

Supplementary information for:

Synthesis and structure of a new thiazoline-based palladium(II) complex that promotes cytotoxicity and apoptosis of human promyelocytic leukemia HL-60 cells.

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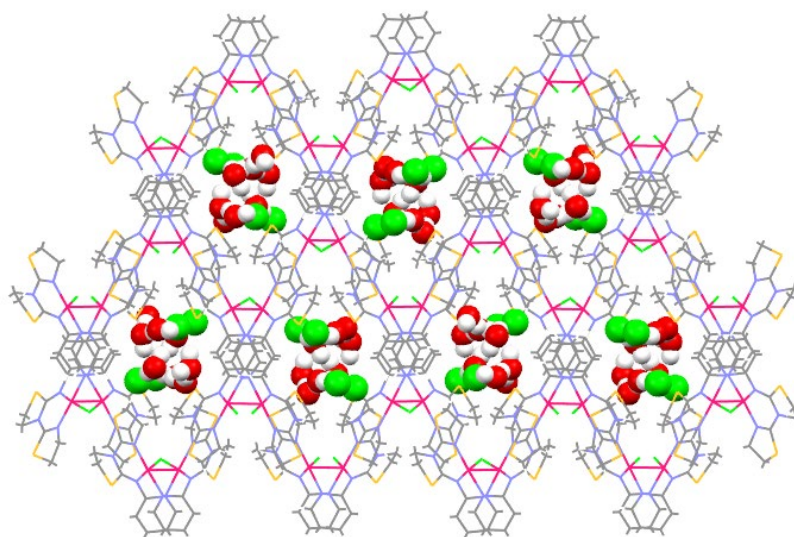


Figure S1. 3D supramolecular metal organic matrix along a-axis in PdPyTT.

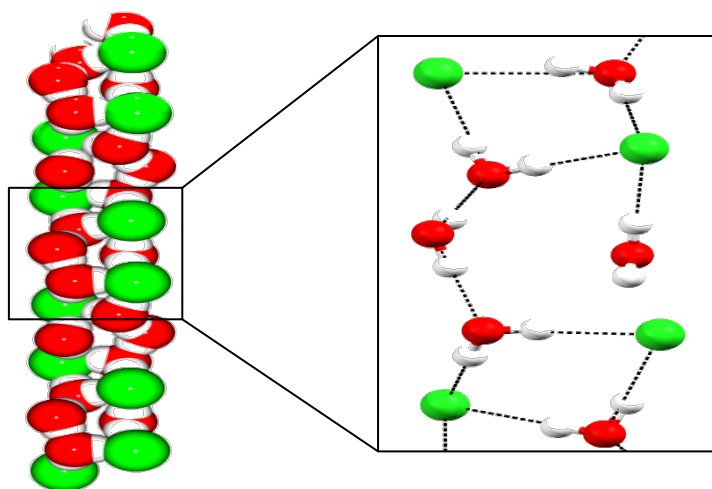


Figure S2. *Hydrogen bonds inside water-chloride chains in PdPyTT.*

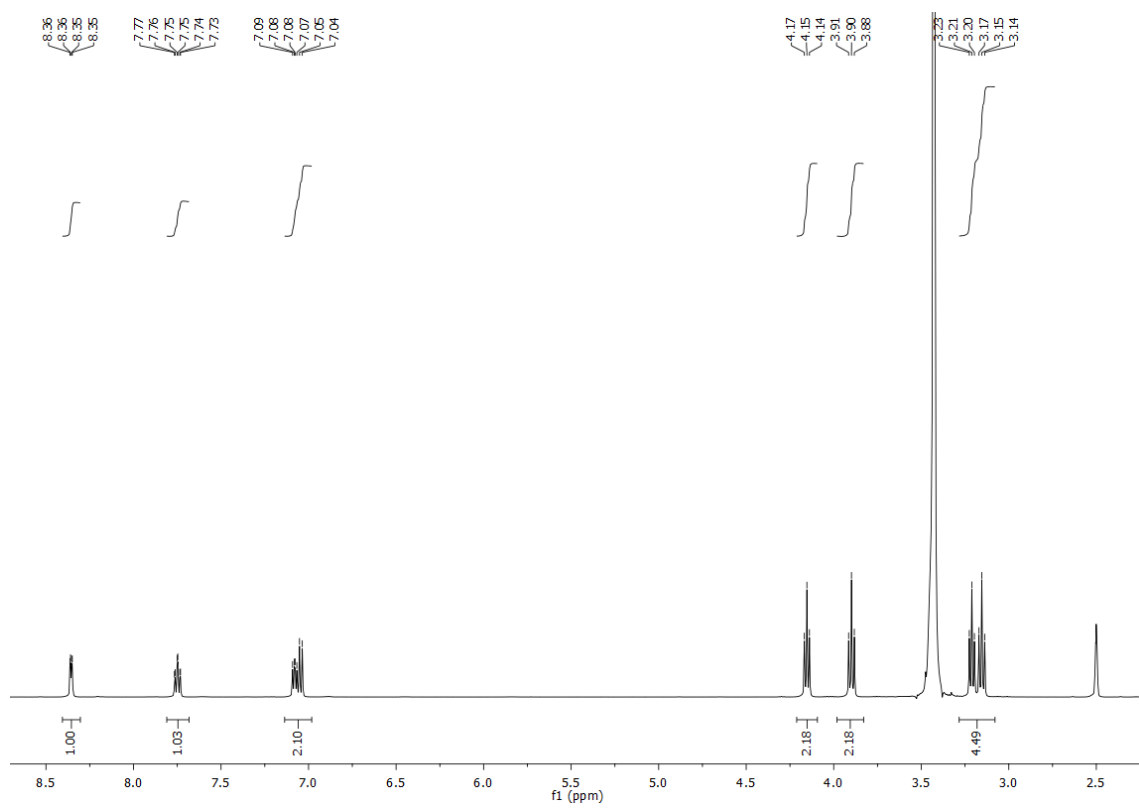


Figure S3. ^1H NMR spectrum of PyTT in $\text{DMSO-}d_6$.

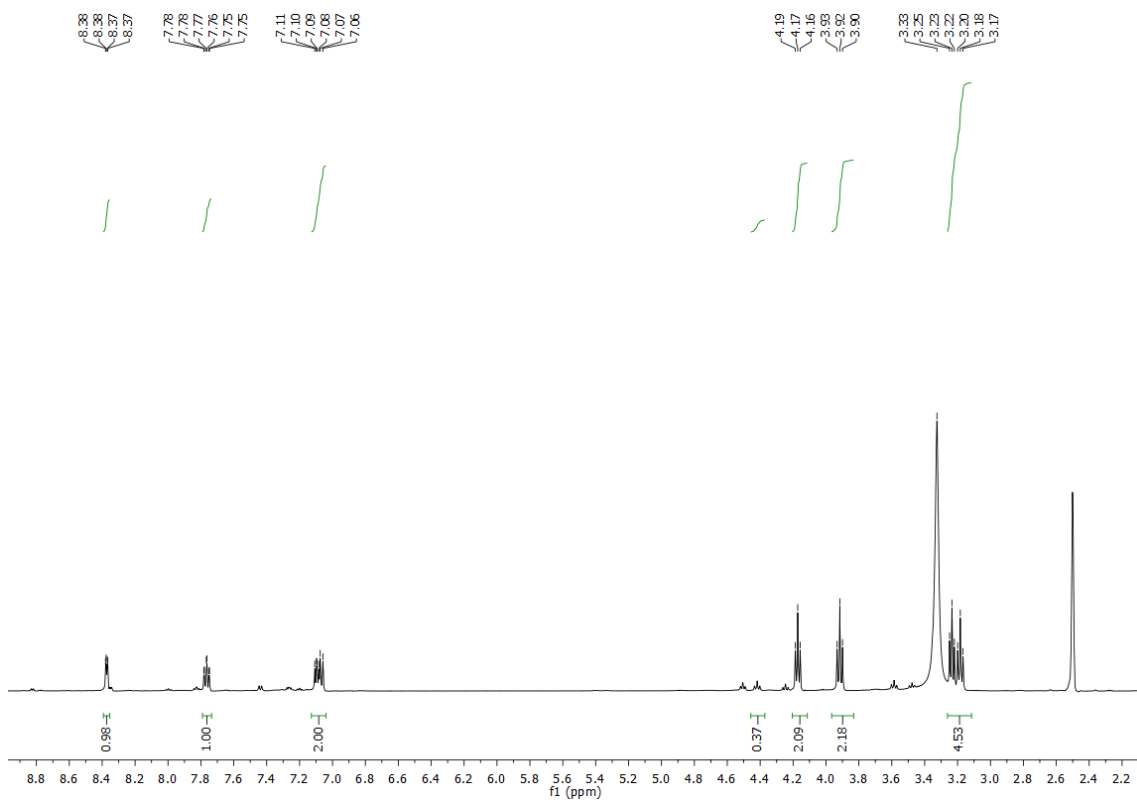


Figure S4. ^1H NMR spectrum of $[\text{Pd}_2\text{Cl}_2(\text{PyTT})_2]\text{Cl}_2 \cdot 4\text{H}_2\text{O}$ in $\text{DMSO-}d_6$.

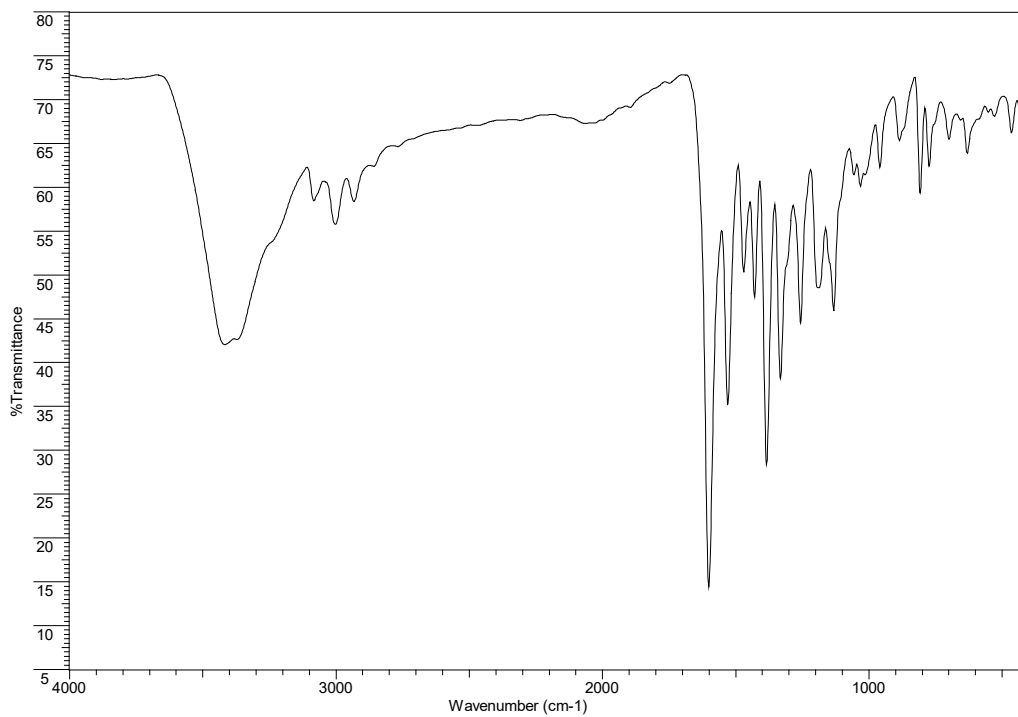


Figure S5. IR spectrum of $[\text{Pd}_2\text{Cl}_2(\text{PyTT})_2]\text{Cl}_2 \cdot 4\text{H}_2\text{O}$ in $4000\text{-}400\text{ cm}^{-1}$ region.

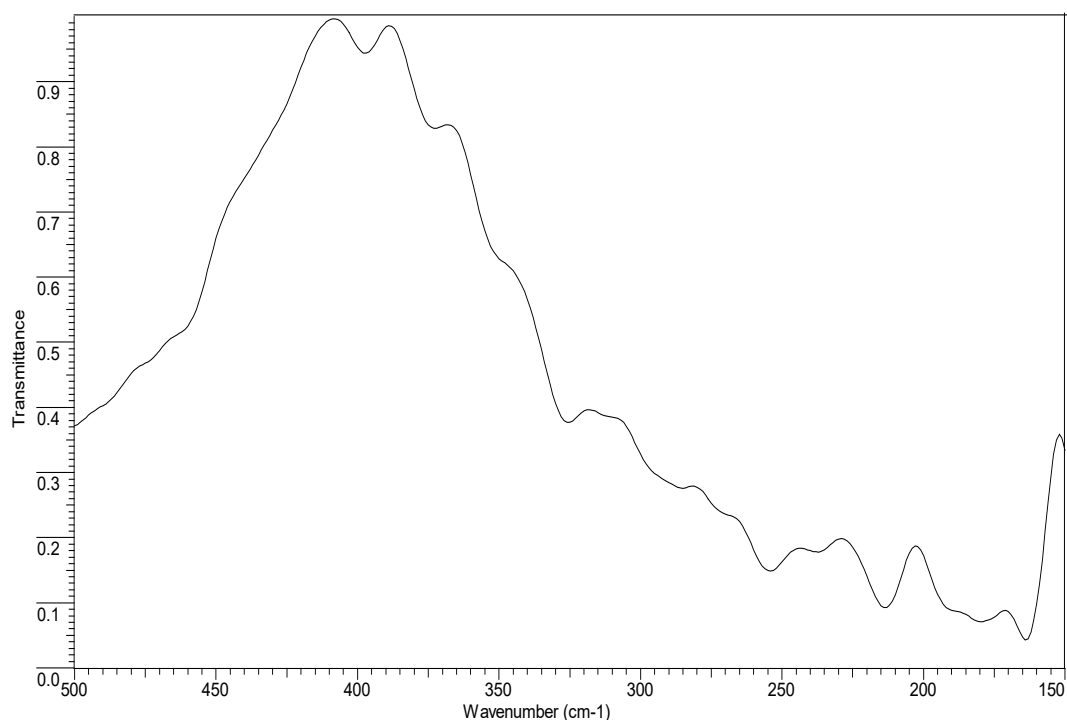


Figure S6. IR spectrum of $[Pd_2Cl_2(PyTT)_2]Cl_2 \cdot 4H_2O$ in $500-150\text{ cm}^{-1}$ region.

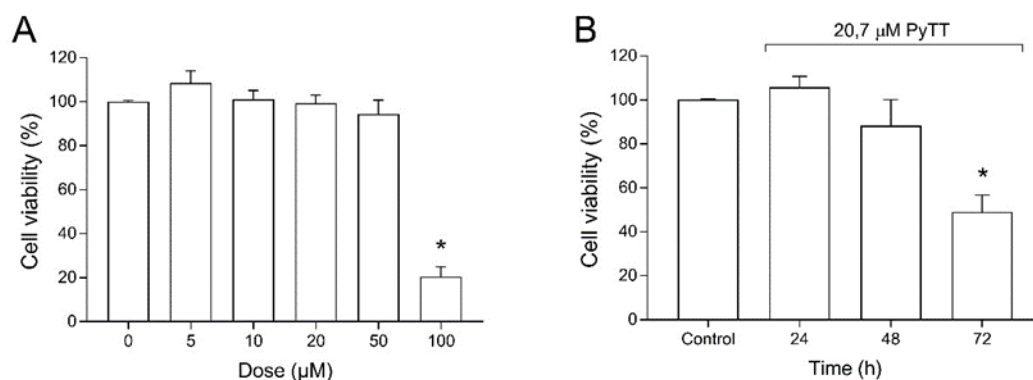


Figure S7: Dose- and time-response curves of the palladium complex ligand on cell viability in HL-60 cells. (A) Cells were treated for 24 hours with increasing concentrations (5, 10, 20, 50 and 100 μM) of free ligand (PyTT), or the vehicle (DMSO, control). (B) Cells were treated with 20.7 μM PyTT, or the vehicle (DMSO, control) for 24, 48 and 72 hours. Cell viability was evaluated by means of the MTS assay. Values are presented as means \pm S.D. of 6 independent experiments and expressed as percentage of control values (untreated cells). * $P < 0.05$ compared to control values.

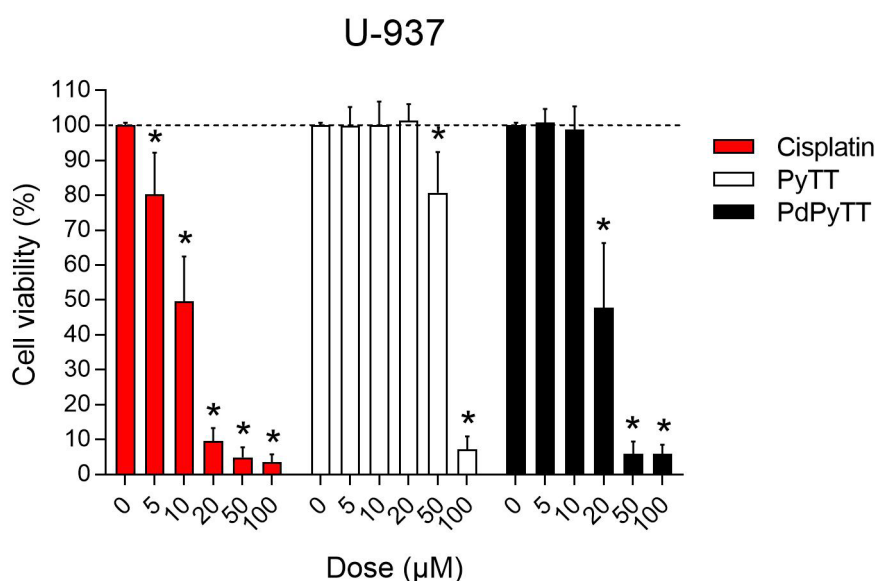


Figure S8: Dose-response curve of chemotherapeutics on cell viability of U-937 cells. (A) U-937 cells were treated for 24 hours with increasing concentrations (5, 10, 20, 50 and 100 μM) of cisplatin, the free ligand PyTT, the Pd(II) complex (PdPyTT), or the vehicle (DMSO, control). Values represent means \pm S.D. of 6 independent experiments and are expressed as percentage of control values. * $P < 0.05$ compared to their corresponding control values.

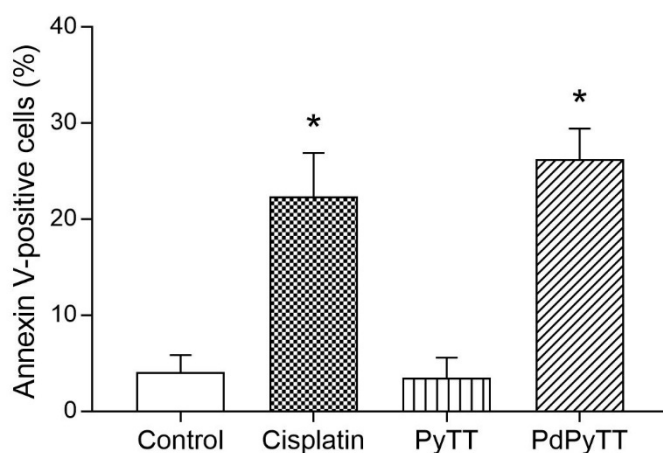


Figure S9: Effects of chemotherapeutics on phosphatidylserine externalization in HL-60 cells. Cells were treated with 11.3 μM cisplatin, 20.7 μM PyTT, 20.7 μM PdPyTT, or the vehicle (DMSO, control) for 24 hours. The phosphatidylserine externalization was determined by double-staining cells with annexin V-FITC/PI and analysed by flow cytometry. As indicated in material and methods, annexin V-positive cells were considered apoptotic cells. Histograms show percentages of apoptotic cells. Values represent means \pm S.D. of 6 independent experiments. * $P < 0.05$ compared to control values.

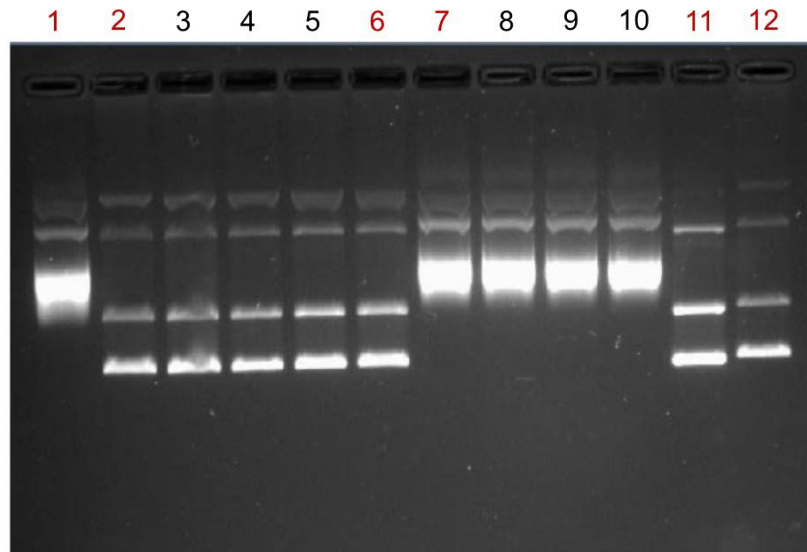


Figure S10: Full-length agarose gel corresponding to Figure 8A. Lane 1: supercoiled pBR322; lane 2: supercoiled pBR322 + wheat germ Topoisomerase I; lane 6: supercoiled pBR322 + wheat germ Topoisomerase I + 20.7 μ M PdPyTT; lane 7: supercoiled pBR322 + 20.7 μ M PdPyTT; lane 11: relaxed pBR322; lane 12: relaxed pBR322 + wheat germ Topoisomerase I. All other lanes (3-5, 8-10) correspond to different organometallic complexes that are still under evaluation in our lab for their potential therapeutic effects.

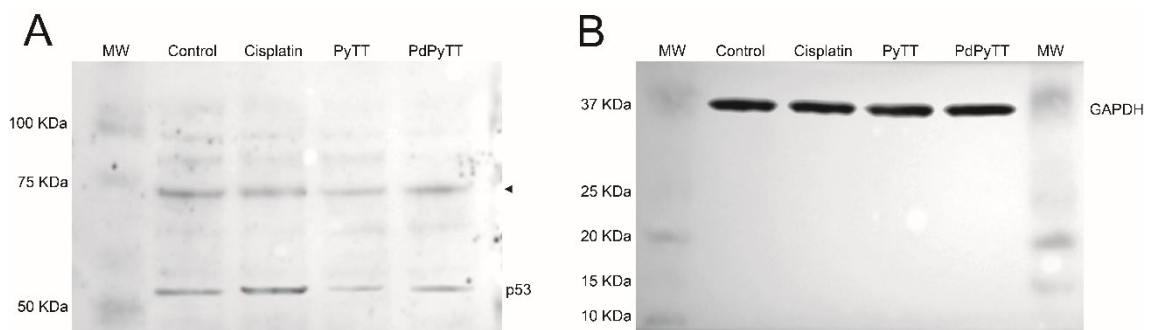


Figure S11: Full-length blots corresponding to Figure 8B. p53 (A) and GAPDH (B) protein expression levels. Arrowheads indicate unspecific bands. MW: molecular weight marker.

Table S1. *Hydrogen-bond parameters for PdPyTT*

D-H \cdots A	Position of A	H \cdots A	A-D (Å)	A \cdots H-D (°)
Hydrogen bonds inside water-chloride chain				
O(1W)-H(1W) \cdots Cl(5)	-x+2,-y+1,-z+1	2.27(2)	3.226(2)	178(1)
O(1W)-H(2W) \cdots Cl(7)	-x+1,-y+1,-z+1	2.39(3)	3.313(2)	163(2)
O(2W)-H(4W) \cdots Cl(6)	-x,-y,-z+1	2.25(2)	3.165(2)	161(1)
O(2W)-H(3W) \cdots Cl(8)	x-1,y,z	2.39(3)	3.293(2)	158(2)
O(3W)-H(5W) \cdots Cl(6)	-x+1,y+1/2,-z+1/2	2.28(3)	3.234(2)	175(2)
O(3W)-H(6W) \cdots Cl(7)	x,y,z	2.26(3)	3.172(2)	158(2)
O(4W)-H(8W) \cdots Cl(6)	-x+1,y+1/2,-z+1/2	2.23(4)	3.173(2)	169(3)
O(4W)-H(7W) \cdots Cl(7)	x,y,z	2.202(7)	3.144(3)	168.4(4)
O(5W)-H(10W) \cdots Cl(5)	x,-y+1/2,z+1/2	2.26(3)	3.216(3)	175(2)
O(5W)-H(9W) \cdots Cl(8)	x,y,z	2.21(2)	3.162(2)	170(2)
O(6W)-H(12W) \cdots Cl(5)	x,-y+1/2,z+1/2	2.23(4)	3.183(3)	171(3)
O(6W)-H(11W) \cdots Cl(8)	x,y,z	2.23(2)	3.129(3)	156(1)
O(7W)-H(14W) \cdots O(4W)	x,-y+1/2,z+1/2	1.90(2)	2.838(3)	167(2)
O(7W)-H(13W) \cdots O(5W)	x-1,y,z	1.92(3)	2.849(3)	163(3)
O(8W)-H(15W) \cdots O(3W)	x,y,z	1.90(3)	2.831(4)	164(3)
O(8W)-H(16W) \cdots O(6W)	x,-y+1/2,z-1/2	1.89(2)	2.844(4)	177(2)
Interactions between water-chloride chain and cationic complexes				
C(8)-H(8) \cdots Cl(5)	x-1,-y+1/2,z+1/2	2.795(1)	3.551(3)	139.2(2)
C(8A)-H(8A) \cdots Cl(6)	x,-y+1/2,z+1/2	2.693(1)	3.469(3)	141.4(2)
C(3C)-H(3C1) \cdots Cl(6)	-x+1,-y,-z+1	2.777(1)	3.625(3)	146.5(2)
C(3B)-H(3B1) \cdots Cl(6)	-x+1,-y,-z+1	2.924(1)	3.674(3)	135.0(2)
C(11B)-H(11B) \cdots Cl(7)	x,y,z	2.843(1)	3.618(3)	141.6(2)
C(6A)-H(6A1) \cdots Cl(7)	-x+1,-y+1,-z+1	2.741(1)	3.508(3)	136.4(2)
C(6)-H(6A) \cdots Cl(8)	x-1,y,z	2.659(1)	3.431(3)	136.8(2)
C(11C)-H(11C) \cdots Cl(8)	-x+2,-y,-z+1	2.708(1)	3.504(3)	144.1(2)
C(3A)-H(3A1) \cdots O(6W)	x-1,-y+1/2,z-1/2	2.590(2)	3.325(4)	132.7(2)
C(11)-H(11) \cdots O(6W)	-x+1,+y+1/2,-z+1/2	2.757(2)	3.553(3)	144.2(2)
C(3A)-H(3A2) \cdots O(1W)	-x+1,-y+1,-z+1	2.702(2)	3.647(4)	164.7(2)
C(11A)-H(11A) \cdots O(7W)	x,y,z	2.699(2)	3.493(3)	143.9(2)
C(5B)-H(5B2) \cdots O(7W)	-x+1,-y,-z+1	2.459(2)	3.269(3)	140.8(2)
C(11A)-H(11A) \cdots O(4W)	x,-y+1/2,z+1/2	2.686(2)	3.468(3)	142.3(2)
C(6B)-H(6B2) \cdots O(4W)	-x+1,+y-1/2,-z+1/2	2.932(2)	3.693(4)	136.1(2)
C(11B)-H(11B) \cdots O(4W)	x,y,z	2.791(2)	3.570(3)	142.0(2)
C(3C)-H(3C2) \cdots O(4W)	x,-y+1/2,z+1/2	2.749(2)	3.457(4)	130.3(2)
C(5B)-H(5B2) \cdots O(7W)	-x+1,-y,-z+1	2.459(2)	3.269(2)	140.8(2)
C(8B)-H(8B) \cdots O(1W)	x,-y+1/2,z-1/2	2.704(2)	3.462(3)	139.3(2)
C(8B)-H(8B) \cdots O(5W)	-x+2,-y,-z+1	2.637(2)	3.415(3)	141.5(2)

C(5C)-H(5C2)···O(8W)	x,y,z	2.427(3)	3.253(4)	142.8(2)
C(8C)-H(8C)···O(2W)	x+1,-y+1/2,z-1/2	2.615(2)	3.308(3)	131.7(2)
C(8C)-H(8C)···O(3W)	x,y,z	2.626(2)	3.425(3)	144.4(2)
Interactions between cationic complexes				
C(5)-H(5B)···Cl(3)	x-1,y,z	2.738(1)	3.518(3)	138.0(2)
C(5A)-H(5A2)···Cl(1)	-x+1,-y+1,-z+1	2.779(1)	3.478(3)	129.5(2)
C(5B)-H(5B1)···Cl(4)	-x+1,-y,-z+1	2.789(1)	3.601(3)	141.7(2)
C(5C)-H(5C1)···Cl(2)	x+1,y,z	2.735(1)	3.345(3)	121.5(2)
C(10)-H(10)···Cl(8)	-x+1,y+1/2,-z+1/2	2.970(1)	3.683(3)	134.6(2)
C(10A)-H(10A)···Cl(4)	x,-y+1/2,z+1/2	2.955(1)	3.595(3)	127.3(2)
C(10B)-H(10B)···Cl(1)	x,-y+1/2,z-1/2	2.813(1)	3.604(3)	143.6(2)
C(10C)-H(10C)···Cl(2)	x+1,-y+1/2,z-1/2	2.797(1)	3.621(3)	148.2(2)
C(9)-H(9)···S(1)	x,-y+1/2,z+1/2	2.921(1)	3.627(3)	133.8(2)
C(9B)-H(9B)···S(1A)	-x+1,y-1/2,-z+1/2	2.857(1)	3.491(3)	126.5(2)
C(9B)-H(9B)···S(2A)	x,-y+1/2,z-1/2	2.900(1)	3.480(3)	121.8(2)
C(9C)-H(9C)···S(2)	x+1,-y+1/2,z-1/2	2.916(1)	3.490(3)	121.3(2)