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Supplementary Information (ESI)

Enhanced binding of 5,10,15,20 (N-Propyl-4-Pyridyl) Porphyrine (TPrPyP4): a comparative study with TMPyP4

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Materials and Methods

Oligonucleotides and porphyrins

The oligonucleotides were obtained from Microsynth (Switzerland). They were purified by PAGE using a denaturing 20% gel (acrylamide: bisacrylamide, 19:1) in TBE, 7 M urea, 55°C. The bands were excised from the gel and eluted in water. The DNA solutions were filtered (Ultrafree-DA, Millipore) and precipitated. The concentration of each DNA was determined from the absorbance at 260 nm in milli Q water, using as extinction coefficients 7500, 8500, 15000 and 12500 M⁻¹cm⁻¹ for C, T, A and G, respectively. Porphyrins TMPyP2 (P2), TMPyP3 (P3) were purchased from Porphyrin Systems (Lübeck, Germany), TMPyP4 (P4) from Sigma (Milan, Italy).

Polymerase stop assay

Single-stranded DNA fragments with a number of nt between 79-82, containing in the middle a quadruplex forming G-rich element from the murine *KRAS* or human *HRAS* promoters, were used as templates in the Taq polymerase primer-extension reactions. The DNA sequences have been purified by PAGE under denaturing conditions. The template (25 nM) was mixed with the ³²P-labelled primer (25 nM), in the presence or absence of porphyrins (P2, P3, P4, PP4 or 4) or phthalocyanines (1-4), in 25 mM KCl, Taq buffer 1X and incubated overnight at 37°C. The primer extension reactions have been carried out for 1h, by adding 10 mM DTT, 100 mM dATP, dGTP, dTTP, dCTP and 3.75U of Taq polymerase (Euro Taq, Euroclone, Milan). The reactions were stopped by adding an equal volume of stop buffer (95% formamide, 10mM EDTA, 10mM NaOH, 0.1% xylene cyanol, 0.1% bromophenol blue). The products were separated on a 12% polyacrylamide sequencing gel prepared in TBE1X, 8 M urea. The gel was dried and exposed to autoradiography. Standard dideoxy sequencing reactions were performed to detect the exact positions in which DNA polymerase was arrested.





Figure S_1 : Putative structure of the G4-DNA formed by the murine GA-element, as determined by DMS-footoprinting (Nucleic Acids Res, 2006, 34, 2536).



Figure S₂: Structures of pentaphyrins TMPyP2 (P2) and TMPyP3 (P3).



Titration TMPyP4 (P4) with duplex d28R

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0,9 PP4_6 microM PP4_28Rdup_42nM PP4_28Rdup_126nM 0,8 PP4_28Rdup_164nM PP4 28Rdup 202nM 0,7 PP4_28Rdup_241nM PP4_28Rdup_279nM 0,6 PP4_28Rdup_318nM PP4_28Rdup_356nM PP4_28Rdup_414nM 0,5 PP4_28Rdup_471nM PP4 28Rdup 529nM 0,4 PP4_28Rdup_586nM PP4_28Rdup_644nM PP4_28Rdup_720nM 0,3 PP4_28Rdup_762nM PP4_28Rdup_804nM 0,2 PP4_28Rdup_871nM PP4 28Rdup 972nM 0,1 PP4_28Rdup_1098nM PP4_28Rdup_1266nM PP4_28Rdup_1518nM 0 -PP4_28Rdup_1854nM 300 350 400 450 500 PP4_28Rdup_2274nM

Titration TPrPyP4 (PP4) with duplex d28R

Figure S₃: UV-vis titrations of porphyrins 6 µM TMPyP4 and TPrPyP4 with duplex d28R [5'-GGGAGGGAGGGAGGGAGGGAGGGAGGGAGGGAGGGA-3'CCCTCCCTCCCTCCCTCCCTCCCTCCCTCCCT] in 50 mMTris-HCl, pH 7.4, 100 mM NaCl.



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Figure S₄: (A) Binding curves for P4 and PP4 on a logarithm scale showing that P4 has more affinity for duplex d28 than quadruplex q28R, whereas PP4 shows similar affinity for duplex and quadruplex; (B) Binding curve on a linear scale and best fitted to a 1:1 binding model. P4-d28: K_d =5.7x10⁻⁷±5.7x10⁻⁷; P4-q28R: K_d =1.5x10⁻⁶±5.0x10⁻⁸; PP4-d28: K_d =4.1x10⁻⁷±4.5x10⁻⁷; P4-q28R: K_d =6.4x10⁻⁷±6.6x10⁻⁸.



Figure S₅: Typical Scatchard plots for duplex d28R and porphyrins P4 and PP4. Titrations performed in 50 mM Tris-HCl pH 7.4, 100 mM KCl, quartz cuvette, 0.5 cm pathlength, 28R concentration 6 μ M.



Pentaphyrine

Supplementary Material (ESI) for Chemical Communications # This journal is (c) The Royal Society of Chemistry 2010 Figure S₆: (Up) Porphyrine (DIGPor) used in this study. [P Alzeer, N Roth, NW Luedtke. ChemComm 2009, 15, 1970; A Membrino, M Paramasivam, S Cogoi, J Alzeer, NW Luedtke, LE Xodo 2010, ChemComm 46, 625] ; (Down) Pentaphyrins used in this study [C Comuzzi, S Cogoi, M Overhand, GA Van der Marel, HS Overkleeft, LE Xodo. J Med Chem. 2006, 49, 196.