

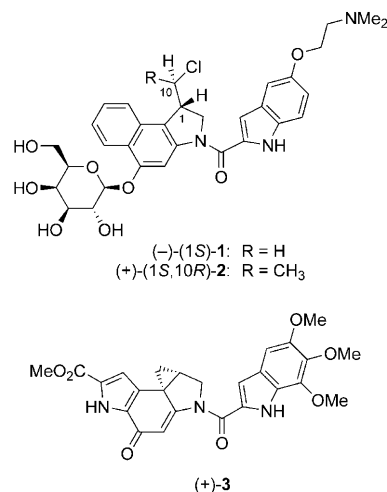
Glycosidic Prodrugs of Highly Potent Bifunctional Duocarmycin Derivatives for Selective Treatment of Cancer**

Lutz F. Tietze,* J. Marian von Hof, Michael Müller, Birgit Krewer, and Ingrid Schuberth

Dedicated to Professor Horst Kunz on the occasion of his 70th birthday

The development of selective chemotherapeutics for treatment of cancer is an important current area of medicinal research. One approach for the reduction of the frequently dose-limiting adverse effects of the currently available cytostatics is the antibody-directed enzyme prodrug therapy (ADEPT),^[1,2] in which in a binary approach a conjugate of a tumor-specific antibody and an enzyme in combination with a hardly toxic prodrug is used. By means of the enzyme in the conjugate the prodrug is selectively cleaved in the tumor tissue leading to the formation of a highly cytotoxic compound. The advantages of the ADEPT approach compared to the use of conjugates of antibodies and toxins lies in the far better diffusion of the low-molecular-weight active compound formed in the malignant tumor, the killing also of those cancer cells to which the antibody does not bind, and the enzymatic approach, which gets along with a comparatively low amount of the antibody. One problem of the prodrugs previously developed for ADEPT was that their cytotoxicity was frequently only insignificantly lower than that of the active compounds formed from them (QIC₅₀ value; QIC₅₀ = IC₅₀ of the prodrug/IC₅₀ of the prodrug in the presence of the enzyme). Furthermore, the efficacy of the active compound formed was frequently not adequate. We have therefore suggested as minimal requirement a QIC₅₀ value of > 1000 and an IC₅₀ value of the active compound formed of < 10 nM.^[3]

With the compounds (–)-(1*S*)-**1** and (+)-(1*S*,10*R*)-**2** derived from seco analogues of the natural cytostatic duocarmycin SA ((+)-**3**;^[4] Scheme 1) we have recently succeeded in developing prodrugs, which with QIC₅₀ values of 3500 and 4800, respectively, and IC₅₀ values of the active



Scheme 1. Glycosidic prodrugs (–)-(1*S*)-**1** and (+)-(1*S*,10*R*)-**2**, derived from the natural cytostatic antibiotic duocarmycin SA ((+)-**3**).

compounds formed of 16 and 750 pM, respectively, best met the requirements of the previously named criteria.^[5,6]

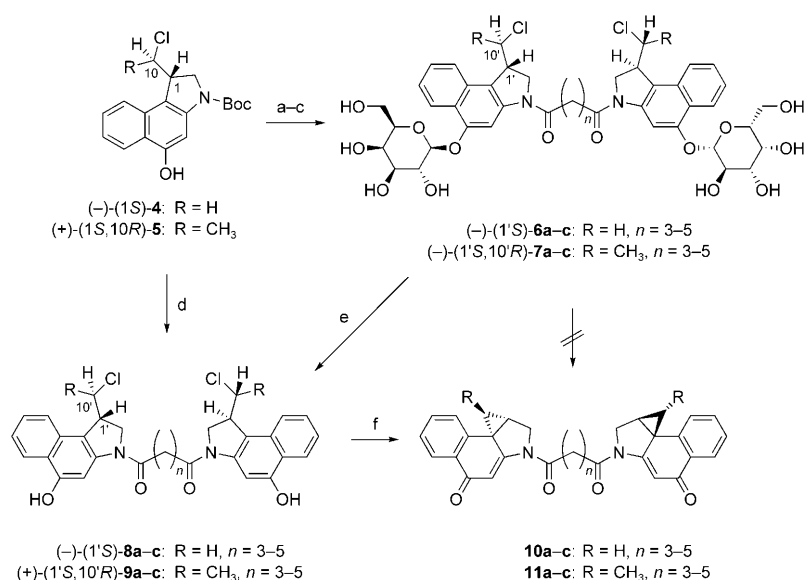
Herein we describe new prodrugs and the active compounds derived from them, which with QIC₅₀ values of up to almost 1000000 and IC₅₀ values of the active compounds formed of up to about 100 fM put all previous results well into the shade.^[3,5–7] The new compounds contain the same pharmacophore as (–)-(1*S*)-**1** and (+)-(1*S*,10*R*)-**2**, but in this case, however, two of each are coupled through a dicarboxylic acid. The length of the dicarboxylic acid used here has a considerable influence on the biological activity of the prodrugs and the resulting active compounds.^[8]

The preparation of the new glycosides (–)-(1'*S*)-**6a–c** (*n* = 3–5) and (–)-(1'*S*,10'*R*)-**7a–c** (*n* = 3–5) was carried out starting from the enantiomerically pure *N*-Boc-protected seco compounds (–)-(1*S*)-**4**^[6,9] and (+)-(1*S*,10*R*)-**5**,^[5,10] respectively, by glycosidation using the trichloroacetimidate method,^[11] subsequent BF₃·OEt₂-mediated *tert*-butyloxycarbonyl(Boc) deprotection and coupling of the secondary amines formed with the respective dicarboxylic acid chlorides (Scheme 2). Solvolytic cleavage of the acetyl protecting groups by using the method applied by Zemlén and Pacsu finally afforded the galactosides (–)-(1'*S*)-**6a–c** (*n* = 3–5) and (–)-(1'*S*,10'*R*)-**7a–c** (*n* = 3–5) in yields of 27–55% over three steps.^[12] The corresponding secodugs (–)-(1'*S*)-**8a,c** (*n* = 3 and 5) and (+)-(1'*S*,10'*R*)-**9a–c** (*n* = 3–5) were obtained in 65–80% yield from (–)-(1*S*)-**4** and (+)-(1*S*,10*R*)-**5**, respectively by cleavage of the Boc protective

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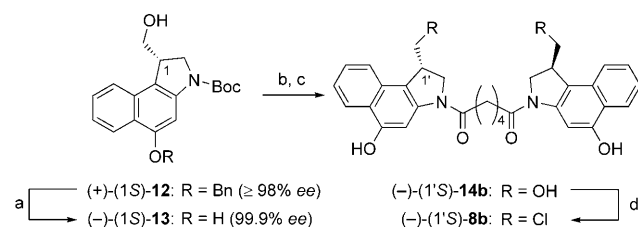
Supporting information for this article is available on the WWW under <http://dx.doi.org/10.1002/anie.201002502> or from the author.



Scheme 2. Synthesis of the prodrugs (a–c) and seco-drugs (d; not for (-)-(1'S)-8b), as well as the enzymatic activation of the prodrugs to the active compounds (e, f): a) α -trichloroacetimidate of D-galactose, BF₃·OEt₂ (0.5 equiv), CH₂Cl₂, MS (4 Å), -10°C, 3.5 h, then BF₃·OEt₂ (3.0 equiv), RT, 5.5 h; b) ClCO(CH₂)_nCOCl (n = 3–5), NEt₃, DMF, RT, 20 h, then preparative HPLC: Kromasil 100 C18 (250 × 20 mm, 7 μ m), A = H₂O, B = MeOH, gradient(A/B) = 30:70 → 0:100 in 6.5 min; c) NaOMe/MeOH, RT, 2 h, then preparative HPLC: Kromasil 100 C18 (250 × 20 mm, 7 μ m), A = H₂O, B = MeOH, gradient(A/B) = 70:30 → 0:100 in 15 min; d) 4 N HCl/EtOAc, RT, 3 h, then ClCO(CH₂)_nCOCl (n = 3–5), pyridine, DMF, RT, 20 h, then preparative HPLC: Kromasil 100 C18 (250 × 20 mm, 7 μ m), A = H₂O, B = MeOH, gradient(A/B) = 30:70 → 0:100 in 6.5 min; e) glycoside cleavage of the prodrugs to the seco-drugs by β -D-galactosidase; f) in situ Winstein cyclization of the seco-drugs to the active compounds under physiological conditions. DMF = N,N-dimethylformamide.

group and reaction with the respective dicarboxylic acid chlorides. Surprisingly the preparation of the seco-drug (-)-(1'S)-8b (n = 4) was not successful in this way neither in acceptable yields nor in pure form.

Thus for (-)-(1'S)-8b an alternative synthetic route was developed starting from (+)-(1S)-12,^[6,13] a precursor of (-)-(1S)-4 (Scheme 3). First cleavage of the benzyl group was carried out in a transfer hydrogenation that led to the formation of the diol (-)-(1S)-13,^[13] which after cleavage of the Boc protective group was coupled with adipoyl chloride. In this way a mixture of (-)-(1'S)-14b and differently acylated



Scheme 3. Alternative synthetic route to (-)-(1'S)-8b: a) Pd/C/ NH₄HCO₂, THF, RT, 4.5 h, then crystallization from n-hexane/EtOAc, 61%; b) 4 N HCl/EtOAc, RT, 100 min, then ClCO(CH₂)₄COCl, pyridine, DMF, RT, 19 h; c) NaOMe/MeOH, DMF, RT, 4 h, 65% over two steps; d) PPh₃ resin, DMF/CH₂Cl₂ (10:1), RT, 7.5 h, then crystallization from DMF/CH₂Cl₂, 19%. Bn = Benzyl.

products was obtained which, however, could be transformed completely into the desired diamide (-)-(1'S)-14b by solvolysis in the presence of catalytic amounts of NaOMe in MeOH. In the last step the desired chloride (-)-(1'S)-8b was formed from (-)-(1'S)-14b in an Appel reaction. Overall, (-)-(1'S)-8b was obtained in pure form in 8% yield over four steps from (+)-(1S)-12 by subsequent crystallization (N,N-dimethylformamide/dichloromethane).

The determination of the in vitro cytotoxicity of the new compounds was carried out in a clonogenicity assay based on the HTCFA (Human Tumor Colony Forming Ability) test on human bronchial carcinoma cells of the line A549, which reflects the proliferation ability of individual cells. For the prodrugs (-)-(1'S)-6a-c and (-)-(1'S,10'R)-7a-c the investigations (Table 1) gave practically identical cytotoxicities in the presence of β -D-galactosidase as were obtained for the corresponding seco-drugs (-)-(1'S)-8a-c and (+)-(1'S,10'R)-9a-c. The reversible detoxification thus established forms the basic requirement for an application within the framework of the ADEPT concept. In agreement with earlier results the seco-drugs that contained the pharmacophoric anti-methyl-seco-CBI unit ((+)-(1'S,10'R)-9a-c) showed a less pronounced cytotoxicity than the analogous compounds with a seco-CBI unit ((-)-(1'S)-8a-c).^[5,6] In addition the cytotoxicity of the seco-drugs within these two homolo-

gous series each fell within the sequence $n = 3 > 5 > 4$. Highlighted is the prodrug (-)-(1'S)-6a (n = 3), which has a QIC₅₀ value of 970000 and in the presence of β -D-galactosidase a cytotoxic activity in the femtomolar range (IC₅₀ = 0.15 pM).

Under physiological conditions the seco-drugs are transformed in situ into the corresponding active compounds 10a-c and 11a-c (Scheme 2) with a spirocyclopropyl group as in (+)-3. By using the seco-drugs generated from (-)-(1S)-1 and (+)-(1S,10R)-2 and related compounds we had been able to show by mass spectrometry on oligonucleotides and CD spectroscopy on living cells that the resulting active compounds are embedded very rapidly into the minor grooves of the DNA and subsequently alkylate more slowly an adenine residue sequence specifically.^[14] The cytotoxic activity of the active compounds released from (-)-(1S)-1 and (+)-(1S,10R)-2 is thus presumably attributable to a stabilization of the double strand of the DNA, whereby the alkylation causes a fixation of the molecules in the minor grooves. In contrast the formation of DNA intra- or DNA interstrand cross-links (ICLs) without incorporation into the minor grooves of the DNA could be held responsible for the comparably high cytotoxicity of the new seco-drugs (-)-(1'S)-8a-c and (+)-(1'S,10'R)-9a-c and of other comparable bifunctional active compounds and hybrid structures,^[8,15–17,18] since these substances have no DNA-binding unit for a

Table 1: Total yields and results of the HTCFA assay for the investigation of the cytotoxicity of the new prodrugs and secoodrugs against human bronchial carcinoma cells (A549).^[a]

Compound	R	n	Yield [%] ^[b]	β -D-Galactosidase ^[c]	IC ₅₀ [pM]	QIC ₅₀
(-)-(1'S)- 6a	H	3	51	—	1.46×10^5	9.7×10^5
(-)-(1'S)- 6a	H	3		+	0.15	
(-)-(1'S)- 8a	H	3	76	—	0.11	
(-)-(1'S)- 6b	H	4	44	—	1.29×10^5	2.2×10^4
(-)-(1'S)- 6b	H	4		+	5.8	
(-)-(1'S)- 8b	H	4	< 6 (8 ^[d])	—	9.0	
(-)-(1'S)- 6c	H	5	55	—	1.80×10^5	1.4×10^5
(-)-(1'S)- 6c	H	5		+	1.3	
(-)-(1'S)- 8c	H	5	65	—	1.0	
(-)-(1'S,10'R)- 7a	CH ₃	3	30	—	3.26×10^5	2.2×10^3
(-)-(1'S,10'R)- 7a	CH ₃	3		+	1.5×10^2	
(+)-(1'S,10'R)- 9a	CH ₃	3	80	—	1.6×10^2	
(-)-(1'S,10'R)- 7b	CH ₃	4	27	—	5.3×10^6	1.9×10^3
(-)-(1'S,10'R)- 7b	CH ₃	4		+	2.86×10^3	
(+)-(1'S,10'R)- 9b	CH ₃	4	76	—	3.14×10^3	
(-)-(1'S,10'R)- 7c	CH ₃	5	35	—	3.83×10^7	1.5×10^5
(-)-(1'S,10'R)- 7c	CH ₃	5		+	2.6×10^2	
(+)-(1'S,10'R)- 9c	CH ₃	5	79	—	1.7×10^2	

[a] The incubation of the cells (A549) with the respective compound was carried out for 24 h at 37°C; after subsequent cultivation for 12 days the relative clone formation rate was determined compared relative to untreated controls. The experiments were carried out as replicates (at least three) of duplicates. β -D-Galactosidase: *Escherichia coli*, 4 U mL⁻¹. [b] Each total yield based on (-)-(1S)-**4** or (+)-(1S,10R)-**5**. [c] Addition of β -D-galactosidase. [d] Total yield based on (+)-(1S)-**12**.

noncovalent interaction with the DNA as have duocarmycin SA ((+)-**3**) and the cytotoxic active substances generated from (-)-(1S)-**1** and (+)-(1S,10R)-**2**.

To estimate the reactivity of the new bifunctional secoodrugs towards double-stranded DNA and to investigate the structure–activity relationships, the compounds (-)-(1'S)-**8a** ($n=3$), (-)-(1'S)-**8c** ($n=5$), and (+)-(1'S,10'R)-**9a–c** ($n=3–5$) were incubated with different double-strand oligodeoxynucleotides in aqueous solution (1 % DMSO) for 24 h at 37°C and then analyzed with high-resolution electrospray ionization Fourier transform ion cyclotron resonance mass spectrometry (ESI-FTICR-MS).^[14] The ESI-FTICR-MS experiments showed no significant formation of ICLs with any of the compounds investigated. An analysis by circular dichroism gave no indication of a characteristic interaction of the active compounds with the oligodeoxynucleotides.^[14b] It is therefore unlikely that alkylation and cross-linking of double-stranded DNA is responsible for the high cytotoxicity of these new compounds.

In conclusion, the glycosidic prodrugs (-)-(1'S)-**6a–c** ($n=3–5$) and (-)-(1'S,10'R)-**7a–c** ($n=3–5$), which are constructed from seco analogues of the pharmacophoric units of duocarmycin SA ((+)-**3**) coupled with one another through dicarboxylic acids, are excellently suited for use within the scope of the ADEPT concept. Worthy of note is (-)-(1'S)-**6a** ($n=3$), which with a QIC₅₀ value of almost 1 000 000 and an IC₅₀ value of the underlying active compound of 0.11–0.15 pM exceeds by far all previously known compounds. In addition, the biological activity of the compounds is most probably not attributable to DNA intra- or DNA interstrand cross-linking.

The cause of the high cytotoxicity is presumably another, as yet unknown mechanism.

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