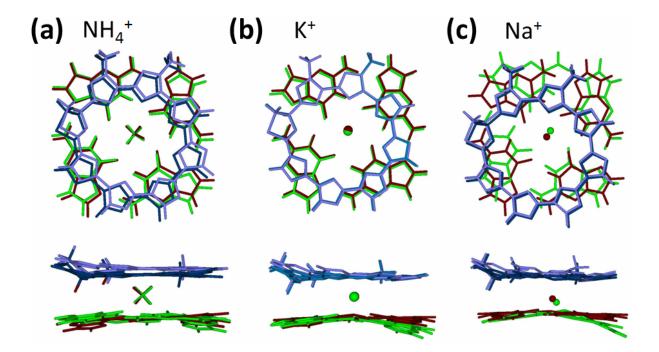
Cation Involvement in Telomestatin Binding to G-Quadruplex DNA

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Supporting information



Supporting Information Figure S1: Manual superimposition of optimized geometry of the ternary complexe telomestatin - cation - G-quartet obtained using B3LYP and MPWB1K functionals (top view and side view). (a) NH₄⁺, (b) K⁺, (c) Na⁺. The B3LYP optimized structures are in green (G-quartet) and purple (telomestatin). The MPWB1K optimized structures are in red (G-tetrad) and blue (telomestatin).

Supporting Information Table S1: Comparison of the geometries optimized with the different hybrid functional: root mean square distances (RMSD) of two-by-two comparisons (in \mathring{A}).

	Cation + telomestatin complex	Cation + telomestatin + G-quartet complex	
B3LYP vs BHHLYP			
NH ₄ ⁺	0.045	0.265	
K ⁺	0.045	0.131	
Na ⁺	0.000	0.856	
B3LYP vs MPWB1K			
NH ₄ ⁺	0.085	0.350	
K ⁺	0.083	0.276	
Na ⁺	0.069	0.726	
MPWB1K vs BHHLYP			
NH ₄ ⁺	0.054	0.302	
K ⁺	0.055	0.188	
Na ⁺	0.069	0.573	

Supporting Information Table S2: Interaction energies of NH_4^+ , K^+ and Na^+ with telomestatin alone, telomestatin + one G-quartet, computed with DFT, 6-31G(d,p) basis set, and three different hybrid functionals: B3LYP, BHandHLYP, and MPWB1K. The calculation of the interaction energy of the cation in the complex (BSSE corrected) has been calculated using the following equations:

$$\Delta E_{int} \!\!=\!\! E_{(telomestatin+cation)} - E_{(telomestatin)} - E_{(cation)} - E_{(BSSE)}$$

$$\Delta E_{int} \!\!=\!\! E_{(G4+telomestatin+cation)} - E_{(G4+telomestatin)} - E_{(cation)} - E_{(BSSE)}$$

	Cation interaction energy with telomestatin alone (kcal/mol)	Cation interaction energy with telomestatin + G-quartet (kcal/mol)		
B3LYP				
NH ₄ ⁺	-84.9	-115.8		
K ⁺	-82.7	-119.9		
Na ⁺	-93.8	-141.2		
BHandHLYP				
NH ₄ ⁺	-87.4	-120.7		
K ⁺	-86.0	-117.9		
Na ⁺	-95.4	-147.0		
MPWB1K				
NH ₄ ⁺	-88.1	-121.6		
K ⁺	-86.1	-124.3		
Na ⁺	-95.6	-145.6		