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Trehalose-based Janus cyclooligosaccharides: "Click" synthesis and DNA-directed assembly into pH-sensitive transfectious nanoparticles

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A convergent preparation of Janus molecular nanoparticles by thiourea-"clicking" of α, α' -trehalose halves has been implemented; the strategy let access macrocyclic derivatives with seggregated cationic and lipophilic domains that in the presence of DNA undergo pH-dependent self-assembling into lamellar superstructures, as established by electrochemical, spectroscopical (SAXS), microscopical (TEM) and computational techniques, that mediate transfection *in vitro* and *in vivo*.

The "Janus" term was initially adopted from the Roman mythology by materials scientists to denote particles with a three-dimensionally dissymmetrical distribution of physical/chemical features.¹ Because of their morphological and chemical asymmetry, Janus particles can assemble into unique superstructures that could not be obtained by the assembly of homogeneous particles.² The notion has been further extended to describe the entities exhibiting dichotomy in general, from covalent structures to supramolecular assemblies and nanocomposites. Among them, molecular Janus architectures are particularly interesting. Their decreased sizes of typically only a few nanometers and the possibility of precise chemical tailoring make them ideally suited for the incorporation of functional and structural hierarchy. However, most of the molecular Janus entities reported so far are based on dendritic³ or polymeric scaffolds⁴ with high degree of conformational flexibility. Alternatively, connecting together hydrophobic and hydrophilic 3D molecular frameworks with precisely defined chemical structures, sizes, shape symmetry and surface functional

been used to generate new monodisperse Janus-like topologies.⁵ The examples on record include snowman or dumbbell-like nanostructures obtained from polyhedral oligomeric silsesquioxane (POSS), C₆₀ fullerene. polyoxometalate (POM), cyclodextrin (CD) or calixarene (CA) building blocks that still keep a relatively high degree of conformational flexibility and in which the spatial separation between the components fluctuates depending on the nature of the tether.⁶ "Monomeric" Janus-like MNPs, better mimicking "hard" Janus objects, can be accessed by selective facial functionalization when the starting precursor exhibit intrinsic molecular anisotropy, as it is the case for CD and CA platforms.⁷ Recently, Chiara and coworkers⁸ have reported an elegant strategy to break the symmetry and create a "perfect" Janus feature in a cubic POSS precursor by exploiting the geometrical complementarity of the cube face with a conveniently tetra-functionalized partner. In any case, with few exceptions such divergent approaches require multiplestep synthesis involving delicate control over reaction conditions and nontrivial purifications to warrant diasterochemical purity. Here we present an alternative convergent tactic based in the multipoint covalent assemblage of dissimilar halves that allows elaborating "perfect" Janus cyclooligosaccharides made of α, α' -trehalose (cyclotrehalans, CTs) whose self-assembly behaviors are dictated by symmetry breaking in both functional groups and supramolecular interactions. The methodology has been applied to the efficient synthesis of MNPs capable of reversibly condensing DNA into self-assembled lamellar nanocomplexes (CTplexes) with transfection-mediating capabilities in vitro and in vivo (Figure 1).

groups, referred to as "molecular nanoparticles" (MNPs), has

Contrary to other macrocyclic MNPs such as CDs or CAs, canonic CTs exhibit identical faces that are brought together after a very efficient macrocyclization step involving a double "click"-type thiourea-forming reaction.⁹ The concave shape of the constitutive α, α' -trehalose moieties, dictated by the concurrence of two exo-anomeric effects at the glycosidic linkages, preorganizes the disaccharide to favour macrocyclic

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over oligomeric structures upon bridging through the primary positions. Thiourea tethering further reinforces macrocyclization efficiency: after the first isothiocyanateamine coupling, the formation of a seven-membered intramolecular hydrogen bond stabilizes the Z,E-rotamer and places the next reacting groups in close proximity to zip the macroring. We envisioned that stacking of amphiphilic molecular Janus CTs composed of hydrophobic and cationic moieties would be prohibited by the repulsive interactions introduced by the charges on the surface. In the presence of an oligonucleotide chain, the interplay of attractive coulombic and hydrophobic interactions may promote DNA-directed assembly into nanocomplexes whose stability would depend on the protonation extent, which may be used for pH-sensitive non-viral gene delivery.



Figure 1. Schematic representation of Janus cyclooligosaccharides obtained by thiourea-"clicking" of α , α '-trehalose halves. In the cartoon, red circles represent cationic groups and blue circles lipophilic substituents. Hierarchical assembly in the presence of plasmid DNA to give nanocomplexes (CTplexes) with a lamellar structure is represented.

To check the above working hypothesis, the diisothiocyanate and diamine precursors **1** and **2**, bearing respectively six hexanoyl tails and six protected cationizable amine groups at the secondary hydroxyls, were synthesized (ESI) and reacted (\rightarrow **3**) to give, after deprotection, the multihead-multitail Janus CT **4** in over 80% yield. Further elaboration of this prototype was performed by thiourea coupling of **4** with the branching building block **5** (\rightarrow **6**), affording the dendritic adduct **7** (Scheme **1**). Both compounds were purposely designed to incorporate structural elements (hexanoyl tails, thiourea Hbond donor centres, multivalent/dendritic amine clusters) previously found advantageous in the design of molecular artificial viruses.¹⁰

The effective charges available for coulombic interaction between the Janus CTs and DNA, determined from ζ -potential measurements, were found to be significantly different from those expected considering the ionizable N and P atoms in each partner. For instance, CT **4** exhibited positive net charges that are around 75% of the nominal ones in the presence of DNA (ESI, Table S1). Contrary to linear double helix DNA, which Page 2 of 4

keeps its negative charge (-2 per base pair) totally available for the gene vector,¹¹ plasmid DNA rendered an available negative charge per bp of around 7% of its nominal value when being compacted by CT **4.** It means that effective charge ratios, ρ_{eff} , are around 10- to 11-folds the nominal charge ratios ρ_{nom} (also named as N/P). These data support that the plasmid remains in a supercoiled conformation at physiological conditions, retaining an important percentage of its cationic sodium counter ions after interaction with the vector.



Circular dichroism spectra of DNA registered in the absence and in the presence of compounds **4** or **7** at pH 7.4 unequivocally evidenced the existence of interactions that distorted the typical B-type structure of uncomplexed DNA, probably to a Z-DNA form. Thus, a concentration-dependent

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decrease in the intensities of the positive and negative bands at 278 and 245 nm, arising from stacking interactions between bases and from polynucleotide helicity, respectively, was initially observed (Figure 2A and ESI, Figure S14), with a change in sign at CT:DNA mass molar ratios higher than 2 (Figure 2B).



Figure 2. A, Circular dichroism spectra in the 230-325 nm region for solutions containing Janus CT **4** (0, 0.37, 0.75, 1.12, 1.50, 1.87, 3.74, 5.91 and 9.85 μ M) and DNA fixed 1.23 μ M concentration) in HEPES at 25 °C. B, Ellipticity values as a function of the **4**:DNA mass molar ratios for the maxima of the two bands appearing near 245 nm (open symbols) and 275 nm (filled symbols). C, SAXS diffractograms for **4**:pDNA complexes formulates at several effective charge ratios (ρ_{eff}). D, TEM micrograph of **4**:pDNA complexes formulated at N/P 10.

Small-angle X-ray scattering (SAXS) diffractograms (Intensity vs q factor) of 4:pDNA complexes at several effective charge ratios (ρ_{eff} = 5, 10, 41 and 81, i.e. N/P ratios 0.45, 0.91, 3.7 and 7.4, respectively) showed three peaks that indexed well to a lamellar lyotropic liquid crystal phase, L_{α} , regardless of ρ_{eff} (Figure 2C). This lamellar arrangement, known to be correlated with potentially high transfection efficiencies,¹² arises from self-assembling of the Janus CT molecules into lipidic bilayers in the confined space between quasi-parallel DNA supercoils, with thicknesses represented by d_m, and d_w, respectively, being d = d_m + d_w . The characteristic interlayer distances, d, directly related to the q factor (d = $2\pi n/q_{hkl}$, n is the diffraction order) remains constant with ρ_{eff} at 5.2 ± 0.1 nm (Figure 2C and ESI, Figure S13). Considering these d values and the fact that pDNA supercoils needs around $d_w\ {\sim}2.5$ nm to be sandwiched by CT bilayers, it can be deduced that the thickness of the bilayer, $d_{\rm m},$ must be around 2.7 nm. TEM micrographs (Figure 2D) of the CTplexes also showed

aggregates with a clear multilamellar pattern, in full agreement with SAXS results.

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To get a deeper insight into the interactions that govern the hierarchical process that leads to CTplex formation, the stability of a head-to-head dimer of 4, the smallest unit of the bilayer arrangement, was first studied by molecular mechanics (MM) in explicit water. The minima binding energy (MBE) structure was initially obtained by sequentially approaching two Janus CT molecules in the corresponding perhydrochloride form, with their center of mass on the y axis in a coordinate system (ESI, Figure S16). The binding energy profile corresponded to a structure in which the lipidic hexanoyl chains significantly intercrossed and the cationic arms onto the opposite face of the Janus macrocycle adopted an open bouquet disposition to avoid electrostatic repulsions and steric clashes (ESI, Figure S17). The MBE structure was then placed between two symmetrically located and oriented B DNA helix fragments containing twelve nucleotides (CGCGAATTCGCG) and the oligonucleotide chains were approached simultaneously in steps of 1 Å (Figure 3) through the major groove. The binding energies obtained by this procedure, which were rather favorable, led to an efficient packing. Remarkably, the distance between the centers of mass of the nitrogen clusters in the CTs, which would represent a measurement of the CT bilayer thickness, was reduced from near ~3.3 nm for the isolated CT dimer to ~2.6 nm for the most stable structure of the CTplex, very well fitting the above SAXS experimental data. The most stable CTplex structure generated (Figure 3) was used as the starting conformation for 1.0 ns MD simulations, which confirmed the stability of the nanocomplex through the trajectory (ESI, Figures S20 and S21). Interestingly, when calculations were conducted on a fully cationized form of 4 (by removing the chloride anions) the dimer became unstable. We speculated that increasing protonation in the acidic environment of the endosomes after CTplex cellular uptake will then destabilize the Janus CT bilayers, contributing both to endosome disruption and intracellular DNA release, thereby facilitating the transfection process.13



Figure 3. Coordinate system used for DNA fragments approaching the MBE structure of the CT **4** dimer along the *y* coordinate by the major groove (left) and MBE structure of the resulting CT:DNA complex (right).

Electrophoretic mobility experiments in agarose gel followed by visualization with the fluorescent intercalating agent GelRed[®] confirmed that full pDNA complexation was achieved with both Janus CTs **4** and **7** at N/P ratios >1 (ρ_{eff} > 10). After treatment with a nuclease and dissociation with sodium

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dodecylsulfate (SDS), intact DNA could be recovered from CTplexes, demonstrating that the DNA cargo was protected from the environment following self-assembling (ESI, Figure S22). CTplexes formulated with 4 and 7 at N/P ratios 5 and 10, (hydrodynamic diameter 123 to 230 nm; ζ-potential +18 to +30 mV. See ESI, Table S3), promoted transfection in African green-monkey epithelial kidney COS-7 cells (Figure 4A) and human hepatocellular carcinoma HepG2 cells (ESI, Figure S22) with efficiencies that were comparable to those obtained with polyplexes formulated with branched poly(ethyleneimine) (bPEI), a commonly used positive control for nonviral gene delivery, even in the presence of serum. Noteworthy, no toxicity was observed for any CTplex formulation, compared with 60-70 % cell viability for bPEI polyplexes. The nanocomplexes formulated with 4 and 7 at N/P 10 were next injected systemically into mice, and their activity was compared with control PBS and naked DNA. The results, based on luciferase reporter gene expression, indicated that 24 h after the intravenous administration of the CTplexes transfection occurs mainly in the liver and lung (Figure 4B), with negligible luminescence detected in other organs. Increases in transfection levels relative to naked DNA were above one and two orders of magnitude for 4: pDNA and 7:pDNA CTplexes, respectively. The advantages of the convergent synthetic methodology in terms of versatility and ease of manipulation of the precursors should now allow the preparation of a larger collection of monodisperse Janus MNPs for structure/self-assembling/biological activity relationship studies in view of developing optimized nanodevices programmed for biomedical applications.



Figure 4. A, Transfection efficiency in COS-7 cells for CTplexes formulated with Janus CTs 4 or 7 and the luciferase-encoding reporter gene pCpG-hCMV-SPEC-eFLuc at N/P ratios 5 and 10 in the absence and presence of 10% fetal bovines serum (FBS). Data obtained with bPEI polyplexes (N/P= 5 and 10, 10% FBS) under identical conditions are included for comparison. The data represent the mean \pm SD of three wells and are representative of three independent determinations. B, Gene expression conducted *in* vivo after intravenous administration of 60 µg of pCpG-hCMV-SPEC-eFLuc formulated as CTplexes with 4 and 7 at a N/P 10. Bars represent the mean \pm SD (n=7 animals).

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