

Structural Design and Synthesis of Bimodal PNA That Simultaneously Binds Two Complementary DNAs To Form Fused Double Duplexes

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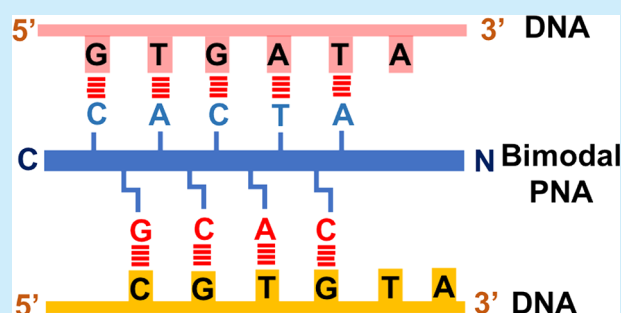


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ABSTRACT: Bimodal PNAs are new PNA constructs designed to bind two different cDNA sequences synchronously to form double duplexes. They are synthesized on solid phase using sequential coupling and click reaction to introduce a second base in each monomer at C_α via alkyltriazole linker. The ternary bimodal PNA:DNA complexes show stability higher than that of individual duplexes. Bimodal PNAs are appropriate to create higher-order fused nucleic acid assemblies.



Peptide nucleic acids (PNA) are designed acyclic analogues of DNA endowed with an ability to recognize and bind complementary DNA/RNA sequences to form PNA:DNA and PNA:RNA duplexes.¹ The achiral PNA backbone with repeating units of aminoethylglycine (*aeg*) has nucleobases (A/T/C/G) connected via tertiary amide group (Figure 1a).² The structural geometry of *aeg* backbone in PNA ensures matching of the interbase distance in PNA with that of adjacent bases on the sugar–phosphate backbone of DNA/RNA,³ leading to sequence-specific complementary base pairing between the PNA and the DNA/RNA complements

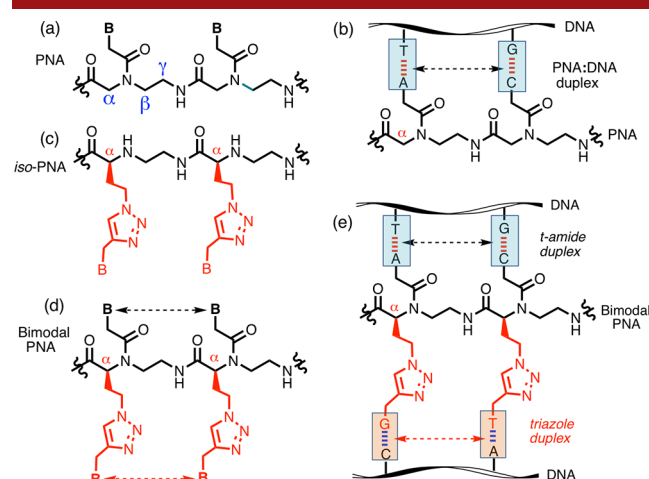


Figure 1. Structures of (a) *aeg*-PNA, (b) PNA:DNA duplex, (c) *iso*-PNA, (d) bimodal PNA (*bm*-PNA), and (e) bimodal PNA double duplex. B = T/A/G/C.

(Figure 1b). PNA:DNA duplexes have strong binding and superior stability, compared to that of natural DNA:DNA duplexes.⁴ This, along with unique strand invasion properties,⁵ has allowed PNA to be used for several biotechnological applications in diagnostics⁶ and antisense therapeutics.⁷ The simplicity of PNA construct provides abundant opportunities for rational design of its structure to improve its properties⁸ and cellular uptake.^{9,10}

Substitutions on *aeg* backbone at C_ω , C_β , and C_γ does not seriously impair the hybridization with cDNA/cRNA and, depending on the nature of substitution, PNAs acquire additional functions.^{10,11} This manuscript describes the design of *iso*-PNA (Figure 1c) carrying nucleobases (A/G/C/T) on side chain at C_α linked through a triazole ring, instead of t-amide link. The interbase distance in *iso*-PNA is similar to that in original PNA and can be structurally fused with original PNA (Figure 1a) to generate “bimodal PNA” composed of two faces (Figure 1d). This equips a single PNA strand to recognize and concurrently bind two different DNA strands to produce double duplexes (Figure 1e) with a common PNA backbone.

Synthetic “Janus bases” known in literature recognize two natural bases, one from each face¹² and PNAs composed from such synthetic bases have innovative applications.¹³ The

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bimodal PNA “*bm*-PNA” (Figure 1d) described here are conceptually different with PNA backbone itself acquiring Janus character. *bm*-PNA is designed to bind cDNA/RNA/PNA as single strands or two strands simultaneously through canonical base pairing. It is demonstrated that *bm*-PNA forms double duplexes by simultaneous binding of two complementary DNA strands (Figure 1e) and the stability of the DNA:*bm*-PNA:DNA ternary complex is enhanced over that of isolated duplexes.

The target *iso*-PNA 1 and *bm*-PNA 2 (Figure 2) have nucleobases attached to C_α(S) of each glycol component on

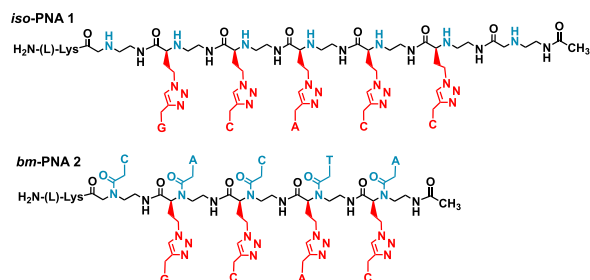


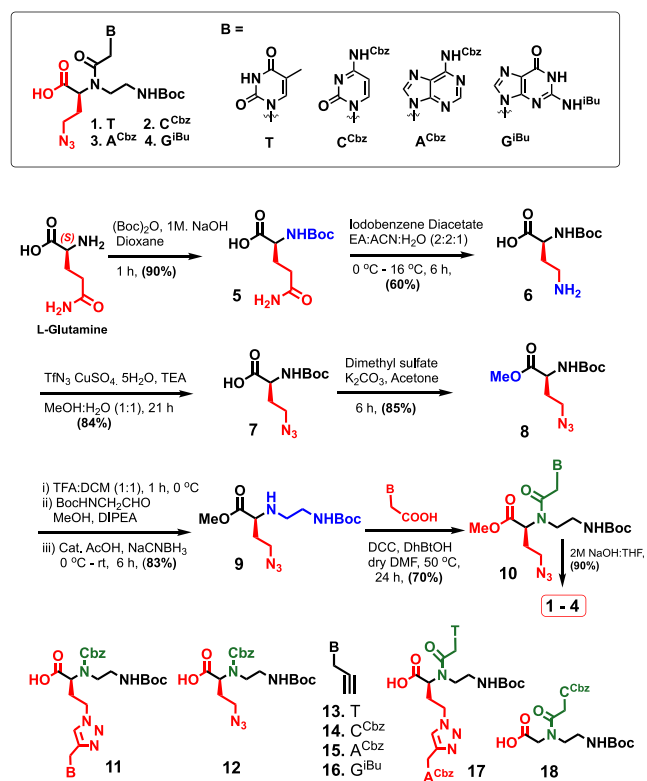
Figure 2. Structures of *iso*-PNA 1 and *bm*-PNA 2.

aeg backbone through an ethyl triazole spacer (Figure 2). In *bm*-PNA 2 this is in addition to the standard t-amide linked nucleobases as in PNA. The triazole side chain at C_α design was chosen more due to ease of synthesis than any structural or functional considerations. *iso*-PNA 1 (Figure 2) bearing only nucleobases anchored on C_α side chain through triazole was used to test the ability of this new PNA analogue for base pairing with cDNA. The monomers 1–4 possessing C_α(S)-ethylazido side chain for subsequent click reaction were synthesized from easily available L-glutamine through intermediate steps via compounds 5–10 (Scheme 1). The monomers 11 (B = C^{Cbz}/A^{Cbz}/G^{iBu}) needed for assembling *iso*-PNA 1 were synthesized from azide 12 and base propynes 14–16 by click chemistry. The bimodal PNA monomer 17 having 2 bases per *aeg* unit was made by coupling 1 with 15. The *aeg*-PNA-C monomer 18 was made by standard procedures.¹⁴

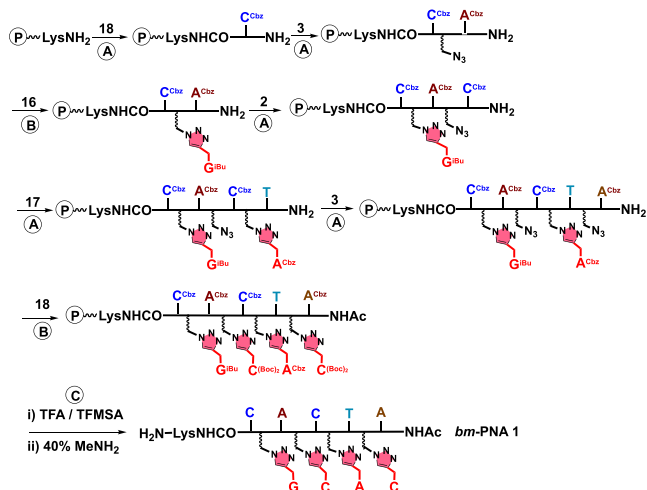
bm-PNA 2 with mixed sequence of five bases on t-amide side and four bases on triazole side was assembled by solid-phase synthesis on MBHA resin (Scheme 2), using well established protocols.¹⁴ The first coupling on resin was done with standard PNA monomer 18 to avoid possible steric effects with modified monomers. Each coupling of C_α-ethylazido PNA monomers (1–3) on resin was followed by click reaction with appropriate base propynes (14–16). At an intermediate step, the preclicked bimodal PNA monomer 17 having T at the t-amide side and A on the triazole side was used to test strategy for future development of a general approach for synthesis of any mixed *bm*-PNA oligomers. The *iso*-PNA 1 was similarly synthesized on solid phase using monomers 11 (B = C^{Cbz}/A^{Cbz}/G^{iBu}). After the solid-phase synthesis, the resin bound *iso*-PNA 1 and *bm*-PNA 2 product oligomers were deprotected and cleaved from the resin, purified by reverse-phase HPLC and characterized by MALDI-TOF spectra (see the Supporting Information).

The successful formation of antiparallel PNA:DNA triazole duplexes from *iso*-PNA 1 and *bm*-PNA 2 (Figures 3A–E) was shown by sigmoidal transitions observed in a temperature–UV absorbance plot^{14,15} (see Figure 4). The melting point (*T*_m)

Scheme 1. Structures of Monomers 1–4 and Their Synthesis



Scheme 2. Solid-Phase Synthesis of Bimodal PNA *bm*-PNA 2^a



^a Monomer coupling: (A/T/G/C; 3 equiv), dry DMF (500 μ L) HOBt (3 equiv), HBTU (3 equiv), and DIPEA (3 equiv); © Click reaction: resin (10 mg) in dry DMF:pyridine (1:1), alkyne (9 mg, 6 equiv), CuI (12 mg, 18 equiv), ascorbic acid (3 mg, 5 equiv), DIPEA (12 μ L); and © Cleavage: resin 10 mg, thioanisole (20 μ L), 1,2-ethanedithiol (8 μ L), TFA (200 μ L), TFMSA (16 μ L). Compound numbers at each step are taken from Scheme 2. For details regarding the reaction procedures, see the Supporting Information.

values extracted from the peak in the first derivative curves are shown in Figure 5. The *iso*-PNA 1:DNA 1 duplex (Figure 3A) showed *T*_m of 42.8 °C, just for a 5-mer duplex (Figure 5A). Since it has cationic amino backbone, it is possible that it can also bind anionic DNA due to electrostatic interactions. To

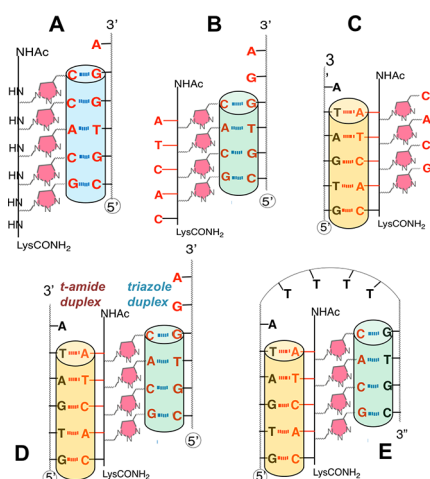


Figure 3. *iso*-PNA 1 and *bm*-PNA 2 duplexes: (A) *iso*-PNA 1: DNA 1; (B) *bm*-PNA 2:DNA 1; (C) DNA 2:*bm*-PNA 2; (D) DNA 2:*bm*-PNA 1:DNA 2 open double duplex and (E) DNA 3_{bp}:*bm*-PNA 2 hairpin double duplexes.

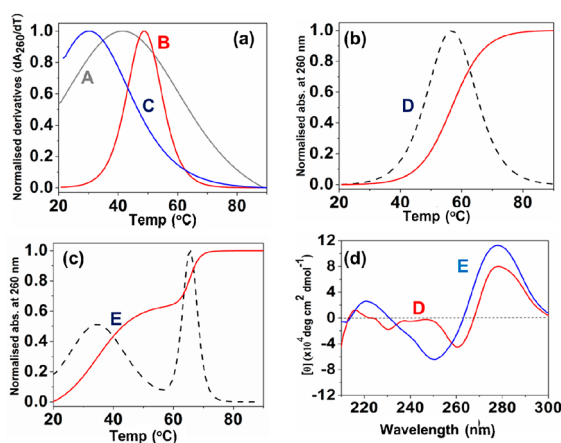


Figure 4. UV-melting plots and CD of duplexes. (a) A, *iso*-PNA 1:DNA 1; B *bm*-PNA 2:DNA 1; and C DNA 2:*bm*-PNA 2. (b) D, DNA 2:*bm*-PNA 2: DNA 1 and (c) E DNA 3_{bp}:*bm*-PNA 2 (red line represents normalized melting curves and black dashed line represents first derivative plots). (d) CD profiles of duplexes D and E.

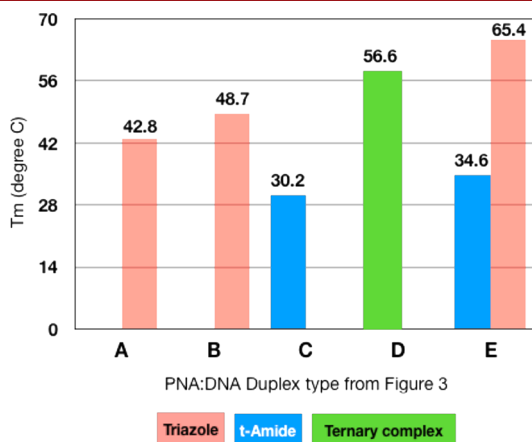


Figure 5. T_m s for various bimodal PNA:DNA duplexes and double duplexes, A–E correspond to complexes as in Figure 3. Values accurate to ± 1.0 °C. *iso*-PNA 1:DNA 1 T_m = 24.7 °C, Mismatch DNA sequence: DNA 1_m = 5'-CGCGGA-3'.

test the sequence specificity of *iso*-PNA 1 binding to DNA 1, its complementation was done with DNA 1_m (see legend in Figure 5) to yield duplex *iso*-PNA 1:DNA 1_m, which has a C-T mismatch and exhibited a significantly lower T_m (24.7 °C). This suggests that *iso*-PNA 1:DNA 2 duplex arises from specific base pairing and not mere electrostatic interactions.

The *bm*-PNA 2 formed corresponding triazole duplex *bm*-PNA 2:DNA 1 (Figure 3B) with T_m of 48.7 °C (Figure 5B) that is slightly higher than that of *iso*-PNA 1:DNA 1 triazole duplex. *bm*-PNA 2 also gave t-amide face duplex DNA 2:*bm*-PNA 2 (Figure 3C) with T_m = 30.2 °C (Figure 5C), which is lower than that of its triazole duplex. Thus, the designed *bm*-PNA 2 forms complementary base-paired duplexes independently from both triazole and t-amide faces. The triazole face duplexes A and B (Figure 3) result from new isomeric bimodal PNA structure, in which the nucleobases are attached at C_{4'} quite different from that of standard PNA.

The designed bimodal PNA *bm*-PNA 2, in the presence of DNA 1 and DNA 2 yielded the ternary complex DNA 2:*bm*-PNA 2:DNA 1 (Figure 3D) made from two duplexes. This complex showed a single melting transition (Figure 4b) with T_m of 56.6 °C, which is significantly higher than the constituent isolated duplexes DNA 2:*bm*-PNA 2 and *bm*-PNA 2:DNA 1 (Figure 5). The double duplex does not show a sequential melting (as in DNA triplexes¹⁶), but displays simultaneous dissociation of both DNA strands from *bm*-PNA 2 similar to PNA₂:DNA triplex.^{2,17} The bimodal PNA backbone being common to both duplexes couples the unwinding of one duplex to influence that of other duplex, causing melting of both duplexes to be synchronous and not sequential. The coexistence of duplexes with common backbone in a single complex strengthens their mutual stability, impacting dissociation of both duplexes. These results provide proof of concept that as per the design, bimodal PNA concomitantly binds to two DNA strands that are complementary to triazole and t-amide face.

Creation of double duplex from *bm*-PNA 2, DNA 1 and DNA 2 (Figure 3D) is a termolecular reaction requiring entropic cost. The two complementary DNA segments were connected by a tetranucleotide TTTT yielding single DNA 3_{bp}. This can now bind both sides to yield hairpin double duplex DNA 3_{bp}:*bm*-PNA 2 (Figure 3E) as a binary complex. This hairpin double duplex interestingly displayed two well-resolved transitions with T_m values of 34.6 and 65.4 °C (Figure 4c), in contrast to open double duplex that showed a single transition. The two transitions in hairpin duplex 3E arise either from sequential melting of t-amide and triazole duplex, or the early transition from a premelting conformational change, followed by simultaneous disassociation of both strands. Notably, both observed transitions in hairpin duplex 3E are stabilized, compared to isolated duplexes 3B and 3C and that of ternary complex 3D. Note that in isolated duplexes and open double duplex 3D, both duplexes are in antiparallel orientation. In hairpin double duplexes, the triazole duplex segment is in parallel orientation and hence stabilities are not directly comparable. The PNA:DNA transitions are generally known to be broad due to slow association–disassociation kinetics,¹⁴ and in bimodal PNA duplex and double duplexes, they are even more broad.

The CD spectra of open double duplex D and hairpin double duplex E are shown in Figure 4d. The profiles resemble that observed for PNA:DNA duplexes,¹⁸ indicating similar overall conformational features. Both D and E show a positive

Table 1. Thermodynamic Parameters from ITC^a

duplex	<i>bm</i> -PNA:DNA	−Δ <i>G</i> (kcal/mol)	−Δ <i>H</i> (kcal/mol)	−Δ <i>S</i> (cal/(mol K))	<i>K</i> _D (× 10 ^{−6} M)
B	<i>bm</i> -PNA 2:DNA 1 (triazole)	6.67	33.7	94.0	8.78
C	DNA 2: <i>bm</i> -PNA 2 (t-amide)	6.52	26.1	68.0	11.4
D	DNA 2: <i>bm</i> -PNA 2:DNA 1	6.22	37.9	109.7	19.5
E	DNA 3 _{hp} : <i>bm</i> -PNA 2	7.49	24.8	60.0	2.11

^aFor experimental details, see Figures S10–S12 in the Supporting Information.

CD band at ~280 nm, with a slight difference in negative band position: 260 nm for the open duplex **D** and 250 nm for hairpin duplex **E**. The CD band intensities were also higher for hairpin duplex **E**, compared to open duplex **D**, suggesting a better stacking in the hairpin duplex. The temperature-dependent CD of hairpin double duplex **E** (Figure S9 in the Supporting Information) showed slight shifts in maximum at 280 nm, accompanied by decreased intensity with an isosbestic point at ~265 nm, indicating alteration in conformation during melting. The stacking of triazole rings concomitant with base pairing may provide additional stability in bimodal PNA duplexes. The results establish formation of double duplexes from *bm*-PNA 2 and biophysical mechanisms of melting of bimodal PNAs need further study.

The thermodynamic parameters for formation of duplexes and double duplexes in bimodal PNAs were sought from ITC (see Table 1, as well as Figures S10–S12 in the Supporting Information). The experiment with bimodal PNA double duplex **D** was performed by titrating an equimolar mixture (in terms of the number of bases/binding sites) of DNA 1 and DNA 2 into *bm*-PNA 2. The dissociation constants (*K*_D) indicated the binding of triazole duplex **B** to be slightly stronger than t-amide duplex **C**, but association is entropically less favored (more negative). The stacking of triazole rings concomitant with canonical base pairing contributes to higher enthalpy, providing additional stability in the formation of *bm*-PNA 2 duplex **B**.

Formation of termolecular double duplex **D**, is accompanied by an entropic cost (more negative entropy), but offset by having a higher enthalpic contribution. Notably, the hairpin duplex **E** is most favored in terms of lowest *K*_D and lowest (−Δ*S*) and highest (−Δ*G*) among different complexes. The negative Δ*G* for bimodal PNA:DNA duplexes in the range of −6.0 kcal/mol to −7.0 kcal/mol was similar to that reported for PNA:DNA duplexes.¹⁹

In summary, this manuscript presents new concept of bimodal PNA that can concurrently hybridize with two cDNAs. In the isomorphous PNA backbone *iso*-PNA 1, nucleobases anchoring on C_α(S) glycol side chain successfully form triazole DNA duplex. The base pairing in this new isomorphous PNA duplex is likely to be Watson–Crick (WC) type, since it shows sequence specificity with lower *T*_m in mismatched duplex. The bimodal *bm*-PNA 2 designed to simultaneously bind two complementary DNAs forms open and hairpin double duplexes from t-amide and triazole faces. These coexist in a ternary complex with stability higher than the individual binary duplexes. The double duplex dissociates in a single step similar to PNA₂:DNA triplexes,¹⁷ while the hairpin double duplex shows distinct premelting conformational change before disassociation.¹⁶ This arises from the fact that a single common backbone hosts two duplexes. Based on choice of sequences (polypurines/polypyrimidines/self-complementary) on either side, bimodal PNAs can lead to fused duplexes, triplexes, and tetraplexes (G_n/C_n) with cDNA. They

can be engineered to yield supramolecular functional nanostructures for probable applications in PNA material science.²⁰ Distant biotechnological applications include simultaneous targeting of two genes or micro RNA structures²¹ for therapeutic purposes, and, as equivalents of short DNA staples, they can be active in programmed folding of plasmids in DNA origami.²²

■ ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acs.orglett.0c01950>.

Synthesis procedures, characterization data (NMR, MS of all new compounds, HPLC, MALDI TOF of PNA oligomers), UV-melting curves, CD spectra, ITC data (PDF)

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Notes

The authors declare no competing financial interest.

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