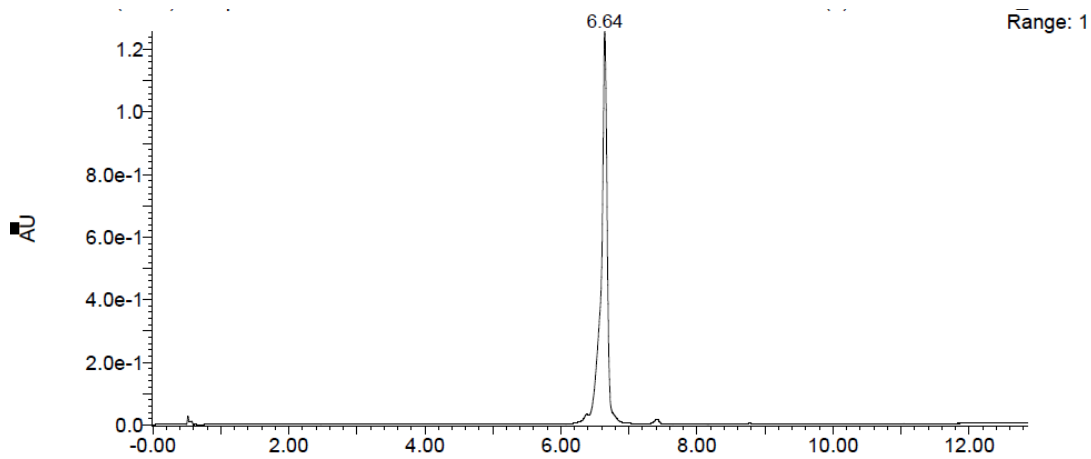


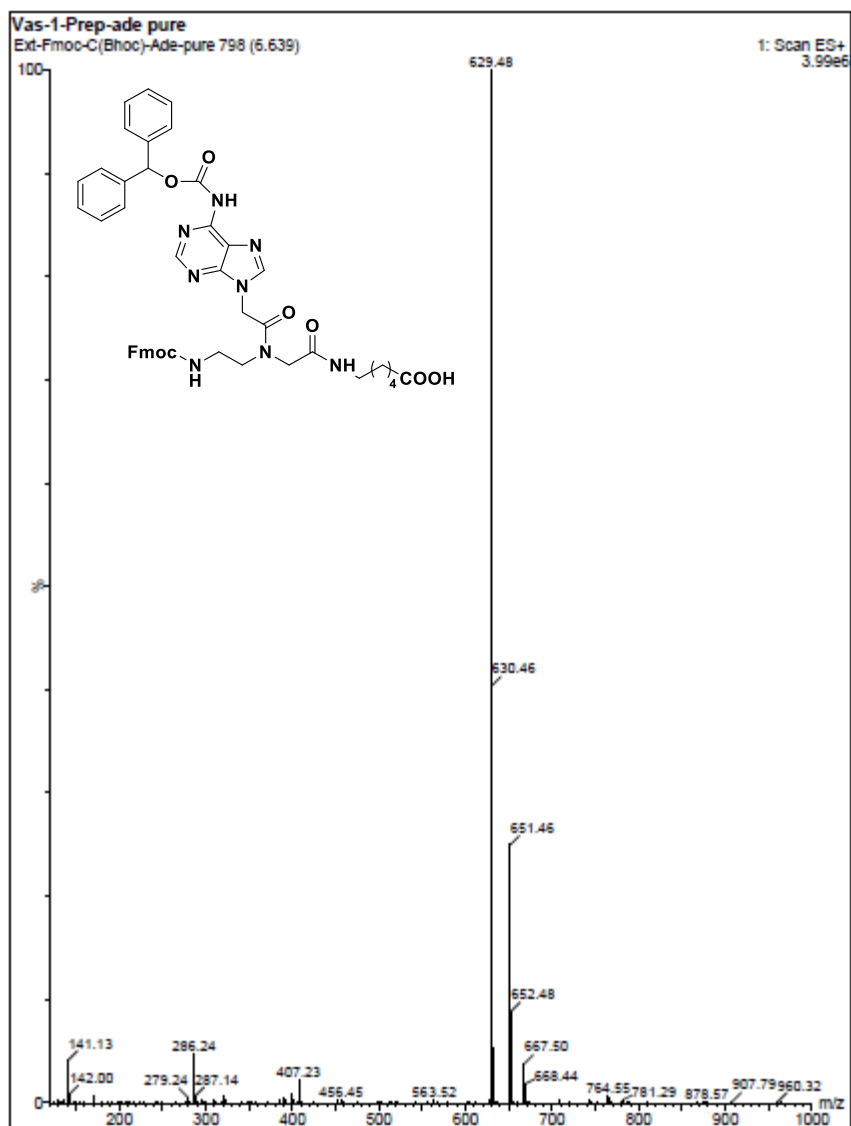
1 SUPPLEMENTARY INFORMATION

2  
3 **Mechanically Rigid Supramolecular Assemblies Formed from an Fmoc-Guanine**  
4 **Conjugated Peptide Nucleic Acid**

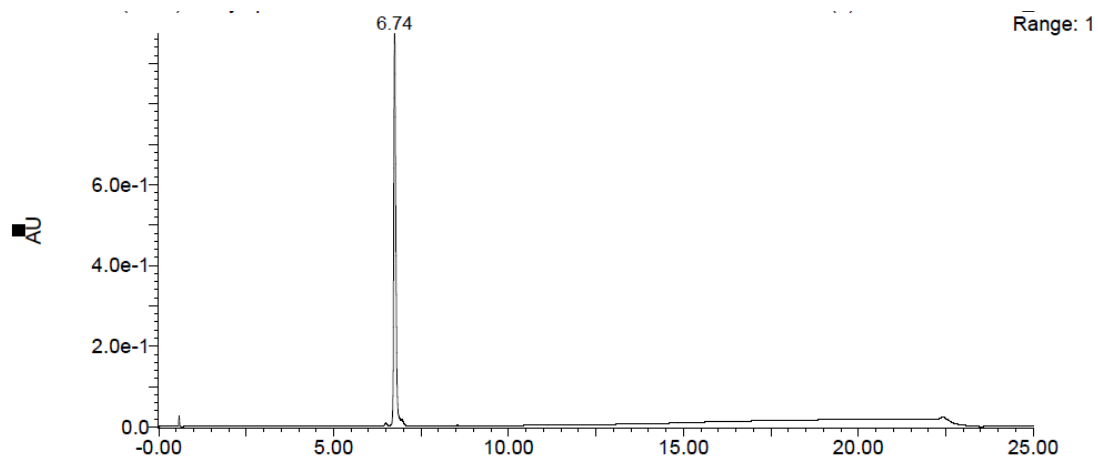
5  
6 *Vasantha Basavalingappa<sup>1</sup>, Santu Bera<sup>1</sup>, Bin Xue<sup>2</sup>, Ido Azuri<sup>3</sup>, Yiming Tang<sup>4</sup>, Kai Tao<sup>1</sup>, Linda*  
7 *J. W. Shimo<sup>5</sup>, Michael R. Sawaya<sup>6</sup>, Sofiya Kolusheva<sup>7</sup>, David S. Eisenberg<sup>6</sup>, Leeor Kronik<sup>3</sup>, Yi*  
8 *Cao<sup>2</sup>, Guanghong Wei<sup>4</sup>, Ehud Gazit\*<sup>1</sup>*



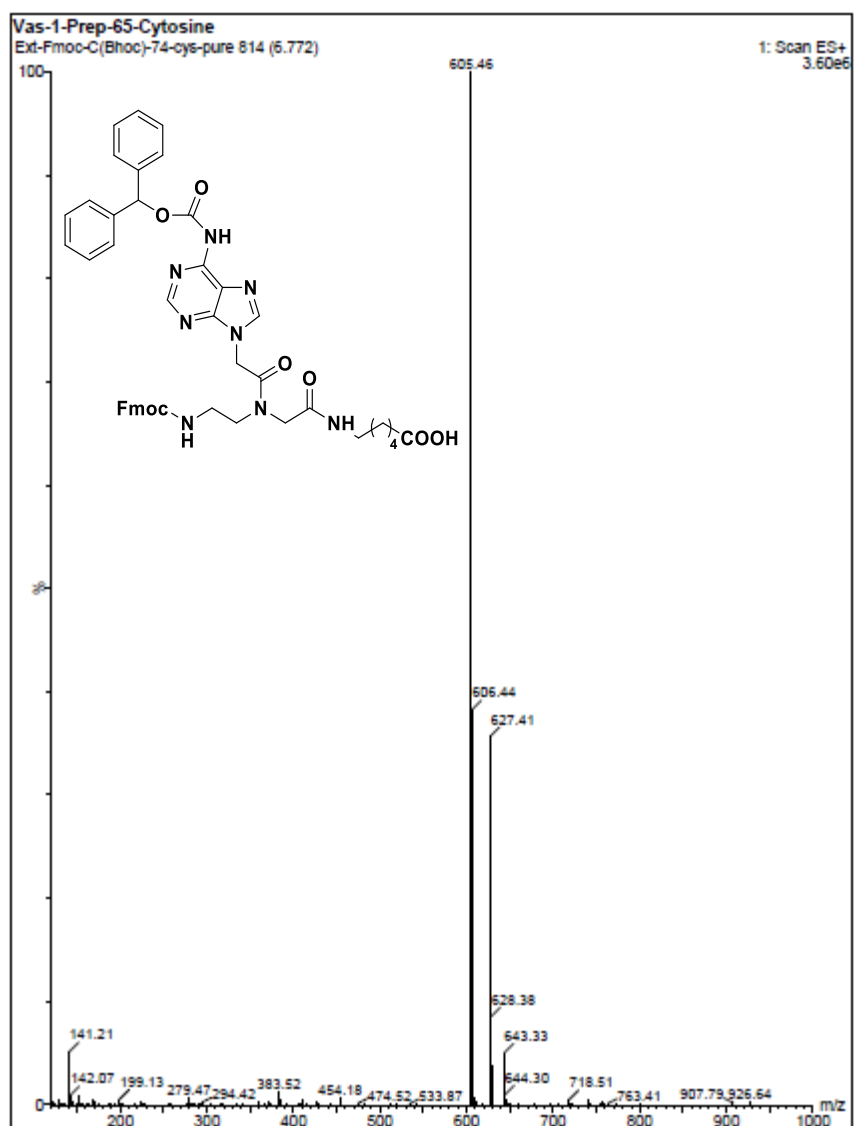
Supplementary Figure 1. LC profile of the pure adenine PNA-derivative.



Supplementary Figure 2. Mass spectrum of the pure adenine PNA-derivative.

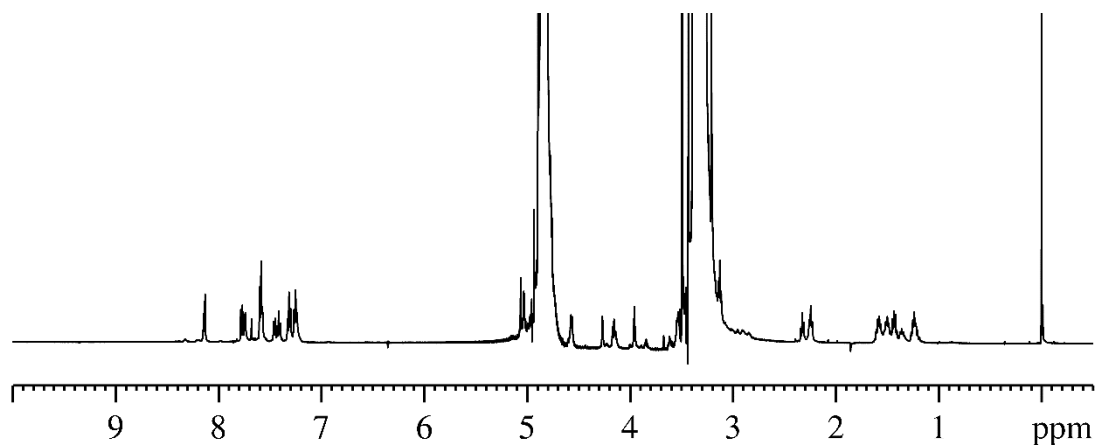


3 **Supplementary Figure 3.** LC profile of the pure cytosine PNA-derivative.



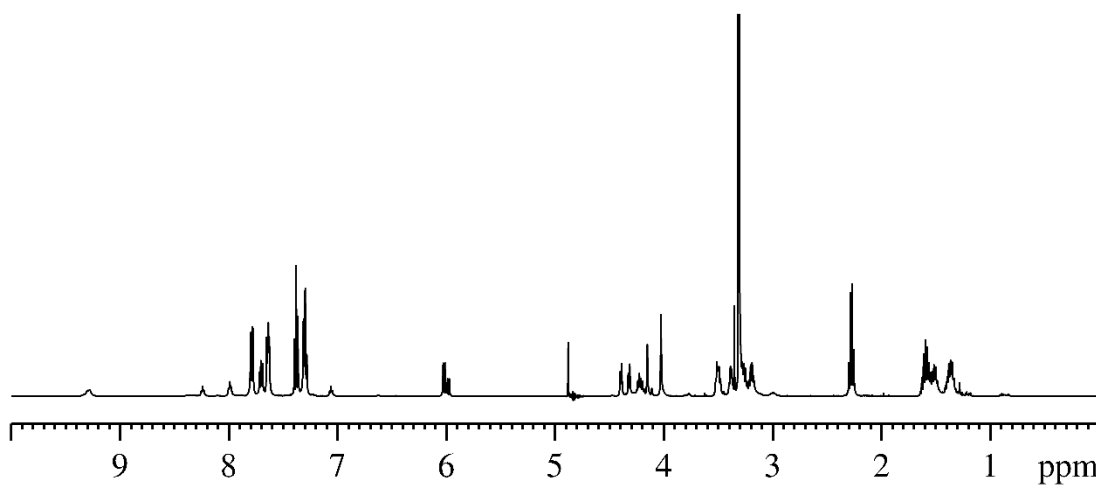
23 **Supplementary Figure 4.** Mass spectrum of the pure cytosine PNA-derivative.

24



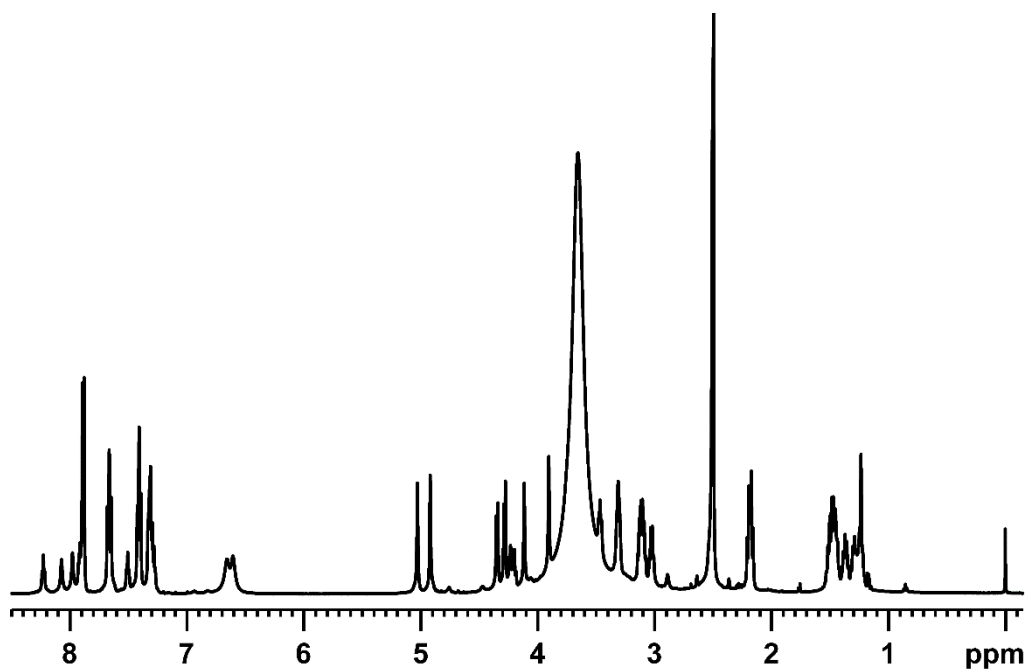
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2 **Supplementary Figure 5.** A 500 MHz  $^1\text{H}$  NMR spectrum of the adenine PNA-derivative in  
3 DMSO- $\text{d}_6$  and MeOH.



4

5 **Supplementary Figure 6.** A 500 MHz  $^1\text{H}$  NMR spectrum of the cytosine PNA-derivative in  
6 MeOH- $\text{d}_3$ .

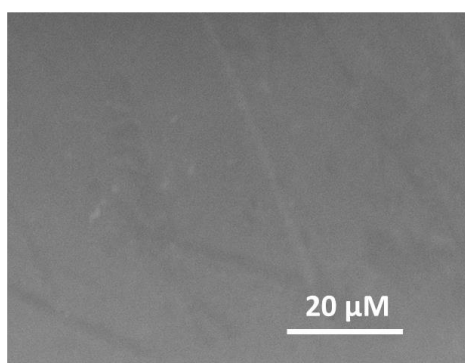


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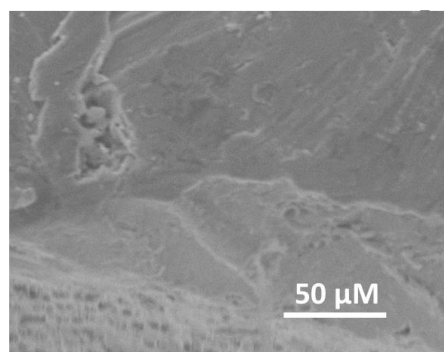
2 **Supplementary Figure 7.** A 500 MHz  $^1\text{H}$  NMR spectrum of the PNA-quadruplex in  
3 DMSO- $\text{d}_6$ .

4

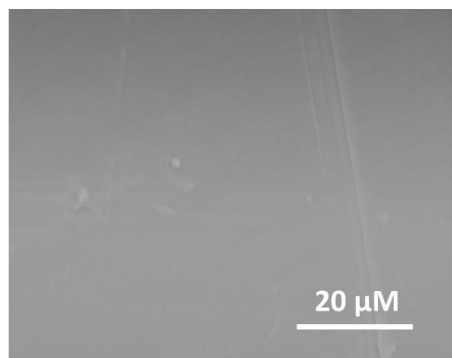
a)



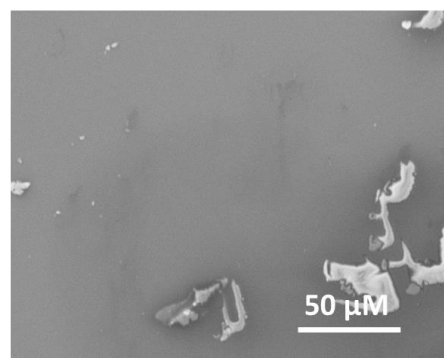
b)



c)

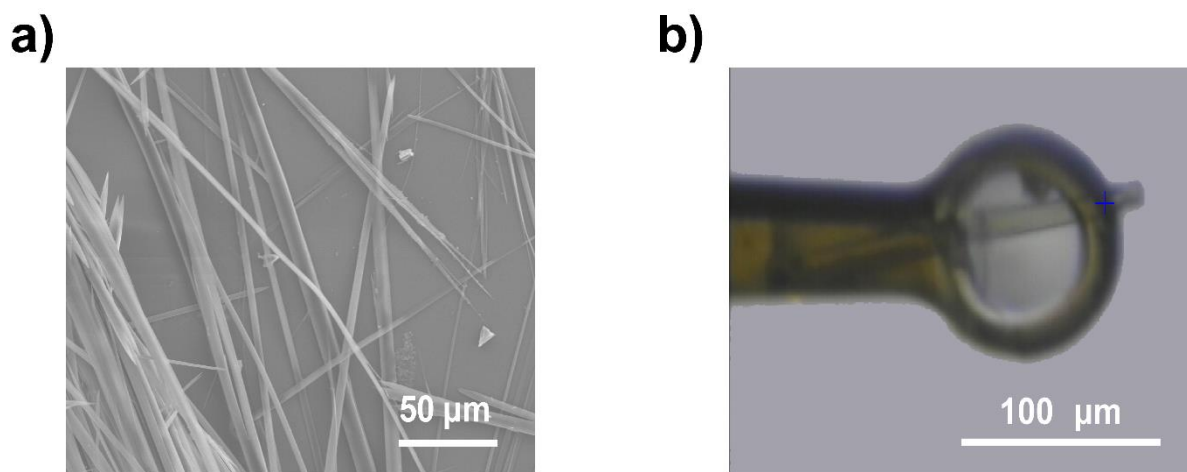


d)

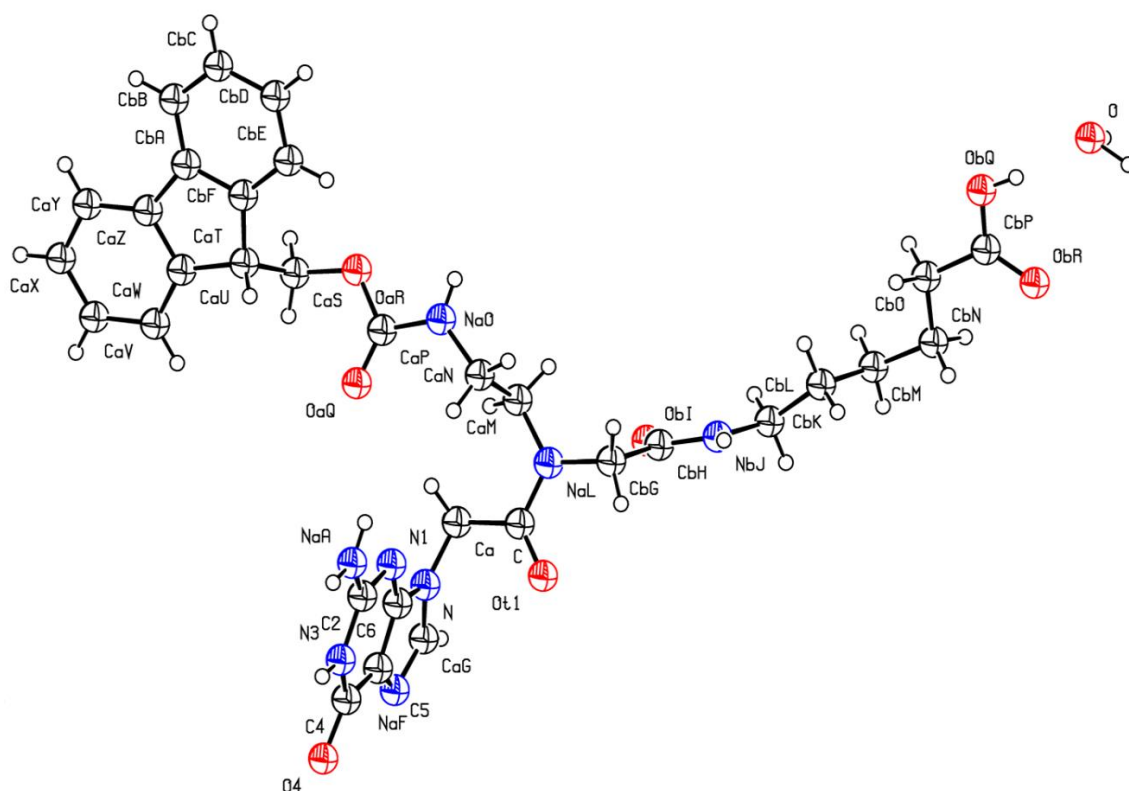


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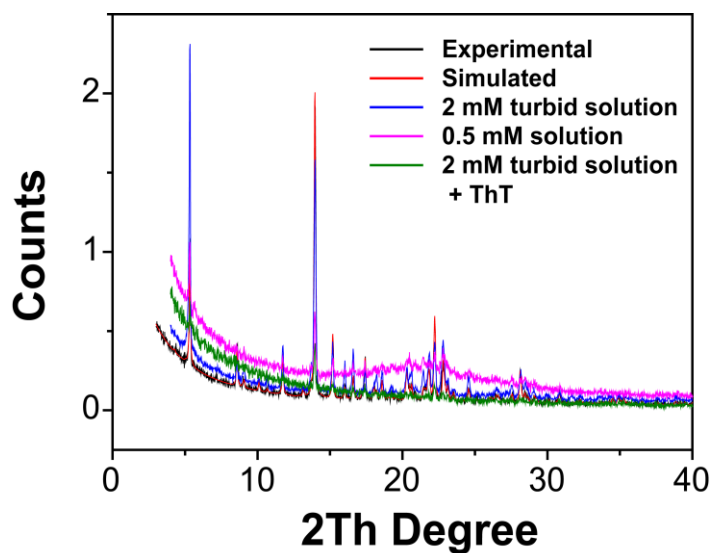
1 **Supplementary Figure 8.** Morphological characterization of adenine and cytosine PNA-  
2 derivatives. **(a, b)** SEM images of the adenine PNA-derivative at **(a)** 0.1 and **(b)** 0.5 mg/mL.  
3 **(c, d)** SEM images of the cytosine PNA-derivative at **(c)** 0.1 and **(d)** 0.5 mg/mL.



4  
5 **Supplementary Figure 9.** Images of the PNA-quadruplex crystals. **(a)** SEM image. **(b)** View  
6 of the PNA-quadruplex crystal mounted and used for data collection.

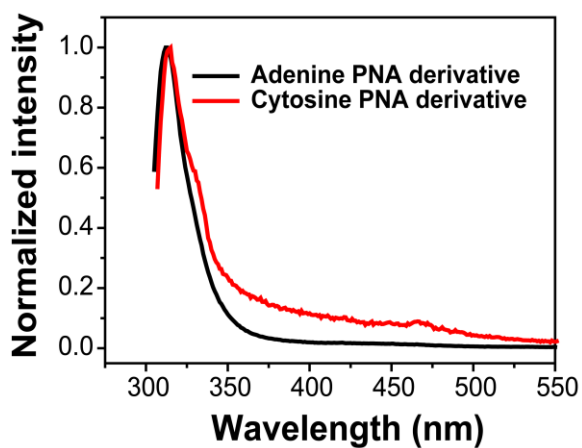


7  
8 **Supplementary Figure 10.** ORTEP diagram of the PNA derivative in 50% probability  
9 ellipsoids, obtained from single crystal X-ray structure.



1

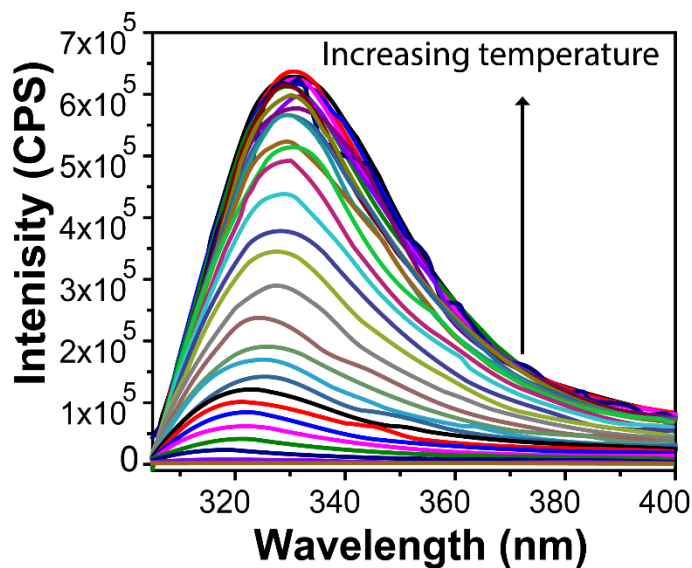
2 **Supplementary Figure 11.** Comparison of PXRD profiles of various self-assembled structural  
 3 forms of PNA-quadruplex.



4

5 **Supplementary Figure 12.** Fluorescence emission spectra for adenine and cytosine PNA-  
 6 derivative at 2 mM.

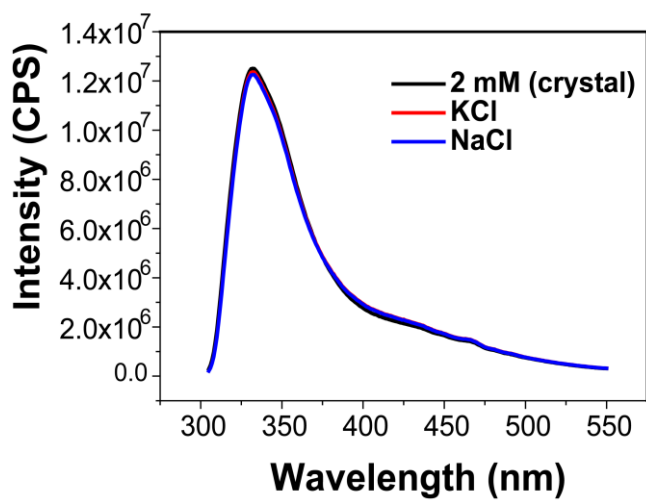
7



1

2 **Supplementary Figure 13.** Temperature dependence of the fluorescence emission peak at 330  
 3 nm, obtained from the PNA-quadruplex solution upon excitation at  $\lambda_{\text{ex}}=290$  nm.

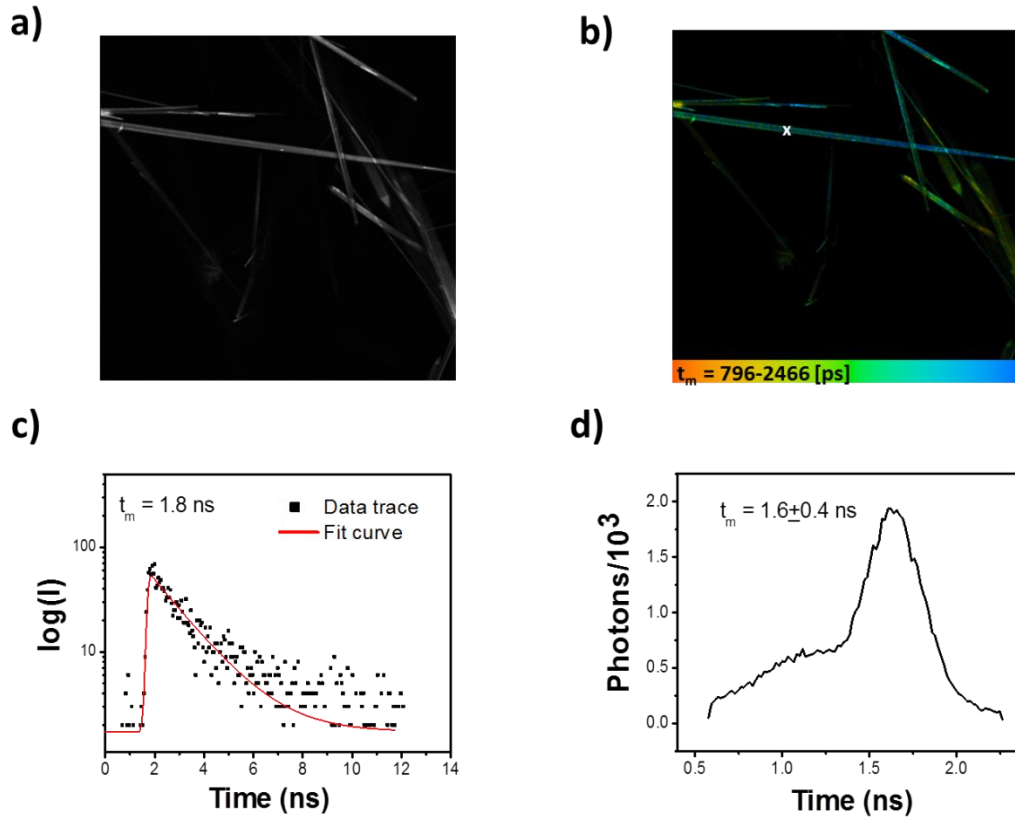
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5

6 **Supplementary Figure 14.** Fluorescence emission spectra of PNA-quadruplex in the presence  
 7 of 100 mM Na<sup>+</sup> or K<sup>+</sup> ions.

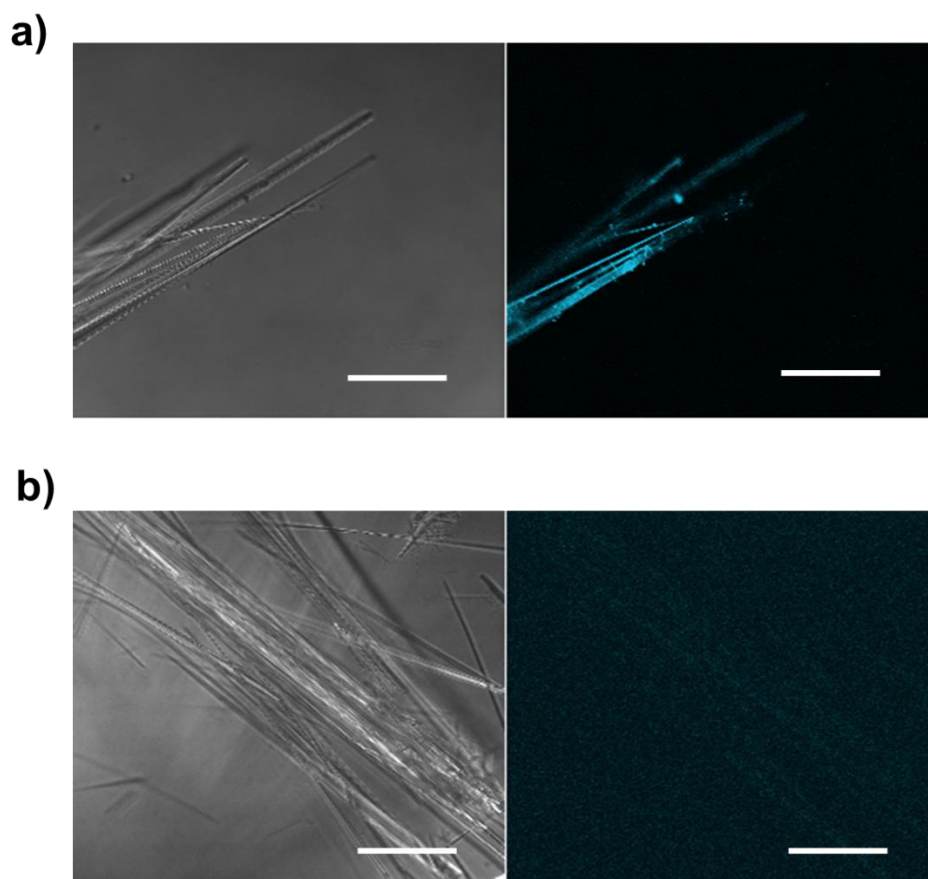




1

2 **Supplementary Figure 15.** FLIM characterization of PNA-quadruplex. (a) Bright field and  
 3 (b) Fluorescence image of the PNA-quadruplex crystals. (c) Representative fluorescent  
 4 emission decay curve at the white point in the FLIM image, showing a lifetime of  $\sim 2.0$  ns. (d)  
 5 Statistical fluorescence lifetime distribution of the crystals.

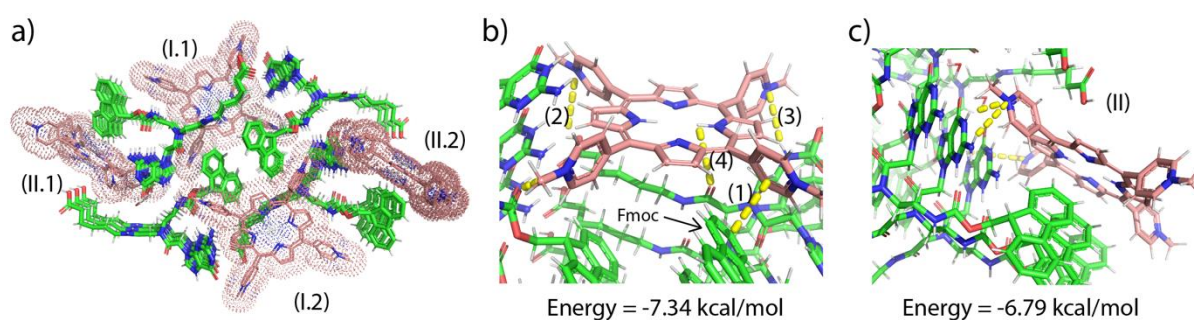
6



1

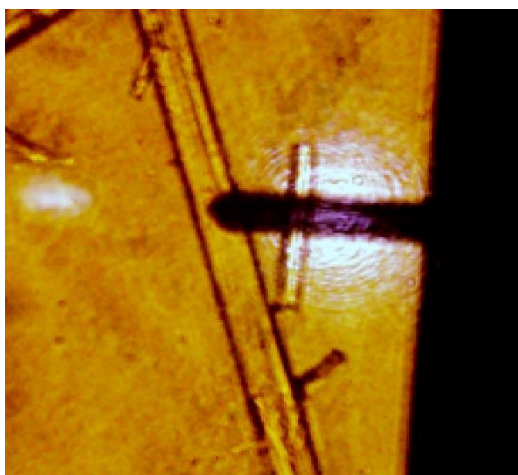
2 **Supplementary Figure 16.** Confocal fluorescence microscopy images of the PNA-quadruplex  
 3 (a) with ThT, (b) without ThT. (left panel: bright field; right panel: fluorescent image,  $\lambda_{\text{exc}} =$   
 4 458 nm, scale bar 50  $\mu\text{M}$ ).

5



6

7 **Supplementary Figure 17.** Results of molecular docking of TMPyP4 to quadruplex inspired  
 8 PNA-assemblies. (a) Binding sites of the first thirty energy-optimized docking pattern. The  
 9 conformations are clustered into two groups: I.1 and I.2 are equivalent due to the crystal  
 10 symmetry, so are II.1 and II.2. (b-c) Interactions responsible for molecule binding in the first  
 11 (b) and the second (c) conformation clusters.



1

2 **Supplementary Figure 18.** Typical IT-AFM image. The crystals were fixed to the surface of  
 3 the quartz substrate and the cantilever was moved to the crystal surface while monitoring under  
 4 an optical microscope.

5 **Supplementary Table 1.** Crystallographic and computed lattice parameters for the PNA-  
 6 quadruplex crystals, in Å. Values in parentheses indicate the relative error in the computed  
 7 lattice parameters, with respect to experiment.

	$\vec{a}$	$\vec{b}$	$\vec{c}$
Experimental	4.74	32.86	20.48
TS-vdW DFT	4.66 (-1.7%)	32.63 (-0.7%)	20.33 (-0.7%)

8

9 **Supplementary Details 1. Calculation of elastic constant tensor.** 6 different distortions were  
 10 applied to the unit cell, where, 1-6 are xx, yy, zz, yz, xz, and xy in the Voigt notation. These  
 11 are:

$$\begin{aligned}
 & \begin{pmatrix} 1 + \delta_1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}, \quad \begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 + \delta_2 & 0 \\ 0 & 0 & 1 \end{pmatrix}, \quad \begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 + \delta_3 \end{pmatrix}, \quad \begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & \delta_4 \\ 0 & 0 & 1 \end{pmatrix}, \quad \begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ \delta_5 & 0 & 1 \end{pmatrix}, \\
 & \begin{pmatrix} 1 & \delta_6 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}
 \end{aligned}$$

13

14 For each distortion, 5 strain ( $\delta$ ) values (0,  $\pm 0.005$ ,  $\pm 0.01$ ) were applied to calculate stress. The  
 15 elastic tensor was obtained from numerical fitting of the obtained stress-strain curves.

16

1 **Supplementary Table 2.** PNA-quadruplex data collection and refinement statistics.

<b>Data collection</b>	<b>PNA-quadruplex</b>
Space group	P2 <sub>1</sub> /c
Cell dimensions	
<i>a</i> , <i>b</i> , <i>c</i> (Å)	4.74, 32.9, 20.5
<i>α</i> , <i>β</i> , <i>γ</i> (°)	90.0, 91.9, 90.0
Resolution (Å)	0.80 (0.82-0.80)
<i>R</i> <sub>sym</sub>	0.048 (0.139)
<i>I</i> / <i>σI</i>	17.2 (5.0)
CC <sub>1/2</sub>	99.9 (98.8)
Completeness (%)	89.4 (72.9)
Redundancy	4.6 (3.1)
<b>Refinement</b>	
Resolution (Å)	0.80
No. reflections	5271
<i>R</i> <sub>work</sub> / <i>R</i> <sub>free</sub>	0.080/0.085
No. atoms	
PNA-2 (including hydrogen)	83
Water	1
B-factors (Å <sup>2</sup> )	
PNA-2	4.6
Water	5.7
R.m.s deviations	
Bond lengths (Å)	0.022
Bond angles (°)	2.2

2 \*Highest resolution shell is shown in parenthesis.