



Enhancement of Ricin Toxin A Chain Immunotoxin Activity: Synthesis, Ionophoretic Ability, and *In Vitro* Activity of Monensin Derivatives

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ABSTRACT. Site-selective toxin delivery was achieved by coupling monoclonal antibody to the A chain subunit of ricin (RTA-IT). The cell-killing potency of RTA-IT can be drastically increased *in vitro* by using ionophores such as monensin. To reduce the intrinsic toxicity of monensin and to enhance its *in vitro* and *in vivo* activity, we synthesized 7 derivatives characterized by different lipophilicities. These derivatives were also analyzed for ionophoretic activity on intact cells, toxicity, and RTA-IT-enhancing activity. Two different RTA-IT were assayed on a human leukemia cell line. A correlation between lipophilicity, ionophoretic activity, and RTA-IT enhancement was observed. The compounds with the highest polar charge showed low intrinsic toxicity, revealed moderate ionophoretic activity, and were able to enhance RTA-IT only at high concentrations, whereas more lipophilic compounds (with a C28 tail or a phenyl group) showed significant ionophoretic activity and good enhancing properties. *BIOCHEM PHARMACOL* 52;1:157–166, 1996.

KEY WORDS. immunotoxin; monensin; ricin A chain; enhancer; drug targeting

Tumor-selective cytotoxic conjugates are produced by chemically cross-linking bacterial toxins (e.g., diphtheria toxin, *Pseudomonas* exotoxin A) or plant toxins (e.g., ricin, abrin) to vehicle molecules (e.g., antibodies, ligands) that bind target antigens or receptors at the cell surface. The potential of MAb-ligand-toxin conjugates (IT) for tumor immunotherapy has been established in experimental models, as well as in clinical trials [1].

Ricin is a toxic plant glycoprotein formed by two subunits linked by a disulphide bond: the A chain, RTA§, which catalytically inactivates the protein synthesis of the target cell; and the B chain (RTB), which binds nonselectively to cell surface carbohydrate structures and helps RTA translocation to the cytosol of the target cell [1].

IT made with monoclonal antibody (MAb) linked to RTA have been extensively studied *in vitro* and in animal models [2–7], and their antitumor potential clinically investigated [8–13]. Although RTA-IT show high target-cell selectivity and low *in vivo* toxicity [8, 9, 11–13], their antitumor potency is variable, depending on the internalization rate, intracellular distribution, target-cell type, and other parameters [14–22]. As a result, the clinical trials

conducted with RTA-IT have met with limited success. The cell-killing potency of RTA-IT, however, can be drastically increased *in vitro* by using lysosomotropic amines (e.g., ammonium chloride, chloroquine), ionophores (e.g., monensin, nigericin), calcium antagonists (perhexiline, verapamil) or lysosomal enzyme inhibitors (e.g., leupeptin, pepstatin) in combination [14–22].

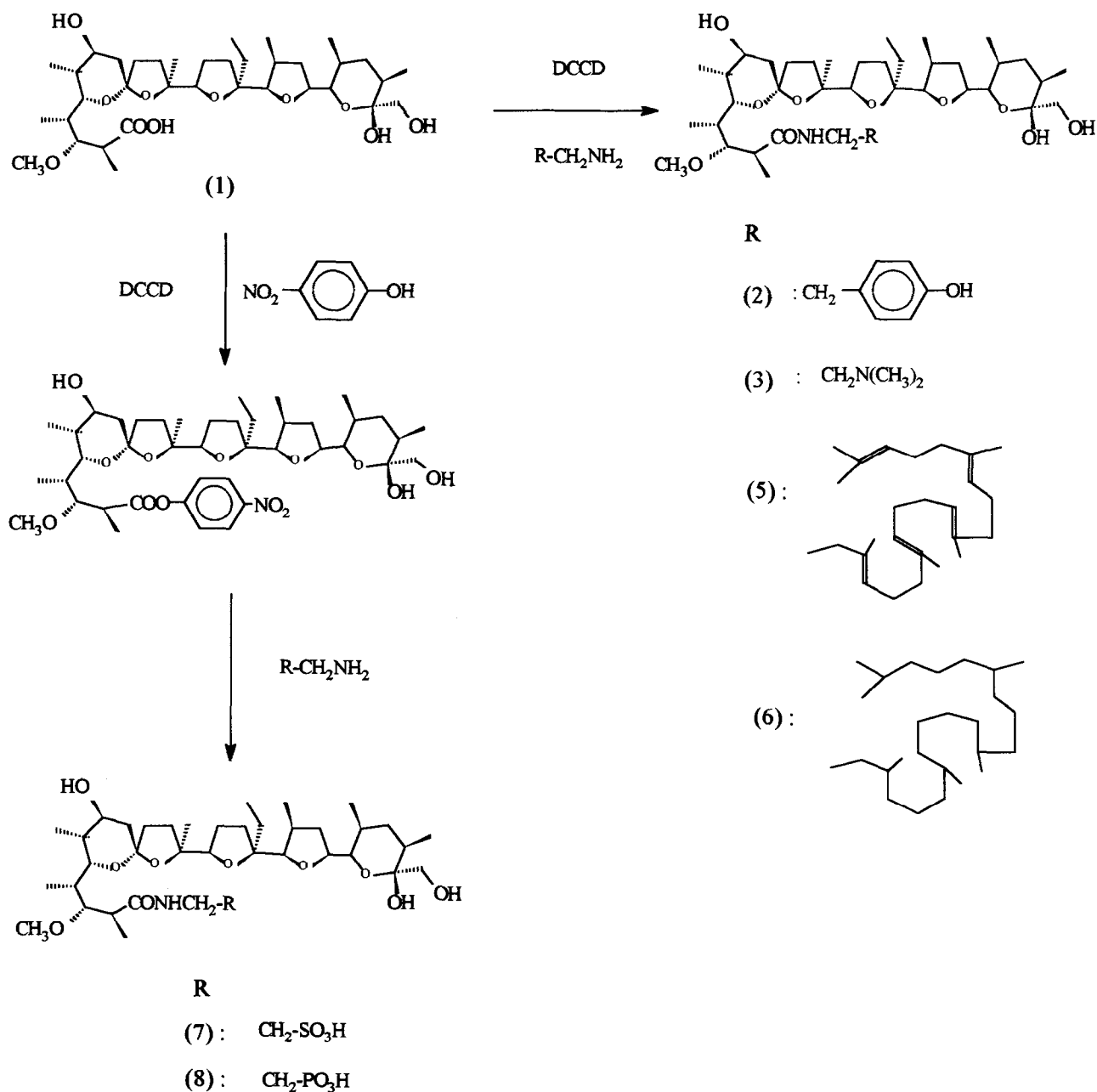
Monensin (1, Scheme 1), a well-characterized metabolite of *Streptomyces cinnamonensis*, is a monovalent carboxylic polyether ionophore able to regulate Na^+/H^+ exchange across the cell membrane [23, 24].

Monensin potentiates the cytotoxic effect of IT directed against different cell surface structures and entering the cell via different pathways [25, 26]. Several other properties make monensin a potential candidate for the enhancement of RTA-IT *in vivo*: a) it potentiates RTA-IT at nanomolar concentrations in cell lines of different histotypes [25, 26]; b) it can be delivered *in vivo* encapsulated in liposomes or cross-linked to carrier proteins (e.g., HSA) to decrease its clearance rate *in vivo* [3, 27]; and c) large amounts of monensin can be tolerated *in vivo* in animals following i.v. or i.p. inoculation [2, 3, 27, 28]. However, attempts at monensin and HSA-linked monensin application *in vivo* have highlighted several obstacles hampering its use in the clinic as a potentiator of RTA-IT. Among these: 1. fast clearance of unconjugated monensin; 2. blockage of its enhancing effect by serum factors [29, 30]; and 3. higher toxicity when injected intrathecally [2]. New monensin-derived mol-

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§ Abbreviations: RTA, ricin A chain; IT, immunotoxin; Tfn, transferrin; HSA, human serum albumin; FITC, fluorescein isothiocyanate; DCCD, dicyclohexylcarbodiimide.

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Scheme 1. Synthetic route to monensin amide derivatives. Monensin acid was condensed with primary amino compounds in the presence of DCCD to provide monensyl tyramide (2), dimethylaminomethyl amide (3), squalene amide (5), and squalane amide (6). Alternatively, to provide monensyl taurinamide (7) or ciliatinamide (8), monensin acid was first condensed with 4-nitro-phenol in the presence of DCCD, and the 4-nitro-phenolate derivative then reacted with the amino compounds.

ecules with properties apt to overcome some of the present limitations are, therefore, needed.

Treatment of cells with monensin leads to endosome and Golgi vacuolization [28, 31] which, in turn, affects the recycling of internalized molecules [32]. The molecular mechanisms by which monensin potentiates RTA-IT and accelerates their cell entry from intracellular compartments [33] have yet to be elucidated. Several hypotheses have been put forward to explain these aspects: 1. monensin may interact with intracellular membranes, facilitating the

translocation of toxin molecules across the lipid bilayer; and 2. the ionophoretic properties of monensin may be required for toxin passage to the cytosol.

To tackle some of the questions related to the hypotheses mentioned above, we set out to study the ability of monensin analogues of increasing lipophilicity to enhance the cell-selective cytotoxicity of IT directed against two cell surface structures expressed by human leukemia cells, the transferrin receptor (TfnR) and the CD5 antigen. Modulation of lipophilicity was achieved by condensing the car-

boxyl group of monensin with several amines. For comparison, derivatives with higher polarity and, thus, increased solubility in water, were also prepared. The availability of a set of monensin derivatives also allowed us to approach some aspects pertaining to the structure-function relationship of monensin molecules.

MATERIALS AND METHODS

Chemicals

All chemicals were of reagent grade and were obtained from either Aldrich Chimica or Bracco-Merck (Milan, Italy). Monensin was purchased from Fluka Chemical Co. (Milan, Italy). ^1H NMR spectra were recorded on a Jeol EX-400. Mass spectra were obtained on a VG Analytical 7070 EQ-HF spectrometer. IR were recorded on a Perkin-Elmer 781. Microanalyses performed on an elemental analyser 1106 (Carlo Erba Strumentazione) were within $\pm 0.4\%$ of theoretical values.

Squalene Amine:

(4E,8E,12E,16E)-4,8,13,17,21-pentamethyl-4,8,12,16,20-docosapentaenylamine

NH_3 (ca. 5 mL) was liquefied at -80°C and anhydrous methanol (10 mL) previously cooled at -50°C was, then, added with continuous stirring. A solution of HCl in anhydrous methanol was added in drops up to pH 6.0; during this addition, a white precipitate formed. NaBH_3CN (314 mg, 5 mmol) was then added, followed by 1,1',2-tris-nor-squalene aldehyde (1.92 g, 5 mmol). The reaction mixture was allowed to reach -25°C and left 12 hr at this temperature. It was, then, extracted with diethyl ether (100 mL) after addition of water (100 mL), dried over anhydrous sodium sulphate, and evaporated to dryness under vacuum. The resulting oil was purified by flash chromatography using diethyl ether/light petroleum 70:30, then diethyl ether and, finally, diethyl ether/methanol 95:5, to give squalene amine (482 mg, 25% yield). Anal. C, H, N. IR (CHCl_3) ν_{max} 2970, 2920, 2870, 1450, 1380 cm^{-1} . ^1H -NMR (CDCl_3) δ : 1.55–1.64 (m, 20 H, allylic CH_3 and $\text{CH}_2\text{CH}_2\text{N}$), 1.96–2.05 (m, 18 H, allylic CH_2), 2.66 (m, 2H, CH_2N), 5.02–5.21 (m, 5H, vinylic CH). HRMS: found M^+ , 385.3698. $\text{C}_{27}\text{H}_{47}\text{N}$ requires M , 385.3708; C, H, N.

Squalene Amine:

4,8,13,17,21-pentamethyldocosylamine

Pd/C (10% Pd; 15 mg) was added to absolute ethanol (100 mL) under stirring, and the suspension was, then, put under H_2 atmosphere. Squalene amine (250 mg, 0.65 mmol) was added and left for 24 hr under H_2 atmosphere under stirring. The suspension was filtered to remove the catalyst and concentrated under vacuum. The crude product was purified by flash chromatography with diethyl ether and then diethyl ether/methanol 95:5 to remove impurities and, finally, with methanol to give 236 mg (92% yield) as a col-

orless viscous oil. Anal. C, H, N. IR (CHCl_3) ν_{max} 2960, 2920, 2860, 1460, 1380 cm^{-1} . ^1H -NMR (CDCl_3) δ : 0.79–0.86 (m, 20 H, CH_3 and $\text{CH}_2\text{CH}_2\text{N}$), 0.98–1.40 (m, 33 H, alkylic CH_2 and CH), 2.59 (broad t, 2H, CH_2N). HRMS: found M^+ , 395.4494. $\text{C}_{27}\text{H}_{57}\text{N}$ requires M , 395.4491; C, H, N.

Monensin Acid (1)

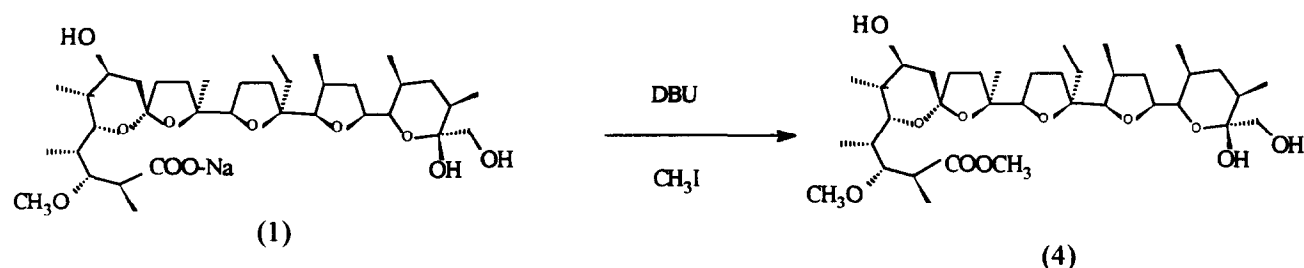
Sodium monensin was dissolved in hot methanol and the solution filtered. Monensin was, then, precipitated by adding water to the cooled methanolic solution. This purification step was repeated twice. The white salt obtained was dried under vacuum (yield 83–87%). It was, then, dissolved in distilled chloroform and stirred vigorously for 1 hr at room temperature with a layer of aqueous perchloric acid (1 M) to convert the sodium salt into the acid form of monensin. The chloroform layer was washed with distilled water until the washings were neutral, and then evaporated to dryness. The obtained crystals were dried under vacuum and stored at -20°C .

To synthesize the reactive intermediate monensin 4-nitrophenolate, 500 mg of monensin acid (0.745 mmol) were dissolved in 10 mL of freshly prepared dry tetrahydrofuran (THF) and maintained at 5°C . 4-nitrophenol (160 mg, 1.15 mmol) and DCCD (250 mg, 1.2 mmol), dissolved in 3 mL of THF, were added and the reaction was maintained at 5°C for 24 hr. The precipitated dicyclohexylurea was removed by centrifugation and the reaction mixture purified by flash chromatography (2×20 cm SiO_2 Merck) using ethyl acetate/methylene chloride 15:85 and, then, 30:70. The fractions containing monensin 4-nitrophenolate were collected and dried under vacuum to obtain 450 mg of white crystalline product. Yield 76%. $R_f = 0.38$ in $\text{EtOAc-CH}_2\text{Cl}_2$ (30:70). IR (KBr) ν_{max} cm^{-1} : 1765, 1525, 1350, 2950. ^1H -NMR (400 MHz, CDCl_3) δ : 8.27 (d, $J = 9.06$ Hz, H arom), 7.37 (d, $J = 9.06$ Hz, H arom), 4.48 (d, $J = 8.8$ Hz OH 2nd), 4.38 (br, OH 3rd), 4.31 (dt, $J = 3.1, 8.1, 20\text{H}$), 4.11 (dd, $J = 2.2, 9.5, 5\text{H}$), 3.94 (d, $J = 4.4, 17\text{H}$), 3.85 (m, 21H), 3.83 (br, 7H), 3.76 (t, $J = 4.9, 3\text{H}$), 3.56 (dd, $J = 5.5, 9.2, 13\text{H}$), 3.45 (br, 26H), 3.39 (s, 35H), 2.98 (dq, $J = 4.2, 6.9, 2\text{H}$), 2.65 (br, OH 1st), 0.85–2.30 pattern of 45 protons. FAB-MS m/z : 814.4354 ($M + \text{Na}$) $^+$, Calcd for $\text{C}_{42}\text{H}_{65}\text{NaNO}_{13}$: 814.4356; C, H, N, O.

Monensin Methyl Ester (4)

To synthesize this compound, the procedure described by Tohda *et al.* [34] was followed using methyl iodide and 1,8-diazabicyclo[5.4.0]undecene (Scheme 2).

The tyramide (2), N,N -dimethylaminoethylamide (3), squalene amide (5), and squalane amide (6) derivatives of monensin were prepared using essentially the same procedure as in Scheme 1: Monensin acid (200 mg, 0.3 mmol) was stirred at 5°C for 30 min in 2 mL of dry dimethylformamide (DMF) or THF and, then, DCCD (64 mg, 0.31 mmol) was added. After being stirred for 1 hr, the reaction



Scheme 2. Synthetic route to monensin methylester derivative (4). Monensin sodium salt was reacted with methyl iodide and 1,8-diazabicyclo[5.4.0]undecene.

was treated with 0.4 mmol of the corresponding amine: tyramine, N,N-dimethylethylenediamine, squalene, and squalene amine dissolved in 1 mL of dry DMF. Stirring was continued at 5°C for 24 hr. The mixture was evaporated under vacuum and the residue suspended in ethyl acetate and filtered off. To remove all the precipitated dicyclohexylurea, this step was repeated 3 times. The crude product was purified by flash chromatography on 230–400 mesh silica gel.

Monensin Tyramide (2)

Elution mixture: methylene chloride/ethyl acetate 85:15 and, then, 5:95 to give 170 mg of pure **2**. $R_f = 0.28$ in CH_2Cl_2 -MeOH (90:10). IR (CHCl_3) $\nu_{\max} \text{ cm}^{-1}$: 1650 ($\text{C}=\text{O}$). $^1\text{H-NMR}$ (400 MHz, CDCl_3) δ : 7.03 (d, $J = 8.24$ Hz, H arom), 6.79 (d, $J = 8.40$ Hz, H arom), 4.25 (dt, $J = 3.1, 8.1, 20\text{H}$), 4.11 (dd, $J = 2.2, 9.5, 5\text{H}$), 3.97 (d, $J = 4.4, 17\text{H}$), 3.85 (m, 21H), 3.65 (t, $J = 4.9, 3\text{H}$), 3.56 (dd, $J = 5.5, 9.2, 13\text{H}$), 3.40 (br, 26H), 3.30 (s, 35H), 2.83 (dq, $J = 4.0, 6.5, 2\text{H}$), 2.65 (t, $J = 5.2$ Hz, CH-Ph), 0.85–2.30 pattern of 45 protons. FAB-MS m/z : 812.501 ($M + \text{Na}$)⁺, Calcd for $\text{C}_{44}\text{H}_{71}\text{NaNO}_{11}$: 812.4925; C, H, N, O.

Monensin Dimethylaminoethylamide (3)

Reaction solvent THF: Elution mixture ethyl acetate/methanol/ammonia 97.5:2:0.5 and, then, 96:3:1 to give 167 mg of pure **3**. $R_f = 0.22$ in EtOAc-MeOH-NH₃ (70:28:0.2). IR (CHCl_3) $\nu_{\max} \text{ cm}^{-1}$: 1650. $^1\text{H-NMR}$ (400 MHz, CDCl_3) δ : 4.41 (dt, $J = 3.1, 8.1, 20\text{H}$), 4.2 (dd, $J = 2.2, 9.5, 5\text{H}$), 4.01 (d, $J = 3.4, 17\text{H}$), 3.91 (br, 7H), 3.85 (m, 21H), 3.60 (dd, $J = 5.5, 8.9, 13\text{H}$), 3.42 (s, 35H), 3.38 (t, $J = 7.5$ Hz, CH-NCO), 2.90 (dq, $J = 3.6, 7.0$ Hz, 2H), 0.85–2.30 pattern of 45 protons. FAB-MS m/z : 763.498 ($M + \text{Na}$)⁺, Calcd for $\text{C}_{40}\text{H}_{72}\text{NaN}_2\text{O}_{10}$: 763.508; C, H, N, O.

Monensin Squalene Amide (5)

Elution mixture: methylene chloride/methanol 99:1 and, then, 96:4 to give 150 mg of pure **5**. $R_f = 0.58$ in CH_2Cl_2 -MeOH (90:10). IR (CHCl_3) $\nu_{\max} \text{ cm}^{-1}$: 1650. $^1\text{H-NMR}$ (400 MHz, CDCl_3) δ : 5.1 (m, vinyl CH-Squalene), 4.41 (dt, $J = 3.0, 8.3$ Hz, 20H), 4.2 (dd, $J = 2.0, 9.4$ Hz, 5H), 3.85 (m, 21H), 3.42 (s, 35H), 2.95 (m, 2H), 0.85–2.30 pattern of

45 protons, 1.99 (m, methylene CH-Squalene), 1.6 (s, methyl CH-Squalene). FAB-MS m/z : 1060.9010 ($M + \text{Na}$)⁺, Calcd for $\text{C}_{63}\text{H}_{107}\text{NaNO}_{10}$: 1060.7792; C, H, N, O.

Monensin Squalene Amide (6)

Elution mixture: methylene chloride/methanol 99:1 and, then, 96:4 to give 110 mg of pure **6**. $R_f = 0.68$ in CH_2Cl_2 -MeOH (90:10). IR (CHCl_3) $\nu_{\max} \text{ cm}^{-1}$: 1650. $^1\text{H-NMR}$ (400 MHz, CDCl_3) δ : 4.41 (dt, $J = 3.0, 8.3$ Hz, 20H), 4.2 (dd, $J = 2.0, 9.4$ Hz, 5H), 3.85 (m, 21H), 3.42 (s, 35H), 2.95 (m, 2H), 0.85–2.30 pattern of 45 protons, 1.25 (m, methylene CH-Squalene), 0.87 (m, methyl CH-Squalene). FAB-MS m/z : 1070.8110 ($M + \text{Na}$)⁺, Calcd for $\text{C}_{63}\text{H}_{117}\text{NaNO}_{10}$: 1070.8575; C, H, N, O.

Monensin Taurine Amide (7)

To a solution of 400 mg of monensin 4-nitrophenolate (0.5 mmol) in 5 mL of dry pyridine were added 160 mg of taurine (2-aminoethanesulfonic acid) (1.28 mmol), previously dissolved in 2.3 mL of distilled water. The solution was maintained vigorously stirred for 48 hr at room temperature. After evaporation under vacuum, the unreacted taurine was filtered off in ethyl acetate and the crude product purified by flash chromatography using methylene chloride/ethyl acetate 95:5 and, then, methylene chloride/ethyl acetate/methanol 55:30:15. The white powder was, then, crystallized from acetone/ethyl acetate to give 383 mg of pure **7**. M.p. 161–162°C. $R_f = 0.0.8$ in CH_2Cl_2 -MeOH (90:10). IR (KBr) $\nu_{\max} \text{ cm}^{-1}$: 1650 ($\text{C}=\text{O}$), 1255 (SO_2 asym), 1050 (SO_2 sym). $^1\text{H-NMR}$ (400 MHz, d_6 -DMSO) δ : 4.35 (dt, $J = 3.1, 7.8, 20\text{H}$), 4.15 (dd, $J = 2.5, 8.9, 5\text{H}$), 3.95 (br, 7H), 3.83 (m, 21H), 3.35 (s, 35H), 3.11 (m, CH-taurin), 0.85–2.30 pattern of 45 protons. FAB-MS m/z : 823.397 ($M + 2\text{Na}$)⁺, Calcd for $\text{C}_{38}\text{H}_{67}\text{Na}_2\text{NO}_{13}\text{S}$: 823.4128; C, H, N, O, S.

Monensin Ciliatine Amide (8)

To a solution of 400 mg of monensin 4-nitrophenolate (0.5 mmol) in 10 mL of dry pyridine were added 160 mg of ciliatine (2-aminoethylphosphonic acid) (1.28 mmol) previously dissolved in 4.5 mL of distilled water. The solution was vigorously stirred for 5 days at room temperature. After

evaporation under vacuum, the unreacted ciliatin was filtered off in ethyl acetate and the crude product was purified by 5 preparative TLC plates (20 × 20 cm × 1 mm, Bracco-Merck, Milan, Italy), eluent methylene chloride/methanol 80:20 to give 85 mg of pure **8**. $R_f = 0.18$ in CH_2Cl_2 -MeOH (80:20). IR (CHCl_3) ν_{max} cm^{-1} : 1650 (C=O), 1280 (PO). $^1\text{H-NMR}$ (400 MHz, d_6 -DMSO) δ : 4.28 (dt, $J = 3.1, 7.8$, 20H), 4.20 (dd, $J = 2.5, 8.9$, 5H), 4.05 (d, $J = 3.4$, 17H), 3.91 (br, 7H), 3.85 (m, 21H), 3.60 (dd, $J = 5.5, 8.9$, 13H), 0.85–2.30 pattern of 45 protons. $^{32}\text{P-NMR}$ δ : 19.7, disacc. dt $J = 17.09, 21.9$ Hz. FAB-MS m/z : 823.412 ($M + 2\text{Na-H}$)⁺, Calcd for $\text{C}_{38}\text{H}_{68}\text{Na}_2\text{NO}_{13}\text{P}$: 823.4228; C, H, N, O, P.

LIPOPHILICITY (R_m) DETERMINATION. The lipophilic character of the synthesized compounds was determined by a chromatographic R_m method [35, 36]. Solutions of compounds were spotted 1 cm above the bottom of silanized silica gel TLC plates (20 × 20 cm, RP-2, Bracco-Merck). The plates were developed with various mixtures of water-methanol (from 50 to 90% in methanol). Then, the plates were dried and sprayed with a 4% solution of vanillin in methanol and 2.5% concentrated sulphuric acid. After a few minutes of 80°C, reddish spots appeared. The R_m were calculated from the following expression: $R_m = \log[(1/R_f) - 1]$ where R_f is the distance travelled by the compound divided by the distance travelled by the solvent front (18 cm). The values reported in Fig. 1 represent the mean of 4 different chromatographic runs for each monensin derivative.

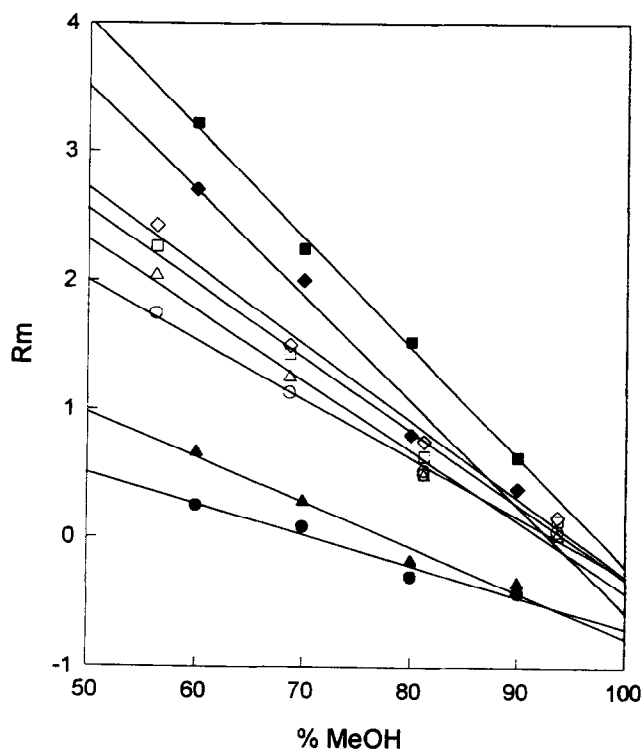


FIG. 1. Linear relationship between R_m values and methanol concentration in the mobile phase. The values reported represent the mean of 4 values: **1** (○), **2** (◇), **3** (△), **4** (□), **5** (■), **6** (◆), **7** (●), **8** (▲).

Biological

CELLS. The Jurkat, a human T-lymphoblastoid cell line CD5+, was used. Jurkat cells were maintained by serial passage in RPMI 1640 (Seromed, Berlin, Germany) + 10% fetal calf serum (Seromed) and gentamicin at 37°C in 5% CO₂ in humidified atmosphere.

IMMUNOTOXINS. ST.1 (Fab')₂-RTA (henceforth designated ST.1-RTA for brevity) IT to the CD5 differentiation antigen of human T lymphocytes was kindly supplied by Dr. P. Casellas (Sanofi Recherche, Montpellier, France). The preparation and properties of this IT have been described in detail previously [37]. Transferrin (Tfn)-RTA IT was synthesized and purified as described [38]. In all cytotoxicity assays, the concentration of IT is expressed as protein concentration.

PROTEIN SYNTHESIS ASSAYS. Protein synthesis measurements were performed to assay the intrinsic toxicity of monensin and its derivatives and to evaluate their RTA-IT enhancing activity.

a) Toxicity: protein synthesis was assayed by incubating 5×10^4 Jurkat cells in 50 μL leucine-free RPMI, 5% fetal bovine serum in 96-well flat-bottomed microtitration plates (Costar, Cambridge, MA, U.S.A.). From a stock solution of 5 mM of monensin in DMSO, 10-fold dilutions were prepared. DMSO (1% final concentration) alone had no effect on cell culture. Monensin and its derivatives were, then, added to Jurkat cell microcultures and target cells incubated for 22 hr at 37°C in 5% CO₂ in a humidified atmosphere in 100 μL final volume. Final concentrations ranged from 500 μM to 0.5 nM. Under these conditions, in the absence of monensin or derivatives, the cells proliferate monoexponentially. One μCi of L-[U-¹⁴C]-Leucine (316 mCi/mmol, DuPont, Boston, MA, U.S.A.) in 10 μL of RPMI was, then, added. After 2 hr, the cells were harvested onto glass fiber filters, washed with water, and dried. Radioactivity incorporated by the cells was, then, measured in a beta-spectrometer. The results are expressed as a percentage of the incorporation of control mock-treated cultures. The cytotoxicity of monensin or its derivatives was compared by taking the molar concentration required to inhibit 50% protein synthesis (IC₅₀).

b) Enhancement of RTA-IT cytotoxicity: Jurkat target cells (5×10^4 /well) were treated with various concentrations of different RTA-IT for 22 hr in the presence or in the absence of monensin or its derivatives, used at the highest nontoxic concentration. Protein synthesis was, then, measured as described above. Raw data expressed in cpm were also processed as above.

CYTOFLUOROMETRY. Tfn and anti-TfnR antibodies are internalized to low pH intracellular compartments [39]. We took advantage of the pH-dependent fluorescence of fluorescein isothiocyanate (FITC)-labelled proteins [40] to investigate *in vivo* the ionophoretic activity of monensin and its derivatives.

Fluorescence emission of FITC-labelled molecules inter-

nalized to low pH compartments is decreased with respect to that of noninternalized surface-bound FITC-labelled molecules. Addition of monensin following internalization into acidic compartments leads to an increase of FITC emission due to monensin-driven neutralization of low pH intracellular organelles [41].

Tfn was labelled with FITC following described procedures [41]. An antiFITC mouse MAb (clone FL-D6) was from Sigma (St. Louis, MO, U.S.A.).

Jurkat cells (3×10^5 in 50 μ L RPMI 1640) were incubated for 1 hr at 4°C with a saturating amount of FITC-labelled Tfn. The cells were, then, washed twice with a cold solution of bovine serum albumin (BSA) 0.2% in phosphate saline buffer (0.1 M, pH 7), to remove unbound Tfn-FITC, and resuspended in 200 μ L cold RPMI medium. The cells were, then, incubated at 37°C for 30 min to allow internalization of surface-bound molecules. The samples were, then, placed at 4°C and treated for 30 min with monensin or its derivatives at various concentrations (range 0.5–50 μ M). AntiFITC MAB (1:100 final dilution) was also added to quench the fluorescent emission of non-internalized surface-bound FITC-labelled molecules. Under these conditions, only an intracellular fluorescent emission could be detected. In control mock-treated samples, monensin or its derivatives were omitted and only the diluent (DMSO) was added. Other control samples were treated as above, except that the temperature was kept at 4°C throughout washings and incubations. The fluorescence intensity expressed by the cells was, then, evaluated by running cell samples through an Epic-Profile II (Coulter, Hialeah, FL, U.S.A.) cytofluorometer with an exciting wavelength of 488 nm at 15 mW power. Fluorescence intensity values are expressed as mean fluorescence intensity (MFI). The difference in MFI between values obtained from samples treated with monensin or its derivatives and untreated samples (Δ MFI) was taken as a measure of their ionophoretic activity.

Experiments in Tables and Figs. are representative of 3–4 experiments; values represent the average of duplicate or triplicate samples (<10% standard error).

RESULTS AND DISCUSSION

In an effort to obtain new RTA-IT enhancers with high activity and to investigate the possible correlation between the structure of monensin and its potentiating activity, seven structural analogues were synthesized (Schemes 1 and 2) and assayed: two compounds with an increased negative polarity (sulphate and phosphate group), and one with an ionizable positive charge (dimethylamino group). Three compounds with more lipophilic groups: a rigid phenyl group, and two long-chain structures of 28 carbon atoms: more rigid for the tris-nor squalene, more flexible for the tris-nor squalane. For comparison, we also prepared a methyl ester derivative of monensin to reduce the polarity of its carboxyl group.

Monensin (1), obtained by treatment of commercially available sodium salt with perchloric acid, was condensed with several substituted primary amines in the presence of DCCD to provide monensylamide compounds (3–6) (Scheme 1). Due to the complete insolubility of taurine and ciliatine in organic solvents, a different synthetic approach involving the reactive 4-nitrophenolate ester was chosen to obtain more polar derivatives (7, 8) (Scheme 1).

Lipophilicity

The partitioning of monensin and related derivatives between a stationary and polar mobile phase is a function of their structure. Thus, using the procedure described by G. L. Biagi *et al.* [35, 36] a relative lipophilicity factor, R_m , can be determined. Under these conditions, R_m values higher than that of monensin are suggestive of higher lipophilicity. From Fig. 1, it appears that only compounds 7 and 8 are more polar than monensin; all others are more lipophilic, in an increasing order, from 2 to 6.

Toxicity

The cytotoxicity of monensin and its derivatives was tested on the human T-lymphoblastoid Jurkat cell line (Table 1). Considering the IC_{50} values, only compounds 7 and 8, endowed with a highly negative charge, show a considerable reduction in toxicity, being 114–207-fold less toxic than the parent drug. Derivatives 2, 3, and 4 are also less toxic (4–7-fold reduction in toxicity), whereas the more lipophilic squalene and, in particular, squalane derivatives were as toxic as monensin. Thus, the relatively higher toxicity of the derivatives correlates with a higher lipophilicity (Fig. 1 and Table 1) and the relatively higher toxicity of more apolar compounds might, thus, be a result of a facilitated interaction with the cell membranes. The reduced toxicity of the synthesized compounds with respect to the parent drug allowed us to test higher concentrations of monensin derivatives in RTA-IT potentiation assays.

Enhancement of RTA-Immunotoxins by Monensin and its Derivatives

To evaluate the ability of monensin derivatives to enhance RTA-ITs, several concentrations of the various drugs and

TABLE 1. Toxicity of monensin derivatives on Jurkat cell line*

Compound	IC_{50} (μ M)
1	1.4 ± 0.1
2	7.5 ± 0.5
3	5.3 ± 0.2
4	10 ± 1.1
5	4.6 ± 0.6
6	1.5 ± 0.5
7	290 ± 3.1
8	160 ± 4.2

* Jurkat cells were incubated with different concentrations of monensin or its derivatives for 22 hr. Protein synthesis inhibition was, then, measured, based on L-[U- 14 C]-Leucine incorporation, and IC_{50} values were determined.

different concentrations of IT were added to Jurkat cell microcultures.

Two different ITs directed against different antigens expressed by the Jurkat cells were tested: one composed of a monoclonal antibody ST.1 (antiCD5) conjugated with RTA, and the other of human transferrin linked to RTA. The antiCD5 RTA-IT was chosen for our study because of its relevance in the treatment of human diseases. In fact, antiCD5 IT have been used *ex vivo* for bone marrow purging, and CD5 IT-based therapy has also been extensively investigated clinically in a broad spectrum of pathologies [42, 43]. Similarly, Tfn IT has been proposed for the treatment of solid tumors of various histotypes, particularly those affecting the central nervous system (CNS) [42, 43]. Furthermore, inasmuch as Tfn is not species-specific [27], it allows the use of a large panel of target human, rat, and mouse cell lines for comparisons.

The optimal dose of **1** to be used in *in vitro* experiments is 50 nM, as reported in previous publications [2, 27]. At higher doses of **1**, no increase in RTA-IT enhancement was observed, and a monensin-dependent toxicity began to appear (not shown). Figure 2 shows the potentiation of the cytotoxic activity of ST.1-RTA and of free RTA by **1**. As shown in Fig. 2, the cell-selective cytotoxicity mediated by ST.1-RTA was increased 1140-fold, whereas the noncell-selective toxicity of unconjugated RTA was increased only 10-fold. These data can be compared to the cytotoxicity of native ricin. These results suggest that both cell surface binding and, as reported by Griffin *et al.* [44], the presence of a reducible disulphide bond between RTA and the car-

rier molecule are necessary for monensin effect. Due to the different toxicity shown by the various derivatives, several concentrations were assayed in each case in an RTA-IT dose-response assay.

The cytotoxicity of the two IT in the absence of potentiators is reported in Table 2: ST.1-RTA showed low cytotoxicity ($IC_{50} = 1140$ pM), whereas Tfn-based IT was 50-fold more active ($IC_{50} = 26$ pM). This allowed us to correlate the activity of the enhancers on a rapid internalization, rapid recycling model (Tfn) in comparison with a slower one (antiCD5).

Overall, the data reported in Table 2 illustrate uniform patterns as regards the RTA-IT enhancing activity of monensin and its derivatives: in general, the extent of RTA-IT potentiation depends on the type of immunotoxin used for all drugs (monensin included) and reflects the cytotoxic activity of RTA-IT observed in the absence of enhancers.

In particular: 1) Compounds **2**, **5**, and **6** showed an RTA-IT potentiating activity comparable to **1**. Compound **6** showed a significantly higher (2-fold) potentiating activity in comparison with the parent drug. However, it must be noted that the RTA-IT enhancement observed with compounds **2**, **5**, and **6** was achieved at a 10-fold higher concentration than **1**, whereas the most polar compounds **7** and **8** required a 1000-fold higher concentration. A higher concentration could be used in this case due to their lower intrinsic toxicity. As previously described, **1** cannot be tested at concentrations higher than 50 nM, due to its high intrinsic toxicity.

2) Derivatives with a positive ionizable charge (e.g., compound **3**) or with a reduced negative charge (e.g., compound **4**) showed no RTA-IT enhancing activity at the concentrations tested.

The mechanism of potentiation of toxin conjugates by monensin has not yet been fully elucidated. Monensin affects cells in many ways, apparently as a consequence of profound changes in the cytoplasmic Na^+/K^+ balance [31].

When RTA-IT are directed against target cells, the most important consequence of the presence of monensin is that the conjugate tends to accumulate in endocytic vacuoles, which appear as enlarged vesicles and in a finely structured tubovesicle system, whereas entry and accumulation of the RTA-IT into the lysosomes is considerably reduced [45]. These phenomena are probably essential to prevent RTA degradation and allow maximal cytotoxic efficacy.

Coated vesicles and endosomal vacuoles are known to be the site of dissociation of Tfn from its receptor [39]. Therefore, analogous to Tfn, a surface-endosome-surface routing can be suggested for Tfn-RTA. In the case of antiCD5 RTA-IT, however, it has been shown that the cell intoxication process involves a surface-Golgi-surface recycling route [14]. The different intracellular routing of the two RTA-IT used in our experiments may account for the greater cytotoxicity of Tfn-RTA in the absence of potentiators. On the other hand, inasmuch as monensin functions intracellularly by altering the Golgi, it could be spec-

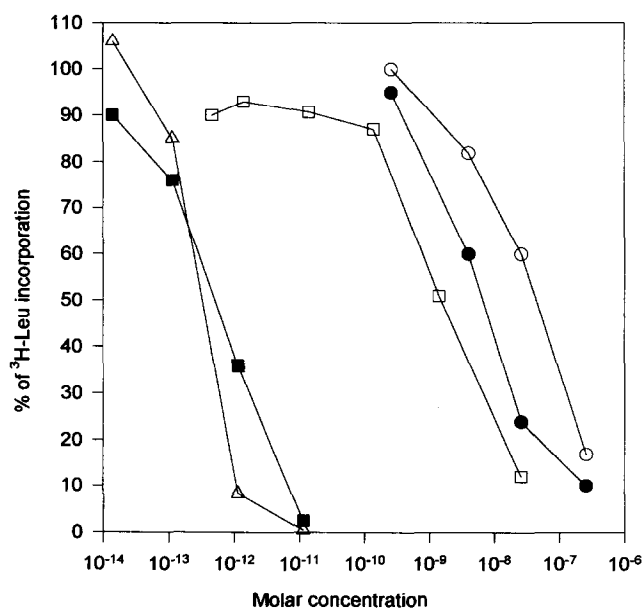


FIG. 2. Enhancement of ST.1-RTA and RTA cytotoxicity by monensin. Jurkat cells were incubated for 22 hr with either ST.1-RTA IT (squares) or ricin A chain (circle) in the absence (open symbols) or in the presence (filled symbols) of 50 nM monensin. Protein synthesis was, then, measured. For comparison, cytotoxicity of native ricin is presented (triangle).

TABLE 2. Comparative RTA immunotoxin enhancement of Jurkat cell line by monensin and derivatives*

Immunotoxin (Enhancer concentration (μ M))	IC ₅₀ (pM)/Fold-Potentiation			
	0.05	0.5	5	50
ST.1-RTA (1140 pM)†				
+1	0.46/2478			
+2	2.8/407	0.44/2590		
+3	840/1.3	800/1.0		
+4	1100/1.0	810/1.4		
+5	240/5.0	0.4/2590		
+6	1.7/670	0.28/4071		
+7	810/1.0	710/1.0	500/2.1	1.8/633
+8	1100/1.0	750/1.5	0.9/1267	0.29/3930
Tfn-RTA (26 pM)†				
+1	0.18/144			
+2	26/1.0	0.16/162		
+3	26/1.0	26/1.0		
+4	26/1.0	26/1.0	26/1.0	
+5	15/1.7	0.17/153		
+6	26/1.0	0.1/260		
+7	26/1.0	26/1.0	0.4/65	0.2/130
+8	15/1.7	15/1.7	13/1.8	0.24/108

* Cytotoxicity assay on Jurkat cells was induced by incubation with different concentrations of RTA-ITs and drugs for 22 hr. Protein synthesis inhibition was then measured based on L-[U-¹⁴C]-Leucine incorporation and IC₅₀ values were determined. †IC₅₀ values of IT without addition of enhancers.

ulated that the greater potentiation observed with anti-CD5-RTA is due to the preferential delivery of ST.1 immunotoxin to the Golgi or to the Golgi-related compartments. However, it must be noted that, at the low concentrations used in our assays, monensin seems unable to alter the pH of intracellular compartments [29].

Ionophoretic Activity

The cytofluorometric assays employed by us rendered possible the evaluation of the ionophoretic activity of monensin and its derivatives in intact cells. In fact, the difference in MFI between treated samples vs control samples (i.e., Δ MFI) is due to their ability to neutralize the low pH of intracellular compartments. It must be mentioned, however, that the ionophoretic activity of the various compounds could be detected with a cytofluorometric assay only at concentrations higher than those required to achieve RTA-IT enhancement in protein synthesis inhibition assays. The cytofluorometric assay was carried out with Tfn-FITC because Tfn is directed to low pH prelysosomal compartments [46] and its internalization cycle proceeds with fast kinetics [39]. As illustrated in Fig. 3, monensin showed the highest pH neutralizing effect, plateauing at 20–50 μ M. Among the derivatives, those endowed with higher lipophilic properties (i.e., 5 and 6) and 2 displayed a relatively higher ionophoretic activity (Δ MFI > 1), whereas the negatively charged 7 and 8 analogues, as well as 4, showed a lower ionophoretic activity (Δ MFI < 1). Thus, a correlation was found between ionophoretic activity and lipophilic properties of the various compounds assayed. Lipophilicity is known to play an important role in penetration of various compounds into the cell membranes;

in particular, lipophilic derivatives of ionophores allowed a more rapid transport of metal ions [47]. In our case, we can correlate the Rm values and the ionophoretic activity expressed as Δ MFI. It is likely that compounds 5 and 6 (approximately 3 times more lipophilic than monensin) and 2 (characterized by a rigid phenyl group) are able to increase ion traffic and, therefore, display relatively greater ionophoretic properties in the cytofluorometric assay. It must be noticed, however, that, in our assay, monensin displayed the greatest ionophoretic activity and, therefore, its lipophilicity/size ratio may be the most advantageous in performing a rapid ion transport, as evidenced by cytofluorometry.

Furthermore, a direct correlation between cell toxicity and ionophoretic activity can be appreciated in Fig. 3 and Table 1. In particular, the low Δ MFI shown by compounds 7 and 8 parallels the strong reduction in toxicity observed with monensin. This may be due to a reduced solubility in the cell membranes and/or to less efficient Na⁺ ion transport by these compounds.

How ionophoretic activity and lipophilicity relate to the ability to potentiate RTA-IT is still under investigation.

As shown in Fig. 3 and Table 2, we found that a correlation also exists between the ionophoretic properties of the various compounds and their RTA-IT potentiating effect. Cell intoxication by RTA is facilitated by neutral or slightly alkaline pH [48]. It is, therefore, likely that compounds showing higher ionophoretic activity and, thus, greater pH neutralizing properties also display greater RTA-IT enhancing activities.

In conclusion, we show here that various structural analogues of the RTA-IT enhancer monensin may be less toxic than the parent drug, but also display variable enhancing

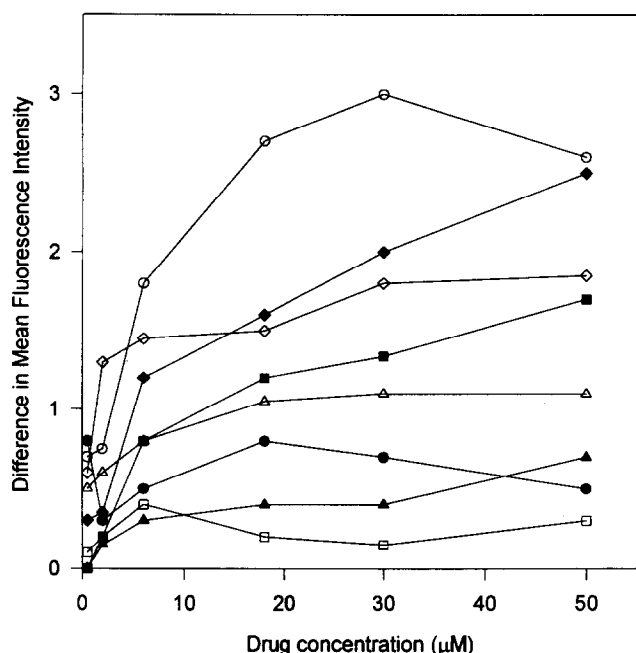


FIG. 3. The ionophoretic activity of monensin and its derivatives was evaluated on living cells by cytofluorometry. The assay measured the ability of monensin or its derivatives to counteract the low pH-induced quenching of internalized Tfn-bound FITC. Results are expressed as difference in mean fluorescence intensity (Δ MFI) between untreated samples and samples treated with Tfn-FITC in the presence of monensin or its derivatives. Compounds are represented as those in Fig. 1.

properties. This variability may be related principally to the structure of derivatives, but may also be influenced by the different intracellular routing of IT. Although no enhancers with greater activity than monensin are produced, these preliminary studies provide some guidance for the development of further monensin derivatives.

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