

# Structure-activity relationship for natural tetracenomycin X congeners

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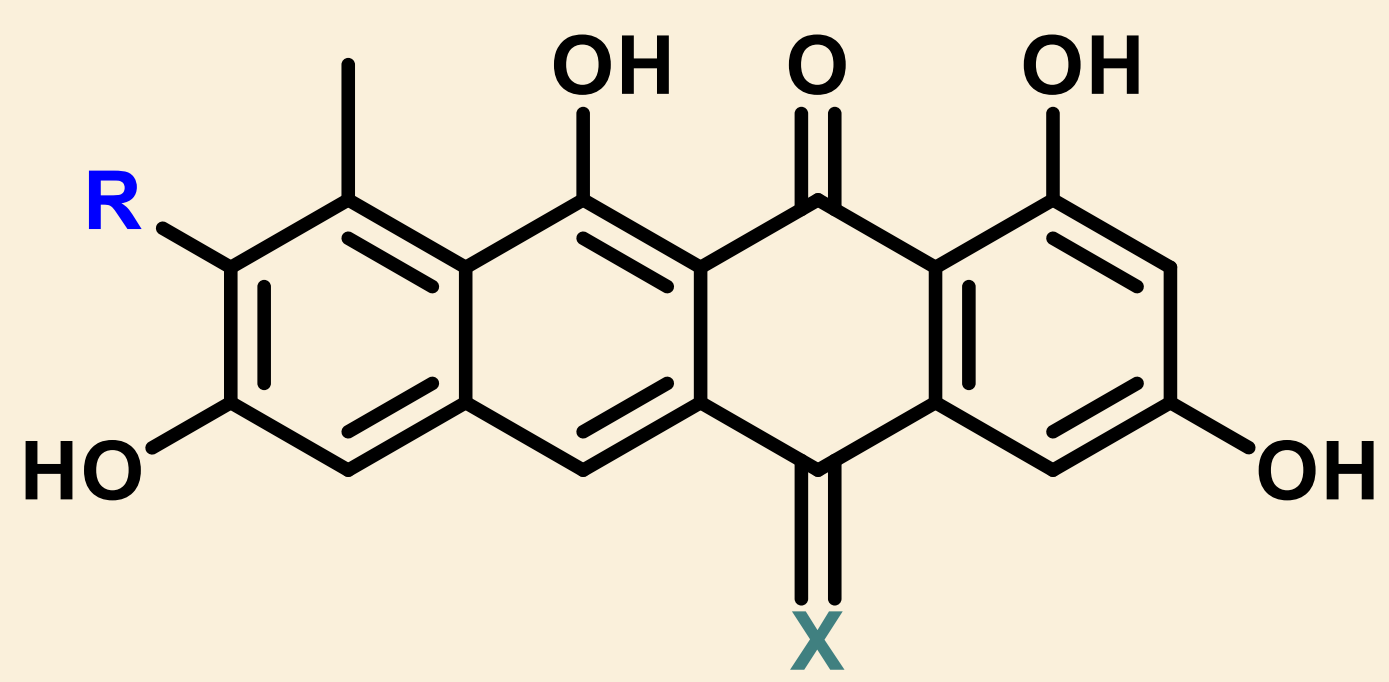
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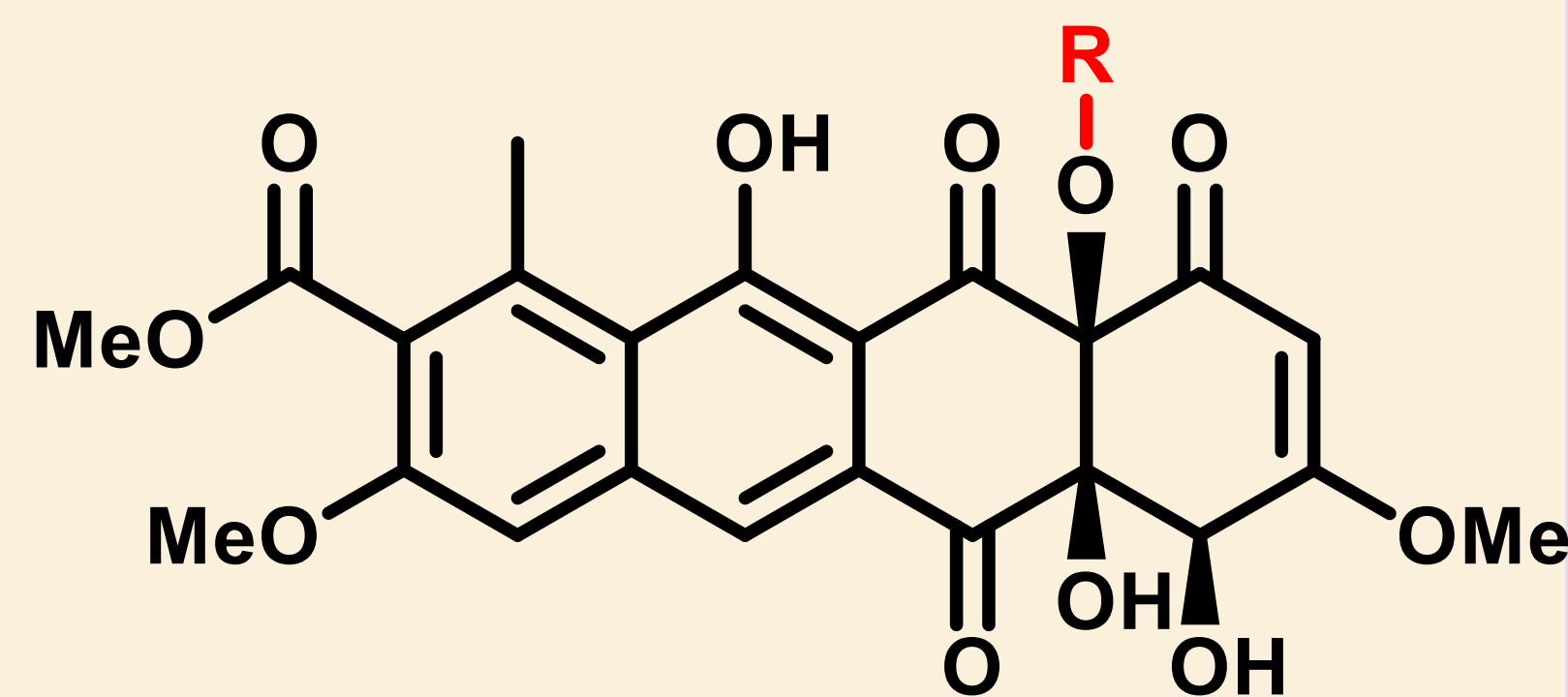
## Tetracenomycin antibiotic family



R = CO<sub>2</sub>Me, X = O, tetracenomycin A2

R = H, X = O, tetracenomycin D

R = CO<sub>2</sub>H, X = H<sub>2</sub>, tetracenomycin F1



R = H, tetracenomycin C

R = Me, tetracenomycin X

**Tetracenomycins (Tcm)** are aromatic polyketides with four fused rings.

Despite structural similarity, only TcmX and TcmC have distinctive 4a,12a-deoxygenation dramatically disrupting the planar structure.

Recently TcmX was shown to have the **unique binding site in the large ribosomal subunit**.

## Isolation of the novel natural teracenomycins

*Amycolatopsis* sp. strain A23

**Cultivation**

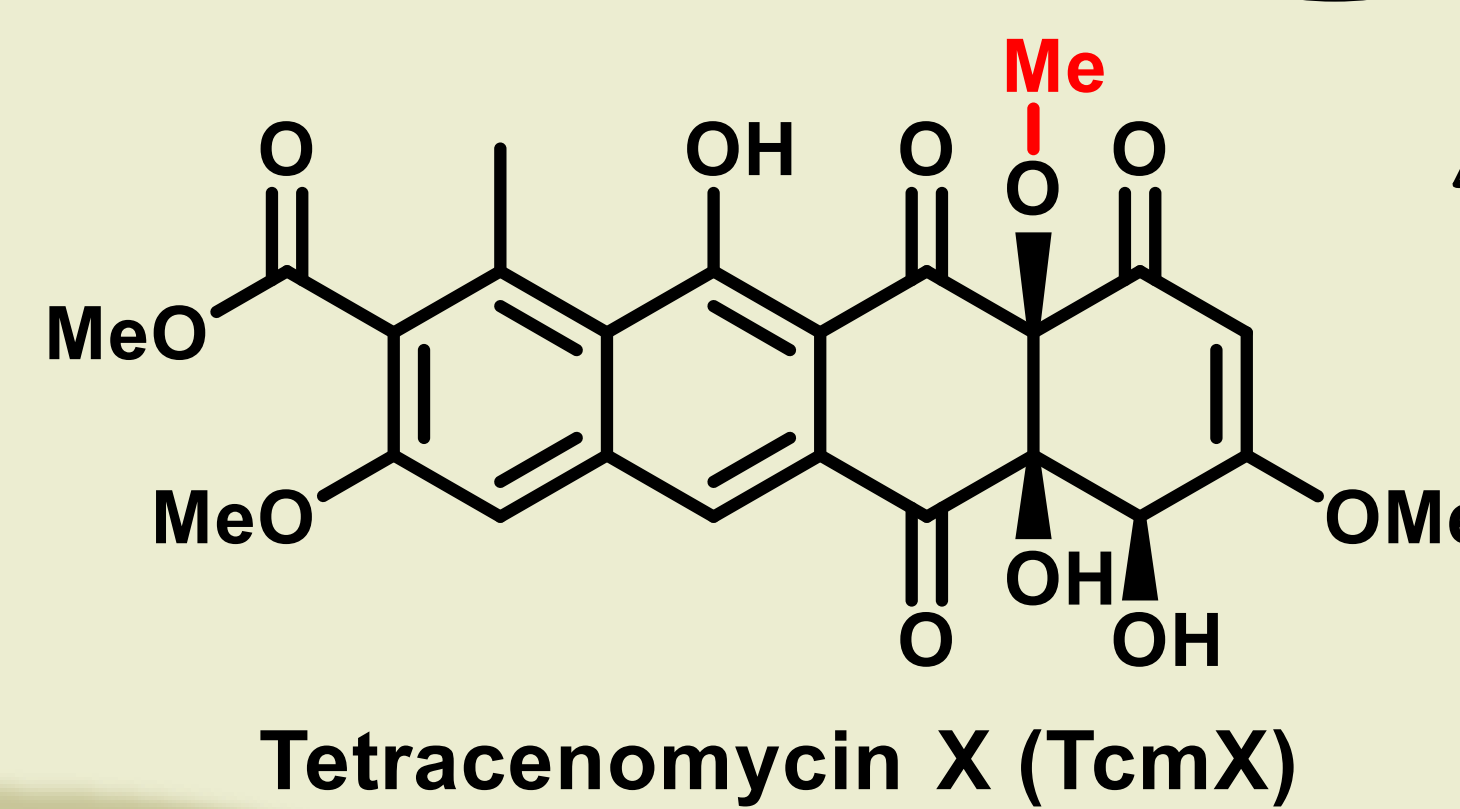


**Extraction**



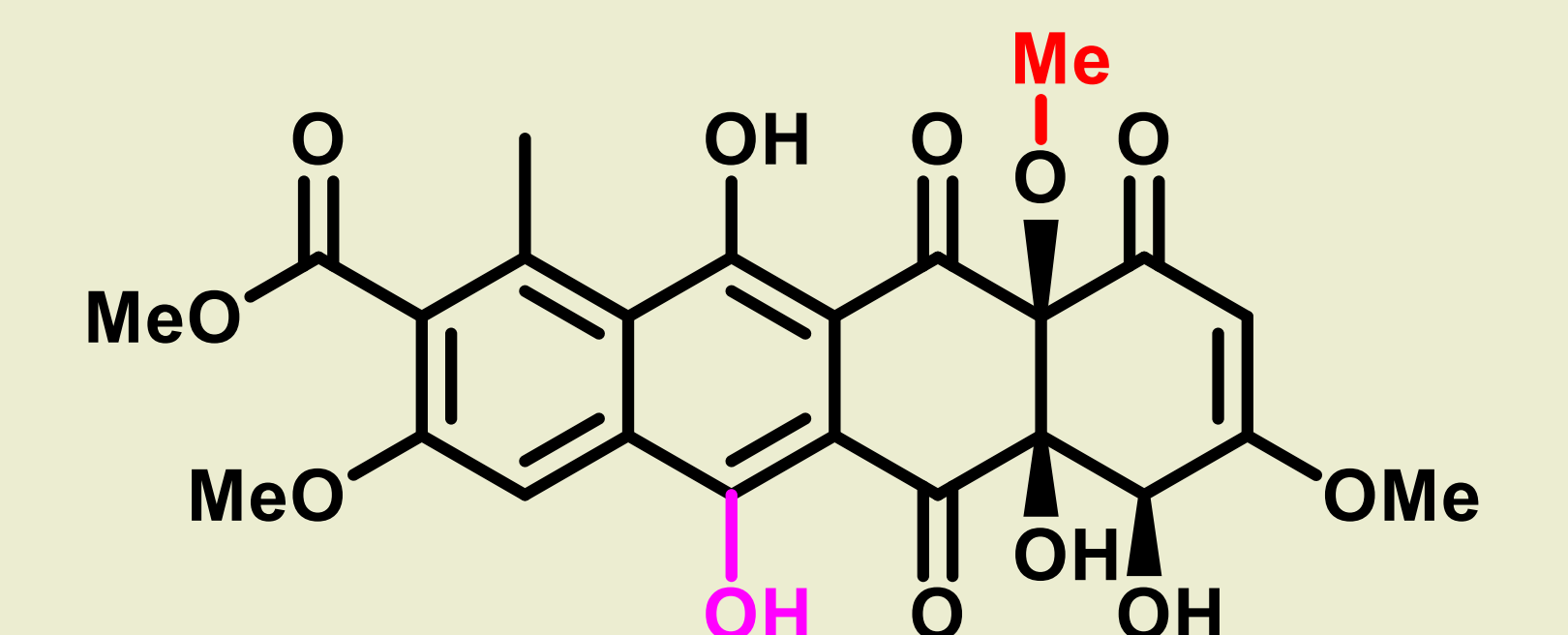
**Chromatography**

2 novel Tcm congeners

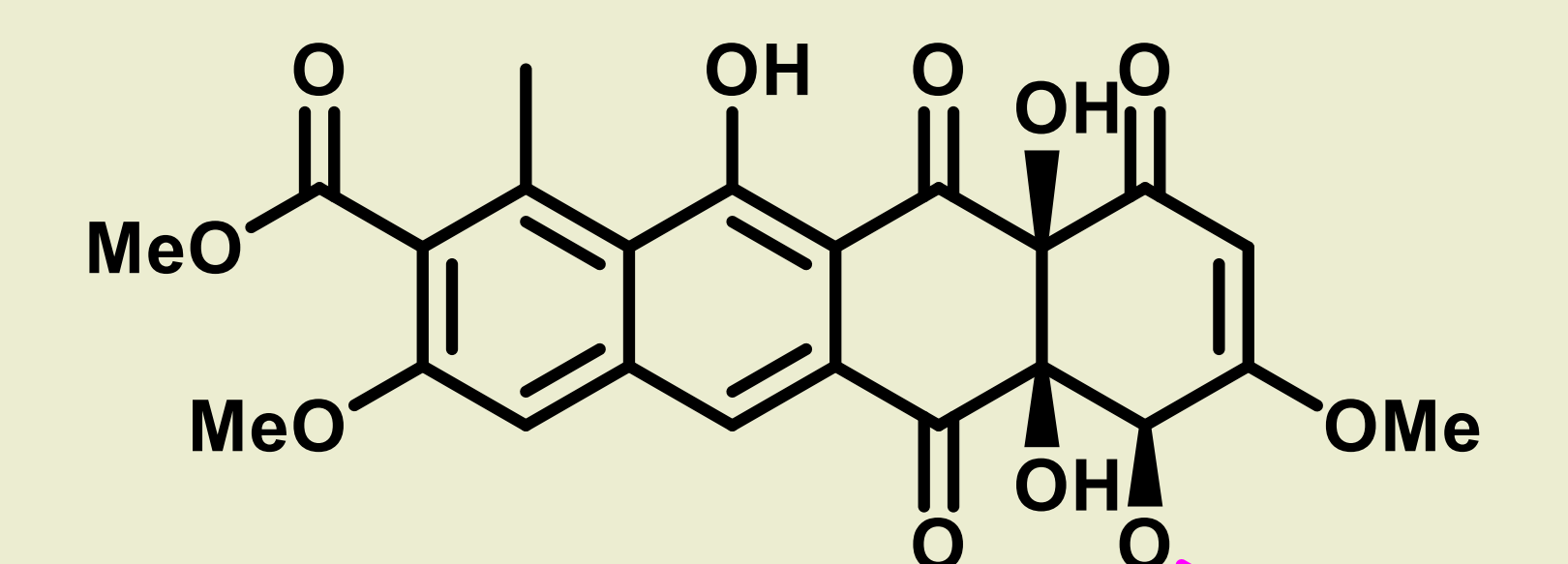


Tetracenomycin X (TcmX)

this work



6-OH-Tetracenomycin X (6-OH-TcmX)



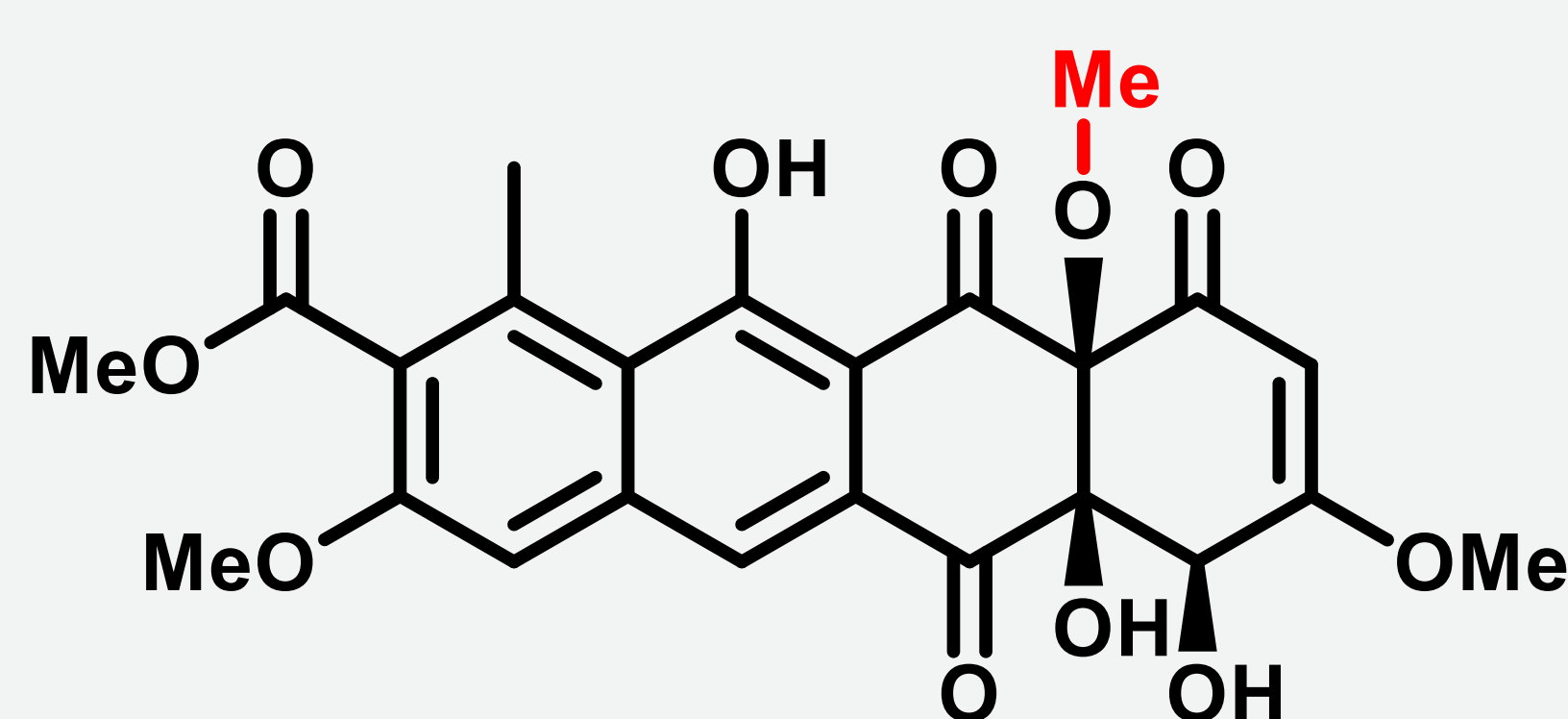
O<sup>4</sup>-Me-Tetracenomycin C (O<sup>4</sup>-Me-TcmC)

STATE OF ART

STRUCTURE ELUCIDATION

## Antibacterial properties

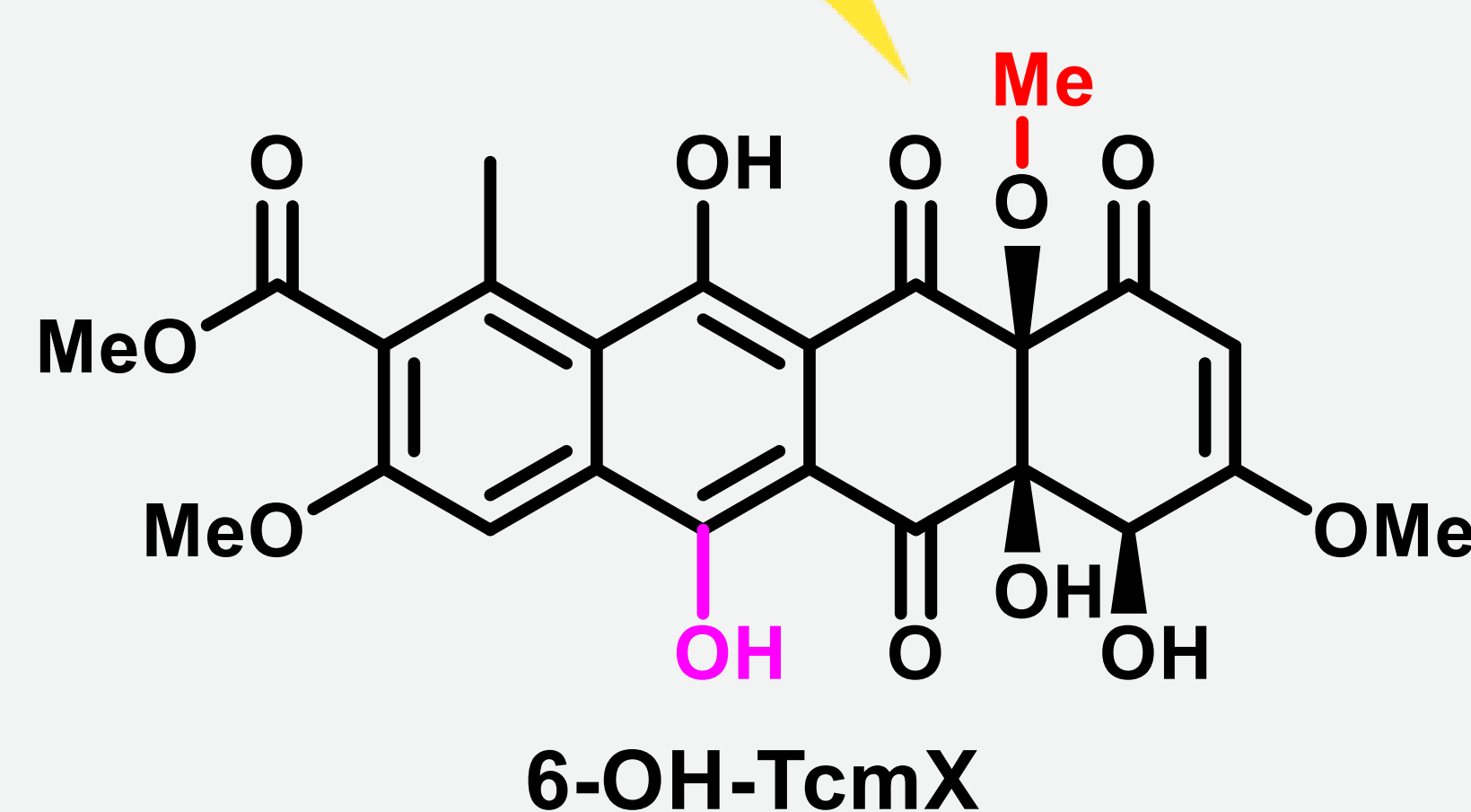
### Translation inhibition



TcmX

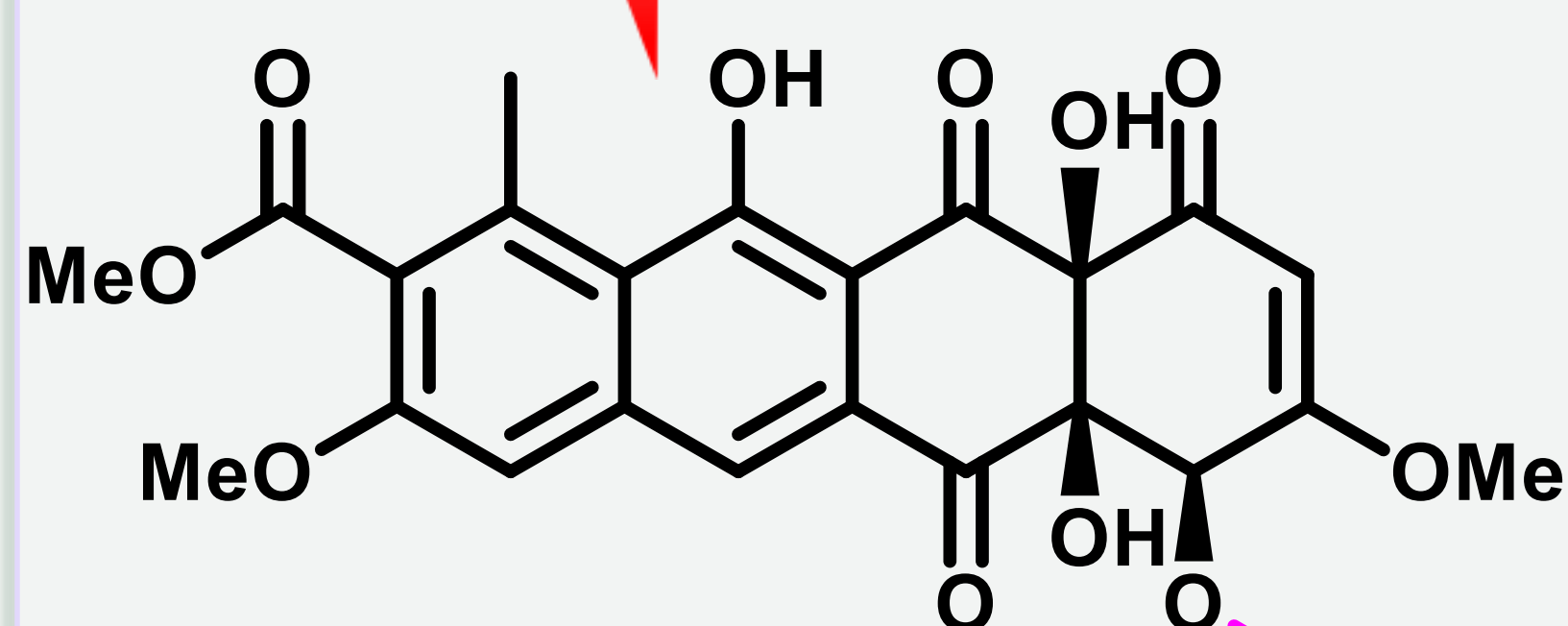
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High antibacterial activity  
Potent translation inhibition



6-OH-TcmX

Reduced antibacterial activity  
Potent translation inhibition



O<sup>4</sup>-Me-TcmC

Inactive  
No translation inhibition

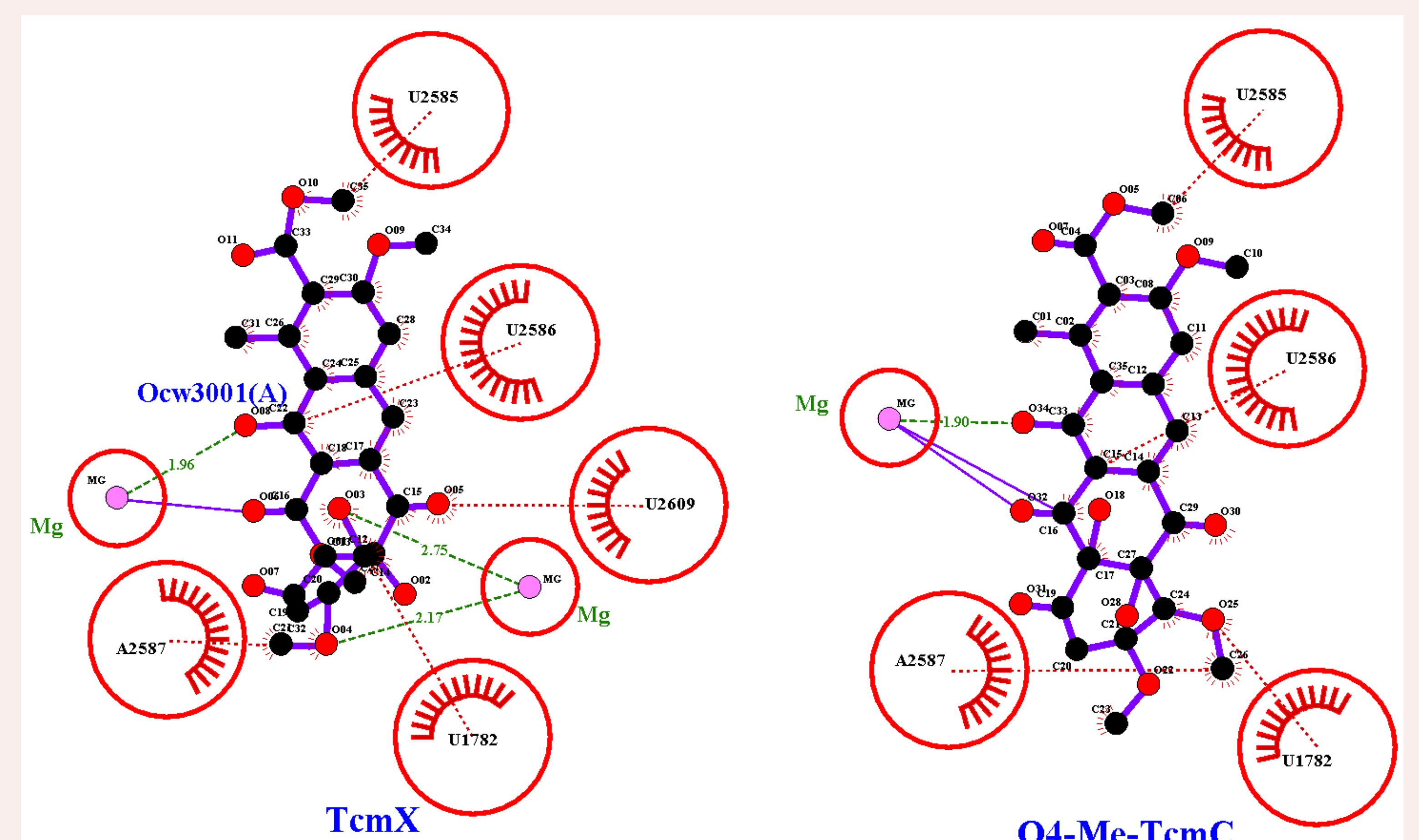
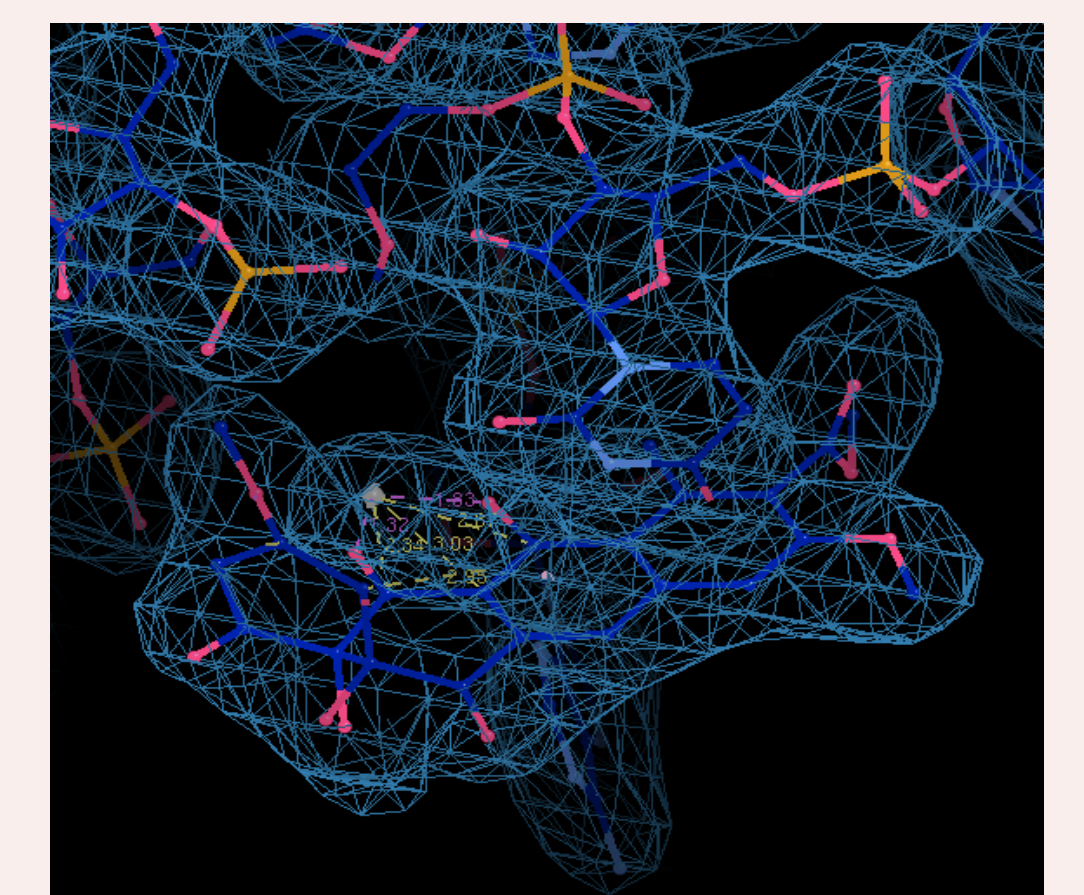
6-hydroxylation impairs activity of 6-OH-TcmX, but **does not interfere with key target interaction sites**

BIOLOGICAL EVALUATION

RIBOSOME BINDING

## Schematic map on the ribosome

O<sup>4</sup>-Me-TcmC fit to cryo-EM TcmX-70S ribosome (PDB 6Y69)



O<sup>4</sup>-Me-TcmC exhibits only a single coordination with magnesium.

Methylation in O<sup>4</sup> position prevents hydrogen bonding, further supporting the idea, that this site is **essential for the ribosome binding**