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PAPER

Efficient and versatile COMU-mediated solid-phase submonomer synthesis of arylopeptoids (oligomeric N-substituted aminomethyl benzamides)[†]

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The development of a highly efficient methodology for solid-phase synthesis of *para-* and *meta-*arylopeptoids (oligomeric *N*-substituted aminomethyl benzamides) with free acids or free amides at the *C*-terminus is described. The arylopeptoids were synthesised by means of a convenient submonomer protocol in which the arylopeptoid residues were created in an iterative manner on the growing chain using an acylation–substitution cycle. The uronium salt COMU was found to be the most efficient reagent for ensuring fast and clean couplings of the benzoic acid building blocks.

Introduction

A group of foldamers which has received ever increasing interest during the last two decades is "one-way sequence" aromatic oligoamides comprised of aromatic amino acids.¹ While the amide core structures of these synthetic oligomers are inherently only distantly related to biopolymers such as peptides, they may still be brought to mimic the side chain presentation and structure of such biopolymers and may therefore for example find use as proteomimetics.² These aromatic oligoamides may be divided into two distinct groups: (i) aromatic oligoamides with free amide protons (Fig. 1a) and (ii) *N*-alkylated aromatic oligoamides (Fig. 1b).

In the first group, oligoamides that contain no aliphatic carbons in the backbone are by far the most widely studied (Fig. 1a). Examples include benzanilides by the groups of Hamilton,³ Gong,⁴ Kilbinger,⁵ Li,⁶ Wilson,⁷ Clayden,⁸ and others;⁹ pyridylamides by the Hamilton group;^{3a-b,10} and quinoline and naphthyridine oligoamides by the Huc group.^{11,12} The syntheses of these oligomers pass *via* acylation of comparatively less reactive aromatic amines and coupling using acid chlorides is the most commonly used synthetic pathway, primarily in solution,^{3,4,5c,6-8,9a-c,e,11,10} but also on solid-phase.^{5a-b,9d} Accordingly, the only reports of synthesis using free acid building blocks

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Fig. 1 Aromatic oligoamides with free amide protons (a) and *N*-alkylated aromatic oligoamides (b).

and peptide-type coupling methods include the use of BOP^{9b} or PyBOP¹² in solution-phase. Although less well studied, several reports on aromatic oligoamides with free amide protons and aliphatic carbons in the backbone have been disclosed: oligoamides of aminomethyl-phenyl acetic acids by the Amblard group,¹³ oligoamides of aminophenoxy acetic acids by the Akazome group,¹⁴ aminomethyl-pyridylamides by the Huc group,¹⁵ and aminomethyl-benzamides by the Hamilton group.¹⁶ Since the syntheses of these oligomers proceed *via* more reactive aliphatic intermediates, activation as acid chlorides is not necessary and they have exclusively been synthesised using peptide-type coupling

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reagents such as BOP,¹³ EDC,¹⁴ HBTU,¹⁵ in solution-phase and PyBOP¹⁶ on solid-phase. A common trait in this first group of aromatic oligoamides is that the formation of secondary structures may be promoted by the presence of intramolecular hydrogen bonding involving the free amide protons analogous to what is observed in peptides. Accordingly, the presence of spherand or crown ether-like structures,⁶ crescent or helical structures,^{4,11-14} and rod-like structures,^{5a,c,7a} as well as the use as α -helix mimetics,^{3a-b,7a,9a-b,10} has been reported.

Compared to aromatic oligoamides with free amide protons, Nalkylated aromatic oligoamides remain far less studied (Fig. 1b). Most of the efforts concerning N-alkylated oligoamides have again been concentrated around the synthesis and study of oligomers with no aliphatic carbons in the backbone, including N-alkylated benzanilides by the groups of Yokozawa,¹⁷ Kilbinger,¹⁸ Wilson,¹⁹ and Clayden;8a and N-alkylated naphthanilides by the Clayden group.^{8a} These oligomers were exclusively synthesised using activation as acid chlorides either in solution-phase^{8a,17,19b-c} or on solid-phase.^{18a-b,19a} Intriguingly, although they are deprived of free amide protons, the presence of crescent or helical structures¹⁷ as well as the use as α -helix mimetics^{19a} has still been reported. Furthermore, in connection with their pioneering work on the development of peptoids (oligomeric N-substituted glycines),²⁰ Zuckermann and co-workers²¹ briefly mentioned the synthesis of a few pentamers of para-arylopeptoids²² (oligomeric N-substituted aminomethyl benzamides) in a series of patents from the mid to late 90's. The oligomers were synthesised on a Rink amide polystyrene resin using a submonomer method unique to peptoids and the benzoic acid building blocks were activated as their corresponding anhydrides using 0.5 equiv DIC. In 2007, Lokey and Combs then reported the synthesis of a few tetramers and pentamers of paraor meta-arylopeptoids using a Wang resin and activation of the benzoic acid building blocks with 1.0 equiv DIC.23 This method was, however, not adaptable to synthesis on a Rink amide resin and provided mediocre yields.

We believe that these arylopeptoids may represent a new group of oligomers with folding propensities as a result of their relatively rigid aromatic amide core structure even though their backbones are deprived of free amide protons. Importantly, they may be developed to retain the favorable characteristics of peptoids such as large potential for diversity and straightforward synthesis using submonomer methods. Thus, based on convenient methods we have previously developed for solutionphase synthesis of peptoid residues,²⁴ we recently published the first solution-phase synthesis of para- and meta-arylopeptoids.²⁵ Shorter oligomers were synthesised using highly convenient and efficient submonomer protocols where each arylopeptoid residue is created in an iterative manner on the growing chain using an acylation-substitution cycle. Peptide coupling methods were studied to access longer oligomers and we furthermore undertook the first conformational studies of arylopeptoids. Thus, the *cis/trans* isomerism of arylopeptoid backbones was studied by means of NMR and we found that increasing the bulkiness of the side chains would greatly favor the cis amide bond conformation. Although this solution-phase methodology is highly useful for gram scale synthesis of lead compounds, the development of efficient and generally applicable solid-phase methodologies for synthesis of large and diversified libraries of arylopeptoids is highly desirable. Thus, herein we describe the development of a

versatile COMU-based solid-phase methodology that meets these demands.

Results and discussion

Solution-phase studies

In order to establish the initial conditions for optimisation of solidphase submonomer synthesis of arylopeptoids, we first screened a series of coupling and substitution conditions on a small scale in solution (see ESI† for details). As a model system for conducting these studies we used conversion of *para*-monomer p-1,²⁵ carrying the moderately bulky isopropyl side chain, into dimer $p-4^{25}$ via bromomethyl intermediate p-2 or chloromethyl intermediate p-3(Scheme 1).



Scheme 1 Model system for solution-phase studies. Key: (a) XCH₂ArCOOH, coupling reagent, DIPEA, solvent, rt, further conditions in Table 1. (b) *i*PrNH₂, solvent, further conditions in Table 2.

In our preceding paper, we screened a selection of coupling reagents for peptide-type coupling of arylopeptoid trimers.²⁵ We found the new uronium salt COMU²⁶ (derived from Oxyma) to be by far the most efficient coupling reagent and we therefore started the present study with the use of COMU. Thus, 1 M solutions of 4-(bromomethyl)benzoic acid or 4-(chloromethyl)benzoic acid (3 equiv) in DMF were preactivated with COMU (3.5 equiv) and DIPEA (7 equiv) at room temperature, and were then reacted with monomer *p*-1. The reactions were monitored by analytical HPLC (see ESI[†] for details) and while the conversion of *p*-1 into bromomethyl intermediate p-2 gave rise to a complex reaction mixture (Table 1, entry 1) the conversion of *p*-1 into chloromethyl intermediate p-3 proceeded cleanly in 2.5 min (Table 1, entry 2). Indeed, we found that while 4-(chloromethyl)benzoic acid remains reasonably pure in DMF in the presence of DIPEA (99.6% purity after 5 min; 98.5% purity after 15 min), 4-(bromomethyl)benzoic acid polymerises quickly under the same conditions (<50% purity after 5 min). These polymerisation reactions presumably occur too fast for COMU activation to compete properly and a complex product mixture is obtained. The use of 4-(bromomethyl)benzoic acid as a building block for solid-phase submonomer synthesis of arylopeptoids under basic conditions was therefore abandoned.

Entry	X, XCH ₂ C ₆ H ₄ COOH (equiv)	Coupling reagent (equiv)	DIPEA (equiv)	Solvent	Pre-activation (min)	Reaction time ^{<i>a</i>} (min)	Comments
1	Br (3.0)	COMU (3.5)	7.0	DMF	5	<2.5	Complex mixture
2	Cl (3.0)	COMU (3.5)	7.0	DMF	5	2.5	Clean reaction to p-3
3	Cl (3.0)	COMU (3.5)	7.0	NMP	5	5	Clean reaction to p-3
4	Cl (3.0)	HOTU (3.5)	7.0	NMP	5	10	Clean reaction to p-3
5	Cl (3.0)	HATU (3.5)	7.0	NMP	5	20	Clean reaction to p-3
6	Cl(3.0)	PyBroP (3.5)	7.0	NMP	5	30	Clean reaction to p-3
7	Cl(3.0)	BOP (3.5)	7.0	NMP	5	>30	93% conv. after 30 min
8	Cl(3.0)	HBTU (3.5)	7.0	NMP	5	>30	90% conv. after 30 min
9	Cl(3.0)	PyBOP(3.5)	7.0	NMP	5	>30	85% conv. after 30 min
10	Cl (6.0)	DIC (3.0)	7.0	NMP	5	>30	93% conv. after 30 min
11	Cl (3.0)	DIC (3.0)	0	NMP	5	>30	83% conv. after 30 min
^a Reactio	ons monitored by analytical	HPLC.					

 Table 1
 Optimisation of coupling step (a)

Next, we found that although the conversion of p-1 into p-3would proceed slightly slower when using the less toxic NMP as solvent (5 min, entry 3), the reaction would proceed just as cleanly as in DMF. A representative selection of other coupling agents were then screened (entries 4-11). Although most of these couplings proceeded with the same purity as when COMU was used, none of the alternative coupling reagents could compete with COMU with respect to reaction times. Thus, compared to COMU, HOTU would double the reaction time (10 min, entry 4), while couplings using HATU and PyBroP would need 20 min and 30 min to reach completion, respectively (entries 5 and 6). Furthermore, couplings using BOP, HBTU and PyBOP did not reach complete conversion within 30 min (entries 7-9). Interestingly, both activation of 4-(chloromethyl)benzoic acid as the corresponding anhydride using 0.5 equiv DIC, as reported in the patents of Zuckermann and co-workers,²¹ and coupling using 1.0 equiv DIC as reported by Lokey,²³ were found to be much less efficient than COMU couplings (93% and 83% conversion after 30 min, respectively, entries 10 and 11). Having established the combination of 4-(chloromethyl)benzoic acid and COMU as the optimal set of conditions for peptide type coupling synthesis of *p*-3, we then proceeded to investigate the ensuing substitution reaction with isopropylamine, yielding dimer p-4 (Table 2).

 Table 2
 Optimisation of substitution step (b)

Although we had by now excluded the use of bromomethyl intermediate p-2, an initial solvent scan was performed using this more reactive intermediate. Thus, p-2 was reacted with 10 equiv isopropylamine as 1 M solutions in different solvents at room temperature (Table 2, entries 1-6) and the evolution of the reactions was followed using analytical HPLC (see ESI[†] for details). We found the substitution reaction to occur the fastest in DMSO (<2 min, entry 6) followed by DMF and NMP (3 min, entries 4-5). When transferring these conditions to the corresponding chloromethyl intermediate p-3, the reaction times were as expected drastically increased (entries 7-9) with DMSO still offering the fastest reaction time (3 h, entry 9). By increasing the concentration of isopropylamine and the temperature, the reaction time could be reduced (entries 10-12) and the reaction would be complete within a reasonable 30 min when using 20 equiv isopropylamine at 2 M concentration in DMSO at 50 °C (entry 12). The reaction could also be performed in DMSO-water 9:1 although the reaction time increased slightly to 45 min. This result indicated that concentrated aqueous solutions could be used as a source of small volatile amines if needed. We also tested the use of microwave irradiation (95 °C) and found that the reaction times could be drastically reduced to 5 min or less (entries 14-16). Overall, these solutionphase investigations showed that the optimisation of solid-phase

Entry	Х	<i>i</i> PrNH ₂ (equiv/[c])	Solvent	Conditions	Reaction time ^a
1	Br	10.0/1.0	MeOH	rt	4 h
2	Br	10.0/1.0	THF	rt	1.5 h
3	Br	10.0/1.0	CH ₂ Cl ₂	rt	1.5 h
4	Br	10.0/1.0	NMP	rt	3 min
5	Br	10.0/1.0	DMF	rt	3 min
6	Br	10.0/1.0	DMSO	rt	<2 min
7	Cl	10.0/1.0	NMP	rt	16 h
8	Cl	10.0/1.0	DMF	rt	12 h
9	Cl	10.0/1.0	DMSO	rt	3 h
10	Cl	20.0/2.0	DMSO	rt	2 h
11	Cl	20.0/2.0	DMSO	40 °C	45 min
12	Cl	20.0/2.0	DMSO	50 °C	30 min
13	Cl	20.0/2.0	DMSO-water 9:1	50 °C	45 min
14	Cl	10.0/1.0	DMSO	MW, 95 °C	5 min
15	Cl	20.0/2.0	DMSO	MW, 95 °C	<2.5 min
16	Cl	20.0/2.0	DMSO-water 9:1	MW, 95 °C	<2.5 min

" Reactions monitored by analytical HPLC.



Scheme 2 Model system for optimisation of solid-phase synthesis. Key: (a) ClCH₂ArCOOH (1.2 equiv, 0.14 M), DIPEA (6.0 equiv), CH₂Cl₂, rt, 1 h. (b) *i*PrNH₂ (20 equiv, 2.0 M), DMSO, further conditions in Table 3. (c) ClCH₂ArCOOH, COMU, DIPEA, further conditions in Table 3. (d) BzCl (4.0 equiv, 1.0 M), DIPEA (8.0 equiv), CH₂Cl₂, rt, 15 min. (e) HFIP-CH₂Cl₂ 1 : 4, rt, 1 h.

submonomer synthesis of arylopeptoids should be based on the use of COMU-mediated coupling of chloromethylbenzoic acid building blocks and substitution reactions with primary amines in DMSO either at 50 °C or by the use of microwave activation.

Solid-phase studies

Optimisation of solid-phase submonomer synthesis of arylopeptoids was performed on a 2-chlorotrityl chloride polystyrene resin with a listed loading of 1.50 mmol g⁻¹. As a direct continuation of the solution-phase studies we chose to use para-trimer p-5 with isopropyl side chains as a model target (Scheme 2). We found that a constant loading of 1.23 ± 0.03 mmol g⁻¹ was obtained when the 2-chlorotrityl chloride polystyrene resin was reacted with 4-(chloromethyl)benzoic acid (1.2 equiv with respect to the listed loading) in the presence of DIPEA (6.0 equiv) in CH₂Cl₂ at rt for 1 h (see ESI[†] for details). A slightly lower loading of 1.10 mmol g⁻¹ was obtained when shortening the reaction time to 30 min, while longer reaction times were avoided in order to minimise the risk of polymerisation of 4-(chloromethyl)benzoic acid. The synthesis of trimer *p*-5 was then carried out by following the iterative coupling-substitution cycle illustrated in Scheme 2 followed by capping with benzoyl chloride and cleavage with HFIP. Based on the solution-phase studies, the reaction times

of the substitution steps and coupling steps were set to 1 h and 20 min, respectively, and we were satisfied to find that trimer p-5 could be isolated in 71–74% crude yield and an excellent 96% crude purity when otherwise adapting the conditions found in the previous solution-phase studies (Table 3, entry 1). We furthermore obtained identical crude yields and purities when conducting the coupling steps at higher dilution (0.5 M) which would prove to be useful when synthesising longer oligomers (vide infra). Increasing the amount of COMU-activated 4-(chloromethyl)benzoic acid added in the coupling steps to 5 equiv did not improve the crude yield or purity of the synthesis (entry 2). The substitution steps could also be performed using microwave irradiation at 95 °C which would furnish trimer p-5 in comparable crude yields and purities (entry 3). However, although the reaction time of the substitution steps could be reduced to as little as 2.5 min, we found heating to 50 °C using conventional heating blocks to be more convenient since the entire arylopeptoid synthesis could be completed without changing the reaction syringe. Otherwise, specialised equipment would be required. Attempts to change the initially selected reaction times did not improve the synthesis either. Thus, reducing the reaction time of the substitution steps to 15 min (entry 4) or increasing the reaction time of the coupling steps to 1 h (entry 5) would in both cases decrease the crude purity slightly to 93%. Furthermore, reducing the reaction time of the

 Table 3
 Optimisation of substitution step (b) and coupling step (c)

Entry	Substitution step (b)	Coupling step (c)					
	(temp/time)	ClCH ₂ ArCOOH (equiv/[c])	COMU (equiv)	DIPEA (equiv)	Time (min)	Crude yield (%)	Crude purity (%) ^a
1	50 °C/1 h	3.0/0.5-1.0	3.5	7	20	71–74	96
2	50 °C/1 h	5.0/1.0	6	12	20	76	96
3	MW 95 °C/2.5–10 min	5.0/1.0	6	12	20	73-75	95
4	50 °C/15 min	3.0/1.0	3.5	7	20	74	93
5	50 °C/1 h	3.0/1.0	3.5	7	60	77	93
6	50 °C/1 h	3.0/1.0	3.5	7	5	69	82

" Determined by analytical HPLC.



Scheme 3 Solid-phase submonomer synthesis of model arylopeptoid hexamers with free acids at the *C*-terminus. Key: (a) ClCH₂ArCOOH (1.2 equiv, 0.14 M), DIPEA (6.0 equiv), CH₂Cl₂, rt, 1 h. (b) R-NH₂ (20 equiv, 2.0 M), DMSO, 50 °C, 1 h. (c) ClCH₂ArCOOH (3.0 equiv, 1.0 M), COMU (3.5 equiv), DIPEA (7.0 equiv), rt, 20 min. (d) BzCl (4.0 equiv, 1.0 M), DIPEA (8.0 equiv), CH₂Cl₂, rt, 15 min. (e) HFIP–CH₂Cl₂ 1:4, rt, 1 h. See Table 4 for yields and purities.

coupling step to 5 min would decrease the crude yield to 69% and at the same time decrease the crude purity to 82% (entry 6). Overall, the conditions listed in entry 1 were chosen as the optimal conditions.

Solid-phase synthesis of arylopeptoids

With an optimised method for solid-phase submonomer synthesis of arylopeptoids in hand, we then first synthesised a range of representative model para-hexamers p-6 and meta-hexamers m-6 with free acids at the C-terminus (Scheme 3 and Table 4). The syntheses were performed on the 2-chlorotrityl chloride polystyrene resin used in the above studies using the optimised conditions. We were satisfied to find that hexamers p/m-6a with isopropyl side chains, p/m-6c with 4-phenylbutyl side chains and p/m-6d with 2-morpholinoethyl side chains were all isolated in good crude purities (78-85%, entries 1-2 and 5-8). Whereas hexamer p-6a unfortunately proved too insoluble to be purified, the remaining hexamers could easily be obtained in 39-58% yield and >99% purity after preparative HPLC. For comparison, Lokey and Combs reported the synthesis of similar N-acetyl capped para- and meta-arylopeptoid tetramers with 2-morpholinoethyl side chains in 20% and 37% yield, respectively, and the synthesis of similar N-acetyl capped para- and meta-arylopeptoid pentamers with 4-phenylbutyl side chains in 12% yield.²³ Our method thus provides longer oligomers in greatly increased yields and is furthermore more efficient when installing more bulky side chains such as the isopropyl side chain. We used a concentrated aqueous solution as the source of ethylamine for the substitution steps in the synthesis of hexamers p/m-6b which carry ethyl side chains (entries 3 and 4). Although we had shown in the solution-phase studies that the substitution step would proceed more slowly in the presence of water, only a slight decrease in crude purity to 71% was observed. Hexamers *p***-6b** and *m***-6b** were isolated in comparable crude yields which were slightly lower than the above-mentioned hexamers. Hexamer p-6b unfortunately also proved too insoluble to allow for purification, but *m*-6b was accordingly isolated in a slightly lower yield than the above-mentioned hexamers after purification (33%, >99% purity, entry 4). This slight decrease in yield is presumably a result of slow ester hydrolysis of the arylopeptoid chains from the 2-chlorotrityl linker during the substitution reactions. Hexamer p-6e with pyridinylmethyl side chains was also synthesised efficiently although the crude purity was reduced to 52% and the yield of the purified product was only 21% (>99% purity, entry 9). Synthesis of hexamers carrying highly bulky tert-butyl side chains was likewise attempted but failed to yield anything but complex product mixtures.

 Table 4
 Results for solid-phase submonomer synthesis of model arylopeptoid hexamers with free acids at the C-terminus

Entry	Hexamer	Crude yield (%)	Crude purity (%) ^a	Purified yield (%) ^b
1	<i>p</i> -6a	69	78	c
2	<i>m</i> -6a	91	80	55
3	<i>p</i> -6b	57	71	c
4	<i>m</i> -6b	54	71	33
5	<i>p</i> -6c	66	80	39
6	<i>m</i> -6c	68	81	39
7	<i>p</i> -6d	86	85	58
8	<i>m</i> -6d	87	81	41
9	<i>p</i> -6e	72	52	21

^{*a*} Determined by analytical HPLC. ^{*b*} Yield of product isolated in >99% purity after preparative HPLC. ^{*c*} Preparative HPLC hindered due to very low solubility.



Scheme 4 Solid-phase submonomer synthesis of model arylopeptoid hexamers with free amide at the *C*-terminus. Key: (a) piperidine–NMP 1:4, rt, 2 and 15 min; then ClCH₂ArCOOH (3.0 equiv, 1.0 M), COMU (3.5 equiv), DIPEA (7.0 equiv), rt, 30 min. (b) R-NH₂ (20 equiv, 2.0 M), DMSO, 50 °C, 1 h. (c) ClCH₂ArCOOH (3.0 equiv, 1.0 M), COMU (3.5 equiv), DIPEA (7.0 equiv), rt, 20 min. (d) BzCl (4.0 equiv, 1.0 M), DIPEA (8.0 equiv), CH₂Cl₂, rt, 15 min. (e) TFA–water 95: 5, rt, 2 h. See Table 5 for yields and purities.

An important point for us was to develop the first efficient and versatile solid-phase methodology for submonomer synthesis of arylopeptoids that could generate both oligomers with free acids at the C-terminus and oligomers with free amides at the Cterminus. Indeed, Lokey and Combs reported that their method was not adaptable to the use of Rink amide resins,23 while synthesis of arylopeptoids with free acids at the C-terminus was not demonstrated by Zuckermann and coworkers.²¹ We were therefore satisfied to find that our methodology was also adaptable to synthesis of oligomers with free amides at the Cterminus using a Rink amide polystyrene resin with a listed loading of 0.74 mmol g⁻¹ (Scheme 4 and Table 5). Compared to synthesis using the 2-chlorotrityl chloride polystyrene resin, the only changes to the syntheses were (1) the initial attachment reactions were carried out for 30 min using 3 equiv of COMUactivated chloromethylbenzoic acids with respect to the listed resin loading and (2) the final cleavage was effected using TFA-water 95:5 instead of HFIP. Hexamers p/m-7a with isopropyl side chains, p/m-7c with 4-phenylbutyl side chains and p/m-7d with 2-morpholinoethyl side chains were all isolated in good crude purities (71-84%, entries 1-2 and 4-7) and good purified yields after preparative HPLC (46–59%, >99% purity). These results are comparable to the results obtained above in the synthesis of the

analogous arylopeptoids with free acids at the C-terminus using the 2-chlorotrityl linker. The two linkers thus perform equally well with these side chains when applying the standard conditions we have developed. Furthermore, when using the Rink amide linker, we did not observe a relative decrease in the yield of p-7b which carries ethyl side chains even though a concentrated aqueous solution was again used as the source of ethylamine (54% vield, >99% purity, entry 3). This supports the assumption that the relative reduction in yield of p/m-6b is caused by slow hydrolysis of the arylopeptoid chains from the 2-chlorotrityl linker during the substitution steps. Our results compare favourably with the method reported in the patents of Zuckermann and coworkers where similar, though N-uncapped, para-arylopeptoid pentamers were obtained in 60-76% crude yield and 80-90% crude purity.²¹ Installing pyridinylmethyl side chains again resulted in a reduced crude purity of 51% and we accordingly obtained p-7e in only 22% yield (>99% purity, entry 8). We furthermore synthesised hexamers p-7f and p-7g which carry 4-aminobutyl side chains and carboxymethyl side chains, respectively (entries 9-10). These side chains were installed using tert-butyl (4-aminobutyl)carbamate and tert-butyl aminoacetate in the substitution steps, respectively, conveniently unmasking the side chain functionalities during the cleavage step.

 Table 5
 Results for solid-phase submonomer synthesis of model arylopeptoid hexamers with free amides at the C-terminus

Entry	Hexamer	Crude yield (%)	Crude purity (%) ^{<i>a</i>}	Purified yield (%) ^b
1	<i>p</i> -7a	127	80	55
2	<i>m</i> -7a	110	71	58
3	<i>p</i> -7b	120	82	54
4	<i>p</i> -7c	110	78	46
5	<i>m</i> -7c	104	84	53
6	<i>p</i> -7d	93	79	58
7	<i>m</i> -7d	115	81	59
8	<i>p</i> -7e	118	51	22
9	<i>p</i> -7f	77	77	49
10	<i>p</i> -7g	109	52	31

" Determined by analytical HPLC. " Yield of product isolated in >99% purity after preparative HPLC.

In order to firmly underline the utility of the present method we then easily synthesised the longest arylopeptoid constructs demonstrated to date: the two hetero-dodecamers p-8 and m-8 shown in Scheme 5, which were obtained in 25% and 27% overall yield, respectively, after preparative HPLC (>99% purity). The syntheses were carried out as outlined in Scheme 4 with the only modifications being that coupling steps beyond the hexamer stage as well as the final capping step were carried out at higher dilution (0.5 M) as a result of the expanding of the resin. The dodecamers were obtained in highly satisfactory 62% and 58% crude purities, respectively.



m-8: 27%, >99% purity (58% crude purity)

Scheme 5 Model hetero-dodecamers *p*-8 and *m*-8.

Importantly, we observed that the solubility of the synthesised arylopeptoid oligomers in different solvents would vary greatly as a function of the side chain decorations, the substitution patterns of the backbone and the nature of the C-terminus. Thus, arylopeptoids that carried hydrophobic side chains such as ethyl, isopropyl or 4-phenylbutyl were only soluble in organic solvents such as methanol or CH_2Cl_2 and the presence of only small amounts of water would be tolerated before cloudy mixtures formed. On the contrary, arylopeptoids that carried hydrophilic side chains such as pyridinylmethyl, 2-morpholinoethyl or 4aminobutyl were generally highly water soluble. Furthermore, we noted that *meta*-arylopeptoids were generally more readily soluble than *para*-arylopeptoids and that arylopeptoids with free amides at the C-terminus were likewise more readily soluble than arylopeptoids with free acids at the C-terminus. Overall, these observations lend promise that arylopeptoids may easily be adapted for a plethora of uses in chemical biology and medicine.

Conclusions

We have developed a highly efficient methodology for solidphase synthesis of *para-* and *meta-*arylopeptoids (oligomeric *N*-substituted aminomethyl benzamides). The oligomers were synthesised by means of a highly versatile submonomer protocol in which the arylopeptoid residues are created in an iterative manner on the growing chain using an acylation–substitution cycle. The difficult acylation steps of the secondary amines were conveniently carried out using free acid building blocks reminiscent of peptide synthesis, and we found the new uronium salt COMU to be the most efficient reagent for ensuring fast and clean couplings. The present methodology is the first that has been shown to be adaptable to synthesis of arylopeptoids with both free acids and free amides at the *C*-terminus. We found the solubility of arylopeptoids in different solvents to vary greatly as a function of the side chain decorations, the substitution patterns of the backbone and the nature of the *C*-terminus, lending promise that arylopeptoids may easily be adapted for a plethora of uses within for example chemical biology and medicine. The developed solidphase methodology will allow for synthesis of highly diversified libraries of arylopeptoids and we will continue to explore all aspects of arylopeptoids and will report our findings in due course.

Experimental Section

General experimental methods

CH₂Cl₂ used as solvent in attachment and capping reactions was dried over 4 Å molecular sieves. All other chemicals and solvents obtained from commercial sources (Alfa Aesar, Fluka, Merck and Sigma-Aldrich) were used as received. 2-Chlorotrityl chloride copoly(styrene-1% DVB) resin (100-200 mesh) with a listed loading of 1.50 mmol g⁻¹ and Rink amide copoly(styrene-1% DVB) resin (100–200 mesh) with a listed loading of 0.74 mmol g⁻¹ were purchased from Merck. 2 M solutions of primary amines in DMSO used in the substitution steps were made from the neat, free amines except in the case of ethylamine where a saturated aqueous solution was used as a source of the amine. Melting points were determined on a Mettler Toledo MP70 melting point system and are referenced to the melting points of benzophenone and benzoic acid. NMR spectra were recorded on a Bruker Avance 300 MHz spectrometer. Chemical shifts are referenced to the residual solvent peak and J values are given in Hz. The following multiplicity abbreviations are used: (s) singlet, (q) quartet, (m) multiplet, and (br) broad. Where applicable, assignments were based on COSY, HMBC, HSQC and J-mod-experiments. IR spectra were recorded on a Shimadzu FTIR-8400S spectrometer equipped with a Pike Technologies MIRacle[™] ATR and wavenumbers (v) are expressed in cm⁻¹. HRMS of all arylopeptoids except for *p*-7f and *p*-7g were recorded on a Micromass LCT apparatus equipped with an AP-ESI probe calibrated with Leu-Enkephalin. HRMS of p-7f and *p*-7g were recorded on a Micromass Q-Tof Micro (3000V) apparatus. Model system monomer p-1 and dimer p-4 were synthesised as previously described.²⁵ See ESI† for details concerning analytical and preparative HPLC as well as for the synthesis of halomethyl intermediates *p*-2 and *p*-3.

General procedure for attachment to trityl resin and synthesis of first arylopeptoid residue

2-Chlorotrityl chloride resin (100 mg, 1.50 mmol g⁻¹; thus 0.150 mmol) was washed with CH₂Cl₂ (2 × 2 mL). The resin was swelled in CH₂Cl₂ (2 mL) for 5 min and drained. To a suspension of 3- or 4-(chloromethyl)benzoic acid (30.7 mg, 0.18 mmol) in CH₂Cl₂ (1.3 mL) at rt was added DIPEA (0.126 mL, 0.72 mmol). The resulting solution was added to the resin and the resulting mixture was agitated for 1 h at rt after which time a loading of 1.23 ± 0.03 mmol was obtained (see ESI[†]). The resin was drained

Downloaded by John Rylands University Library on 25 September 2011 Published on 12 August 2011 on http://pubs.rsc.org | doi:10.1039/C1OB05729D and washed with CH_2Cl_2 (3 × 2 mL) and DMSO (3 × 2 mL). A solution of the secondary amine (20 equiv, 2 M) in DMSO (1.3 mL) was added to the resin and the mixture was agitated at 50 °C for 1 h. The resin was drained and washed with DMSO (3 × 2 mL) and NMP (3 × 2 mL).

General procedure for attachment to rink amide resin and synthesis of first arylopeptoid residue

Rink amide resin (162 mg, 0.74 mmol g⁻¹; thus 0.120 mmol) was washed with CH_2Cl_2 (2 × 2 mL) and NMP (5 × 2 mL). Piperidine– NMP 1 : 4 (1.0 mL) was added and the resin was agitated for 2 min and drained. Further piperidine–NMP 1 : 4 (1.0 mL) was added and the resin was agitated for 15 min, drained and washed with NMP (5 × 2 mL). To a solution of 3- or 4-(chloromethyl)benzoic acid (64 mg, 0.38 mmol) in NMP (0.38 mL) at rt was added COMU (187 mg, 0.44 mmol) and then DIPEA (0.157 mL, 0.90 mmol). After agitating for 5 min, the resulting deep red mixture was added to the resin was drained and washed with NMP (5 × 2 mL) and DMSO (3 × 2 mL). A solution of the secondary amine (20 equiv, 2 M) in DMSO (1.3 mL) was added to the resin and the mixture was agitated at 50 °C for 1 h. The resin was drained and washed with DMSO (3 × 2 mL) and NMP (3 × 2 mL).

General procedure for arylopeptoid residue synthesis

To a solution of 3- or 4-(chloromethyl)benzoic acid (64 mg, 0.38 mmol) in NMP (0.38 mL) at rt was added COMU (187 mg, 0.44 mmol) and then DIPEA (0.157 mL, 0.90 mmol). After agitating for 5 min, the resulting deep red mixture was added to the resin and the resulting mixture was agitated for 20 min at rt. The resin was drained and washed with NMP (5×2 mL) and DMSO (3×2 mL). A solution of the secondary amine (20 equiv, 2 M) in DMSO (1.3 mL) was added to the resin and the mixture was agitated at 50 °C for 1 h. The resin was drained and washed with DMSO (3×2 mL) and NMP (3×2 mL). This general procedure was repeated until the desired oligomer length was reached.

General procedure for capping with benzoyl chloride

The resin was washed with CH_2Cl_2 (3 × 2 mL). To a solution of benzoyl chloride (0.060 mL, 0.52 mmol) in CH_2Cl_2 (0.52 mL) at rt was added DIPEA (0.181 mL, 1.04 mmol). The solution was added to the resin and the resulting mixture was agitated for 15 min at rt. The resin was drained and washed with CH_2Cl_2 – DIPEA 4:1 (3 × 1 mL, only for arylopeptoids that contain side chains with basic functionalities), CH_2Cl_2 (3 × 2 mL), NMP (3 × 2 mL) and CH_2Cl_2 (3 × 2 mL).

General procedure for cleavage from trityl resin

The resin was cleaved in HFIP/CH₂Cl₂ 1:4 (1 mL) with agitation for 1 h. The resin was drained and washed with CH₂Cl₂ (5 × 2 mL). The solvents were evaporated under reduced pressure and the residue was evaporated with CH₂Cl₂ (3 × 5 mL) yielding the crude product which was purified by preparative HPLC (see ESI[†] for details).

General procedure for cleavage from rink amide resin

The resin was cleaved in TFA–water 95:5 (2 mL) with agitation for 2 h. The resin was drained and washed with TFA (2 mL) and CH_2Cl_2 (5 × 2 mL). The solvents were evaporated under reduced pressure and the residue was evaporated with CH_2Cl_2 (3 × 5 mL) yielding the crude product which was purified by preparative HPLC (see ESI† for details).

Arylopeptoid trimer *p*-5. Colorless foam (52 mg, 65%, >99% purity). mp = 87–90 °C. ¹H NMR (300 MHz, CDCl₃): δ = 9.85–9.30 (br s, 1H, COO*H*), 8.12–7.92 (m, 2H), 7.62–6.98 (m, 15H), 4.88–4.44 (br s, 6H, 3×CONC*H*₂Ar), 4.38–4.03 (br m, 3H, 3×CONC*H*(CH₃)₂), 1.42–0.94 (m, 18H) ppm. ¹³C NMR (75 MHz, CDCl₃): δ = 172.7, 172.6, 170.5 (4C_q), 145.0, 140.6, 136.4, 134.9 (7C_q), 130.3, 129.5, 128.6, 127.0, 126.8, 126.5, 126.1 (17CH), 51.0 (br, 3CH, 3×CONCH(CH₃)₂), 43.5 (br, 3CH₂, 3×CONCH₂Ar), 21.3 (6CH₃) ppm. *v*_{max}/cm⁻¹ (ATR) 2980 (CH), 1713 (C=O), 1627 (C=O), 1464 (CH), 1439, 1409, 1339 (OH), 1174 (CN), 1061. HRMS (TOF MS ES⁺) calcd for C₄₀H₄₆N₃O₅ [M + H]⁺ *m*/*z* 648.3432, found 648.3425.

Arylopeptoid hexamer *p*-6a. Preparative HPLC of the crude product hindered due to very low solubility. Crude *p*-6a: Pale rose foam (97 mg, 69%, 78% purity). mp = 223–228 °C. ¹H NMR (300 MHz, CDCl₃–MeOD 1:1): δ = 8.00–7.92 (m, 2H), 7.54–7.10 (m, 27H), 4.78–4.40 (br s, 12H, 6×CONC*H*₂Ar), 4.24–4.04 (br m, 6H, 6×CONC*H*(CH₃)₂), 1.32–1.00 (36H, m) ppm. *v*_{max}/cm⁻¹ (ATR) 2968, 2929, 1628 (C=O), 1437, 1404, 1338 (OH), 1175 (CN), 1054, 842. HRMS (TOF MS ES⁺) calcd for C₇₃H₈₅N₆O₈ [M + 2H]²⁺ *m/z* 587.3248, found 587.3250.

Arylopeptoid hexamer *m*-6a. Colorless foam (82 mg, 55%, >99% purity). mp = 80–83 °C. ¹H NMR (300 MHz, CDCl₃): δ = 10.06–9.68 (br s, 1H, COO*H*), 8.10–7.83 (m, 2H), 7.65–7.02 (m, 27H), 4.84–4.40 (br s, 12H, 6×CONC*H*₂Ar), 4.30–3.88 (br m, 6H, 6×CONC*H*(CH₃)₂), 1.42–0.84 (m, 36H) ppm. ¹³C NMR (75 MHz, CDCl₃): δ = 172.6 (7C_q), 139.6, 139.5, 136.7, 136.6, 130.2 (13C_q), 131.8, 129.5, 128.8, 128.6, 128.5, 128.1, 128.0, 126.1, 124.6 (29CH), 51.0 (br, 6CH, 6×CONC*H*(CH₃)₂), 43.4 (br, 6CH₂, 6×CON*C*H₂Ar), 21.3 (12CH₃) ppm. *v*_{max}/cm⁻¹ (ATR) 2980, 1713 (C=O), 1627 (C=O), 1464 (CH), 1447, 1410, 1338 (OH), 1178 (CN), 1062. HRMS (TOF MS ES⁺) calcd for C₇₃H₈₅N₆O₈ [M + 2H]²⁺ *m*/*z* 587.3248, found 587.3257.

Arylopeptoid hexamer *p***-6b.** Preparative HPLC of the crude product hindered due to very low solubility. Crude *p***-6b**: Pale rose foam (77 mg, 57%, 71% purity). mp = 123–128 °C. ¹H NMR (300 MHz, CDCl₃/MeOD 1:1): δ = 8.05–7.96 (m, 2H), 7.52– 7.10 (m, 27H), 4.85–4.48 (br m, 12H, 12×CONC*H*₂Ar), 3.60–3.18 (br m, 12H, 12×CONC*H*₂CH₃), 1.30–1.01 (m, 18H) ppm. v_{max}/cm^{-1} (ATR) 2973, 2915, 1716 (C=O), 1627 (C=O), 1464 (CH), 1429, 1312 (OH), 1262, 1175 (CN), 1093, 836. HRMS (TOF MS ES⁺) calcd for C₆₇H₇₄N₆O₈ [M + 2H]²⁺ *m/z* 545.2779, found 545.2775.

Arylopeptoid hexamer *m*-6b. Colorless oil (45 mg, 33%, >99% purity). ¹H NMR (300 MHz, CDCl₃): δ = 8.11–7.78 (m, 2H), 7.60–7.14 (m, 27H), 4.88–4.63 (br s, 7.20H, 6×CONCH₂Ar, major rotamer), 4.63–4.36 (br s, 4.80H, 6×CONCH₂Ar, minor rotamer), 3.62–3.33 (m, 4.80H, 6×CONCH₂CH₃, minor rotamer), 3.33–3.04 (m, 7.20H, 6×CONCH₂CH₃, major rotamer), 1.32–0.94

(m, 18H) ppm. ¹³C NMR (75 MHz, CDCl₃): $\delta = 172.2$ (7C_q), 137.8, 137.5, 136.4, 136.0 (13C_q), 129.7, 129.1, 128.9, 128.6, 126.4, 125.6 (29CH), 51.9 (br, 2.40CH₂, 6×CONCH₂Ar, minor rotamer), 47.1 (br, 3.60CH₂, 6×CONCH₂Ar, major rotamer), 43.3 (br, 3.60CH₂, 6×CONCH₂CH₃, major rotamer), 40.1 (br, 2.40CH₂, 6×CONCH₂CH₃, minor rotamer), 13.7, 12.2 (6CH₃) ppm. v_{max} /cm⁻¹ (ATR) 2992, 2933, 1714 (C=O), 1627 (C=O), 1456 (CH), 1418, 1290 (OH), 1192, 1172 (CN), 1076, 980, 795. HRMS (TOF MS ES⁺) calcd for C₆₇H₇₄N₆O₈ [M + 2H]²⁺ m/z 545.2779, found 545.2776.

Arylopeptoid hexamer p-6c. Colorless foam (82 mg, 39%, >99% purity). mp = 51–54 °C. ¹H NMR (300 MHz, CDCl₃): $\delta = 8.58 - 8.18$ (br s, 1H, COOH), 8.11 - 8.00 (m, 2H), 7.48 -6.95 (m, 57H), 4.86-4.62 (br s, 6.60H, 6×CONCH₂Ar, major rotamer), 4.62–4.36 (br s, 5.40H, 6×CONCH₂Ar, minor rotamer), 3.57-3.31 (m, 5.40H, 6×CONCH₂CH₂, minor rotamer), 3.31-3.03 (m, 6.60H, 6×CONCH2CH2, major rotamer), 2.74-2.54 (m, 5.40H), 2.54–2.35 (m, 6.60H), 1.80–1.31 (m, 24H) ppm. ¹³C NMR (75 MHz, CDCl₃): δ = 172.5, 172.2, 171.8 (7C_a), 142.0, 141.4, 138.9, 138.4, 135.8, 135.4, 135.1 (19C_q), 130.5 (2CH), 129.7, 128.5, 128.3, 128.0, 127.2, 127.0, 126.8, 126.4, 125.8 (57CH), 52.2 (br, 2.70CH₂, 6×CONCH₂Ar, minor rotamer), 48.3 (br, 3.30CH₂, 6×CONCH₂CH₂, major rotamer), 47.2 (br, 3.30CH₂, 6×CONCH₂Ar, major rotamer), 44.6 (br, 2.70CH₂, 6×CONCH₂CH₂, minor rotamer), 35.4, 35.0 (CH₂), 28.6, 27.9, 27.5, 26.3 (12CH₂) ppm. v_{max} /cm⁻¹ (ATR) 3025, 2930, 2859, 1717 (C=O), 1632 (C=O), 1455 (CH), 1423, 1410, 1308 (OH), 1260, 1174 (CN). HRMS (TOF MS ES⁺) calcd for $C_{115}H_{121}N_6O_8$ [M + 2H]²⁺ m/z 857.4657, found 857.4657.

Arylopeptoid hexamer m-6c. Colorless foam (84 mg, 39%, >99% purity). mp = 48–51 °C. ¹H NMR (300 MHz, CDCl₃): $\delta = 9.70-9.30$ (br s, 1H, COOH), 8.04–7.70 (m, 2H), 7.48– 6.94 (m, 57H), 4.83–4.59 (br s, 6.60H, 6×CONCH₂Ar, major rotamer), 4.59–4.32 (br s, 5.40H, 6×CONCH₂Ar, minor rotamer), 3.57-3.28 (m, 5.40H, 6×CONCH2CH2, minor rotamer), 3.28-3.00 (m, 6.60H, 6×CONCH₂CH₂, major rotamer), 2.71–2.51 (m, 5.40H), 2.51-2.33 (m, 6.60H), 1.78-1.22 (m, 24H) ppm. ¹³C NMR (75 MHz, CDCl₃): δ = 172.6, 172.2, 171.8 (7C_a), 142.0, 141.4, 137.6, 137.4, 137.3, 136.4, 136.2, 135.7 (19C_a), 129.7, 129.0, 128.5, 128.3, 127.7, 126.4, 126.2, 125.8, 125.6 (59CH), 52.2 (br, 2.70CH₂, 6×CONCH₂Ar, minor rotamer), 48.4 (br, 3.30CH₂, 6×CONCH₂CH₂, major rotamer), 47.4 (br, 3.30CH₂, 6×CONCH₂Ar, major rotamer), 44.7 (br, 2.70CH₂, 6×CONCH₂CH₂, minor rotamer), 35.4, 35.0 (6CH₂), 28.5, 27.9, 27.6, 26.4 (12CH₂) ppm. v_{max} /cm⁻¹ (ATR) 3025, 2933, 2860, 1714 (C=O), 1630 (C=O), 1453 (CH), 1416, 1301 (OH), 1195, 1171 (CN). HRMS (TOF MS ES⁺) calcd for $C_{115}H_{121}N_6O_8$ [M + 2H]²⁺ m/z 857.4657, found 857.4650.

Arylopeptoid hexamer *p*-6d. Colorless foam isolated as the corresponding TFA-salt (164 mg, 58%, >99% purity). mp = 62–65 °C. ¹H NMR (300 MHz, MeCN-*d*₃): δ = 11.30–9.40 (br s, 7H, 6×CF₃COO*H* and COO*H*), 8.00–7.91 (m, 2H), 7.64–7.08 (m, 27H), 4.83–4.44 (m, 12H, 6×CONC*H*₂Ar), 4.10–3.89 (m, 12H), 3.88–3.66 (m, 24H), 3.66–3.45 (m, 12H), 3.40–3.21 (m, 12H, 6×CONC*H*₂CH₂), 3.20–2.91 (m, 12H) ppm. ¹³C NMR (75 MHz, MeCN-*d*₃): δ = 174.8, 174.2 (6C_q), 167.9 (C_q), 161.0 (q, *J* = 36.1 Hz, 6C_q, 6×CF₃COOH), 142.6, 139.5, 136.0, 135.5, 130.9 (13C_q), 131.0

(2CH), 131.2, 129.4, 128.4, 128.0 (27CH), 116.0 (q, J = 290.9 Hz, 6C_q, 6×CF₃COOH), 64.7 (12CH₂), 55.8 (6CH₂, 6×CON*C*H₂CH₂), 53.7 (6CH₂, 6×CON*C*H₂Ar), 53.1 (12CH₂), 41.3 (6CH₂) ppm. $v_{\rm max}/{\rm cm^{-1}}$ (ATR) 3001, 2930, 2858, 1669 (C=O), 1635 (C=O), 1458 (CH), 1418, 1179 (CN), 1129 (CO), 1088 (CN), 938, 798. HRMS (TOF MS ES⁺) calcd for C₉₁H₁₁₈N₁₂O₁₄ [M + 4H]⁴⁺ *m/z* 400.7217, found 400.7208.

Arylopeptoid hexamer *m*-6d. Colorless foam isolated as the corresponding TFA-salt (119 mg, 41%, >99% purity). mp = 61-64 °C. ¹H NMR (300 MHz, MeCN- d_3): δ = 7.94–7.75 (m, 2H), 7.54-6.50 (m, 34H, Ar-H, 6×CF₃COOH and COOH), 4.80-4.39 (m, 12H, 6×CONCH₂Ar), 4.12–3.87 (m, 12H), 3.87–3.39 (m, 36H), 3.39–3.16 (m, 12H, 6×CONCH₂CH₂), 3.16–2.92 (m, 12H) ppm. ¹³C NMR (75 MHz, MeCN- d_3): $\delta = 174.6, 174.4, 174.2, 174.2$ $(6C_{q}), 168.0 (C_{q}), 160.8 (q, J = 36.4 Hz, 6C_{q}, 6 \times CF_{3}COOH), 137.7,$ 137.6, 136.6, 136.5, 136.4, 135.6, 132.2 (13C_q), 132.7, 131.4, 130.4, 130.3, 130.2, 129.9, 129.8, 129.4, 128.6, 127.9, 127.5, 127.2, 125.4, 125.3, 124.8 (27CH), 117.1 (q, J = 290.3 Hz, $6C_{q}$, $6 \times CF_{3}$ COOH), 64.7, 64.3 (12CH₂), 56.2, 56.1, 55.9, 55.6 (6CH₂, 6×CONCH₂CH₂), 53.7 (6CH₂, 6×CONCH₂Ar), 53.2 (12CH₂), 41.5, 41.0, 40.8 (6CH₂) ppm. v_{max}/cm⁻¹ (ATR) 3013, 1671 (C=O), 1633 (C=O), 1463 (CH), 1417, 1269 (OH), 1184 (CN), 1138 (CO), 1088 (CN), 986, 1798. HRMS (TOF MS ES⁺) calcd for $C_{91}H_{118}N_{12}O_{14}$ [M + 4H]⁴⁺ m/z 400.7217, found 400.7217.

Arylopeptoid hexamer p-6e. Pale yellowish foam isolated as the corresponding TFA-salt (58 mg, 21%, >99% purity). mp = 60-63 °C. ¹H NMR (300 MHz, MeCN- d_3): $\delta = 8.80-8.63$ (m, 6H), 8.63-8.54 (m, 6H), 8.51-8.39 (br s, 7H, 6×CF₃COOH and COOH), 8.36-8.24 (m, 6H), 7.92-7.74 (m, 8H), 7.47-7.07 (m, 27H), 4.84–4.65 and 4.65–4.44 (2×br s, 2 × 12H, 6×CONC H_2 Ar and 6×CONCH₂Pyr) ppm. ¹³C NMR (75 MHz, MeCN- d_3): δ = 173.3, 172.9 (6C_q), 167.7 (C_q), 161.3 (q, J = 36.0 Hz, 6C_q, 6×CF₃COOH), 146.1 (6CH), 142.3 (6CH), 141.6 (6CH), 139.4, 138.9, 136.4, 135.8 (19C_q), 130.9 (2CH), 129.5, 128.6, 128.1, 127.8, 127.6 (27CH), 127.7 (6CH), 117.3 (q, J = 290.9 Hz, 6C_q, $6 \times CF_3$ COOH), 53.9 and 47.1 (2 × 6CH₂, 6×CONCH₂Ar and $6 \times \text{CON}C\text{H}_2\text{Pyr}$ ppm. $v_{\text{max}}/\text{cm}^{-1}$ (ATR) 3060, 3028, 1675 (C=O), 1634 (C=O), 1458 (CH), 1410, 1177 (CN), 1135 (CN), 995, 796. HRMS (TOF MS ES⁺) calcd for $C_{91}H_{81}N_{12}O_8$ [M + 3H]³⁺ m/z 489.8761, found 489.8763.

Arylopeptoid hexamer *p*-7a. Pale rose foam (77 mg, 55%, >99% purity). mp = 229–232 °C. ¹H NMR (300 MHz, CDCl₃): δ = 7.82–7.65 (m, 2H), 7.55–6.92 (m, 27H), 6.84–6.16 (m, 2H, CONH₂), 4.82–4.37 (br s, 12H, 6×CONCH₂Ar), 4.34–4.02 (br m, 6H, 6×CONCH(CH₃)₂), 1.42–0.92 (m, 36H) ppm. ¹³C NMR (75 MHz, CDCl₃): δ = 172.6, 172.5, 170.1 (7C_q), 140.8, 136.7, 135.1, 131.3 (13C_q), 129.6, 128.7, 127.9, 127.1, 126.5, 126.2 (29CH), 51.0 (br, 6CH, 6×CONCH(CH₃)₂), 43.5 (br, 6CH₂, 6×CONCH₂Ar), 21.4 (12CH₃) ppm. v_{max} /cm⁻¹ (ATR) 2969, 2929, 1674 (C=O), 1628 (C=O), 1437 (CH), 1405, 1338, 1174 (CN), 1054, 842. HRMS (TOF MS ES⁺) calcd for C₇₃H₈₆N₇O₇ [M + 2H]²⁺ m/z 586.8328, found 586.8336.

Arylopeptoid hexamer *m*-7a. Pale rose foam (82 mg, 58%, >99%). mp = 70–73 °C. ¹H NMR (300 MHz, CDCl₃): δ = 7.93–7.61 (m, 2H), 7.59–6.90 (m, 28H, Ar-*H* and CON*H*H), 6.70–6.47 (br s, 1H, CON*HH*), 4.82–4.34 (br s, 12H, 6×CONC*H*₂Ar), 4.30–3.92 (br m, 6H, 6×CONC*H*(CH₃)₂), 1.42–0.92 (m, 36H) ppm. ¹³C

NMR (75 MHz, CDCl₃): δ = 172.8, 172.7, 170.7 (7C_q), 139.6, 136.4, 132.7 (13C_q), 130.9, 129.5, 128.8, 128.7, 128.6, 128.1, 126.2, 126.1, 124.8, 124.6 (29CH), 51.0 (br, 6CH, 6×CON*C*H(CH₃)₂), 43.5 (br, 6CH₂, 6×CON*C*H₂Ar), 21.3 (12CH₃) ppm. v_{max} /cm⁻¹ (ATR) 2973, 2932, 1678 (C=O), 1624 (C=O), 1448 (CH), 1410, 1338, 1175 (CN), 1061, 793. HRMS (TOF MS ES⁺) calcd for C₇₃H₈₆N₇O₇ [M + 2H]²⁺ *m*/*z* 586.8328, found 586.8329.

Arylopeptoid hexamer p-7b. Pale rose foam (70 mg, 54%, >99% purity). mp = 67-70 °C. ¹H NMR (300 MHz, CDCl₃): $\delta = 7.86-7.79$ (m, 2H), 7.51–7.09 (m, 27H), 6.95–6.32 (m, 2H, $CONH_2$), 4.90–4.65 (br s, 7.20H, 6×CONCH₂Ar, major rotamer), 4.65-4.41 (br s, 4.80H, 6×CONCH₂Ar, minor rotamer), 3.65-3.38 (m, 4.80H, 6×CONCH₂CH₃, minor rotamer), 3.38–3.11 (m, 7.20H, 6×CONCH2CH3, major rotamer), 1.33-0.95 (m, 18H) ppm. ¹³C NMR (75 MHz, CDCl₃): δ = 172.4, 171.9, 170.0 (7C_a), 139.0, 138.6, 135.9, 135.2, 131.8 (13C_a), 129.7, 128.5, 128.1, 126.9, 126.3 (29CH), 51.9 (br, 2.40CH₂, 6×CONCH₂Ar, minor rotamer), 47.0 (br, 3.60CH₂, 6×CONCH₂Ar, major rotamer), 43.3 (br, 3.60CH₂, 6×CONCH₂CH₃, major rotamer), 40.2 (br, 2.40CH₂, 6×CONCH₂CH₃, minor rotamer), 13.7, 12.1 (6CH₃) ppm. v_{max}/cm⁻¹ (ATR) 2973, 1675 (C=O), 1627 (C=O), 1430 (CH), 1410, 1286, 1176 (CN), 1095, 1074, 980. HRMS (TOF MS ES⁺) calcd for $C_{67}H_{75}N_7O_7 [M + 2H]^{2+} m/z$ 544.7859, found 544.7861.

Arylopeptoid hexamer p-7c. Colorless foam (95 mg, 46%, >99% purity). mp = 52–55 °C. ¹H NMR (300 MHz, CDCl₃): δ = 7.82–7.72 (m, 2H), 7.46–6.97 (m, 57H), 6.82–6.20 (m, 2H, CON H_2), 4.84–4.60 (br s, 6.60H, 6×CONC H_2 Ar, major rotamer), 4.60–4.36 (br s, 5.40H, 6×CONCH₂Ar, minor rotamer), 3.57-3.31 (m, 5.40H, 6×CONCH₂CH₂, minor rotamer), 3.31-3.05 (m, 6.60H, 6×CONCH₂CH₂, major rotamer), 2.74–2.54 (m, 5.40H), 2.54–2.35 (m, 6.60H), 1.80–1.24 (m, 24H) ppm. ¹³C NMR (75 MHz, CDCl₃): $\delta = 172.4$, 172.1, 171.7 (7C_a), 142.0, 141.4, 138.9, 138.4, 136.0, 135.4, 135.2 (19C_q), 129.6, 128.5, 128.3, 128.1, 128.0, 127.1, 127.0, 126.8, 126.4, 125.8 (59CH), 52.2 (br, 2.70CH₂, 6×CONCH₂Ar, minor rotamer), 48.3 (br, 3.30CH₂, 6×CONCH₂CH₂, major rotamer), 47.2 (br, 3.30CH₂, 6×CONCH₂Ar, major rotamer), 44.6 (br, 2.70CH₂, 6×CONCH₂CH₂, minor rotamer), 35.4, 35.0 (6CH₂), 28.6, 27.9, 27.5, 26.3 (12CH₂) ppm. v_{max} /cm⁻¹ (ATR) 2929, 1675 (C=O), 1629 (C=O), 1453 (CH), 1410, 1309, 1173 (CN). HRMS (TOF MS ES⁺) calcd for $C_{115}H_{123}N_7O_7 [M + 2H]^{2+} m/z 856.9737$, found 856.9747.

Arylopeptoid hexamer *m*-7c. Colorless foam (109 mg, 53%, >99% purity). mp = 54–57 °C. ¹H NMR (300 MHz, CDCl₃): δ = 7.84–7.56 (m, 2H), 7.56–6.92 (m, 57H), 6.22–5.54 (m, 2H, CON*H*₂), 4.81–4.56 (br s, 6.90H, 6×CONC*H*₂Ar, major rotamer), 4.56–4.32 (br s, 5.10H, 6×CONC*H*₂Ar, minor rotamer), 3.56–3.29 (m, 5.10H, 6×CONC*H*₂CH₂, minor rotamer), 3.29–3.03 (m, 6.90H, 6×CONC*H*₂CH₂, major rotamer), 2.71–2.54 (m, 5.10H), 2.54–2.33 (m, 6.90H), 1.75–1.23 (m, 24H) ppm. ¹³C NMR (75 MHz, CDCl₃): δ = 172.3, 172.0, 171.6, 169.6 (7C_q), 142.0, 141.4, 137.8, 137.5, 136.8, 136.6, 136.1 (19C_q), 129.6, 129.0, 129.0, 128.5, 128.3, 126.5, 126.2, 125.8, 125.5 (59CH), 52.2 (br, 2.55CH₂, 6×CONC*H*₂Ar, minor rotamer), 48.4 (br, 3.45CH₂, 6×CONC*H*₂Ar, major rotamer), 44.6 (br, 2.55CH₂,

6×CON*C*H₂CH₂, minor rotamer), 35.4, 35.0 (6CH₂), 28.6, 28.0, 27.7, 26.4 (12CH₂) ppm. v_{max}/cm^{-1} (ATR) 3023, 2929, 2858, 1675 (C=O), 1631 (C=O), 1453 (CH), 1415, 1301, 1200, 793. HRMS (TOF MS ES⁺) calcd for C₁₁₅H₁₂₃N₇O₇ [M + 2H]²⁺ *m/z* 856.9737, found 856.9743.

Arylopeptoid hexamer p-7d. Colorless foam isolated as the corresponding TFA-salt (159 mg, 58%, >99% purity). mp = 69-72 °C. ¹H NMR (300 MHz, MeCN- d_3): $\delta = 11.50-9.60$ (br s, 6H, 6×CF₃COOH), 7.85-7.76 (m, 2H), 7.55-7.14 (m, 27H), 7.09-6.25 (m, 2H, CONH₂), 4.77-4.47 (m, 12H, 6×CONCH₂Ar), 4.08-3.89 (m, 12H), 3.89-3.65 (m, 24H), 3.65-3.47 (m, 12H), 3.37-3.18 (m, 12H, 6×CONCH₂CH₂), 3.18–2.94 (m, 12H) ppm. ¹³C NMR (75 MHz, MeCN- d_3): $\delta = 174.9$, 174.4, 174.2, 169.6 (7C_a), 160.8 $(q, J = 36.3 \text{ Hz}, 6C_q, 6 \times CF_3 COOH), 141.1, 139.5, 135.5, 134.0$ (13C_q), 131.2, 129.4, 129.0, 128.4, 128.0, 127.8 (29CH), 117.2 (q, J = 290.6 Hz, $6C_q$, $6 \times CF_3$ COOH), 64.7 (12CH₂), 55.9 (6CH₂, 6×CONCH₂CH₂), 53.7 (6CH₂, 6×CONCH₂Ar), 53.1 (12CH₂), 41.3 (6CH₂) ppm. v_{max}/cm⁻¹ (ATR) 2874, 1675 (C=O), 1631 (C=O), 1466 (CH), 1181 (CN), 1129 (CO), 1088, 984, 798. HRMS (TOF MS ES⁺) calcd for $C_{91}H_{119}N_{13}O_{13}$ [M + 4H]⁴⁺ m/z 400.4757, found 400.4759.

Arylopeptoid hexamer m-7d. Colorless foam isolated as the corresponding TFA-salt (162 mg, 59%, >99% purity). mp = 57-60 °C. ¹H NMR (300 MHz, MeCN- d_3): $\delta = 8.60-7.80$ (br s, 6H, 6×CF₃COOH), 7.80-7.65 (m, 2H), 7.55-7.13 (m, 28H, Ar-H and CONHH), 6.60-6.35 (br s, 1H, CONHH), 4.80-4.39 (m, 12H, 6×CONCH₂Ar), 4.10–3.90 (m, 12H), 3.90–3.37 (m, 36H), 3.37-3.15 (m, 12H, 6×CONCH₂CH₂), 3.15-2.92 (m, 12H) ppm. ¹³C NMR (75 MHz, MeCN- d_3): $\delta = 175.0, 174.5, 174.1, 169.9$ $(7C_q)$, 160.7 (q, J = 36.7 Hz, $6C_q$, $6 \times CF_3 COOH$), 160.5, 160.0, 137.7, 137.5, 136.7, 136.6, 136.5, 135.7, 135.2 (13C_q), 131.5, 131.4, 129.4, 127.9, 127.5, 127.4, 127.2, 125.5, 125.2, 125.0 (27CH), 117.1 $(q, J = 290.1 \text{ Hz}, 6C_a, 6 \times CF_3 \text{COOH}), 64.7 (12 \text{CH}_2), 56.0, 55.9,$ 55.5 (6CH₂, 6×CONCH₂CH₂), 53.7 (6CH₂, 6×CONCH₂Ar), 53.2 $(12CH_2), 41.5, 41.1, 41.0 (6CH_2) \text{ ppm. } v_{\text{max}}/\text{cm}^{-1} (\text{ATR}) 2973, 1668$ (C=O), 1634 (C=O), 1464 (CH), 1418, 1178 (CO), 1137 (CN), 1088, 983, 798. HRMS (TOF MS ES⁺) calcd for C₉₁H₁₁₉N₁₃O₁₃ [M $+ 4H^{4+} m/z 400.4757$, found 400.4764.

Arylopeptoid hexamer *p*-7e. Pale yellowish foam isolated as the corresponding TFA-salt (58 mg, 22%, >99% purity). mp = 62–65 °C. ¹H NMR (300 MHz, MeCN-*d*₃): δ = 8.90–8.73 (m, 6H), 8.73–8.62 (m, 6H), 8.50–8.20 (m, 6H), 7.00–7.81 (m, 8H), 7.60–7.20 (m, 27H), 6.85–6.25 (m, 8H, 6×CF₃COO*H* and CON*H*₂), 4.95–4.76 and 4.76–4.54 (2×br s, 2 × 12H, 6×CONC*H*₂Ar and 6×CONC*H*₂Pyr) ppm. ¹³C NMR (75 MHz, MeCN-*d*₃): δ = 173.2, 172.9, 169.4 (7C_q), 161.2 (q, *J* = 36.2 Hz, 6C_q, 6×CF₃COOH), 146.0 (6CH), 142.4 (6CH), 141.6 (6CH), 139.4, 138.9, 136.4, 135.8, 134.3 (19C_q), 130.9, 129.5, 128.9, 128.6, 128.3, 128.2, 127.6 (35CH), 117.3 (q, *J* = 290.7 Hz, 6C_q, 6×CF₃COOH), 54.0 and 47.0 (2 × 6CH₂, 6×CONCH₂Ar and 6×CONCH₂Pyr) ppm. *v*_{max}/cm⁻¹ (ATR) 3086, 1673 (C=O), 1634 (C=O), 1461 (CH), 1410, 1185, 1134, 994, 796. HRMS (TOF MS ES⁺) calcd for C₉₁H₈₂N₁₃O₇ [M + 3H]³⁺ *m*/*z* 489.5481, found 489.5472.

Arylopeptoid hexamer *p*-7f. Colorless foam isolated as the corresponding TFA-salt (119 mg, 49%, >99% purity). mp = 78–81 °C. ¹H NMR (300 MHz, MeOD): δ = 7.98–7.82 (m, 3H), 7.57–7.12 (m, 26H), 4.90–4.71 (br s, 5.00H, 6×CONCH₂Ar, minor

rotamer), 4.71–4.48 (br s, 7.00H, 6×CONCH₂Ar, major rotamer), 3.60-3.41 (m, 7.00H, 6×CONCH₂CH₂, major rotamer), 3.41-3.22 (m, 5.00H, 6×CONCH2CH2, minor rotamer), 3.05-2.85 (m, 7.00H), 2.85–2.65 (m, 5.00H), 1.82–1.30 (m, 24H) ppm. ¹³C NMR (75 MHz, MeOD): $\delta = 174.7$, 174.4, 174.2, 174.1, 171.9, 171.8 $(7C_q)$, 162.3 (q, J = 35.9 Hz, $6C_q$, $6 \times CF_3 COOH$), 142.2, 140.8, 140.2, 137.3, 136.7, 136.5, 134.5, 134.2 (13C_q), 131.1, 129.9, 129.3, 129.1, 128.9, 128.4, 128.2, 127.6 (29CH), 117.9 (q, J = 291.4 Hz, $6C_{q}$, $6 \times CF_{3}COOH$), 53.5 (br, 3.50CH₂, $6 \times CONCH_{2}Ar$, major rotamer), 50.0 (br, 2.50CH₂, 6×CONCH₂CH₂, minor rotamer), 48.9 (br, 2.50CH₂, 6×CONCH₂Ar, minor rotamer), 46.0, 45.9, 45.7 (br, 3.50CH₂, 6×CONCH₂CH₂, major rotamer), 40.4, 40.2 $(6CH_2)$, 26.4, 25.9, 25.6, 25.1 $(12CH_2)$ ppm. v_{max}/cm^{-1} (ATR) 3026, 2938, 2874, 1675 (C=O), 1609 (C=O), 1471 (CH), 1429, 1410, 1200, 1177, 1131, 835, 799. HRMS (TOF MS ES⁺) calcd for $C_{79}H_{105}N_{13}O_7 [M + 2H]^{2+} m/z$ 673.9130, found 673.9119.

Arylopeptoid hexamer p-7g. Colorless foam (47 mg, 31%, >99% purity). mp = 181-184 °C (decomposition). ¹H NMR (300 MHz, MeOD): δ = 7.92–7.82 (m, 2H), 7.52–7.19 (m, 27H), 4.90-4.75 (br s, 5.00H, 6×CONCH₂Ar, minor rotamer), 4.71-4.55 (br s, 7.00H, 6×CONCH₂Ar, major rotamer), 4.21-4.08 (br s, 7.00H, 6×CONCH₂COOH, major rotamer), 4.01-3.87 (br s, 5.00H, 6×CONCH₂COOH, minor rotamer) ppm. ¹³C NMR (75 MHz, MeOD): δ = 175.0, 174.7, 174.5, 172.1, 171.9 (13C_a), 142.0, 141.8, 140.3, 140.2, 136.8, 136.6, 136.2, 136.1, 136.0, 135.9, 135.8, 134.5, 134.3 (13C_q), 131.4, 131.3, 129.9, 129.8, 129.6, 129.5, 129.4, 129.3, 129.2, 128.6, 128.4, 128.3, 128.2, 127.8, 127.6 (29CH), 55.0 (br, 3.50CH₂, 6×CONCH₂Ar, major rotamer), 51.6 (br, 2.50CH₂, 6×CONCH₂COOH, minor rotamer), 50.6 (br, $2.50CH_2$, $6 \times CONCH_2Ar$, minor rotamer), 48.3 (br, $3.50CH_2$, $6 \times CONCH_2 COOH$, major rotamer) ppm. v_{max}/cm^{-1} (ATR) 1730, 1645, 1634 (C=O), 1609 (C=O), 1456 (CH), 1423, 1404, 1260, 1199, 1176 (CN), 1006, 952, 838. HRMS (TOF MS ES⁺) calcd for $C_{67}H_{61}N_7O_{19}$ [M + 2Na]²⁺ *m*/*z* 656.6909, found 656.6875.

Arylopeptoid hexamer p-8. Colorless foam isolated as the corresponding TFA-salt (99 mg, 25%, >99% purity). mp = 80-84 °C. ¹H NMR (300 MHz, MeCN- d_3): δ = 7.83–7.70 (m, 2H), 7.57-6.90 (m, 71H), 5.15-4.28 (m, 30H, CONH₂, 4×CF₃COOH and 12×CONCH₂Ar), 4.10–3.87 (m, 12H), 3.87–3.63 (m, 16H), 3.63-3.46 (m, 8H), 3.45-3.12 (m, 16H), 3.11-2.89 (m, 8H), 2.67-2.24 (m, 8H), 1.69–1.23 (m, 16H), 1.22–0.94 (m, 24H) ppm. ¹³C NMR (75 MHz, MeCN- d_3): $\delta = 174.8, 172.6, 172.3, 169.7 (13C_a),$ 160.5 (q, J = 37.1 Hz, $4C_q$, $4 \times CF_3 COOH$), 143.1, 142.1, 138.3, 137.8, 136.1, 134.9, 133.5 (29C_q), 131.3, 129.5, 129.2, 128.8, 128.3, 128.2, 127.8, 127.6, 127.4, 126.6 (73 CH), 117.0 (q, J = 289.7 Hz, 4C_q, 4×CF₃COOH), 64.8 (8CH₂), 56.3 (4CH₂), 54.1 (4CH₂, 4×CON[2-morpholinoethyl]CH₂Ar), 53.2 (8CH₂), 51.7 (4CH, 4×CONCH(CH₃)₂), 49.9, 46.0 (4CH₂), 48.5 (4CH₂, 4×CON[4phenylbutyl]CH₂Ar), 44.0 (4CH₂, 4×CON[isopropyl]CH₂Ar), 41.7 (4CH₂), 35.9, 35.6 (4CH₂), 30.3, 29.5, 28.9, 28.7 (8CH₂), 21.2 (8CH₃) ppm. *v*_{max}/cm⁻¹ (ATR) 2927, 2864, 1670 (C=O), 1625 (C=O), 1458 (CH), 1410, 1364, 1311, 1266, 1173 (CO), 1138 (CN), 1088, 928. HRMS (TOF MS ES⁺) calcd for C₁₇₉H₂₁₁N₁₇O₁₇ [M + 4H]⁴⁺ m/z 717.6537, found 717.6536.

Arylopeptoid hexamer *m*-8. Colorless foam isolated as the corresponding TFA-salt (108 mg, 27%, >99% purity). mp = 73–77 °C. ¹H NMR (300 MHz, MeCN-*d*₃): δ = 7.85–7.63 (m, 2H), 7.55–

6.90 (m, 71H), 5.05–4.27 (m, 30H, CONH₂, 4×CF₃COOH and 12×CONCH₂Ar), 4.09–3.83 (m, 12H), 3.83–3.37 (m, 24H), 3.37– 3.07 (m, 16H), 3.07–2.87 (m, 8H), 2.65–2.23 (m, 8H), 1.68–1.20 (m, 16H), 1.20-0.90 (m, 24H) ppm. 13C NMR (75 MHz, MeCN d_3): $\delta = 174.4, 172.7, 172.3, 169.9 (13C_a), 160.5 (q, J = 36.8 Hz, J)$ 4C_a, 4×CF₃COOH), 143.0, 141.2, 139.7, 139.0, 137.8, 137.7, 136.4, 135.0 (29C_a), 131.3, 130.2, 129.8, 129.7, 129.6, 129.5, 129.2, 128.8, 127.9, 126.6, 126.4, 125.8 (73 CH), 117.1 (q, J = 288.2 Hz, 4C_a, 4×CF₃COOH), 64.7 (8CH₂), 56.2 (4CH₂), 54.0 (4CH₂, 4×CON[2-morpholinoethyl]CH2Ar), 53.2 (8CH2), 51.8 (4CH, 4×CONCH(CH₃)₂), 49.9, 45.5 (4CH₂), 48.6 (4CH₂, 4×CON[4phenylbutyl]CH₂Ar), 44.0 (4CH₂, 4×CON[isopropyl]CH₂Ar), 41.6 (4CH₂), 35.9, 35.6 (4CH₂), 29.6, 29.4, 28.9, 28.8 (8CH₂), 21.4 (8CH₃) ppm. v_{max} /cm⁻¹ (ATR) 2935, 2859, 1669 (C=O), 1622 (C=O), 1463 (CH), 1414, 1343, 1306, 1269, 1179 (CO), 1136 (CN), 1086, 796. HRMS (TOF MS ES⁺) calcd for $C_{179}H_{211}N_{17}O_{17}$ [M + 4H]⁴⁺ *m*/*z* 717.6537, found 717.6542.

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