Polyphenylene dendrimers as scaffolds for shape-persistent multiple peptide conjugates
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The present work describes synthetic concepts for the coupling of peptides to polyphenylene dendrimers (PPDs). Novel functionalized cyclopentadienones have been synthesized whose Diels–Alder cycloadition with various core molecules leads to polyphenylene dendrimers possessing (protected) amino or carboxyl groups. In addition, the resulting functionalized molecules exhibit the characteristic shape-persistence and monodispersity of PPDs. Their functions have been used for the attachment of polysylene to the dendritic scaffold. Three different methods for the decoration of dendrimers with polypeptides are presented. First, polylsine segments are grafted from the surface of the dendrimers employing α-amino acid N-carboxyanhydride (NCA) polymerization. Second, the C-terminal carboxyl groups of protected polypeptides are activated and then coupled to the amino groups on the surface of the PPD. Finally, cysteine terminated, unprotected peptide sequences are attached to polyphenylene dendrimers utilizing the addition of the sulphydryl group of a cysteine to the maleimide functions on the dendrimer surface. Moreover, Diels–Alder cycloaddition of suitably functionalized cyclopentadienons to a desymmetrized core molecule allows the design of a dendritic scaffold with a specific number of different anchor groups on its periphery. These approaches are important for the tailoring of new, shape-persistent, polyfunctional multiple antigen conjugates.

1. INTRODUCTION

Dendrimers are monodisperse, highly branched macromolecules (1–3). Their synthesis involves either a convergent or divergent protocol, both implicating the creation of a core molecule and an iterative procedure for the construction of the dendrimer. The high number of functional groups and the well-defined structure make dendrimers attractive carriers for the design of nanoparticles with potential uses in clinical and diagnostic applications. Recent advances in the biomedical applications of various dendritic structures are reviewed by Striba et al. (4). The polyphenylene dendrimers (PPDs) described here are unique shape-persistent and polyfunctional molecules bearing topologically isolated, non-interfering functional groups (5). This is essential for the active function of PPDs decorated with short peptide sequences because the bioactivity of the latter is strongly influenced by inter- and intramolecular chain interactions. The attachment of a defined number of oligopeptides onto PPDs has been inspired by the recent achievement of multiple antigen peptides (MAPs) (6).

MAPs are nanoparticles, which, when introduced to an organism, are able to trigger an immune response. In nature, the recognition of foreign bodies (antigens) by antibodies proceeds via the so-called key and lock principle, in which the shape of the antibodies’ receptors is complementary to a given pattern on the surface of the antigen that is to be recognized. Complex antigens have a variety of epitopes on different parts of their surface, each of which can be recognized separately by a different antibody. Antigens of special interest are so-called immunogens that trigger an immune response, i.e., the production of antibodies.

Very often, the identifiable part for the antibody is a polypeptide segment. The attachment of several recognizable polypeptides to a nanoparticle is referred to as a MAP. These structures have proven to exhibit amplified immunogenicity and pave the way toward fully synthetic vaccines (7). Currently, MAPs are constructed by attaching the desired polypeptides to a dendritic matrix composed, most frequently, of branched polylysine. It has been shown that a stronger immunogenic effect may be achieved when the MAP system contains two different antigen peptides (8).

However, the polylsine carriers do not possess a defined spatial structure, which is believed to have an unfavorable effect on their function (6). In fact, inter- and intramolecular interactions of the antigens may modify their secondary structure as well as limit their acces-
sibility. Therefore, the synthesis of nanosized peptide carriers comprising a well-defined structure and shape-persistence may provide improved MAPs. In this respect, the polyphenylene dendrimers are very promising templates due to their structural perfection and precise spatial definition of functional units in the center, in the dendritic scaffold, or at the periphery. Furthermore, they are chemically inert under the conditions of living organisms, which is another important prerequisite for biologically oriented applications.

In addition, the polyphenylene dendrimers can be synthesized not only as all-carbon structures but also using a strongly fluorescent perylenediiimide core (PDI). Recently, we have described several synthetic approaches toward water-soluble and fluorescent PDI chromophores (9). The chromophore adds a further important property, namely, the possibility of facile optical detection of the dendritic conjugate, and such molecules may find diagnostic application. Thus, the aim of the present work is the development of reliable and expandable synthetic routes affording peptide-functionalized polyphenylene dendrimers with prospective application for attachment of antigen peptides onto a shape-persistent, polyphenylene scaffold. A further development is the synthesis of polyphenylene dendrimers bearing two different polypeptides. While we have recently described routes toward the synthesis of novel polyphenylene dendrimers with lysine and glutamic acid moieties anchored on the surface (10), we herein extend this idea by employing different synthetic approaches for the preparation of peptide—PPD conjugates. Polyphenylene dendrimers were functionalized with polylysine following three parallel routes. First, oligosilanes were grafted directly from the surface of polyphenylene dendrimers employing α-amino acid N-carboxyanhydride (NCA) polymerization. Second, protected pentalysine moieties were coupled to PPDs via carbodiimide activation. Third, unprotected hexapeptides, composed of five lysine residues and a C-terminal cysteine, were specifically attached to the polyphenylene scaffold. This was achieved employing a Michael addition of the thiol containing peptides to maleimide functions on the dendrimer periphery. An important prerequisite for the synthesis of polyphenylene dendrimers decorated with two different peptide units is the creation of a core comprising two types of anchor groups, allowing the independent coupling of different peptide sequences.

2. EXPERIMENTAL SECTION

General Information. Unless stated otherwise, all solvents and reagents are commercially available and were used as received. 1H NMR spectra were recorded on Bruker AMX 250, Bruker AMX 300, Bruker Advance DRX 500 or 700 spectrometers. Spectra were referenced to the residual proton signal of the deuterated solvent. Molecular weights were determined with matrix-assisted laser desorption/ionization time-of-flight (MALDI-TOF) or field desorption (FD) mass spectrometry; MALDI-TOF mass spectra were recorded on a Bruker MALDI-TOF mass spectrometer, and FD mass spectra were recorded on VG Instruments ZAB 2 Se-FPD. Peptide syntheses were carried out on an Applied Biosystems peptide synthesizer model 433A. Ultraviolet-visible (UV—vis) spectra and fluorescence spectra were recorded using a Perkin-Elmer Lambda 9 and SPEX Fluorolog 2 spectrophotometer, respectively. Circular dichroism (CD) spectra were measured on a Jasco J-715 spectropolarimeter. Gel permeation chromatography (GPC) was performed at 60 °C with a setup consisting of Waters 510 pump and a series of three styrene-divinylbenzene columns (300 × 8 mm) with pore sizes of 500, 10^5, and 10^6 Å (Polymer Standard Services, Mainz, Germany). The elemental analyses were carried out by the Microanalytical Laboratory of the University of Mainz (Germany).

The amino acids were free acids, and coupling was facilitated by the use of O-benzotriazole-N,N,N',N''-tetramethyluronium hexafluorophosphate (HBTU) and N-hydroxybenzotriazole (HOBt). Purity was checked using analytical reverse-phase high performance liquid chromatography (RP-HPLC, AKTA-Purifier, Amersham-Pharmacia Biotech). The column used was a reverse-phase C8 column (Sephasil peptide C8, volume 4.155 mL, 5 μm particle size). The synthesis of first- and second-generation dendrimers with maleimide functionalities (9a–d and conjugates 38a–d) have been described previously (10). Compounds 1 (11), 2 (12), 3 (13), 4 (14), and 16 (15) were synthesized according published procedures.

Procedures. Compound 7a. 3-(4-Aminophenyl)-2,4,5-triphenylcycloenta-2,4-diene (6a) (1.4 g, 3.55 mmol), N-BOC-γ-aminobutyric acid (2.4 g, 17.71 mmol), N-(3-dimethylaminopropyl)-N-ethylcarbodiimide hydrochloride (EDC) (2.4 g, 17.71 mmol), and DMAP (1.3 g, 10.6 mmol) were dissolved in 50 mL of DMF and stirred for 48 h under an argon atmosphere. Thereafter, DCM (50 mL) was added, and the reaction mixture was washed twice with water. The organic phase was isolated, dried over magnesium sulfate, and filtrated, and the solvent was evaporated. The dried residue was purified by column chromatography with DCM/acetone (10:1) eluent to give 7a as brown powder. Yield: 2.07 g (82.1%). FD (M+): m/z = 585.5; 1H NMR (250 MHz, CDCl3) δ 7.43 (d, 2H, J = 8.48 Hz), 7.32–7.18 (m, 10H, arom.), 6.87 (d, 2H, J = 7.85 Hz), 3.21 (q, 2H, J = 6.28 Hz), 1.44 (s, 9H); 13C NMR (75 MHz, DMSO-d6) δ 199.6, 171.1, 171.0, 155.5, 154.5, 139.6, 139.5, 132.7, 130.7, 130.5, 129.7, 129.6, 128.8, 128.5, 128.0, 127.9, 127.5, 127.4, 126.8, 124.9, 124.1, 118.1, 117.9, 77.4, 33.7, 28.2, 25.5, 11.0; mp 106 °C.

Compound 7b. 3,4-Bis-(4-aminophenyl)-2,5-diphenyl-cycloenta-2,4-diene (6b) (1.1 g, 2.69 mmol), N-BOC-γ-aminobutyric acid (5.46 g, 26.9 mmol), EDC (5.16 g, 26.9 mmol), and DMAP (1.04 g, 8.5 mmol) in 50 mL of DMF. The crude product was purified by column chromatography with DCM/acetone (10:1) eluent to give 7b as brown powder. Yield: 1.54 g (73%). FD (M+): m/z = 782.6; 1H NMR (250 MHz, DMSO-d6) δ 7.43 (d, 2H, J = 8.85 Hz), 7.30–7.43 (m, 10H, arom.), 6.83 (d, 2H, J = 8.53 Hz), 2.92 (q, 2H, J = 6.63 Hz), 2.26 (t, 4H, J = 7.27 Hz), 1.65 (p, 4H, J = 7.27 Hz), 1.35 (s, 18H); 13C NMR (75 MHz, DMSO-d6) δ 199.8, 171.2, 155.8, 139.8, 139.7, 139.0, 132.0, 129.8, 129.7, 128.6, 128.4, 124.9, 124.1, 118.1, 117.9, 77.4, 33.7, 28.2, 25.5, 11.0; mp 106 °C.
Compound 8b. This compound was prepared analogously to 7b using 6b (1.1 g, 2.69 mmol), N-Z-aminobutyric acid (6.39 g, 26.9 mmol), EDC (7.92 g, 26.9 mmol), and DMAP (1.97 g, 16.1 mmol) in 50 mL of DMF. The crude product was purified by column chromatography with DCM/acetonitrile (2:1) as eluent to give 8b as brown powder. Yield: 1.75 g (76.2%). FD (M^+): m/z = 853; 1H NMR (250 MHz, DMSO-d_6) δ 7.44 (d, 4H, J = 8.53 Hz), 7.33–7.13 (m, 24H, arom.), 6.84 (d, 4H, J = 8.53 Hz), 4.99 (s, 4H), 3.02 (q, 4H, J = 6.32 Hz), 2.29 (t, 4H, J = 7.27 Hz), 1.65 (p, 4H, J = 5.95 Hz); 13C NMR (75 MHz, DMSO-d_6) δc 199.8, 171.3, 171.2, 156.3, 154.5, 139.8, 139.7, 137.4, 131.0, 130.0, 128.5, 125.8, 127.9, 127.6, 127.1, 124.6, 118.3, 65.4, 46.4, 33.8, 25.3; mp 248 °C.

Compound 9. This compound was prepared analogously to 7a using 6a (0.7 g, 1.8 mmol), hexanediol acid monomethyl ester (1.44 g, 9 mmol), EDC (1.72 g, 9 mmol), and DMAP (0.66 mmg, 5.4 mmol) in 50 mL of DMF. The crude product was purified by column chromatography with DCM/acetonitrile (50:1) as eluent to give 9 as brown powder. Yield: 0.830 g (85%). FD (M^+): m/z = 542; 1H NMR (250 MHz, CD_2Cl_2) δ 7.51–7.16 (m, 15H, arom.), 6.96 (d, 2H, J = 6.91 Hz), 6.85 (d, 2H, J = 8.53 Hz), 3.63 (s, 3H), 2.49–2.25 (m, 4H), 1.80–1.58 (m, 4H); 13C NMR (75 MHz, CD_2Cl_2) δc 199.9, 179.0, 130.8, 130.7, 129.8, 129.0, 128.6, 128.5, 128.4, 127.9, 119.1, 51.9, 34.1, 24.8, 24.7.

Compound 14a. Compounds 3 (0.1 g, 0.855 mmol) and 7a (0.597 g, 1.02 mmol) were dissolved in 31.5 mL of a oxylene/toluene/DMSO mixture (5/5/2). The resulting solution was heated at 150 °C for 36 h under an argon atmosphere. The reaction mixture was precipitated in petroleum ether, and the crude product was purified by chromatography (silica gel, CH_2Cl_2/acetone 2:1) to give 14a. Yield: 202 mg (70%). MALDI-TOF MS ((M + Na)^+): m/z = 3426; 1H NMR (250 MHz, d_6-DMSO) δ 9.65, 9.59 (2H, NH), 7.91 (s, 4H, arom.), 7.42 (s, 4H, arom.), 7.33–6.66 (m, 98H, arom.), 2.89 (q, 8H, J = 6.63 Hz), 2.87 (sept, 4H, J = 6.32 Hz), 1.71 (q, 4H, J = 7.27 Hz), 1.59 (p, 8H, J = 7.27 Hz), 1.34 (s, 36H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz), 1.34 (s, 72H), 1.02 (d, 24H, J = 5.68 Hz).
4H, 1J = 6.31 Hz), 2.20 (br, 32H), 1.64 (br, 32H), 1.02 (br, 24H); 13C NMR (75 MHz, DMSO-d6) δ 170.6, 156.1, 156.0, 141.5, 141.3, 140.9, 140.3, 139.7, 139.4, 139.0, 138.4, 137.5, 137.2, 136.8, 136.5, 134.6, 134.2, 131.6, 131.2, 130.9, 129.5, 128.3, 127.7, 126.7, 126.3, 117.3, 116.9, 60.2, 60.1, 33.6, 28.9, 25.2, 23.8.

**Compound 18a.** Compound 17a (0.26 g, 0.035 mmol) was dissolved in 6 mL of trifluoroacetic acid and 210 μL of 5.7 M HBr in acetic acid was added. The reaction mixture was stirred at room temperature for 2 h and added to a saturated NaHCO3 solution. The precipitate was filtered and dried under reduced pressure. Yield: 215 mg (97%).

**Compound 18b.** Compound 17b (0.23 g, 0.025 mmol) was dissolved in 6 mL of trifluoroacetic acid and 280 μL of 5.7 M HBr in acetic acid were added. The precipitate was filtered and dried under reduced pressure. Yield: 170 mg (96%).

**Compound 19.** Compounds 2 and 5b (0.408 g, 0.55 mmol) in o-xylene (15 mL) were heated at 160 °C for 24 h under an argon atmosphere. The reaction mixture was concentrated to 2 mL and heated at 160 °C for 24 h under an argon atmosphere. The reaction mixture was concentrated to one-fourth of the initial volume, and the product was purified by precipitation from methanol to give a yellow powder. Yield: 0.130 g (84%). MALDI-TOF MS (M+): m/z = 2875; 1H NMR (250 MHz, CDCl3) δH 7.74–6.21 (m, 134H, arom.), 3.62 (s, 6H), 2.52–2.18 (m, 18H), 1.69–1.55 (m, 8H); 13C NMR (75 MHz, CDCl3) δC 171.0, 168.1, 149.4, 149.0, 144.9, 144.6, 142.9, 142.5, 141.5, 141.3, 141.1, 140.7, 140.4, 140.2, 140.0, 139.7, 139.3, 136.5, 136.3, 135.9, 132.9, 132.3, 132.2, 130.8, 130.4, 130.1, 129.7, 129.2, 128.6, 128.5, 128.4, 128.2, 127.3, 126.7, 126.0, 120.2, 120.0, 118.3, 118.0, 114.2, 113.9, 63.9, 51.9, 37.6, 34.1, 25.3, 24.9.

**Compound 20.** To a solution of 20 (0.100 g, 0.035 mmol) in tetrahydrofuran (2 mL) was added an aqueous solution of LiOH (0.300 mL, 2.3 M), and the mixture was heated at 80 °C for 12 h under an argon atmosphere. The reaction mixture was added to distilled water, and the product was extracted with DCM (3 × 20 mL). The organic phases were collected, washed twice with water, and dried with MgSO4, and the solvent was evaporated to give 22 as yellow powder. Yield: 0.093 g (94%). MALDI-TOF MS: m/z = 2847 g/mol (M+), 2869 g/mol ((M + Na)+).

**Compound 23.** Compound 22 (0.080 g, 0.032 mmol) was dissolved in THF (0.5 mL), and 2 N HCl (1 mL) added. After 10 min, the mixture was cooled with ice and concentrated HCl (0.5 mL) added. The mixture was stirred for 20 min, and the solvents were evaporated under high vacuum. The final product was isolated as a bright brown powder, which was used in the next step without additional purification. Yield: 0.055 g (89%).

**Compound 24.** Compound 23 (0.065 g, 0.034 mmol) and 4-(2,5-dioxopyrrol-1-yl)butanoyl chloride (0.085 g, 0.41 mmol) were dissolved in a mixture of DMF (5 mL) and triethylamine (0.5 mL). The solution was stirred for 6 h at RT under an argon atmosphere. The product was purified by precipitation in methanol to give a dark yellow powder. Yield: 0.080 g (92%). MALDI-TOF MS ((M + Na)+) m/z = 2873 g/mol; 1H NMR (250 MHz, CDCl3) δH 7.56–6.53 (m, 102H, arom.), 3.59–3.48 (m, 8), 2.45–2.12 (m, 16H), 1.95–1.65 (m, 16H); 13C NMR (175 MHz, DMSO-d6) δC 174.1, 170.9, 169.9, 162.2, 143.7, 141.2, 140.2, 140.0, 139.5, 139.0, 136.7, 136.4, 134.3, 132.9, 131.2, 130.0, 129.5, 128.4, 127.6, 126.8, 126.6, 126.2, 125.4, 117.3, 116.9, 36.8, 35.7, 33.4, 30.7, 24.4, 21.4, 23.6.

**Compounds 36a–d.** The PDI dendrimers 15a,b and 18a,b together with 5 equiv of peptide and 20 equiv of HOBt and HBTU were dissolved in a minimum quantity DMF and treated with 50 equiv of diisopropylethylamine (DIPEA). The reaction was stirred for 5 days at room temperature, and the product afterward was precipitated from diethyl ether. The resulting powder contained dendrimer–peptide–conjugate and peptide. Purification by dialysis proved impossible, and thus the mixture was dissolved in TFA and deprotected with HBr in acetic acid by the method described above. The crude product was precipitated from diethyl ether and subsequent ultrafil-
Table 1. Synthesis and Properties of Peptide-Decorated PDI-Dendrimers

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* Notation: x, number of polypeptide chains per molecule; m, number of lysine residues per chain. The polydispersity ($M_w/M_n$) of the peptide-dendrimer conjugates is calculated from the gel permeation chromatography data, obtained in DMF against polystyrene standards.

3. RESULTS AND DISCUSSION

The synthesis of polyphenylene dendrimers commences with a multi-ethynyl-substituted core molecule, namely, tetrakis-(4-ethynylphen-1-yl)-methane (1), its desymmetrized derivative 2, as well as the PDI core 3 (Scheme 1).

We emphasize that the geometry of the core determines the overall shape of the resulting cascade molecules (16). The growth of each dendritic layer is based upon an iterative process including a Diels–Alder reaction and subsequent deprotection of ethynyl functions (5). Scheme 2a introduces a functionalized cyclopentadienone 4 used as $A_2$-type branching agent (16). This molecule can act both as diene and after removal of the silyl protective groups from the ethynyl groups, as dienophile. In the [4+2]-cycloaddition between the cyclopentadienone and a free ethynyl group of the core molecule or the preceding dendrimer generation, a pentaphenylbenzene unit is formed under the extrusion of carbon monoxide (17–19).

3.1. Synthesis of Functionalized Cyclopentadienones. By using prefunctionalized cyclopentadienones 5ab, 7ab, 8ab, 9 in the Diels–Alder reaction, polyphenylene dendrimers with chemical functions are obtained. These functions are either protected amino (5ab, 7ab, 8ab) or carboxyl (9) groups that can be readily converted into versatile anchor groups. The synthesis of the func-
Functionalized cyclopentadienones is depicted in Scheme 2b. The preparation of the imino (5a,b) and amino (6a,b) functionalized cyclopentadienones has been described previously (10).

γ-Aminobutyric acid-functionalized building blocks 7ab, 8ab were obtained by EDC-catalyzed coupling of 6a,b and the corresponding N-protected amino acid derivatives in DMF at room temperature. The products were purified by flash chromatography, and the structures were confirmed by 1H NMR spectroscopy and FD mass spectrometry. Monocarboxy-functionalized cyclopentadienone 9 was synthesized analogously via a carbodiimide induced reaction of 6a with hexanedioic acid monomethyl ester.

3.2. Synthesis of Polyphenylene Dendrimers. For the sake of brevity, the terminology introduced in Figure 1 will be used for the central dendritic polyphenylene units.

Scheme 3 summarizes the synthesis of amino- and maleimide-functionalized dendrimers. We have previously described the preparation of maleimide-decorated polyphenylene dendrimers 13a–d (10). The Diels–Alder reaction of the functionalized cyclopentadienones 5ab with either the core 1 or the octaethynyl-functionalized dendrimer 10 yielded the benzophenoneimino dendrimers 11a–d. Subsequently, the complete cleavage of the protective groups was achieved with HCl in THF at room temperature, and the resulting amino dendrimers 12a–d were reacted without additional purification with 4-(2,5-dioxo-pyrrol-1-yl)-butanoyl chloride to afford polyphenylene dendrimers with different numbers of maleimide anchor groups 13a–d.

The synthesis of first-generation BOC-protected amino-functionalized PDI-dendrimers 14ab proceeded via the Diels–Alder cycloaddition of 7ab with the perylenediimide core 3. The subsequent removal of the BOC groups with trifluoroacetic acid in dichloromethane (20) afforded

Figure 1. Symbols used to represent the polyphenylene scaffold of dendrimers in the schemes hereafter. The abbreviation in the circles shows the dendrimer generation and the central core unit.
the amino compounds 15ab. Employing the reagents 8ab in a Diels–Alder reaction with 16 yielded the PDI dendrimers of the second generation 17ab, which were readily converted to 18ab by treatment with HBr in glacial acetic acid (21). The amino groups of compounds 15ab and 18ab are more stable against oxidation and more accessible for the coupling of polypeptides than amino groups of the aniline derivatives 12a–d.

The synthesis of polyphenylene dendrimers possessing two different chemical functions is depicted in Scheme 4. First, cyclopentadienone 5b was reacted with the diethynyl core 2 to afford compound 19. After deprotection of the ethynyl groups, the subsequent Diels–Alder cycloaddition of 9 produced compound 20. Note that 20 has both (protected) amino and carboxyl anchor groups. The ester groups were converted to free carboxyl groups by LiOH in THF at 80 °C (22), while the deprotection of the amino groups was achieved by HCl in THF. Finally, the maleimide groups were introduced via reaction with 4-(2,5-dioxo-pyrrol-1-yl)-butanoyl chloride in DMF at room temperature to obtain the bifunctional compound 24.

### 3.3. Coupling of Polyllysines to Polyphenylene Dendrimers

Scheme 5 gives an overview of the three synthetic routes toward peptide-decorated PPDs. Here we demonstrate the synthetic versatility of polyphenylene dendrimers as 3D carriers for the preparation of spatially defined bioactive conjugates. Three different strategies for peptide coupling were employed. We emphasize that the developed strategies are versatile and may be used for coupling of various oligopeptide fragments, according to the desired application.

#### 3.3.1. Grafting of Protected Polypeptide from the PPD

The grafting of the polylsine from the surface of the polyphenylene dendrimers 15ab and 18ab was achieved by ring opening polymerization (Scheme 5a) of α-amino acid N-carboxyanhydrides (NCA). The ε-benzyloxy carbonyl-L-lysine N-carboxyanhydride (Lys(Cbz)-NCA) 25 was prepared following a literature procedure (23–25). 15a,b and 18a,b can be regarded as macrorinitiators to which the corresponding amount of Lys(Cbz)-NCA was added. The reaction proceeded in DMF at room temperature for 5 days.

The products 26a–c, 27a–c, 28a–c, 29a–c were precipitated in water. Unreacted monomer was removed.
by extensive washing with Et₂O. Finally, the deprotection of the side chains of the lysine residues yielded 30a-c, 31a-c, 32a-c, 33a-c (Table 1). This reaction scheme permits control of the polypeptide length via the amount of the Lys(Cbz)-NCA added to the reaction. The dendrimers bear different numbers of polypeptide chains of variable length (Table 1). These variable polypeptide lengths were used in the investigation of the secondary structure of the polypeptide shell attached to a polyphenylene core (see section 2.4.). The Cbz-groups of the lysine residues were removed by HBr in glacial acetic acid, followed by precipitation in diethyl ether. To investigate the conjugates in salt-free solutions 30-33a-c were purified via ultrafiltration in milli-Q water.

3.3.2. Attachment of Protected Pentalysine. In this case, the C-terminus of a Cbz-protected pentalysine 34 is coupled to the amino groups on the surface of the PDI dendrimers (Scheme 5b). The peptide was obtained by solid-phase peptide synthesis using standard Fmoc protecting group chemistry (26). The activation of the C-terminal carboxyl group of the peptides was achieved using HOBt, HBTU, and DIPEA as activating agents (27).

The dendrimers 15a,b or 18a,b, peptide 34, HOBT, HBTU, and DIPEA were allowed to react for 5 days in dry DMF under an inert atmosphere at room temperature. After precipitation of the sample in diethyl ether, the crude product was found to contain the target compounds 35a-d and the unreacted polypeptide. Since purification at this stage was not possible, the Z-groups were removed by the same protocol as mentioned in 3.3.1. Subsequent ultrafiltration in water (NMWL 10 000) and freeze-drying yielded the water soluble conjugates 36a-d. The structure and purity of these compounds were verified by means of 1H NMR spectroscopy and gel permeation chromatography. The decoration of PDI dendrimers with preformed polypeptide sequences has the important advantage that no byproducts such as imperfect polypeptide chains occur. Note that the removal of such byproducts in the subsequent purification steps would be rather laborious (28).

3.3.3. Coupling of Unprotected Pentalysine. We also demonstrate a functionalization of the dendrimers with unprotected peptides sequences employing the Michael addition of the sulfhydryl group of terminal cysteine to the maleimide units on the dendrimer surface (Scheme 5c). Hexapeptide 37 is composed of a C-terminal cysteine and five lysine residues. The peptide was prepared via solid-phase peptide synthesis. The reaction of 13a-d with 37 was performed in DMF at room temperature and the products 38a-d were purified by dialysis and high performance liquid chromatography (HPLC). As for the method described in section 3.3.2, the absence of byproducts with wrong peptide sequences is a considerable advantage. Furthermore, in this case one circumvents the deprotection step(s), making this method especially convenient for coupling of polypeptide sequences sensitive to the deprotection procedures. It should be mentioned that the coupling via the SH group of the cysteine residue is not restricted to polypeptides.

3.4. Properties. 3.4.1. Circular Dichroism (CD). The intrinsic properties of proteins are achieved via secondary, tertiary, and quaternary structures. Therefore, a self-organization of the polypeptide shell attached
to a dendrimer core is essential for targeting bioproperties as for example antigen–antibody recognition. This motivated us to study the effect of either solvent or length of the peptides on the secondary structure of attached polylysine moieties.

PPDs bearing polylysine chains longer than six amino acid residues were investigated by CD spectroscopy. It is known that 2,2,2-trifluoroethanol (TFE) (29) as well as high pH (23) induce coil-to-helix transition in polylysine moieties. We employed these protocols to study how the polyphenylene scaffold affects the capability of the attached polypeptides of forming secondary structure. Figure 2 shows the CD spectra of 30c (c = 1.1 μmol/L) as a function of pH (Figure 2a) and concentration of TFE (Figure 2b), respectively. The data are representative of all studied compounds, and the following conclusions can be drawn.

Below pH 9, the polypeptide chains form random coils. Above this pH value, the coils turn into helices. Above pH 10, however, the polypeptides begin to precipitate. Coil-to-helix transformation can be induced by the exchange of the solvent, namely, when the concentration of TFE is above 75% vol.

The coil-to-helix transition with TFE as a solvent is induced because the intramolecular repulsion is suppressed (29–31). The same effect can be obtained by the basicity of the aqueous solution being increased (23). The helix content at pH 9.9 and in 90% TFE was calculated for all polylysine-decorated PPDs according to the method of Chen et al. (32–34). The mean residue ellipticity of a polylysine with 100% helix content at 222 nm was taken to be \(-37 600 \text{ deg cm}^2 \text{ dmol}^{-1}\) (29). The results are summarized in Table 2. For all studied compounds, the helix content at pH 9.9 is about 40% or above. Furthermore, the helix content determined in 90% TFE is higher than that determined at pH 9.9.

The dissociation of polypeptide coiled coils and their organization into single-stranded helices are also evidenced by the ratio of the mean residue ellipticities at 222 and 208 nm, measured at pH 9.9 and in 90% TFE for all studied compounds. As evident from the plot, for polypeptide chains longer than 10 lysine residues, the ratio $[\theta]_{222}/[\theta]_{208}$ remains constant whatever the generation and number of polypeptide chains.

In water the ratio of the mean residue ellipticities at 222 and 208 nm is nearly one and indicates aggregation between the helices, whereas the value of the measurements in 90% TFE is around 0.8 points at dissociation. Shorter chains, comprising less than 10 residues, are less spatially organized and exhibit higher coil content, the latter being evidenced by the minimum in the ellipticity located in the region below 206 nm. An example is presented in Figure 3b where the CD spectra of 30a and 30c are compared.

The main conclusion from the CD data is that polypeptides attached on the dendrimers periphery are capable of forming secondary structures. This finding is essential for the design of synthetic multi-immunogens since the secondary structure of epitopes is responsible for the unique antigen–antibody recognition.

### 3.4.2. UV–Vis Absorption and Fluorescence

The compounds 30a, 31a, 32a, and 33a, each containing a PDI core, were characterized by UV–Vis and fluorescence spectroscopy. As an example, the absorption and emission spectra of 33c, recorded in water and 90% TFE, are shown in Figure 4. Absorption and emission maxima in aqueous solution are at 580 and 614 nm, respectively, while a small bathochromic shift occurs when changing the solvent from water to 90% TFE with maxima at 597 (absorption) and 624 nm (emission).

For all compounds studied, the maxima of absorption are observed at 580–600 nm, being essentially independent of the dendrimer generation and number of peptide residues. Interestingly, the extinction in the less polar solvent water (58 000–88 000 cm$^2$ mol$^{-1}$) is lower than the extinction in TFE solutions (81 000–103 000 cm$^2$ mol$^{-1}$). All of the compounds 31–33 exhibit fluorescence. The emission maxima were detected at wavelengths between 615 and 630 nm upon excitation at 560 nm. The quantum yields in aqueous solutions of all compounds

**Table 2. Helix Content in Percent from the CD Spectra (80 μg/mL) in Water at pH = 9.9 and the Percentage in Comparison with the Measurements in a 90% TFE Solution**

<table>
<thead>
<tr>
<th>sample</th>
<th>$m_{Lys, UV}$</th>
<th>pH</th>
<th>α-helix</th>
<th>$[\theta]<em>{222}^{\text{water}}/[\theta]</em>{222}^{90%\text{TFE}}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>30a</td>
<td>n = 1</td>
<td>9</td>
<td>9.9</td>
<td>43.6</td>
</tr>
<tr>
<td>30b</td>
<td>x = 4</td>
<td>10</td>
<td>9.9</td>
<td>46.4</td>
</tr>
<tr>
<td>30c</td>
<td>n = 1</td>
<td>9</td>
<td>9.9</td>
<td>35.2</td>
</tr>
<tr>
<td>31a</td>
<td>x = 8</td>
<td>76</td>
<td>9.8</td>
<td>47.7</td>
</tr>
<tr>
<td>32a</td>
<td>n = 2</td>
<td>6</td>
<td>9.9</td>
<td>46.4</td>
</tr>
<tr>
<td>32b</td>
<td>x = 8</td>
<td>28</td>
<td>10</td>
<td>44.4</td>
</tr>
<tr>
<td>32c</td>
<td></td>
<td>41</td>
<td>9.9</td>
<td>43.2</td>
</tr>
<tr>
<td>33a</td>
<td>n = 2</td>
<td>14</td>
<td>10</td>
<td>24.8</td>
</tr>
<tr>
<td>33b</td>
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<td>41</td>
<td>9.9</td>
<td>80.6</td>
</tr>
<tr>
<td>33c</td>
<td></td>
<td>89</td>
<td>10.1</td>
<td>43.3</td>
</tr>
</tbody>
</table>

* Notation: $m_{Lys, UV}$, number of lysine residues per chain, calculated by UV absorption of dendrimers.
were low (about 1%). Nevertheless, the optical properties (strong absorption at 580–600 nm and fluorescence in water) of the peptide-decorated PDI dendrimers allow their identification in complex mixtures of biological samples. These features are very important for in vitro/in vivo studies and pharmaceutically oriented applications since at wavelengths greater than 550 nm cells and tissue usually do not show any chromophoric properties.

4. CONCLUSIONS

We have demonstrated the synthesis of new amino acid functionalized cyclopentadienones. Their Diels–Alder reactions with different core molecules lead to polyphenylene dendrimers functionalized with amino or carboxyl groups. Polylsine fragments were then coupled to these dendrimers employing three different methods.

(i) Grafting of the peptides from the surface of the dendrimer by applying the NCA polymerization strategy. This route affords a decoration of PPDs with polypeptides, connected via their C-terminus. Furthermore, the attached polypeptide chains are able to fold into defined secondary structures, an important prerequisite for the preparation of functional bioconjugates.

(ii) Carbodiimide coupling of a protected peptide. The C-terminal carboxyl groups of the peptides are activated employing HBTU/HOBT and are attached to the surface amino groups of the dendrimers.

(iii) Specific attachment of unprotected peptide fragments under mild conditions via Michael addition of a thiol group of the terminal cysteine of the polypeptide to the maleimide of the PPD. This method allows both C- and N-terminal attachment of the polypeptide.

The resulting dendrimers have good solubility in water and combine shape-persistence with a controlled number of surface functions. Additionally, when perylenediimide PPDs are used as cores, absorption and fluorescence techniques can be employed for their identification.

By using an asymmetric core molecule and appropriately functionalized cyclopentadienones in a Diels–Alder reaction, we have obtained PPDs bearing two different functional groups. For example, compound 33, a PPD bearing carboxyl and maleimide functions in isolated surface domains, is a scaffold for the creation of a conjugate possessing two different polypeptide sequences.

We have recorded circular dichroism (CD) spectra of PPDs bearing different numbers of polypeptide chains of various degrees of polymerization. On the basis of these data, we have concluded that at pH 9.9, as well as in the presence of more than 75% TFE, the attached polypeptide is capable of forming an α-helical secondary structure. This fact is of particular importance for the design of MAP since the secondary structure is essential for the intrinsic properties of its distinct sites. In addition, conjugates synthesized with the perylenediimide core strongly absorb at 580–600 nm and are fluorescent in water.

Our synthetic methods are not restricted to oligolysines but may be easily adapted to the substitution of PPD with various polypeptide sequences. This opens the way to biofunctionalized compounds with perspectives as spatially defined MAPs with a hydrophilic shell formed of certain epitopes and chromophor-labeled shape-persistent scaffolds.

ACKNOWLEDGMENT

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LITERATURE CITED


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**Figure 3.** (a) Ratio of the mean residue ellipticities at 222 and 208 nm: $\Theta_{222}/\Theta_{208}$ of different generations: 30 (red down triangles), 31 (black up triangles), 32 (red squares), 33 (blue circles) in 90% TFE (filled symbols) and at pH 9.9 ± 0.1 (empty symbols). (b) CD spectra of 30a and 30c. CD spectra of 30a (circles) and 30c (squares) in 90% TFE (filled symbols) and at pH 9.9 (empty symbols).

**Figure 4.** UV–vis absorbance and fluorescence emission spectra of 33c in 90% TFE and in aqueous solution pH 9.9.


