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Total Synthesis of Tiacumicin B: Implementing H-bond-Directed Acceptor Delivery for Highly Selective β -Glycosylations

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Dedicated to Professor Henri-Philippe Husson

Abstract: We report a total synthesis of tiacumicin B, a natural macrolide whose remarkable antibiotic properties are used to treat severe intestinal infections. The strategy is in part based on our experience of the synthesis of the tiacumicin B aglycone, and on the decisive use of sulfoxides as anomeric leaving-groups in H-bond-mediated Aglycone Delivery (HAD). This new HAD variant permitted highly β -selective rhamnosylation and noviosylation. To increase convergence, the rhamnosylated C1-C3 fragment thus obtained was anchored to the C4-C19 aglycone fragment by adapting the reliable Suzuki-Miyaura cross-coupling used for the aglycone synthesis. Ring-size selective macrolactonization provided a compound engaged directly in the noviolysation step with a virtually total β -selectivity. The final efficient removal of all the protective groups (PGs) provided synthetic tiacumicin B.

Tiacumicin B (Tcn-B, 1) - also known as clostomicin B1, fidaxomicin or lipiarmycin A3 - (Scheme 1), is a molecule isolated for the first time in 1975 from Dactylosporangium aurantiacum, an actinobacterium.^[1] Consisting of an 18-membered macrolactonic core decorated by two rare sugars (D-noviose and D-rhamnose) attached through β glycosidic bonds, displaying 14 stereogenic centers and several polysubstituted alkenes, Tcn-B is therefore one of the most complex antibiotic-macrolides known. A high degree of synthetic difficulty eventually of course arises from such a structural complexity, whose the thorny problem of the 1,2-cis glycosylations.^[2] Resistance to antibiotics has become a serious biomedical risk and a major threat to public health with sever impacts on the economy. In this context the finding of new antibiotics with new biological targets is essential. Tcn-B is one of them and received a fast-track FDA approval in 2011 for the treatment of frequently nosocomial and fatal gut infections associated with Clostridium difficile.[3] Tcn-B eradicates bacteria by inhibiting the RNA polymerase, targeting the "switch-region".^[4] Cross-resistance with other antibiotics is very unlikely,^[5] even with rifamycin because, although close, the domains targeted do not overlap, so that rifamycin-resistant forms of Mycobacterium tuberculosis remain highly sensitive to Tcn-B.^[6] The demand for

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Tcn-B analogues is therefore strong, which implies setting reliable total synthesis pathways. The daunting structure of Tcn-B being in addition very inspiring in the eyes of synthesis chemists, several renowned research groups have embarked in the adventure of its total synthesis.^[7] In 2015 Gademann^[8] and Altmann,^[9] published the firsts syntheses of the aglycone of Tcn-B, Zhu^[10] reporting the synthesis of a diastereomer. Nonetheless, only Gademann had so far been capable of completing the total synthesis of Tcn-B, bringing the following answers to the β -glycosylations problem:^[11,12] a) The Helferich's protocol (activating agent: HgO(excess)/HgBr2(cat))^[13] was used to noviosylate the C4-C13 aglycone fragment (α/β : 1/3, β : 54%), as the cyclic aglycone gave only α adducts whatever conditions used. b) The rhamnosylation was carried out on the macrolide, using an imidate donor (α/β : 1/4, β : 62%). Following similar strategies, Gademann also synthesized three congeners of Tcn-B: tiacumicin A,^[12] and mangrolide A^[14] and D.^[15] In 2019, de Brabander also reported a total synthesis of mangrolide D.^[16] For our part, we reported two related synthetic pathways leading to the aglycone.[17] The original strategy we designed for the synthesis of its C12-C15 diene region resulted in the discovery that Pd-nanoparticles catalyze the Kumada-Corriu reaction of vinylsulfides,^[18] and led us to propose a mechanism for the Grigg's allene/alkyne cross-coupling based on DFT calculations.[19]



Scheme 1. Tiacumicin B (1) and our retrosynthetic analysis

The new total synthesis of tiacumicin B (1) we depict here is based on strategic and methodological innovations that allowed an original and selective assemblage of the three main regions of the molecule (Scheme 1). It naturally relies on our synthesis of the Tcn-B aglycone, whose strategy had been designed with an eye to the total synthesis. Originally, we had imagined glycosylating our aglycone sequentially, a probably viable pathway. However, we finally opted for a less traditional but more convergent retrosynthetic plan in which Tcn-B was disconnected into fragments A, B, and the known 2,^[17a] equivalent in size and complexity. We chose to assemble fragment A together with fragment 2 first, then close the macrolactone, and install fragment **B** at the very end. The Suzuki coupling developed during our aglycone synthesis^[17a] was considered robust enough to allow assembling fragments 2 with fragment A, instead of the small silvlated C1-C3 fragment formerly involved,^[17a] adding thus convergence. This approach reduces risks of failure since rhamnosylation conditions needed for the synthesis of A can be developed using as acceptor the structurally simple C1-C3 fragment. Complete ring-size selectivity had been observed previously during the aglycone macrolactonization. Applied to this total synthesis, this step should provide a monoglycosylated macrolactone bearing at C11 an unprotected OH directly ready for the β -noviosylation step. This exciting scenario was nonetheless uncertain for at least three reasons: a) noviosylation is a very late step, b) a high β -selectivity is required, and c) this macrolactone has been described as a reluctant glycosylation acceptor.^[12]

To prepare fragment **A** we considered using a β -selective glycosylation of acceptor $18^{[20]}$ (Scheme 3) by the phenylthiorhamnosyl donor 16 that bears a picoloyl group (Pico) at *O*-3 (Scheme 2).



Scheme 2. Syntheses of rhamnopyranosyl donors 16 and 17. TFA: trifluoroacetic acid, NapBr: 2-naphthalene-methyl-bromide, LDA: lithium di-*iso*-propylamide, CSA: camphorsulfonic acid, PTSA: paratoluene sulfonic acid, DCC: dicyclohexylcarbodiimide, DMAP: 4-dimethylaminopyridine, *m*-CPBA: 3-chloroperoxybenzoic acid.

As described by Demchenko,^[21] a remotely positioned Pico group can direct, through intermolecular H-bonding, a selective facial attack on the glycosyl donor. We started with the synthesis of the homodichloro orsellinate attached to the rhamnoside. Commercially available orcellinate **3** was dichlorinated, and both phenols were TBS-protected providing **4**. The benzylic methyl group was deprotonated with LDA then methylated giving **5**. Acidic hydrolysis led to deprotected carboxylic acid **6** that upon a treatment with acetone and Tf₂O in trifluoroacetic acid furnished cyclic ester **7**. The remaining free phenol of **7** was protected, supplying 2-naphthalene-methyl (Nap) ether **8**.

We then addressed the rhamnosyl donor synthesis starting from phenyl-2,3,4,6-tetra-*O*-acetyl-1-thio- α -D-manopyranoside **9** (Scheme 2). Diol **10** was obtained through Zemplén deacetylation and 4,6-benzylidene formation. Then **10** was selectively TBS-protected at *O*-3, methylated at *O*-2, and deprotected furnishing triol **11**. Selective tosylation in 2,6-lutidine gave **12**, whose reduction with LAH provided the desired rhamnose derivative **13**. Gademann's conditions allowed assembling ester **8** together with diol **13** giving selectively **14**.^[11] The free phenol of **14** was protected as a Nap ether **15**, and the rhamnoside *O*-3 position was esterified with picolinic acid. This led to the expected donor **16** ready for the glycosylation of alcohol **18**^[22] (Scheme 3).



Scheme 3. Glycosylation conditions and synthesis of fragment A. NIS: *N*iodosuccinimide, DTBMP: 2,6-di-terbutyl-4-methylpiridine, ADMB: 4-allyl-1,2dimethoxybenzene.

Initial glycosylation attempts carried out in 1,2-dichloroethane with dimethyl(methylthio)sulfonium triflate, a classical promoter of Hbond-mediated Aglycone Delivery (HAD),^[21] were disappointingly unsuccessful. First promising coupling results of donor **16** with acceptor **18** were obtained using *N*-iodosuccinimide (1 equiv/donor) and triflic acid (0.92 equiv/donor)^[23] in CH₂Cl₂ at – 40 °C to rt and produced **19** (76%, α/β : 1/4),^[24] whose anomers was separated by prep-HPLC. Seeking much higher β -selectivity, we shifted to sulfoxide **17** featuring an anomeric leaving-group never used before in HAD. In this case, donor **17** (1.7 eq.) was activated using Tf₂O in CH₂Cl₂ at -70 °C in the presence of the acceptor **18**, DTBMP, and ADMB.^[25, 26] We were pleased to find that the desired glycosylated compound **19** was formed with a high facial selectivity (α/β : 1/20), with 64% yield. To verify that this glycosylation well took place through HAD, we made a control experiment using donor **17**, an analog of **17** whose picoloyl was replaced by a benzoyl. Product **19**' was the only one to form here, its α configuration indicating that the picoloyl group on donor **17** could direct remotely the nucleophilic attack to the β -face likely thanks to the formation of a H-bond. The picoloyl was easily removed from **19** with Cu(OAc)₂, and replaced by a TBS to lead to key fragment **A** (**21**).

Our synthesis of aglycone fragment **2** was robust enough to be scaled up.^[17a] We were pleased to see that the cross-coupling of **2** with rhamnoside **21** proceeded cleanly requiring only slight modifications of formerly used the conditions. This convergent step provided ester **22** in a 79% yield (Scheme 4).



Scheme 4. Assemblage of fragments 2 and A, and macrolactonization. MPM: 4-Methoxyphenylmethyl, Ruphos: CAS number: 787618-22-8.

A critically important selectivity was needed to convert 22 into seco-acid 23 since a second ester function was present on the rhamnoside moiety. This was achieved by using Me₃SnOH in toluene at 120 °C giving 23 with yields of 70 to 93%.^[27] With seco-acid 23 in hand, we focused on the macrolactonization expecting again ring-size selectivity. However, the strict transposition of the Yamaguchi conditions^[28] we had used before, led to the desired hemi-glycosylated tiacumicin B 24 with only 23% yield. The Boden-Keck's protocol,^[29] allowed reaching 58% yield, but 24 proved to be a mixture of two products resulting from the isomerization of the C4-C5 alkene. Finally, the Shiina's conditions^[30] allowed a far cleaner and reproducible macrolactonization into 24 with 72% yield, and an isomerization minimized at 15%. Our strategy allowed keeping the OH at C11

free of protective-group throughout this synthesis, so that **24** could be directly engaged in the noviosylation step.

At first, to secure good β -selectivity, we had programmed anchoring the noviose using a silicon tether delivery.^[31] This strategy failed in our case, so we decided implementing again an HAD approach installing a directing picoloyl group at O-3 of the sugar. The required noviosyl donor was prepared from D-arabinose 25 (Scheme 5) which was transformed into lactone 26 through selective acetonide protection, oxidation of the lactal, and silvlation of the remaining OH group. Lactone 26 was treated with MeMgBr, then oxidized into lactone 27. Dibal-H reduction of 27 led to the corresponding lactol whose acetonide protection was removed under mild acidic conditions leading to 28. A thiophenyl group was then installed at the anomeric position giving diol 29, which was protected as bis-dichloroacetyl ester 30. Treatment by (iPrCO)₂O and a catalytic amount of Sc(OTf)₃ in MeCN allowed the direct replacement of the TBS group by an isobutyrate leading to 31.^[32] The two dichloroacetates of 31 were then selectively removed with sym-collidine giving diol 32. Through the intermediate stannylene of 32, and a NapBr/CsF treatment, a Nap PG was introduced at the O-2 position leading to alcohol 33 with an unexpectedly high selectivity.[33]



Scheme 5. Noviosyl donor synthesis. TEMPO: 2,2,6,6-Tetramethyl-1piperidinyloxy, TCCA: trichloroisocyanuric acid.

Alcohol **33** was esterified with picolinic acid using DCC to give sulfide **34**, the desired noviosyl donor. Preliminary trials using (+)menthol as acceptor led predominantly to the β -adduct (α/β : 1/5) in 72 % yield demonstrating the feasibility of this approach. Unfortunately, the reactions with macrolactonic acceptor **24** was unsuccessful, no glycosylated adduct being detected.^[12] The success of the above-mentioned rhamnosylation led us to consider that sulfoxide **35** could be a superiorly reactive donor. Under our activation conditions (Tf₂O activation of **35** in the presence of **24**, DTBMP, ADMB/CH₂Cl₂, -70 °C) we were pleased to isolate the desired noviosylated product **33** with 68% yield, with a virtually total facial selectivity ($\alpha/\beta > 1/20$).^[34]

The very last steps of this total synthesis consisted in removing seven PGs from compound **36**: 2 MPMs, 3 Naps, 1 Pico and 1 TBS.

First, the TBS group located on the rhamnoside moiety was cleaved using HF.NEt₃ giving alcohol **37** (Scheme 5). The 2 MPMs,



Scheme 6. Noviosylation and PGs removal. DDQ: 2,3-dichloro-5,6-dicyano-1,4-benzoquinone.

as well as the Nap located at O-2 on the novioside, were readily oxidized by DDQ at 0 °C leading to tetraol **38** in 3 hours. However, the two Naps on the phenol functions of the rhamnoside proved resistant to these smooth conditions, and an extended reaction time at r.t. produced an intractable mixture of products. Nonetheless, the removal of the Pico was cleanly carried out (Cu(OAc)₂/methanol-CH₂Cl₂, 0 °C) giving **39**. Importantly, this Pico group has to be removed after its neighboring Nap otherwise DDQ would lead to the formation of a 2-naphthylmethylidene bridge over O-2 and O-3 of the novioside. These three operations proved very clean and were performed without intermediate purification giving **39** with an overall 74% yield (94% *per* PG). Finally, we had to address what turned out to be a tricky final problem: the cleavage of the two reluctant Nap groups protecting the phenol functions. Mild Pdcatalyzed hydrogenation conditions failed at being selective.^[35] However, we had previously observed that Suzuki cross-coupling conditions used to create the C3-C4 bond of the aglycone led, when conducted at 80 °C, to a partial loss of the Nap groups of these phenols. Exploiting this, we developed Pd-catalyzed $(Pd_2(dba)_3/4.PPh_3,$ 1,3-dimethylbarbituric conditions acid. pyridine/DMF, 80 °C) that ultimately provided tiacumicin B (1) cleanly and with a good yield. This 4-steps removal of the 7 PGs took place with 55.5% overall yield (91.9% per PG). The physicochemical data of our synthetic Tcn-B are strictly identical to those of the naturally occurring compound; $[\alpha]^{23}_{D} = -5.6 \text{ deg}$ cm³ g⁻¹dm⁻¹ (c= 0.41 g/100 cm³, MeOH), lit.:^[1b] ($[\alpha]^{23}_{D} = -5.5$ deg cm³ g⁻¹dm⁻¹ (c=1.98 g/100 cm³, MeOH).

In summary, we have achieved the total synthesis of tiacumicin B (1), with as salient steps a highly β -selective rhamnosylation, a Suzuki cross-coupling that allowed assembling the rhamnoside 21 with aglycone fragment 2, a ring-size selective macrolactonization, a final and virtually totally selective β -noviosylation of the cyclic aglycone, and a successful removal of all PGs. The remarkable facial selectivity of both glycosylations relies on an H–bond-directed effect of a remote 3-*O*-picoloyl group set on the incoming glycosyl acceptors, and the conjoint use of a phenylsulfoxide leaving-group. We believe that this new variant of the Demchenko procedure will prove useful to address the biological relevance of the carbohydrate moieties of tiacumicin B or other sensitive aglycones through the preparation of a set of glycosylated analogues.

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