

Interaction of a Gold(I) Dicarbene Anticancer Drug with Human Telomeric DNA G-Quadruplex: Solution and Computationally Aided X-ray Diffraction Analysis

Federica Guarra,^{a†} Tiziano Marzo,^{a,b†} Marta Ferraroni,^c Francesco Papi,^c Carla Bazzicalupi,^{*c} Paola Gratteri,^{*d} Gennaro Pescitelli,^a Luigi Messori,^b Tarita Biver^a and Chiara Gabbiani^{*a}

[†]Equally contributed

^a*Department of Chemistry and Industrial Chemistry, University of Pisa, Via G. Moruzzi 13, 56124 Pisa, Italy. E-mail: chiara.gabbiani@unipi.it.*

^b*Laboratory of Metals in Medicine (MetMed), Department of Chemistry "U. Schiff", University of Florence, Via della Lastruccia 3, 50019 Sesto Fiorentino, Italy.*

^c*Department of Chemistry "U. Schiff", University of Florence, Via della Lastruccia 3, 50019 Sesto Fiorentino, Italy. E-mail: carla.bazzicalupi@unifi.it*

^d*Department NEUROFARBA – Pharmaceutical and nutraceutical section; Laboratory of Molecular Modeling Cheminformatics & QSAR, University of Firenze, via Ugo Schiff 6, 50019 Sesto Fiorentino, Firenze, Italy. E-mail: paola.gratteri@unifi.it.*

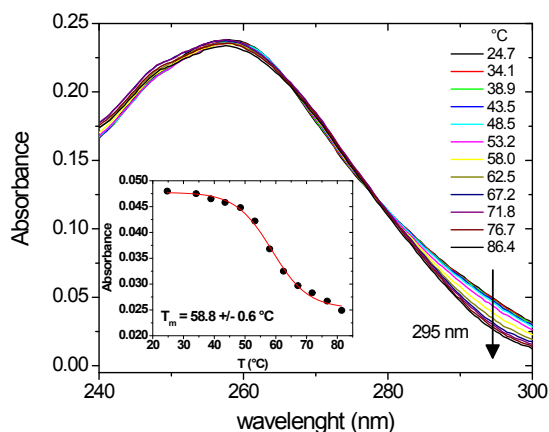


Figure S0. Example of variation of absorbance profile for the $[\text{Au}(\text{NHC})_2]^+/\text{Tel 23}$ system at different temperatures; $[\text{Tel23}] = 1.0 \mu\text{M}$, $r = [\text{Au}(\text{NHC})_2]^+/\text{Tel 23} = 5.0$, 50 mM potassium phosphate buffer, pH = 7.0, cell path 1.0 cm. Inset is the simoidal fit of the absorbance at 292 nm vs. temperature plot which provides the melting temperature, T_m .

Table S1. Data collection and data refinement about the crystal structure of the adduct formed by the complex $[\text{Au}(\text{NHC})_2]^+$ and the human telomeric sequence Tel24.

Data collection	
Wavelength	0.872900 Å
Cell parameters	$a = 36.600$ $b = 71.370$ Å $c = 27.050$ Å $\beta = 92.42^\circ$
Crystal system	monoclinic
Space Group	C 2
Resolution limits	35.62-1.6(1.70-1.60) Å
Number of reflections	48359 (5872)
Rsym	11.0 (265.5)
Multiplicity	5.4 (4.5)
Completeness (%)	97.6 (90.4)
$\langle I / \sigma(I) \rangle$	6.04(0.45)
CC (1/2)	99.2 (22.5)
Refinement	

Resolution range	35.62-2.00Å
Unique reflections (working/free)	4178/517
Rfactor	22.8%
Rfree factor	25.3%
Non-hydrogen atoms	567
Drug atoms	21
Water molecules	39
RMSD bonds	0.004
RMSD angles	1.114

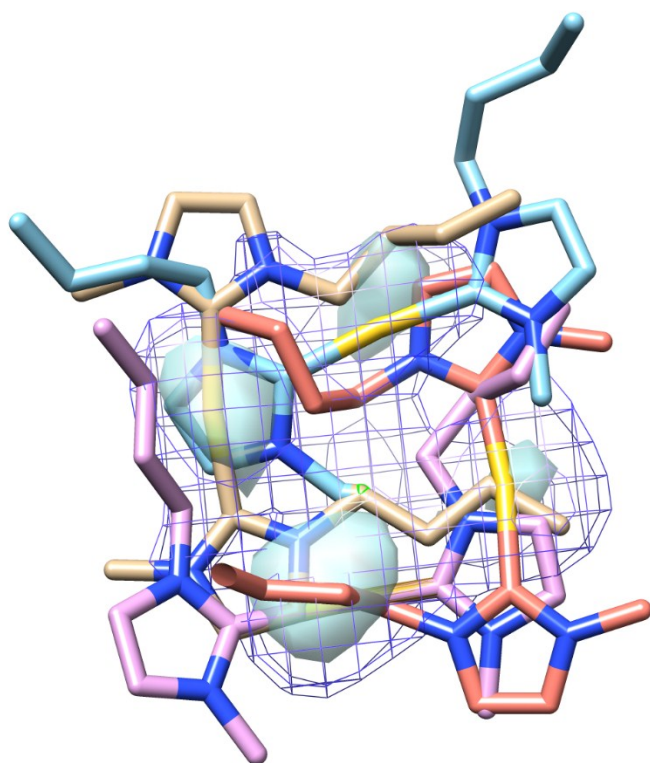


Figure S1 Skeleton of the disordered [Au(NHC)₂]⁺ complex, spread over four crystallographic positions. The following color code has been used to identify each single position: salmon (Au1), tan (Au2), light brown (Au3), light pink (Au4). OMIT electron density map (meshed) and anomalous Fourier difference map for the gold ions (continuous) contoured at 1.5 σ and 3 σ levels, respectively.