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Synthesis of giant globular multivalent glycofullerenes as potent inhibitors in a model of Ebola virus infection

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The use of multivalent carbohydrate compounds to block cell-surface lectin receptors is a promising strategy to inhibit the entry of pathogens into cells and could lead to the discovery of novel antiviral agents. One of the main problems with this approach, however, is that it is difficult to make compounds of an adequate size and multivalency to mimic natural systems such as viruses. Hexakis adducts of [60]fullerene are useful building blocks in this regard because they maintain a globular shape at the same time as allowing control over the size and multivalency. Here we report water-soluble tridecafullerenes decorated with 120 peripheral carbohydrate subunits, so-called 'superballs', that can be synthesized efficiently from hexakis adducts of [60]fullerene in one step by using copper-catalysed azide-alkyne cycloaddition click chemistry. Infection assays show that these superballs are potent inhibitors of cell infection by an artificial Ebola virus with half-maximum inhibitory concentrations in the subnanomolar range.

When compared with monovalent binding¹.

A remarkable example from nature in which multivalency plays a significant role is the interaction between viruses and bacteria with their respective host cells. In particular, the DC-SIGN (dendritic cell-specific intercellular adhesion molecule-3-grabbing non-integrin) receptor is one of the most important pathogen-recognition receptors. This lectin efficiently recognizes, in a multivalent manner, saccharides that contain mannoses and fucoses from glycoproteins (GPs)². In this regard, it is well established that protein-carbohydrate interactions are a key issue in a variety of biological processes, but because the affinity of simple glycans for their respective receptors is often weak, multivalent interactions typically occur. An important challenge for research nowadays is to better understand multivalency and its practical use. In a broader sense, glycobiology is currently a field of research in which chemically inspired approaches and strategies are producing significant advances3.

It is well known, however, that some viruses are able to escape the immune defence by using DC-SIGN as an entry point to infect the cell. Therefore, to inhibit the entry of pathogens by blocking this receptor at the early stages of infection represents a valuable strategy for the design of new antiviral agents. To address this challenge, a variety of different multivalent scaffold architectures have been synthesized, all of them endowed with multiple carbohydrates⁴.

The design of these glycoconjugates typically requires a multivalent central scaffold or core covalently connected to the carbohydrate epitopes that decorate the periphery. Thus, glycoclusters in which the carbohydrate units are connected directly to the core, glycoden-drimers connected through a dendritic structure and glycopolymers that involve a polymeric backbone have been investigated intensively in recent years^{5–9}.

Fullerenes have also been employed as a biocompatible scaffold for the multivalent presentation of ligands, given the possibility of multiple functionalization on their convex surface. In particular, hexakis adducts of [60]fullerene with a Th-symmetrical octahedral addition pattern have a unique three-dimensional structure that allows the introduction of six one-type or mixed-type addends^{10,11}. Such derivatives can be obtained in one synthetic step by the addition of malonates to C_{60} , but because the reaction is very sensitive to steric effects this approach has been limited by the low yields that result when larger malonates are used ¹². Recently, we developed a procedure based on a click-chemistry approach that provides hexakis adducts with 12 alkyne or azide terminal groups in high yields from simple malonates^{13,14}. These hexakis adducts can be functionalized easily and efficiently by using the copper-catalysed azide-alkyne cycloaddition (CuAAC) reaction, which allows the introduction of 12 functional groups simultaneously in a regioselective and efficient way. As these hexakis adducts show an octahedral arrangement of equally separated addends located on the [60]fullerene periphery in a globular topology, they constitute a very attractive platform for the study of multivalent interactions with lectins with important advantages, such as a better biocompatibility of the carbon central core, the globular

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Figure 1 | Synthesis of azide-substituted glycofullerene 9 appended with a long linker and mannose as the carbohydrate unit. Five linear steps are required to prepare compound **9** in a 26% overall yield. Reagents and conditions: (i) ethylmalonyl chloride, pyridine, CH₂Cl₂, 0 to 25 °C, one hour (99%); (ii) C₆₀, I₂, DBU, 25 °C, PhMe, 16 hours (49%); (iii) CBr₄, DBU, 25 °C, ODCB, 72 hours (67%); (iv) CuSO₄·5H₂O, sodium ascorbate, THF/H₂O, 100 °C (MW), two hours (91%); (v) **8** (19 equiv.), CuSO₄·5H₂O, sodium ascorbate, TBAF, THF/H₂O, 80 °C (MW), 1.5 hours (87%).

geometry of the functional groups and an easy and versatile chemical functionalization.

Recently, we reported the syntheses of [60]fullerene hexakis adducts endowed with different carbohydrate units in the periphery. Depending on the starting sugar derivatives employed (monomer, dimer or trimer) 12, 24 and up to 36 monosaccharides were introduced on the periphery of the fullerene central core in a straightforward manner¹⁴⁻¹⁷.

Some of the aforementioned glycofullerenes showed interesting biomedical applications¹⁸. A significant multivalent effect towards various glycosidases was observed in the inhibition profile of a [60]fullerene hexakis adduct endowed with 12 iminosugar units¹⁶. However, fullerene hexakis adducts that bear 12 mannoses on the periphery behave as inhibitors of FimH, a bacterial adhesin¹⁹. Furthermore, studies carried out on hexakis adducts of [60]fullerene endowed with 12, 24 and 36 mannoses have shown a multivalent effect in the interaction with concanavalin A¹⁵, and act as efficient inhibitors of cell infection by Ebola-pseudotyped viral particles²⁰. These preliminary studies reveal that fullerenes are adequate platforms for the multivalent presentation of carbohydrates, and thus pave the way to the covalent linkage of a wide variety of different bioactive molecules in a multivalent manner and to a singular and less-explored globular topology.

In this work, we synthesized molecules with a globular topology formed by making hexakis adducts of [60]fullerene (with the fastest dendrimeric growth reported up to now) through the introduction of 120 sugar units in one synthetic step by using the highly efficient CuAAC click-chemistry methodology. The synthesis of hexakis adducts of [60]fullerene with an octahedral addition pattern has been reported previously. This previous example used macrocyclic bismalonates and enabled the synthesis of heptafullerenes with tunable properties^{21–23}. In contrast, our click-chemistry approach yields tridecafullerenes in which the central [60]fullerene is covalently connected to 12 [60] fullerenes, each endowed, in turn, with ten monosaccharides. To confirm the proposed structures, we carried out a thorough structural study that involved a variety of techniques (infrared, ¹H NMR and ¹³C NMR spectroscopy, dynamic light scattering (DLS), transmission electron microscopy (TEM) and X-ray photoelectron spectroscopy (XPS)). Biological studies revealed that the new giant glycofullerenes exhibit a very strong inhibition of cell infection by Ebola-pseudotyped viral particles with half-maximum inhibitory concentration (IC₅₀) values in the subnanomolar range.

Results and discussion

Synthesis. The synthesis of the glycofullerene superballs was carried out as depicted in Figs 1–3. Although, at first glance, the molecules **17a–17c** may appear difficult to obtain, mainly because of their size and molecular complexity, their synthesis is straightforward and, most importantly, easily reproducible, and it affords the final

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Figure 2 | Synthetic pathway to azide-substituted glycofullerenes 15a and 15b with a short spacer. The peripheral glycofullerene units 15a and 15b were prepared using Bingel conditions and CuAAC click chemistry to conjugate the carbohydrate moieties. Reagents and conditions: (i) CBr₄, DBU, 20 °C, PhMe, 72 hours (49%); (ii) 2-azidoethyl α-D-mannopyranoside for 14a (or 2-azidoethyl α-D-galactopyranoside for 14b), CuBr·S(CH₃)₂, sodium ascorbate, Cu⁰, DMSO, 72 hours (14a, 86%; 14b, 86%); (iii) NaN₃, 70 °C (MW), DMSO, three hours (15a, 84%; 15b, 81%).

compounds in good overall yields. Furthermore, as discussed below, purification of the samples—which includes the required removal of copper for further biological studies—is carried out easily because, after column filtration, compounds **17a** and **17b** are precipitated in the reaction medium and pure samples that show highly reproducible biological assays are obtained.

To study the effect of the size and/or steric congestion of these superballs on biological properties, two different and complementary synthetic strategies were developed simultaneously to obtain final products with different spacers between the central fullerene core and the peripheral carbohydrate-substituted fullerene appendages. In both cases, the synthesis relies on the grafting of clickable A₁₀B macromonomers onto a compact C₆₀ hexa-adduct scaffold that bears 12 terminal alkyne units. The preparation of macromonomer 9 is depicted in Fig. 1. Esterification of alcohol 1 with ethylmalonyl chloride followed by reaction of the resulting malonate (2) with C_{60} under Bingel conditions gave methanofullerene 3. Subsequent treatment of 3 with an excess of malonate 4, CBr₄ and 1,8-diazabicyclo[5.4.0]undec-7-ene (DBU) in o-dichorobenzene (ODCB) gave building block 5 to which the mannose derivative 6 was clicked. Importantly, the trimethylsilyl-protected alkyne unit is not reactive under these conditions and intermediate 7 was thus obtained in a good yield. Finally, compound 7 was desilylated in situ with tetrabutylammonium fluoride (TBAF) to generate the corresponding terminal alkyne, and reaction with a large excess of diazide 8 provided the desired macromonomer 9 in 87% yield.

The synthesis of macromonomers **15a** and **15b** starts from monoadduct **10**, which results from the Bingel nucleophilic cyclopropanation of C_{60} with 6-bromohexyl ethyl malonate (Fig. 2)²⁴. To obtain the [5:1]-hexa-adduct **12**, a tenfold excess of di(pent-4-yn-1-yl) malonate and a ~50-fold excess of carbon tetrabromide were added in the presence of DBU as the base. Flashchromatography purification provided hexa-adduct **12** as a red solid. Different carbohydrate azides (mannose, galactose) were then linked to C_{60} by the CuAAC reaction, with CuBr·S(CH₃)₂ as the catalyst and sodium ascorbate as the reducing agent in the presence of a piece of metallic Cu, to yield derivatives **14a** and **14b**. The nucleophilic substitution of bromine by the azide was carried out with an excess of sodium azide under microwave (MW) heating, and thus gave the versatile building blocks **15a** and **15b**.

The azide-containing macromonomers **9**, **14a** and **14b** were then clicked to the symmetric alkyne derivative **16**¹⁴ under CuAAC conditions. Superballs **17a–17c** substituted with up to 120 monosaccharide units were thus obtained with yields over 70% (Fig. 3). Derivatives **17a** and **17b** were obtained in only three synthetic steps from easily accessible starting materials and the use of protecting groups was avoided.

Although the three tridecafullerenes **17a–17c** contain the same number of monosaccharides, compound **17c** has a larger spacer between the central fullerene moiety and the peripheral carbo-hydrate-functionalized fullerenes. This longer spacer affects the flexibility and the size of the compound and can favour the accessibility and availability of the carbohydrate ligands to interact with the receptor, which could have an important influence on the biological activity²⁰.

However, the aforementioned synthetic approaches enable fullerene derivatives to be synthesized from malonates endowed with alkyl chains of variable length through a Bingel cyclopropanation followed by a subsequent CuAAC reaction. This synthetic strategy affords hexakis adducts with long chains that connect the fullerenes and carbohydrates whereas their direct preparation from suitably functionalized malonates that bear long chains typically occurs with low yields¹⁰.

Characterization of these superballs was carried out by standard spectroscopic techniques. Thus, Fourier transform infrared (FTIR) spectra do not show the presence of either of the azide groups (the typical signal is observed at ~2,097 cm⁻¹) or the alkynes (~2,117 cm⁻¹) (see the Supplementary Information). The molecular ion peak of these compounds could not be detected, but the transfer of glycoclusters of such high molecular weights into the gas



Figure 3 | Syntheses of the tridecafullerenes 17a-17c using CuAAC click chemistry. The core fullerene **16** (endowed with 12 alkyne groups) is joined to the peripheral fullerene-based compounds **9**, **15a** and **15b** by click chemistry. Reagents and conditions: for compounds **17a** or **17b** (i) **15a** or **15b**, CuBr·S(CH₃)₂, sodium ascorbate, Cu(0), DMSO, 25 °C, 48 hours (**17a** (from **15a**), 73%; **17b** (from **15b**), 79%); for compound **17c** (i) **9**, CuSO₄·5H₂O, sodium ascorbate, THF/H₂O, 80 °C (MW), two hours (76%).

phase during matrix-assisted laser desorption/ionization-time-offlight mass spectroscopy analysis is very difficult. Moreover, both the sugars and the fullerene hexa-adduct moieties give rise to a high level of fragmentation²⁵. The unambiguous structural characterization of 17a-17c was, however, greatly facilitated by their high symmetry. Indeed, ¹³C NMR spectroscopy was particularly helpful for the characterization of hexakis adducts of [60]fullerene, as only two signals are usually observed for the sp^2 carbons of C₆₀, which provides evidence for the octahedral symmetry of the fullerene core. As a typical example, the ¹³C NMR spectrum of compound 17a is depicted in Fig. 4. In this case, only two sp^2 carbons are observed in the spectra ($\delta \approx 145.4$ and 141.6 ppm), in which we can also distinguish the two different kinds of triazole rings present in the compound (at $\delta \approx 146.9$ and 123.3 ppm for the carbons of the outer triazole rings and at $\delta \approx 146.4$ and 122.9 ppm for the carbons of the inner triazole rings). In addition, only one signal is detected for all the carbonyl groups ($\delta \approx 163.8$ ppm) and the sp³ carbons of the C_{60} ($\delta \approx 69.3$ ppm), whereas the malonate bridgehead carbons present in the structure are observed at $\delta \approx 45.2$ ppm.

Additional characterization was achieved by DLS analysis (H₂O, 0.01 mg ml⁻¹ and 0.1 mg ml⁻¹), in which, regardless of the

concentration used, we found two or three main size distributions for 17a–17c (Supplementary Fig. 1). This is compatible with a weak aggregation of 17a–17c in water. The first, at around 5–6 nm, must correspond to only one molecule, whereas the second, at ~120–150 nm, and the third, at ≥200 nm, show the aggregation of several molecules. In dimethylsulfoxide (DMSO) (0.1 mg ml⁻¹) (Supplementary Fig. 2), although most of the molecules show no aggregation, the presence of aggregates of different sizes and, especially, of very large aggregates is also detected. The tendency to form aggregates was also confirmed by the broadening of the ¹³C NMR spectrum recorded in D₂O when compared with that recorded in DMSO-d₆ (see the Supplementary Information).

The TEM images of freshly prepared samples reveal the presence of small spherical particles, which correspond to a few or even just one molecule (\sim 4 nm), independently of whether the concentration is 0.01 mg ml⁻¹ or 0.1 mg ml⁻¹, in good agreement with the experimental findings in DLS analyses (Fig. 5 and Supplementary Fig. 3).

XPS was additionally used to confirm the composition of the sugar balls. This surface technique enables the identification of the atoms present on the molecule together with their chemical state and their relative abundance. The survey spectra of superball

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Figure 4 | ¹³C NMR spectrum of tridecafullerene 17a in DMSO-*d*₆. The assignments depict the most representative signals. This spectrum is in full agreement with the T-symmetrical structure of the compound and enables confirmation of the complete functionalization of the alkyne moieties in precursor 16 because of the absence of signals that correspond to the *sp*-hybridized carbons of the alkyne groups.

17a (Supplementary Fig. 4) displays the C 1*s*, O 1*s* and N 1*s* features as expected, with no additional spectroscopy signatures of possible impurities. Moreover, the high-resolution N 1*s* core-level spectrum (Supplementary Fig. 4, inset right) was composed of two different components with a 1:2 ratio of the integrated areas; that located at 400.3 eV is related to one nitrogen atom of the triazole ring (N–N–N) and the other, centred at 398.8 eV, is attributed to the other two nitrogen atoms attached to carbon atoms (C–N)²⁶. The absence of a well-resolved peak around 405.0 eV demonstrates the lack of the electron-deficient nitrogen of the azide group in the final compound^{27,28}. The composition of compounds **17b** and **17c** was also ascertained by their respective XPS analyses, which showed the presence of the expected elements according to their relative abundance (Supplementary Fig. 4 and Supplementary Table 1).

Biological studies. A number of molecules, including DC-SIGN (dendritic cell-specific intercellular adhesion molecule-3-grabbing non-integrin), have been proposed as receptors for the Ebola virus (EBOV)^{29–31}. Although DC-SIGN is not the main receptor in the case of the EBOV, it is thought to play a significant role in the cell entrance of this infectious agent in significant cell populations, such as dendritic cells^{29,32}, and thus facilitate early viral dissemination. Therefore, DC-SIGN can function as a good model for (1) studying the first steps of pathogenesis of EBOV and (2) screening the antiviral strategies based on DC-SIGN-targeting compounds for prevention and treatment purposes. DC-SIGN

recognizes mannosylated and fucosylated oligosaccharides presented in a multivalent manner on the surface of several pathogen envelope GPs. Thus, the preparation of multivalent carbohydrate systems is necessary for an efficient interaction with this receptor as well as for effective competition with the natural ligands. In this study we evaluated the inhibitory effect of giant globular multivalent glycofullerenes in an experiment with the direct infection of Jurkat cells that express the surface receptor DC-SIGN (Jurkat-DC-SIGN) with pseudotyped viral particles that present EBOV-GP. These globular multivalent systems are water soluble and show no cytotoxicity in cell lines, which allows the study of their potential biological function in preventing viral infection. All the multivalent compounds were checked for the possibility of blocking the DC-SIGN receptor in six independent experiments. The results of blocking the DC-SIGN receptor by different compounds are reported as a function of concentration. A 50% inhibition of the infection was calculated with a 95% confidence interval (CI). As a control, infection with DC-SIGN-independent vesicular stomatitis virus envelope GP (VSV-GP)-pseudotyped lentiviral particles²⁹ was performed under the same conditions.

The results obtained in the infection experiment revealed the dependence of the inhibition effect on mannoses. Compound **17b**, which displayed 120 galactoses, as expected, was not able to inhibit the infection process mediated by DC-SIGN. Compounds with 120 mannose-based residues (**17a** and **17c**) showed very strong antiviral activity at picomolar to nanomolar concentrations. Compound **17a** could effectively block EBOV infection at

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Figure 5 | TEM images and DLS analysis of tridecafullerene 17a. These images show small spherical particles with a diameter of around 4 nm, which corresponds to a single molecule. **a**, TEM images of compound **17a** on deposition of a 0.01 mg ml⁻¹ solution in H₂O. **b**, Detail of a particle that apparently corresponds to one molecule. **c**, Width profile of the particle shown in **b**, which has a diameter of ~4 nm in accordance with the DLS data. a.u., arbitrary units



Figure 6 | Biological study of tridecafullerenes 17a-17c. The graphic shows the inhibition of infection with EBOV-GP- or VSV-GP-pseudotyped lentiviral particles of Jurkat DC-SIGN⁺ cells using **17a** (blue), **17b** (green) and **17c** (red). In the *cis*-infection experiments, 2.5×10^5 Jurkat DC-SIGN⁺ cells were challenged with 5,000 TCID of recombinant lentiviral particles. The results represent the mean of six independent experiments ± s.e.m. Compounds **17a** and **17c** show a strong inhibitory activity when EBOV-GP-pseudotyped lentiviral particles are used. No inhibitory activity is detected when VSV-GP-pseudotyped particles (control) are used as the infective agent. Compound **17b** (endowed with galactoses as carbohydrate units) does not show inhibitory activity because it is not able to block the DC-SIGN receptor.

low nanomolar concentrations, with an IC_{50} of 20.37 nM (95% CI = 14.63–28.37 nM). Compound 17c was almost one order of magnitude more potent at inhibiting the infection process, with an IC_{50} of 667 pM (95% CI = 411 pM–1.08 nM) (Fig. 6).

Previous inhibition studies using the same infection model and fullerenes that displayed up to 36 mannoses show relative inhibitory potency (RIP) values at least two orders of magnitude smaller¹⁹. Moreover, huge virus-like particles (VLPs) with a radius of 16 nm and up to 1,640 mannoses³² were 18-fold less potent compared with compound **17c** (see Table 1). These results confirm the efficiency of these systems both to interact with DC-SIGN and to compete with EBOV-GP-pseudotyped particles during their entry into target cells.

The cytotoxic effect of multivalent glycofullerenes was verified by a cell-proliferation assay using the Cell Titer 96 AQueous Non-Radioactive Cell Proliferation Assay (Promega). There were no appreciable cytotoxic effects of compounds 17a–17c at the concentrations used in the infection experiments (Supplementary Fig. 6).

Table 1 | Comparison of IC_{50} and RIP values of different mannosylated multivalent compounds.

Compound	IC ₅₀ (nM)	Mannoses (No.)	RIP*	Reference
17c (120 mannoses)	0.667	120	1.58 × 10 ⁴	This work
VLP (1,620 mannose)	0.91	1,620	8.6 × 10 ²	33
17a (120 mannoses)	20.375	120	5.2 × 10 ²	This work
VLP (540 mannoses)	9.62	540	2.44 × 10 ²	33
C60LL (36 mannoses) †	300	36	1.17 × 10 ²	20
C60 (36 mannoses) [†]	68,000	36	0.5	20
C60 (12 mannoses) [†]	2,000	12	53	20
α-methyl-D-	1.27 ×	1	1	34
mannopyranoside	10 ⁶			

Data obtained from inhibition studies using pseudotyped EBOV particles for the new compounds 17a and 17c in comparison with other carbohydrate multivalent systems previously reported by us; *RIP, calculated as (IC₅₀)mono/IC₅₀*valency ((IC₅₀)mono, IC₅₀ of the monovalent compound; IC₅₀*valency, IC₅₀ of the multivalent compound multiplied by the number of ligands present in the multivalent compound). ¹See Supplementary Fig. 5 for the chemical structure.

Conclusions

In summary, we synthesized giant globular multivalent glycofullerenes in which the central C_{60} core is covalently connected to 12 hexakis adducts of C_{60} to form the first tridecafullerenes reported so far. Each peripheral fullerene is endowed with ten monosaccharides and so a total of 120 carbohydrates decorate the periphery of each molecule. This represents the fastest dendrimer growth ever reported, affording molecular weights as high as 56 kDa.

The syntheses of non-symmetric hexakis adducts (7, 9, 14a, 14b, 15a and 15b) as well as the tridecafullerenes (17a–17c) were accomplished efficiently by using CuAAC click-chemistry reactions. In this way, very sophisticated molecular ensembles were produced in a minimum of synthetic steps, and their apparent structural complexity does not place a limit for their application. Despite the high molecular weights, the new molecules were characterized by standard spectroscopic techniques (FTIR, ¹H NMR and ¹³C NMR spectroscopy) as well as by DLS, XPS and TEM. Interestingly, NMR spectroscopy unambiguously revealed the high degree of symmetry (T_h) in hexakis adducts.

Tridecafullerenes are soluble in water and, therefore, they were tested as globular multivalent systems to inhibit lectin-mediated viral infection processes in cellular assays. In particular, compounds 17a–17c were used to test their ability to inhibit the infection of cells

by an artificial EBOV. These tridecafullerenes were found to block the EBOV infection efficiently in the subnanomolar concentration range. These values surpass by three orders of magnitude (two if the number of mannoses is considered) those exhibited by hexakis adducts endowed with 12 mannoses.

The aforementioned results reveal fullerenes to be appealing platforms for the study of multivalent interactions, particularly because of their biocompatibility and globular presentation. Furthermore, the compounds reported here pave the way to the introduction of dendritic dimers and trimers of monosaccharides as well as to the use of disaccharides to improve significantly the scope of the biological applications of tridecafullerenes.

Methods

Production of recombinant viruses. Recombinant viruses were produced in 293T cells. The viral construction was pseudotyped with the Zaire EBOV envelope GP or VSV-GP and expressed luciferase as a reporter of the infection^{29,33}. One day (18–24 hours) before transfection, 5×10^6 293T cells were seeded onto 10 cm plates. The cells were cultured in DMEM medium supplemented with 10% heat-inactivated fetal bovine serum, 25 mg gentamycin and 2 mM L-glutamine. A few minutes before transfection, the medium on the transfection plates was changed to 9 ml DMEM and chloroquine was added to a 25 µM final concentration. Transfection reactions with all the reagents at room temperature (r.t.) were prepared in 15 ml tubes: 183 μl of 2 M CaCl₂, 500 ng of EBOV-GP or 2 µg of VSV-GP, 21 µg of pNL4-3 luc³⁴ and 1,300 µl of H₂O. Next, 1.5 ml of 2 × HBS (HEPES buffer saline) pH 7.00 was added quickly to the tubes and bubbled for 30 seconds. HBS/DNA solution was gently dropped onto the medium. After eight hours of incubation at 37 °C with 5% CO₂, the medium on the transfection plates was changed to 10 ml DMEM and once again, one day after transfection, to 7 ml DMEM. Transfection supernatants were harvested after 48 hours, centrifuged at 1,200 revolutions per minute for ten minutes at r.t. to remove cell debris and stored frozen at $-80 \degree C^{32,33,35}$.

EBOV infection experiments. Infection was performed on Jurkat cells

(T-lymphocyte cell line) that expressed the receptor DC-SIGN on their surfaces. As the EBOV does not infect T-lymphocytes, its entry is absolutely dependent on DC-SIGN for infection of the Jurkat cells^{29,36}.

Jurkat–DC-SIGN cells (2.5×10^5) were plated into each well of a 96-well plate. Cells were incubated at r.t. for 20 minutes with the carbohydrate-based compounds and then challenged with 5,000 TCID (tissue culture infective dose) of recombinant viruses. After 48 hours of incubation, the cells were washed twice with PBS and assayed with the Luficerase Assay System (Promega).

The range of concentrations tested for compounds 17a–17c was 1 pM to 10 μ M. As a control, an experiment of infection with VSV-GP pseudoviruses was performed under the same conditions. Infection with VSV-GP was independent of the presence of DC-SIGN receptor.

Statistical analysis. The values of percentages of inhibition of the infection presented in Fig. 6 correspond to the mean of six independent experiments, and the error bars correspond to s.e.m. The IC_{50} values were estimated using GraphPad Prism v6.0 with a 95% CI and settings for normalized dose–response curves.

Cytotoxicity assay. The Cell Titer 96 AQueous Non-Radioactive Cell Proliferation Assay (Promega) was used. Jurkat–DC-SIGN cells (5×10^5 cells per well) were seeded into the wells of 96-well plates and left to culture in the presence of different concentrations of glycofullerenes for the time of the infection assay. After 48 hours, the proliferation cell assay was performed. Briefly, 2 ml of MTS (5-(3-carboxymethoxyphenyl)-2-(4,5-dimethylthiazolyl)-3-(4-sulfophenyl) tetrazolium) solution was mixed with 100 µl of phenazine methosulfate (PMS) solution. The mix MTS/PMS in a volume of 20 µl was pipetted into each well of the 96-well assay plate, which contained 100 µl of cells in the culture medium. The plate was incubated for two hours at 37 °C in a humidified 5% CO2 atmosphere. After the incubation period, the absorbance at 490 nm was recorded using an enzyme-linked immunosorbent assay plate reader. As a control of the toxicity, the viability of the cells in the presence of 0.1% Triton-X100 was measured. The results of the assay are presented as the percentage of viable cells, which was calculated from the absorbance at 490 nm as compared with the absorbance shown by cells incubated without the addition of glycofullerenes, which represent 100% cell viability.

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Author contributions

A.M., D.S., I.N., M.H. and K.B. carried out the synthesis and characterization of all the new derivatives. L.R.-P. and J.-S.R. realized and analysed the DLS and TEM. L.R.-P. realized the XPS analyses and contributed to the writing of the paper. J.L. and R.D. realized the biological and cytotoxicity studies. B.M.I., S.P.V., J.R., R.D., J.-F.N. and N.M. designed the project, supervised the work, discussed the data and wrote the manuscript.

Additional information

Supplementary information and chemical compound information are available in the online version of the paper. Reprints and permissions information is available online at www.nature.com/reprints. Correspondence and requests for materials should be addressed to J.R., R.D., J.F.N. and N.M.

Competing financial interests

The authors declare no competing financial interests.