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TARGETING THE G-QUADRUPLEX FOLD OF NUCLEIC ACIDS BY DIFFERENT COMPUTATIONAL DRUG DESIGN APPROACHES

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The G-quadruplex fold is an unusual DNA and/or RNA conformation adopted in physiological and pathological conditions. The hyper-elongation of the human repeated telomeric sequence folding into this conformation is related to the hyperproliferation of neoplastic cells. Recently the rationale of blocking the elongation process has been proposed as new and selective mechanism to develop antitumor agents. The conformational properties of the Gquadruplex target will be discussed as well as the opportunity to carry out rational design of novel antitumor agents based on the stabilization of special this fold. Details about new different computational tools based on approaches developed in our laboratory will be described with some our G-guadruplex case studies.

Information and registration available at www.sardegnaricerche.it





