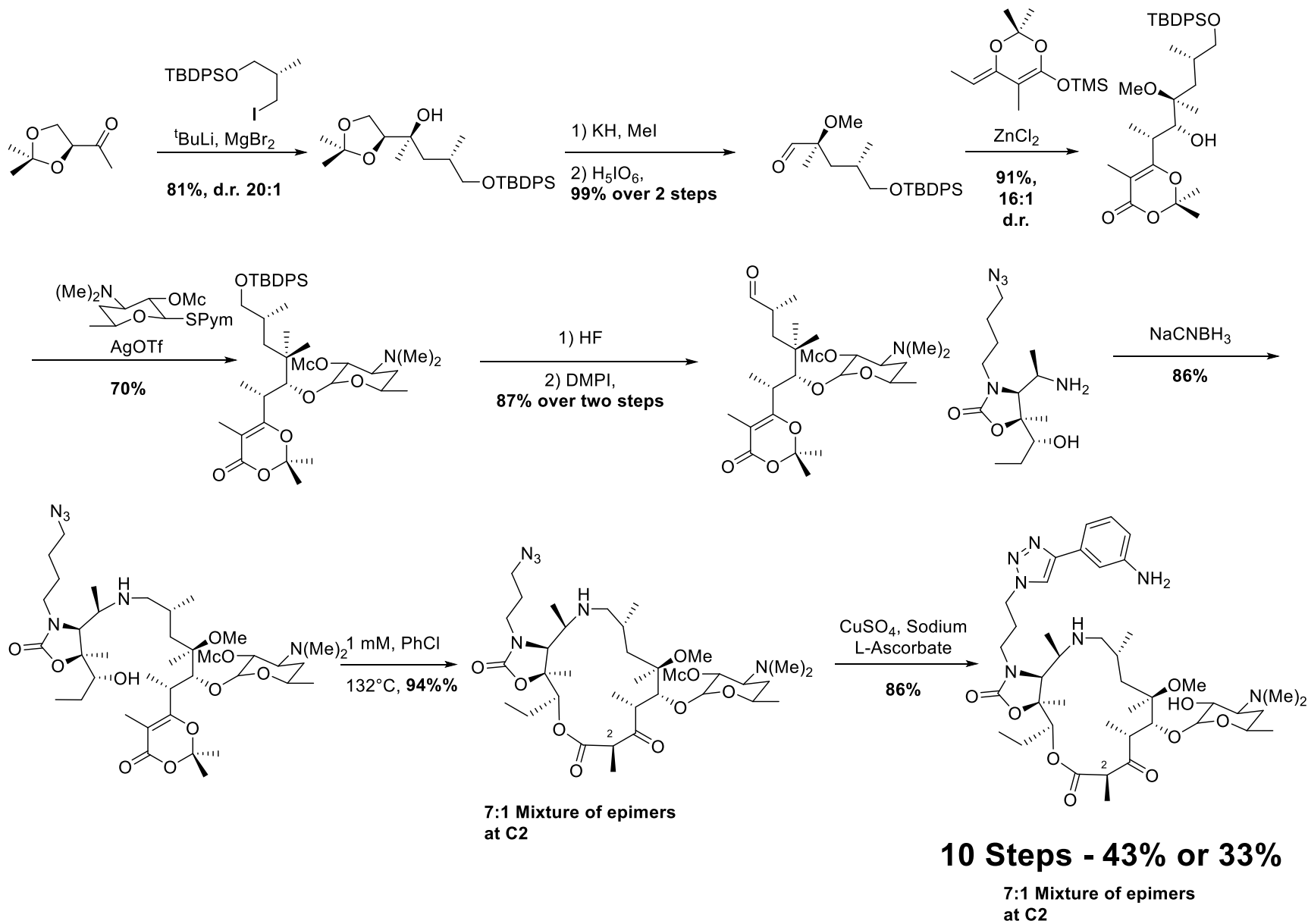
14-Membered Azaketolides 1



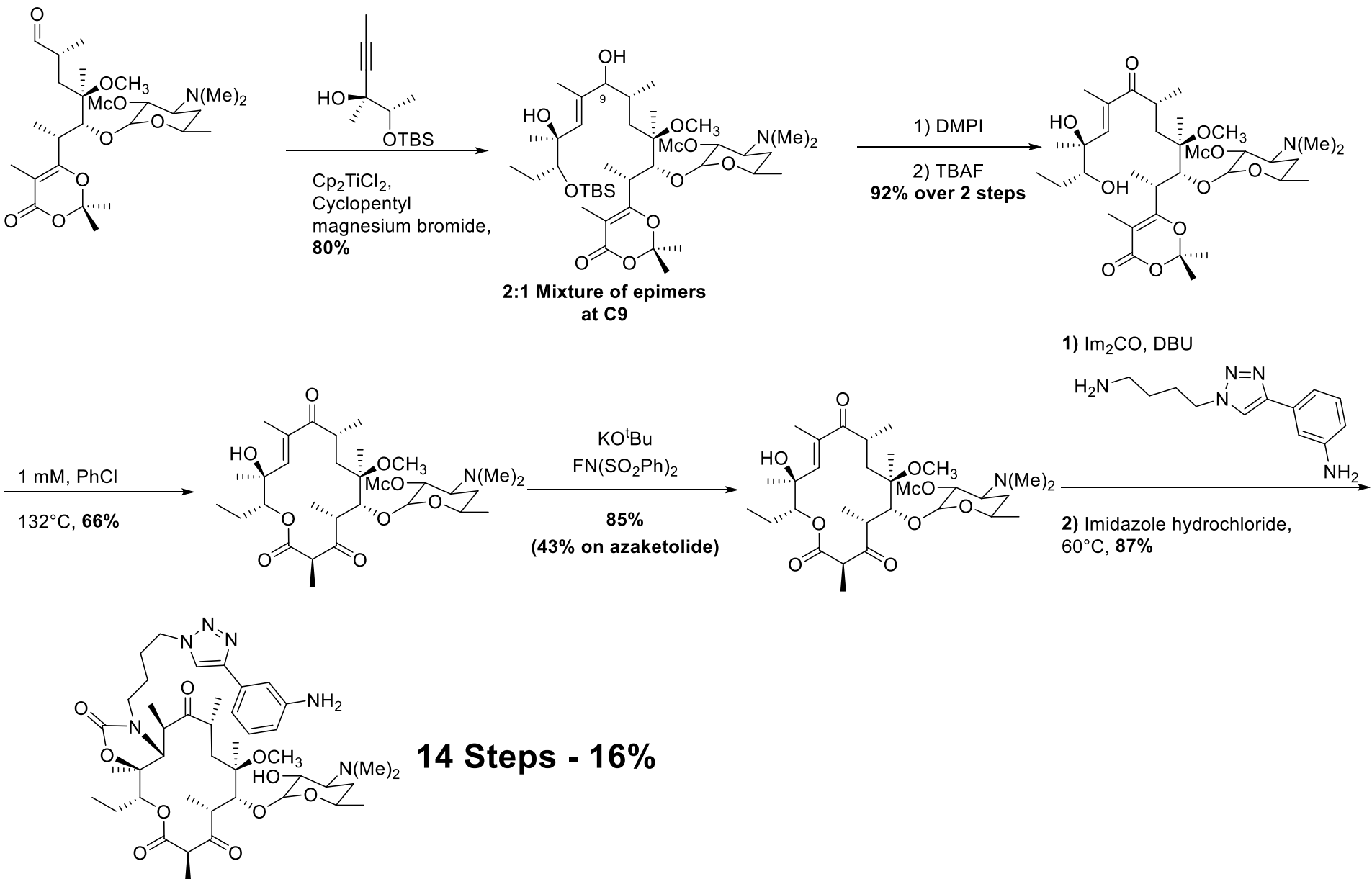
14-Membered Azaketolides 2



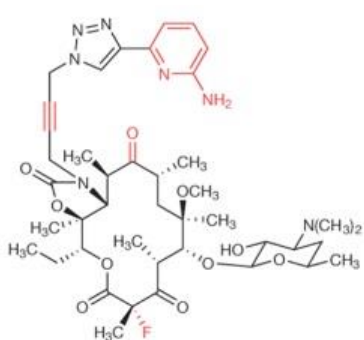
15-Membered Azaketolides



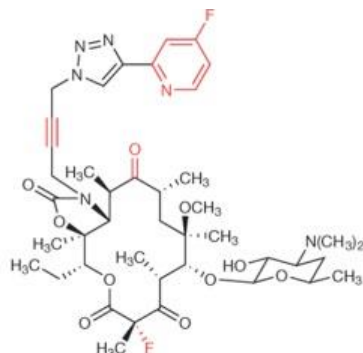
14 Membered Ketolides



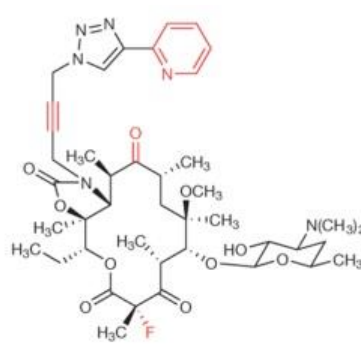
Library construction



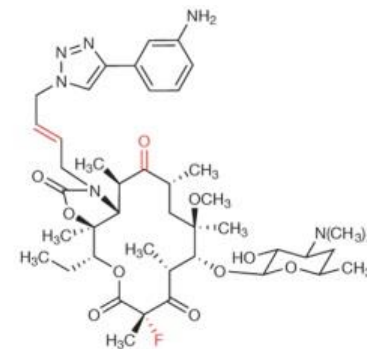
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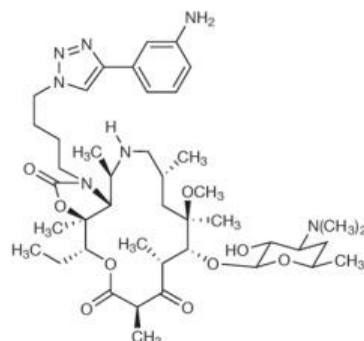
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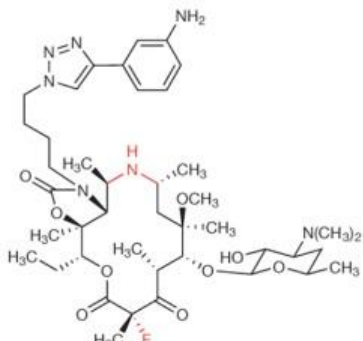
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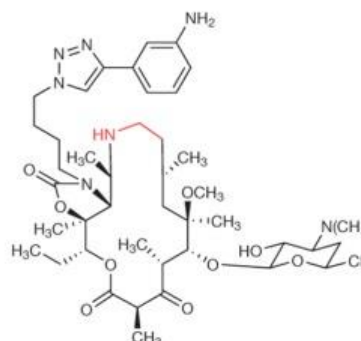
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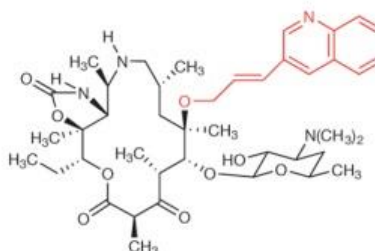
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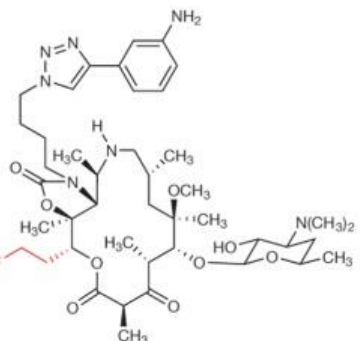
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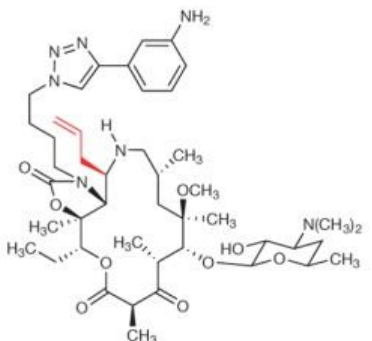
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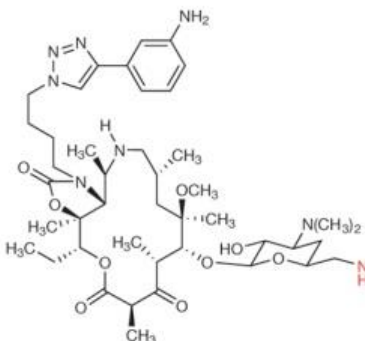
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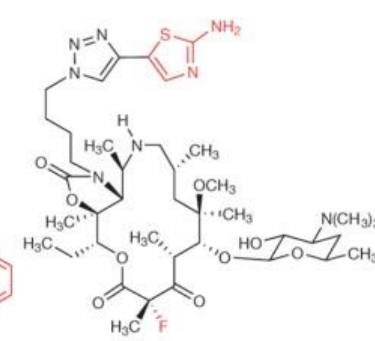
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FSM-11044



FSM-21887



FSM-21760

Microbiological testing

Species	Strain description	Erythro	Azithro	Telithro	Solithro	100573	100563	100490	11563	20707	22391	21397	20919	11453	11044	21887	21760	
Gram-positive	<i>S. aureus</i> ATCC 29213	0.5	1	0.125	0.125	0.06	≤0.03	≤0.03	0.06	0.5	0.25	4	0.25	1	1	8	0.5	
	<i>S. aureus</i> BAA-977; iErmA	>256	>256	0.06	≤0.03	0.06	0.06	0.03	0.06	0.5	0.5	4	0.5	1	1	8	1	
	<i>S. aureus</i> MP513; MRSA; cErmA	>256	>256	256	>64	16	16	64	64	>64	64	64	64	>64	>64	>64	>64	64
	<i>S. aureus</i> NRS384; MRSA; MsrA	64	128	0.125	0.25	0.06	0.125	0.06	0.125	1	1	8	0.5	2	2	16	4	
	<i>S. pneumoniae</i> ATCC 49619	0.03	0.06	≤0.03	≤0.03	≤0.03	≤0.03	≤0.03	≤0.03	≤0.03	≤0.03	0.06	≤0.03	≤0.03	≤0.03	0.06	≤0.03	
	<i>S. pneumoniae</i> UNT-042; ErmB/MefA	>256	>256	0.125	0.25	≤0.03	≤0.03	≤0.03	≤0.03	2	0.125	8	0.5	2	8	1	1	
	<i>S. pyogenes</i> ATCC 19615	≤0.03	≤0.03	≤0.03	≤0.03	≤0.03	≤0.03	≤0.03	≤0.03	≤0.03	≤0.03	0.06	≤0.03	≤0.03	≤0.03	0.06	≤0.03	
	<i>E. faecalis</i> ATCC 29212	1	4	≤0.03	≤0.03	0.03	0.03	0.03	≤0.03	0.125	0.06	0.5	0.25	0.125	0.125	0.5	0.06	
	<i>E. faecalis</i> UNT-047; VRE; ErmB	>256	>256	16	32	1	2	2	4	>64	32	64	>64	>64	64	>64	>64	>64
Gram-negative	<i>H. influenzae</i> ATCC 49247	4	2	2	4	2	2	2	2	2	4	8	4	4	8	16	4	
	<i>A. baumannii</i> ATCC 19606	16	32	4	16	2	8	8	4	4	4	16	16	4	32	32	32	
	<i>K. pneumoniae</i> ATCC 10031	4	2	4	4	2	8	4	4	2	4	8	16	2	8	8	4	
	<i>E. coli</i> ATCC 25922	64	4	16	32	8	16	16	16	4	8	32	4	8	64	16	8	
	<i>P. aeruginosa</i> ATCC 27853	64	64	64	64	16	32	64	32	64	64	64	64	64	>64	>64	>64	64
	MIC colour scale (µg ml ⁻¹)		<0.03	0.03	0.06	0.125	0.25	0.5	1	2	4	8	16	32	64	128	256	>256

- ❖ Screened 305 compounds against a panel of pathogens comprising of Gram + and Gram – stains.
 - ❖ 83% of the candidates showed a MIC of less than 4µg ml⁻¹ against WT *S. pneumoniae*.
- ❖ Most promising candidates were screened against an expanded panel that had developed various antibacterial resistance strategies.
 - ❖ FSM-100573 and 100563 showed were more active than any current clinical macrolide against mutated *S. pneumoniae*. and *Pseudomonas aeruginosa*.

Conclusions

In conclusion the Meyers group have made several contributions with this publication:

- ❖ Designed and executed several efficient and highly convergent routes to three macrolide skeletons.
- ❖ Further functionalised and furnished these skeletons to generate a synthetic library of 300+ macrolides.
- ❖ Demonstrated that these candidates represent possible avenues to new antimicrobial agents in the war against drug resistant bacteria.