



The overview of PNAs

- Discovered in 1991 by Egholm, Nielsen, Berg, and Buchardt
- Synthesis chemistry commercialized in 1993
- More than 1,000 PNA publications in the fields of chemistry, molecular biology, diagnostics, and therapeutics
- A variety of products are on the market today
- A lot of sequence limitations in PNA oligomer synthesis
- Expensive
- PNA comes of age

Characteristics of PNA

- Higher affinity to complementary nucleic acid (DNA, RNA)
- Strong hybridization independent of salt concentration
- Greater specificity and sensitivity of interaction
- Thermal and chemical stability
- Resistance to nucleases and proteases
- Poor cellular uptake

Higher affinity to complementary nucleic acid (DNA, RNA)

			a,t,g,c : PNA A, T, G, C : DNA		
Sequence	T _m [°C]		Buffer		
	DNA	RNA			
H-tgt acg tca caa cta	69.5	72.3	Α		
5'-TGTACGTCACAACTA	53.3	50.6	Α		
H-aca tca tgg tcg	58.7	62.3	В		
5'-ACATCATGGTCG	47.9	44.5	В		
5'-ACATCA tgg tcg	52.6	52.1	В		

 $\begin{array}{l} \mbox{Buffer A: 100 mM NaCl, 10 mM NaH_2PO_4, 0.1 M EDTA, pH 7.0 \\ \mbox{Buffer B: 140 mM KCl, 10 mM NaH_2PO_4, 0.1 M EDTA, pH 7.4 \\ \end{array}$

	bility 15-mer PNA	vs. DNA
NaCI [mM]	PNA/DNA T _m (°C)	DNA/DNA T _m (°C)
0	72	38
100	69	54
140	69	56
1000	65	65
1	0 mM phosphate buffer,	0.1 M EDTA, pH = 7.0
ater specificity	and sensitivity of	of interaction
T _m for single mi	smatch	
15-mer PNA /	DNA 8 ~ 20 °	C

	DNA	PNA
Hybridization affinity with DNA		At least 1 °C higher per base
Hybridization rate with DNA		100 - 5000 times
Salt concentration for hybridization	Dependent	Independent
T _m single mismatch	Lowering 10 °C	Lowering 15 °C
Chemical stability	Unstable in acid and base	Stable
Water solubility	Soluble	Restricted solubility
Maximum base length	No limit	18 bases: aggregation
Required base length for diagnosis	20 - 30	13 - 17
Biological stability	Degradation in nuclease	Stable
Thermal stability	Moderate	Good















	PNA		DNA antiparallel		DNA parallel	
	Seque	Sequence		(2 <i>R</i> ,4S)	(2S,4 <i>S</i>)	(2R,4S)
	I-GTA GAT CAC T-NH(CH ₂) ₂ COOH		43.8		40.3	
	1-GTa GAT CAC T-NH(CH ₂) ₂ COOH (1)		53.8	53.1	50.2	50.2
	H-GTA GAt CAC T-N	-GTA GAt CAC T-NH(CH ₂) ₂ COOH (2)		33.0	34.0	26.2
	H-GTA gAT CAC T-N	-GTA gAT CAC T-NH(CH ₂) ₂ COOH (3)		53.1	58.3	28.3
	H-GTA GAT CAC T-NH(CH ₂) ₂ COOH (4)		55.2	62.3	31.2	34.0
	a, t, g,	and c represent (2S/	R, 4S)-aep	PNA unit		
	T _m (°C) of Mism	atched PNA/DNA	Duplex			
	DNA (5'-3') aegPNA		(2S,4S)-aepPNA		4	
	AGT GAT CCA C	GT GAT CCA C 35.4 (-8.4) (1)		44.4 (-9.4)		
	AGT G <i>T</i> T CTA C	36.8 (-7.0) (2)	26.9	-29.7)		
	AGT GAT ATA C	39.6 (-4.2) (3) 36.8 (-7.0) (4)	24.7	-18.5) -11.6)		













































