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Shortcut to Mycothiol Analogues

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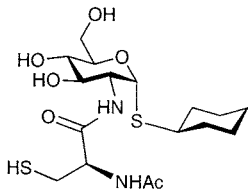
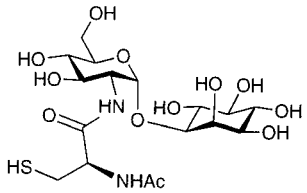
ABSTRACT

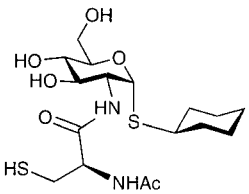


The synthesis of a simplified thioglycosidic analogue (2) of mycothiol (1) is described. Evaluation of 2 against mycothiol S-conjugate amidase from *Mycobacterium tuberculosis* reveals good specific activity (7500 nmol min^{-1} mg-protein^{-1}, vs 14 200 for 1), indicating that 2 can serve as a starting point for antitubercular drug design.

Mycothiol[1] (**1**) is the major low molecular weight thiol found in actinomycetes, including *Mycobacterium tuberculosis*.[2−5] It is thought to protect these organisms against oxidative stress[6−10] and function in the removal of exogenous electrophilic agents.[11−15] The biosynthesis of **1** (Scheme 1) proceeds by way of 1-*O*(2′-acetamido-2′-deoxy-α-D-glucopyranosyl)-D-*myo*-inositol **3**, first by deacetylation to give **4**,[16] and then acylation with L-cysteine under the influence of a ligase[17,18] to provide **5**. A transacetylase[17,19] converts **5** to **1**. Two further pathways involving **1** have been elucidated (Scheme 1). A reductase[6−8] regenerates **1** from the corresponding disulfide, mycothione **(6)**, thus maintaining the reducing intracellular environment. Upon reaction of **1** with electrophiles E^+, the resulting conjugate **7** is cleaved by mycothiol S-conjugate amidase[14,15] into **4** and an *N*-acetylcysteine adduct **8** that is exported from the cell.



1: MSH



2

Drug-resistant tuberculosis now threatens a large portion of earth's population,[20] and the development of new treatments for tuberculosis infection has become a national[21] and international[22] priority. The disruption of enzymatic pathways of mycothiol biosynthesis and/or mycothiol-based detoxifi-

[†] National Institutes of Health.

(1) Systematic name: 1-*O*(2′[*N*-acetyl-L-cysteinyl]amido-2′-deoxy-α-D-glucopyranosyl)-D-*myo*-inositol.

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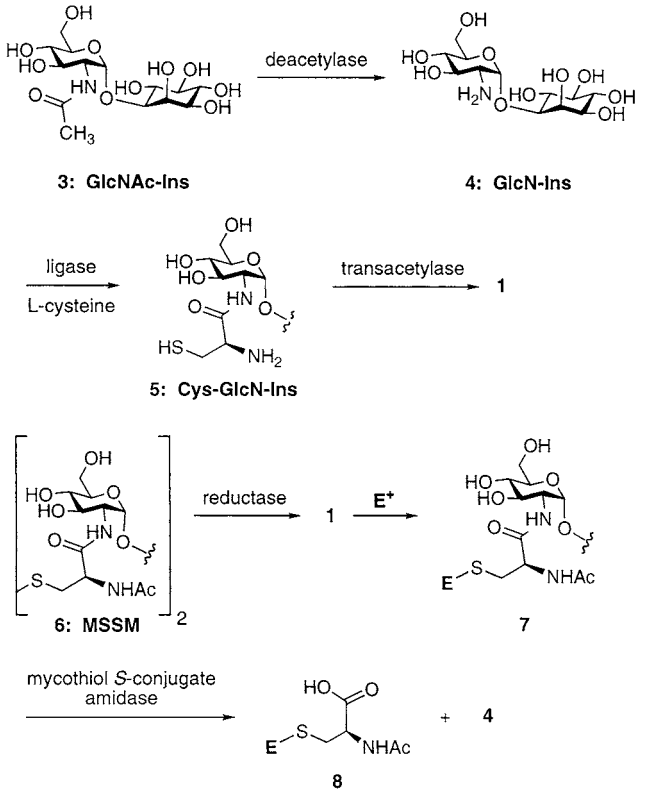
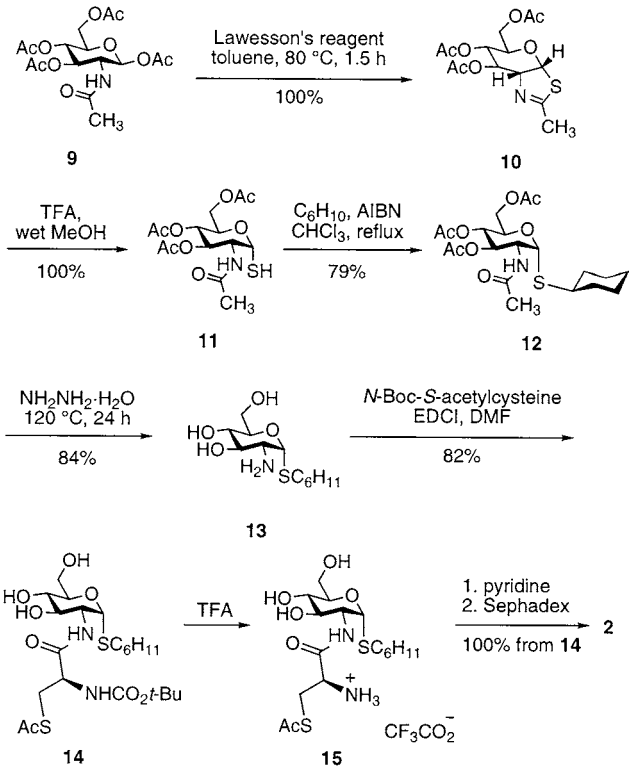
Scheme 1. Enzyme-Mediated Mycothiol Pathways



cation could leave *M. tuberculosis* vulnerable to drugs, oxygen, and other stress factors. The enzymes shown in Scheme 1 accept substrates or produce products that are N-acylated 1-*O*(2′-amino-2′-deoxy-α-D-glucopyranosyl)-D-*myo*-inositols. For this reason, compounds based on the GlcN−Ins substructure that additionally bear groups on N that resemble those of the respective transition states are potential inhibitors for any one, or more than one, of these enzymes. In previous synthetic studies on **1** and related compounds,[6,15,17,23] the preparation of a protected D-*myo*-inositol glycosylation acceptor has required several steps and a resolution, and both the inositol α-glycosylation and N-acylation steps have been problematic. Some of these synthetic difficulties could be dodged if a stripped-down version of the inositol ring could be used as a component of inhibitors. We recently found that a variety of α-GlcNAc thioconjugates can be prepared stereoselectively and in good yield by *S*-derivatization of an α-GlcNAc mercaptan.[24,25] Thioglycosides are generally more resistant to degradation by glycosidases than *O*-glycosides,[26−28] so this approach to inhibitor design combines several possible advantages.

Commercially available 2-acetamido-2-deoxy-*β*-D-glucopyranose tetraacetate (**9**, Scheme 2) was treated with Lawesson's reagent as described previously,[24,26] and the resulting thiazoline **10** was then hydrolyzed to the acetamido mercaptan **11**.[24] Reaction of **11** with cyclohexene under conditions for free radical addition of anomeric mercaptans to alkenes [chloroform as a cosolvent, azobis(isobutyronitrile) as a radical initiator][25] afforded the cyclohexyl thioglycoside **12** with no trace of the corresponding *β*-isomer. Hydrazinolysis[29] of the four acetyls provided aminotriol **13**, and then coupling with *S*-acetyl-*N*-Boc-L-cysteine[30] gave **14** in good

Scheme 2. Synthesis of Simplified Mycothiol Analogue **2**



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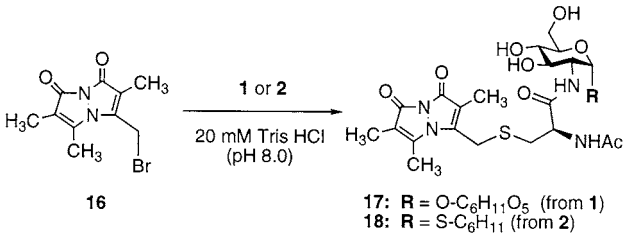
yield and high isomeric purity. The *N*-Boc protecting group was removed by treatment of **14** with neat trifluoroacetic acid, leading to ammonium salt **15**, and then basification with pyridine in the same pot gave the simplified mycothiol analogue **2** as the result of a spontaneous and convenient intramolecular S-to-N acetyl migration.[31]

Evaluation of **2** as a substrate for *M. tuberculosis* mycothiol S-conjugate amidase[14,15] was carried out by prefatory S-alkylation with bromobimane (**16**, Scheme 3) under mildly

Scheme 3. Preparation of Bimane Derivatives



basic conditions.[15] The resulting bimane derivative **18** was subjected to cleavage by the amidase in parallel with mycothiol-bimane **17**, while monitoring formation of the cysteine-*S*-bimane product (see **8**) by fluorescence-detected HPLC assay.[15] Specific activities for **18** and **17** are 7500 and 14 200 nmol min^{-1} mg-$protein^{-1}$, respectively, establishing **18** as a good substrate for this amidase. Neither the inositol hydroxyls nor the glycosidic linking atom (O vs S) plays a major role in enzyme binding. An earlier study[6] had indicated that the inositol ring is not required for reduction of disulfides (see **6**) by the *M. tuberculosis* mycothione reductase. The accumulated information thus suggests that **2**, which dispenses with the inositol hydroxyls and the linking oxygen atom, can serve as a suitable foundation upon which to base inhibitor design.

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Supporting Information Available: Experimental details and spectral characterization for all new compounds and description of the enzyme assay. This material is available free of charge via the Internet at http://pubs.acs.org.

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