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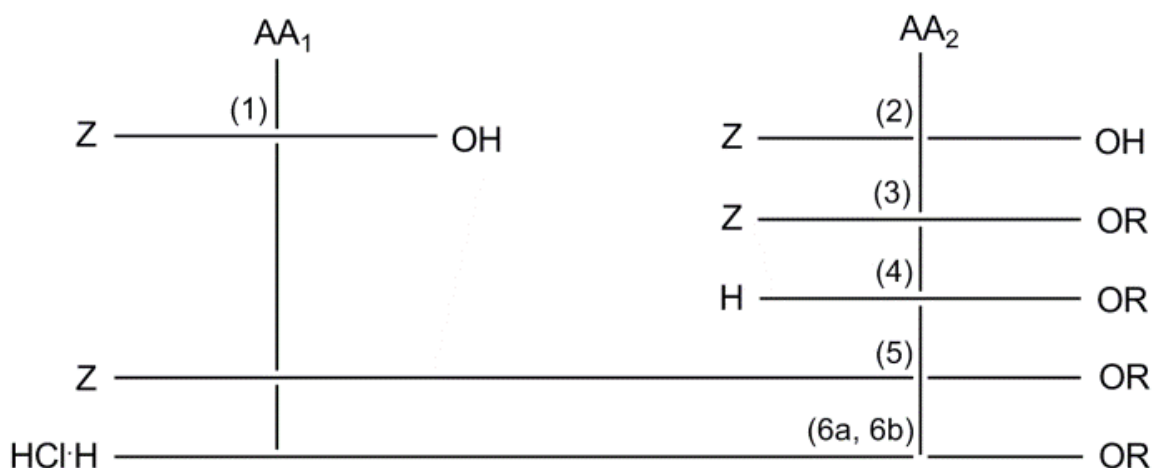
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Scheme S1. Synthetic scheme for the preparation of hydrochloride dipeptide esters P1-P5 (AA_1 = Sar, L/D-Pro; AA_2 =L/D-Ala, Aib; R= OtBu, OTEG). (1) and (2) Z-OSu, triethylamine, CH_3CN/H_2O ; (3) –OtBu: isobutene, H_2SO_4 cat., CH_2Cl_2 ; -OTEG: EDC/DMAP/HOBt* (Figure S1), triethylen glycol monomethyl ether, CH_2Cl_2 ; (4) H_2 , Pd/C (10% w/w), methanol; (5) for the Sar-derivatives isobutyl chloroformate (Figure S2), *N*-methylmorpholine, tetrahydrofurane; for the Pro-derivatives EDC, HOBt, diethylisopropylamine, dichloromethane; (6a) H_2 , Pd/C, methanol; (6b) HCl/diethylether.

*1-ethyl-3-(3-dimethylaminopropyl)carbodiimide/4-dimethylaminopyridine/Hydroxybenzotriazole

Table S1. Thermogravimetric (TG) and differential scanning calorimetric (DSC) data.

Compound	Weight loss (%)		DSC
	Found	Calculated	Peak temperature [°C] (process*)
IT01	67.95	69.61	158.0 (endo); 499.4 (exo)
IT02	67.90	69.61	151.0 (endo); 469.4 (exo)
IT03	72.00	71.38	113.7 (endo); 439.7 (exo)
IT04	71.89	71.38	178.0 (endo); 436.7 (exo)
IT05	73.82	72.61	122.6 (endo); 493.7 (exo); 542.1 (exo)

*endo/exo=endothermic/exothermic

Table S2. Selected IR frequencies of the starting dipeptides P1-P5 and the corresponding gold(III) dithiocarbamate derivatives IT01-IT05.

		Vibrational mode [cm ⁻¹]								
	ν NH	$\nu_{a/s}$ NH ₂ ⁺	ν C=O	amide I	ν N-CSS	Amide II	Amide III	$\nu_{a/s}$ SCS	$\nu_{a/s}$ SAuS	$\nu_{a/s}$ BrAuBr
P1	3332	2763	1735	1669	-	1564	1266	-	-	-
IT01	3317	-	1733	1668	1564 ^a	-	1219	1003/ 567	419/ 380	251/ 229
P2	3332	2766	1735	1669	-	1564	1266	-	-	-
IT02	3311	-	1732	1668	1563 ^a	-	1218	1003/ 566	419/ 380	251/ 228
P3	3430	2744	1733	1678	-	1551	1256	-	-	-
IT03	3341	-	1730	1680	1553 ^a	-	1247	1000/ 535	410/376	252/222
P4	3437	2744	1733	1678	-	1551	1256	-	-	-
IT04	3363	-	1732	1680	1549 ^a	-	1241	991/ 533	410/ 376	251/ 222
P5	3193	2762	1739	1684	-	1554	1256	-	-	-
IT05	3290	-	1735	1683	1576	1549	1253	1022/ 569	412/ 390	250/ 226

^a ν N-CSS and amide II are overlapped.

The main signals generated by the peptide chains are the vibration mode of the amide-proton bond (ν_{N-H}) at 3550-3200 cm⁻¹ (Amide A band), and the three bands characteristic of the C(O)NH function, resulting from the combination of the ν (C=O), δ (N-H) and ν (C-N) vibration modes (referred to as amide I, II and III).

With respect to the formation of the gold(III) dithiocarbamate complexes, there are three main diagnostic regions:

- the 1580-1450 cm⁻¹ region, primarily associated with the ν (N-CSS) vibration mode (“thioureide” band);
- the 1060-940 cm⁻¹ region, associated with the ν (S-C-S) vibration modes;
- the 420-350 cm⁻¹ region, associated with the ν (M-S) vibration modes.

Upon complex formation, the deprotonation step of the HCl·dipeptide to form the corresponding dithiocarbamate ligand causes the disappearance of the signal at 2766-2736 cm⁻¹ of the $\nu_{a/s}$ (NH₂⁺) vibration mode, and the appearance of both the ν (N-CSS) band at 1580-1450 cm⁻¹ (somewhat overlapped to amide II band) and the ν_a (S-C-S) band at 1060-940 cm⁻¹.

As described by Chatt *et al.*,^[39] the dithiocarbamate moiety is characterized by a strong delocalization of the electrons, being described by three main resonance structures (Figure S3). The position of the ν (N-CSS) bands recorded for our compounds is consistent with the presence of a carbon-nitrogen bond order which lies between a single bond (ν =1350-1250 cm⁻¹) and a double bond ($\tilde{\nu}$ =1690-1640 cm⁻¹).^[40] This is in agreement with a relevant contribution to the final structure

of the resonance form III reported in Figure S3, characterized by a symmetric chelated coordination of the dithiocarbamate function to the gold(III) metal center.^[52] Likewise, the presence of a single band for the $\nu_a(\text{S-C-S})$ at 1022-991 and the $\nu_s(\text{S-C-S})$ at 569-533 cm^{-1} validates a bidentate symmetric coordination of the ligand to the metal center (Figure S4).^[53] In the Far FT-IR region, the formation of a symmetrically-chelated dithiocarbamate is further confirmed by the position of the metal-sulfur stretching vibrations $\nu_{a/s}(\text{S-Au-S})$ at 419-410/390-376 cm^{-1} , in agreement with previously reported data.^[52, 54] The bands related to the *cis*-coordinated halides $\nu_{a/s}(\text{Br-Au-Br})$ set at 252-250/229-220 cm^{-1} .^[52, 54-55]

Table S3. ^{13}C NMR spectral data of the starting dipeptides P1-P5 (HCl-H-AA₁-AA₂-OR: AA₁= Sar, L/D-Pro; AA₂=D/L-Ala, Aib; R= -tBu, TEG) and the corresponding gold(III)-dithiocarbamate derivatives IT01-IT05 ([Au^{III}Br₂(dtc-AA₁-AA₂-OR)] (DMSO-d₆, 298 K, 75.48 MHz).

Compound	δ (^{13}C) [ppm]			
	-OR	AA ₁	AA ₂	C-SS
P1	R= tBu	AA₁=Sar	AA₂=L-Ala	-
	27.36(-C(CH ₃) ₃)	32.40(N- CH ₃)	16.75 (β- CH ₃)	
	80.58(-C(CH ₃) ₃)	48.53(N- CH ₂)	48.29 (α-CH)	
IT01*	28.35(-C(CH ₃) ₃)	164.86 (C=O)	171.27 (C=O)	198.2
	82.36(-C(CH ₃) ₃)	40.5 (N-CH ₃)	18.34 (β- CH ₃)	
		55.5 (N- CH ₂)	50.35 (α-CH)	
P2	R= tBu	AA₁=Sar	AA₂=D-Ala	-
	27.28(-C(CH ₃) ₃)	32.41(N- CH ₃)	16.76(β- CH ₃)	
	80.55(-C(CH ₃) ₃)	48.52(N- CH ₂)	48.29(α-CH)	
IT02*	27.69(-C(CH ₃) ₃)	164.87 (C=O)	171.22 (C=O)	197.72
	81.70(-C(CH ₃) ₃)	39.9 (N- CH ₃)	17.67 (β- CH ₃)	
		54.83 (N- CH ₂)	49.68 (α-CH)	
P3	R= tBu	AA₁=L-Pro	AA₂=Aib	-
	27.13(-C(CH ₃) ₃)	23.28 (CH ₂ ⁴)	55.69 C(CH ₃) ₂	
	79.65(-C(CH ₃) ₃)	29.59 (CH ₂ ³)	24.41 (CH ₃)	
IT03*		45.32 (CH ₂ ⁵)	172.15 (C=O)	
		58.07 (CH ²)		
		167.15 (C=O)		
P4	R= tBu	AA₁=D-Pro	AA₂=Aib	-
	27.19(-C(CH ₃) ₃)	23.56 (CH ₂ ⁴)	24.50 (CH ₃)	
	79.65(-C(CH ₃) ₃)	29.67 (CH ₂ ³)	55.70 C(CH ₃) ₂	
IT04*		45.55 (CH ₂ ⁵)	172.06 (C=O)	
		58.26 (CH ²)		
		167.75 (C=O)		
P5	R= tBu	AA₁=L-Pro	AA₂=Aib	-
	27.13(-C(CH ₃) ₃)	23.28 (CH ₂ ⁴)	24.95 (CH ₃)	
	79.65(-C(CH ₃) ₃)	29.59 (CH ₂ ³)	57.44 C(CH ₃) ₂	
IT05*		45.32 (CH ₂ ⁵)	172.96 (C=O)	190.21
		58.07 (CH ²)		
		167.15 (C=O)		

		166.94 (C=O)		
P5	R=TEG	AA₁=Sar	AA₂=Aib	
	57.26 (O-CH ₃)	31.84 (N-CH ₃)	23.39(β-CH ₃)	
	63.19 (CH ₂ ¹)		54.77	
			C(CH ₃) ₂	
			172.52 (C=O)	
	67.44 (CH ₂ ²)	47.94 (N-CH ₂)		
		163.80 (C=O)		
	68.8-70.7 (CH ₂ ^{3,4,5,6})		172.52 (C=O)	
IT05*	58.52 (OCH ₃)	39.86 (NCH ₃)	24.86(β-CH ₃)	197.3
			57.00 C(CH ₃) ₂	
	64.52 (CH ₂ ¹)	54.84 (NCH ₂)	173.63 (C=O)	
	69.25 (CH ₂ ²)	163.54 (C=O)		
	70.8-71.2 (CH ₂ ^{3,4,5,6})			

* acetone-d₆ (recorded in such a solvent due to signal overlapping with DMSO resonances).

Table S4. Summary of X-ray crystallographic data for IT05-pentane.

Empirical formula	C ₂₀ H ₃₉ AuBr ₂ N ₂ O ₆ S ₂
Formula weight	824.44
Colour, habit	Yellow, block
Crystal size, mm	0.07x0.04x0.04
Crystal system	Orthorhombic
Space group	<i>Pbcn</i>
<i>a</i> , Å	22.15(2)
<i>b</i> , Å	9.599(9)
<i>c</i> , Å	25.88(2)
α , deg.	90
β , deg.	90
γ , deg.	90
<i>V</i> , Å ³	5503(8)
<i>Z</i>	8
<i>T</i> , K	293(2)
ρ (calc), Mg/m ³	1.990
μ , mm ⁻¹	8.440
θ range, deg.	1.82 to 20.75
No. of rflcn/unique	9221 / 2636
Goof	1.028
<i>R</i> 1	0.0877
<i>wR</i> 2	0.1774

$$R1 = \frac{\sum ||F_o| - |F_c||}{\sum |F_o|}, \quad wR2 = \frac{[\sum [w(F_o^2 - F_c^2)^2]]}{\sum [w(F_o^2)^2]}^{1/2}, \quad w = 1/[\sigma^2(F_o^2) + (aP)^2 + bP], \quad \text{where } P = [\max(F_o^2, 0) + 2F_c^2]/3$$

Table S5. Selected bond lengths (Å) and angles (°) for IT05-pentane.

Au-Br(1)	2.429(4)	C(1)-N(1)	1.32(3)
Au-Br(2)	2.431(5)	C(4)-O(1)	1.27(3)
Au-S(1)	2.310(9)	C(4)-N(2)	1.33(3)
Au-S(2)	2.306(9)	C(8)-O(2)	1.21(4)
C(1)-S(1)	1.78(3)	C(8)-O(3)	1.36(4)
C(1)-S(2)	1.71(3)		
Br(1)-Au-Br(2)	94.0(2)	S(1)-C(1)-S(2)	108(2)
S(1)-Au-S(2)	75.7(3)	C(1)-S(1)-Au	87(1)
Br(1)-Au-S(1)	96.1(2)	C(1)-S(2)-Au	89(1)
Br(2)-Au-S(2)	94.1(2)		

Table S6. Crystallographic data and structure refinement for the racemate IT03/IT04 [Au^{III}Br₂(dtc-L,D-Pro-Aib-O*t*Bu)] acetone solvate.

Identification code	mc181fl	
Empirical formula	C ₁₄ H ₂₃ AuBr ₂ N ₂ O ₃ S ₂ x C ₃ H ₆ O	
Formula weight	746.33	
Temperature	293(2) K	
Wavelength	0.71070 Å	
Crystal system	Orthorhombic	
Space group	Pbca	
Unit cell dimensions	a = 16.5424(6) Å	α = 90°.
	b = 11.7649(4) Å	β = 90°.
	c = 26.3963(7) Å	γ = 90°.
Volume	5137.2(3) Å ³	
Z	8	
Density (calculated)	1.930 Mg/m ³	
Absorption coefficient	9.024 mm ⁻¹	
F(000)	2864	
Crystal size	0.50 x 0.40 x 0.15 mm ³	
Theta range for data collection	3.09 to 26.37°.	
Index ranges	-20 ≤ h ≤ 20, -14 ≤ k ≤ 14, -25 ≤ l ≤ 27	
Reflections collected	9339	
Independent reflections	3000 [R(int) = 0.0488]	
Completeness to theta = 26.37°	57.1 %	
Absorption correction	Semi-empirical from equivalents	
Max. and min. transmission	1.00000 and 0.20457	
Refinement method	Full-matrix least-squares on F ²	
Data / restraints / parameters	3000 / 158 / 285	
Goodness-of-fit on F ²	1.255	
Final R indices [I > 2σ(I)]	R ₁ = 0.0871, wR ₂ = 0.1526	
R indices (all data)	R ₁ = 0.1151, wR ₂ = 0.1610	
Largest diff. peak and hole	0.783 and -1.024 e.Å ⁻³	

Table S7. Selected bond lengths (Å) and angles (°) for the racemate IT04 [Au^{III}Br₂(dtc-L,D-Pro-Aib-O^tBu)].

Au-Br(1)	2.420(3)	C(1)-N(1)	1.28(2)
Au-Br(2)	2.435(3)	C(6)-O(1)	1.23(2)
Au-S(1)	2.313(6)	C(6)-N(2)	1.32(2)
Au-S(2)	2.307(7)	C(10)-O(2)	1.25(3)
C(1)-S(1)	1.71(2)	C(10)-O(3)	1.29(3)
C(1)-S(2)	1.74(2)		
Br(1)-Au-Br(2)	93.64(12)	S(1)-C(1)-S(2)	110.3(12)
S(1)-Au-S(2)	75.7(2)	C(1)-S(1)-Au	87.1(8)
Br(1)-Au-S(1)	95.77(18)	C(1)-S(2)-Au	86.8(8)
Br(2)-Au-S(2)	94.93(18)		

Table S8. Crystallographic data and structure refinement for Z-L-Pro-Aib-OtBu.

Identification code	Z-L-Pro-Aib-OtBu	
Empirical formula	C ₂₁ H ₃₀ N ₂ O ₅	
Formula weight	390.47	
Temperature	293(2) K	
Wavelength	1.54178 Å	
Crystal system	Orthorhombic	
Space group	P 21 21 21	
Unit cell dimensions	a = 10.050(2) Å	α = 90°.
	b = 10.460(2) Å	β = 90°.
	c = 21.305(3) Å	γ = 90°.
Volume	2239.6(7) Å ³	
Z	4	
Density (calculated)	1.158 Mg/m ³	
Absorption coefficient	0.674 mm ⁻¹	
F(000)	840	
Crystal size	0.50 x 0.35 x 0.20 mm ³	
Theta range for data collection	4.15 to 59.97°.	
Index ranges	-1 ≤ h ≤ 11, 0 ≤ k ≤ 11, 0 ≤ l ≤ 23	
Reflections collected	2158	
Independent reflections	2121 [R(int) = 0.1078]	
Completeness to theta = 59.97°	99.9 %	
Max. and min. transmission	0.8770 and 0.7293	
Refinement method	Full-matrix-block least-squares on F ²	
Data / restraints / parameters	2121 / 4 / 250	
Goodness-of-fit on F ²	0.992	
Final R indices [I > 2σ(I)]	R1 = 0.0611, wR2 = 0.1612	
R indices (all data)	R1 = 0.0774, wR2 = 0.1735	
Absolute structure	-0.1(5)	

parameter	
Largest diff. peak and hole	0.258 and -0.174 e.Å ⁻³

Table S9. Bond lengths [Å] and angles [°] for Z-L-Pro-Aib-OtBu.

C01-C02	1.39
C01-C06	1.39
C01-C07	1.491(5)
C02-C03	1.39
C03-C04	1.39
C04-C05	1.39
C05-C06	1.39
C07-OU	1.421(5)
OU-C0	1.335(5)
C0-O0	1.231(5)
C0-N1	1.325(5)
N1-C1D	1.456(6)
N1-C1A	1.464(5)
C1A-C1	1.523(6)
C1A-C1B	1.528(6)
C1B-C1G	1.480(8)
C1B-C1G'	1.487(8)
C1G-C1D	1.499(8)
C1G'-C1D	1.497(8)
C1-O1	1.230(5)
C1-N2	1.345(5)
N2-C2A	1.444(6)
C2A-C2B2	1.519(6)
C2A-C2B1	1.536(6)
C2A-C2	1.536(6)
C2-O2	1.204(5)
C2-OT	1.335(5)
OT-CT1	1.481(5)
CT1-CT2	1.501(8)
CT1-CT4	1.501(8)
CT1-CT3	1.507(8)
C02-C01-C06	120
C02-C01-C07	122.4(3)
C06-C01-C07	117.6(3)
C01-C02-C03	120
C02-C03-C04	120
C05-C04-C03	120
C04-C05-C06	120
C05-C06-C01	120
OU-C07-C01	109.0(3)
C0-OU-C07	116.0(3)

O0-C0-N1	124.8(4)
O0-C0-OU	123.3(4)
N1-C0-OU	111.8(3)
C0-N1-C1D	122.1(4)
C0-N1-C1A	122.6(4)
C1D-N1-C1A	112.3(3)
N1-C1A-C1	110.3(3)
N1-C1A-C1B	102.9(3)
C1-C1A-C1B	112.2(4)
C1G-C1B-C1A	106.7(5)
C1G-C1B-C1G'	28.6(7)
C1A-C1B-C1G'	103.9(5)
C1B-C1G-C1D	107.9(6)
C1D-C1G'-C1B	107.6(6)
N1-C1D-C1G'	103.9(5)
N1-C1D-C1G	100.9(5)
O1-C1-N2	122.6(4)
O1-C1-C1A	122.6(4)
N2-C1-C1A	114.9(3)
C1-N2-C2A	123.3(3)
N2-C2A-C2B2	111.3(4)
N2-C2A-C2B1	108.1(4)
C2B2-C2A-C2B1	109.4(4)
N2-C2A-C2	112.0(3)
C2B2-C2A-C2	109.9(4)
C2B1-C2A-C2	106.0(3)
O2-C2-OT	124.9(4)
O2-C2-C2A	123.1(4)
OT-C2-C2A	111.8(3)
C2-OT-CT1	120.8(3)
OT-CT1-CT2	109.8(4)
OT-CT1-CT4	108.3(5)
CT2-CT1-CT4	115.2(6)
OT-CT1-CT3	102.1(4)
CT2-CT1-CT3	110.7(6)
CT4-CT1-CT3	109.9(5)

Table S10. Selected torsion angles [°] for Z-L-Pro-Aib-OtBu.

C02-C01-C07-OU	32.7(6)
C06-C01-C07-OU	-147.7(4)
C01-C07-OU-C0	170.2(4)
C07-OU-C0-N1	176.2(5)
OU-C0-N1-C1A	-17.5(6)
C0-N1-C1A-C1	-60.0(5)
N1-C1A-C1-N2	168.8(3)
C1A-C1-N2-C2A	175.7(3)
C1-N2-C2A-C2	-53.1(5)
N2-C2A-C2-OT	-31.8(5)
C2A-C2-OT-CT1	-178.5(4)
C2-OT-CT1-CT2	-59.4(6)
C2-OT-CT1-CT4	67.2(6)
C2-OT-CT1-CT3	-176.9(4)

Table S11. Main absorption band maxima (nm) recorded for IT03 in different solvents.

	λ_{max}/nm		
	Band I	Band II	Band III
DMSO	276	316	378
Bidistilled water	272	310	370
Saline solution 0.9% w/v NaCl	261	312	--
PBS	262	313	375
Cell culture medium + 10% bovine fetal serum	273	307	362

* λ_{max} determined soon after dissolution

Table S12. Inhibition potency of selected compounds incubated (1 h for the Au(III) complexes, and 24 h for Cisplatin and Olaparib[®]) with purified PARP-1. Values expressed in terms of IC₅₀ ± st.dev. (μM).

	IC ₅₀ ± st.dev (μM)
IT03	0.017 ± 0.006
AuL12	0.019 ± 0.004
Cisplatin	12 ± 2
Olaparib [®]	0.00500 ± 0.00001

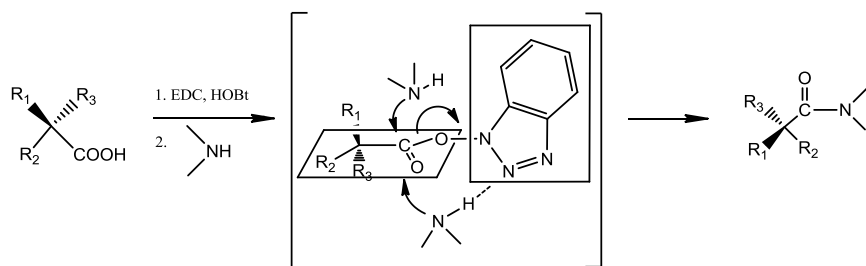


Figure S1. Schematic representation of the peptide bond formation using HOBt/EDC approach.

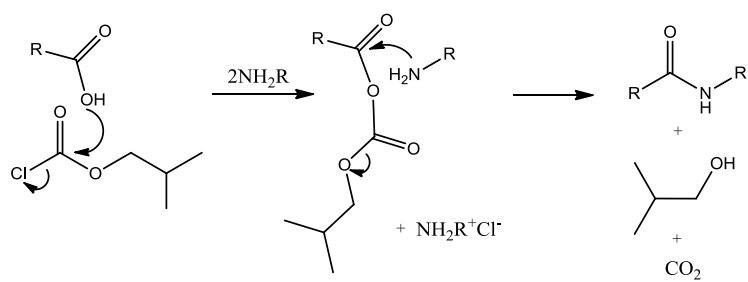


Figure S2. Schematic representation of the peptide bond formation using isobutylchloroformiate as a coupling reagent.

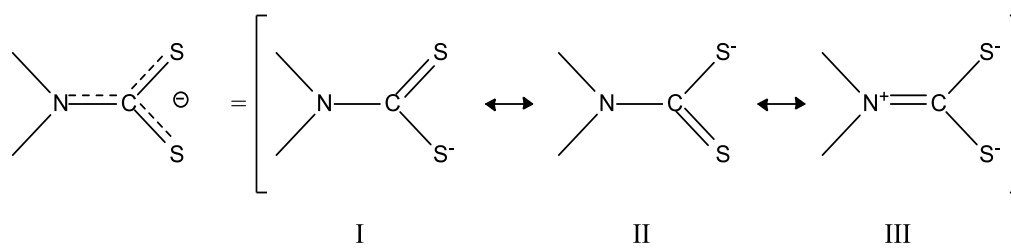


Figure S3. Resonance forms of the dithiocarbamic -NCSS^- moiety.

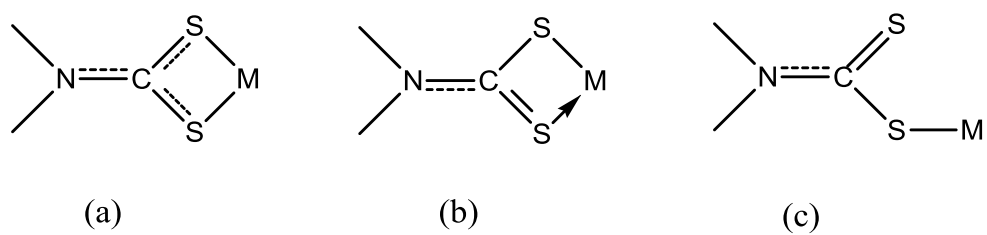


Figure S4. Different ways of metal-sulfur binding in dithiocarbamate complexes: symmetrical bidentate (a), asymmetrical bidentate (b) and monodentate (c).

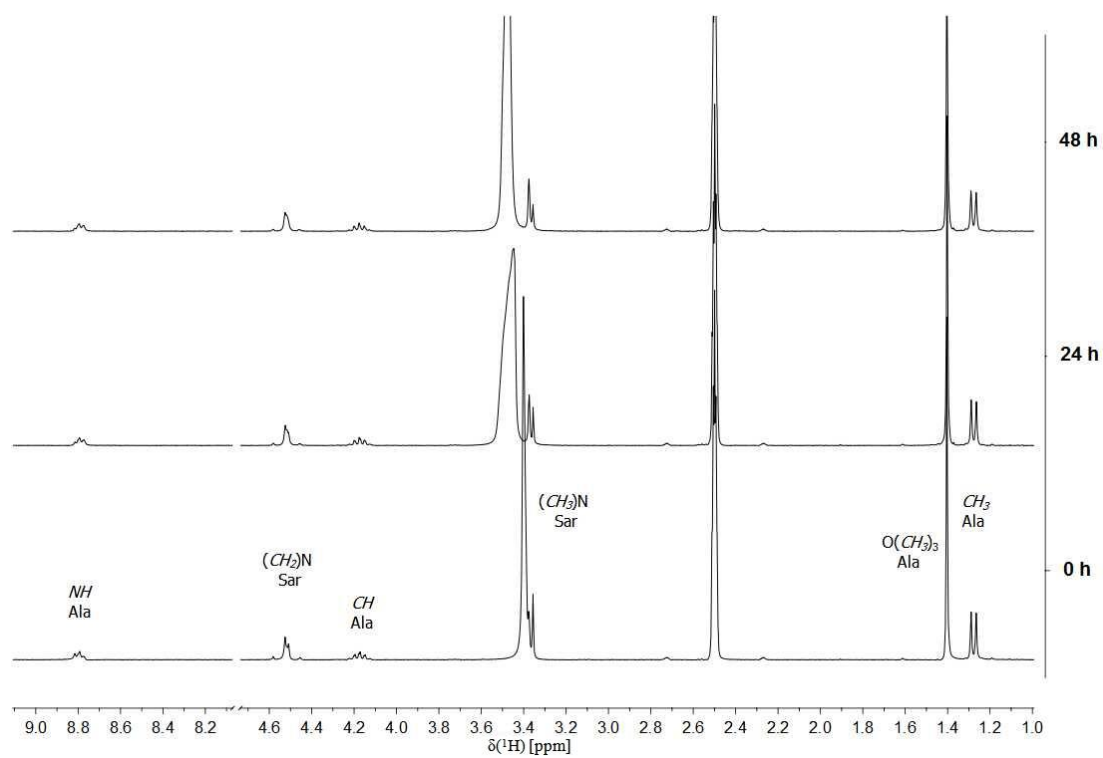


Figure S5. ¹H-NMR spectra of IT01 in DMSO-d₆ over 48 h.

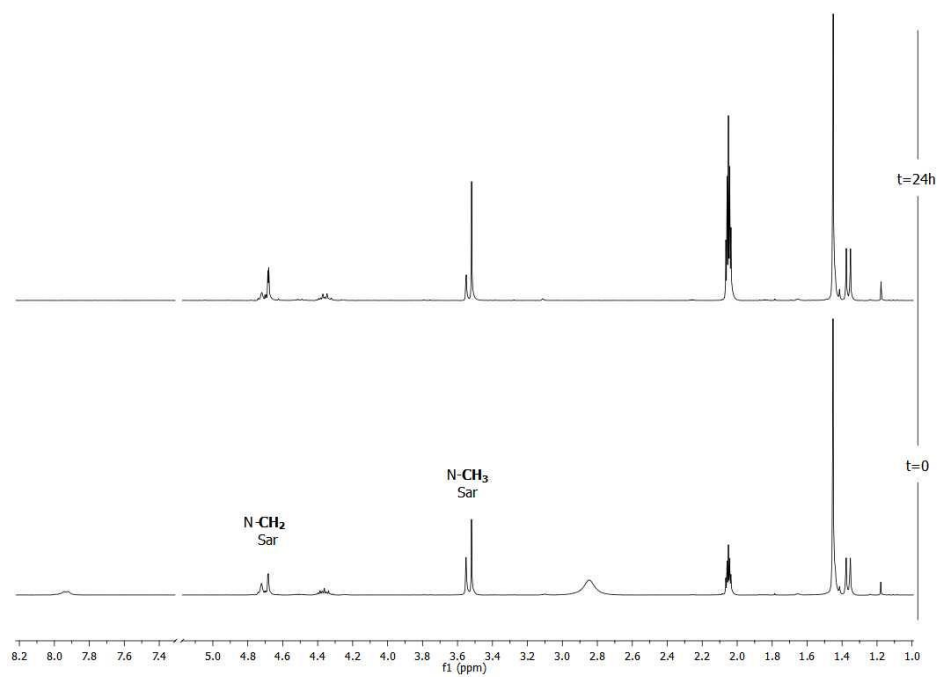


Figure S6. ¹H-NMR spectra of IT01 in acetone-d₆ over 24 h.

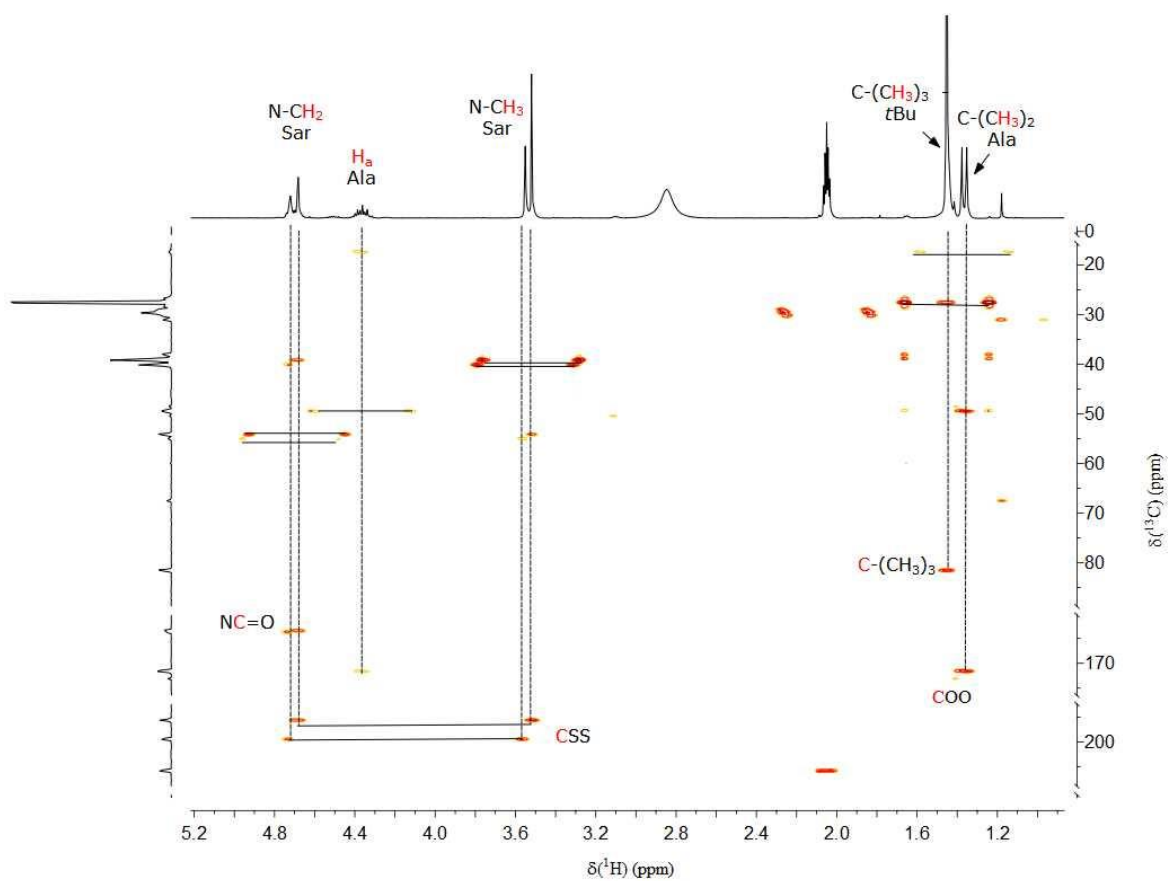


Figure S7. [^1H , ^{13}C]-HMBC spectrum of IT01 in acetone- d_6 .

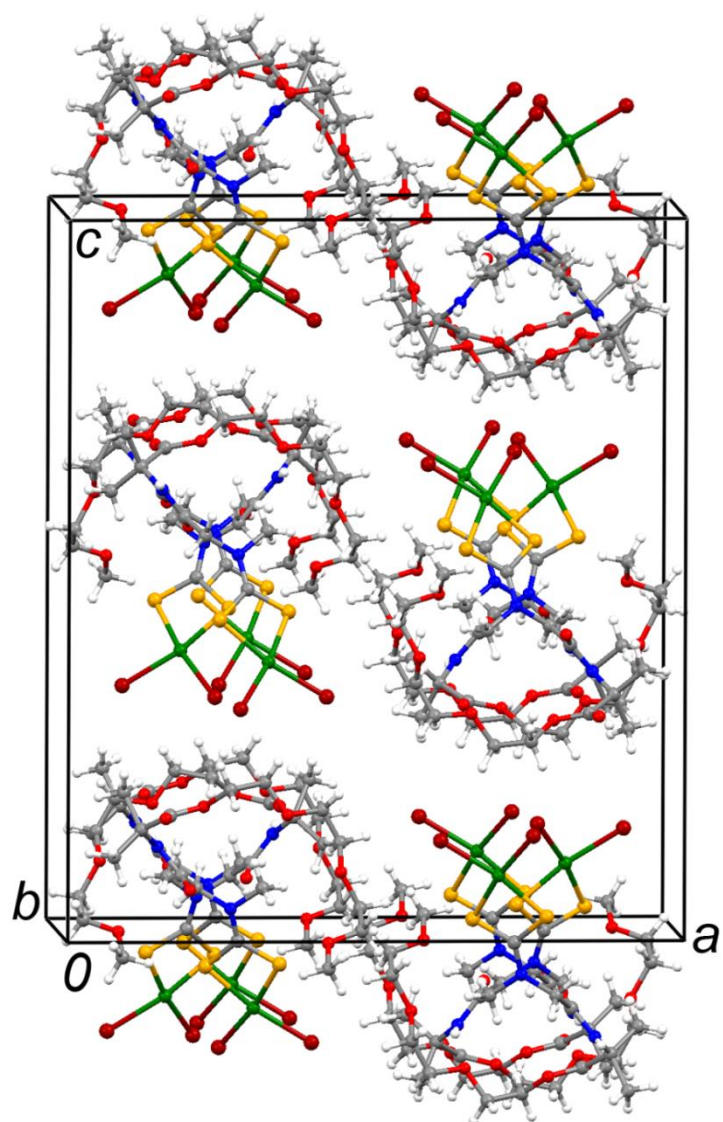


Figure S8. Crystal packing of IT05·pentane viewed along the b crystallographic axis.

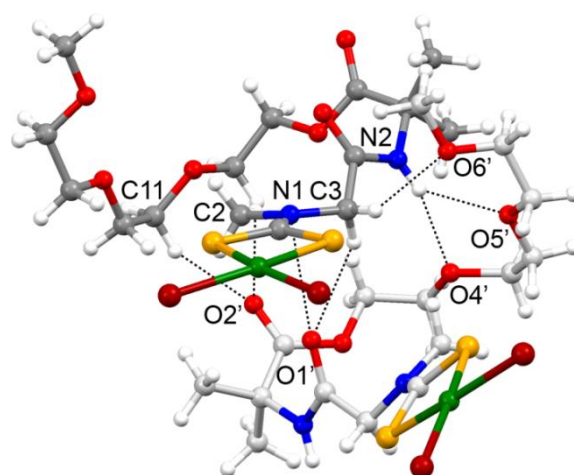


Figure S9. Depiction of intermolecular contacts between two symmetry related molecules of IT05-pentane. Symmetry code ' = $1/2-x; 1/2+y; z$.

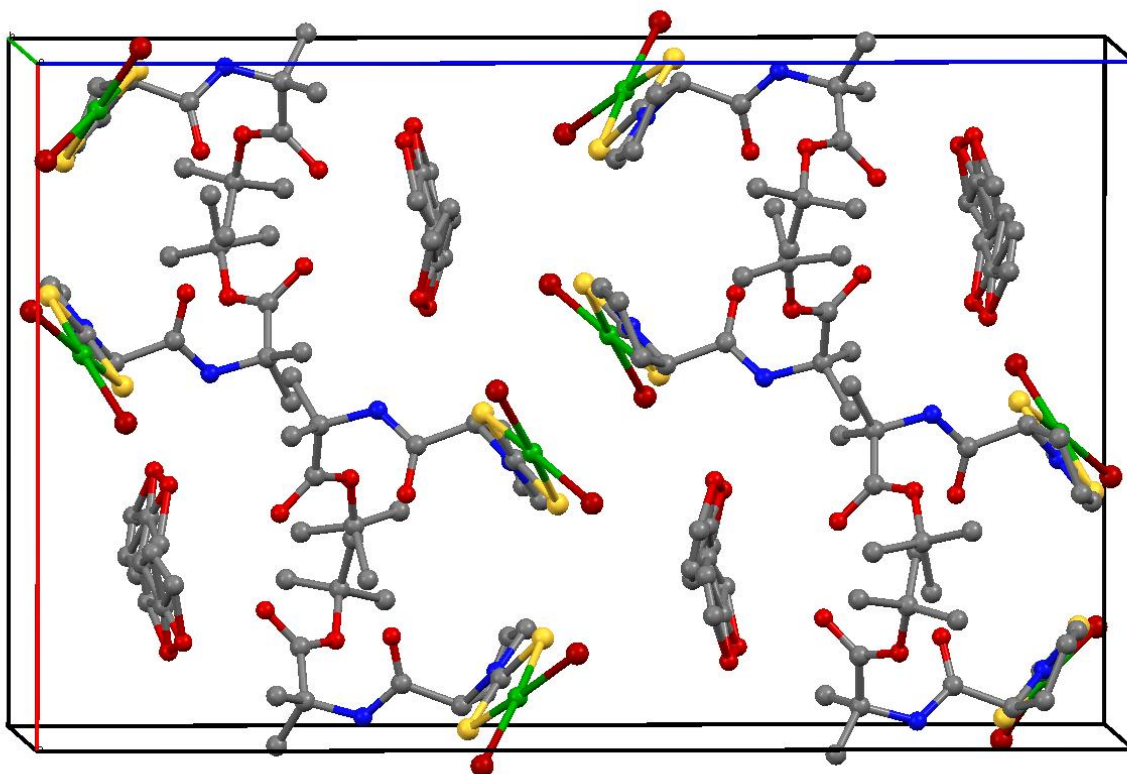


Figure S10. Crystal packing of IT03/IT04 acetone solvate viewed along the b crystallographic axis.

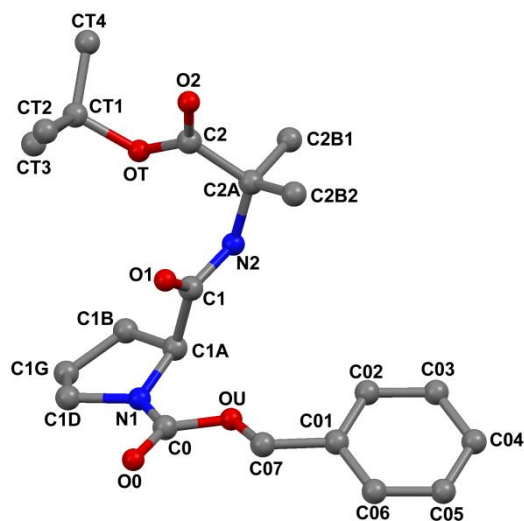


Figure S11. X-ray diffraction structure of Z-L-Pro-Aib-OtBu with atom numbering (only one position for the disordered C1G atom is shown; H-atoms have been omitted for clarity), characterized by a *cis* configuration of the Z-L-Pro urethane bond [the value of the OU-C0-N1-C1A torsion angle being $-17.5(6)^\circ$], a not uncommon observation for urethane-protected Pro derivatives. The peptide backbone features a *semi*-extended conformation for the Pro residue [$\phi, \psi = -60.0(5)^\circ, +168.8(3)^\circ$] while right-handed helical for Aib [$\phi, \psi = -53.1(5)^\circ, -31.8(5)^\circ$]. Such a conformation is closely matched by the L-Pro-Aib segment in the structure of the racemate IT03/IT04. In the packing mode, an intermolecular H-bond is observed between the (peptide) N2-H group and the $(-x, 1/2+y, 3/2-z)$ symmetry equivalent of the (urethane carbonyl) O0=C0 group, linking molecules related by a twofold screw axis along the *b* direction.

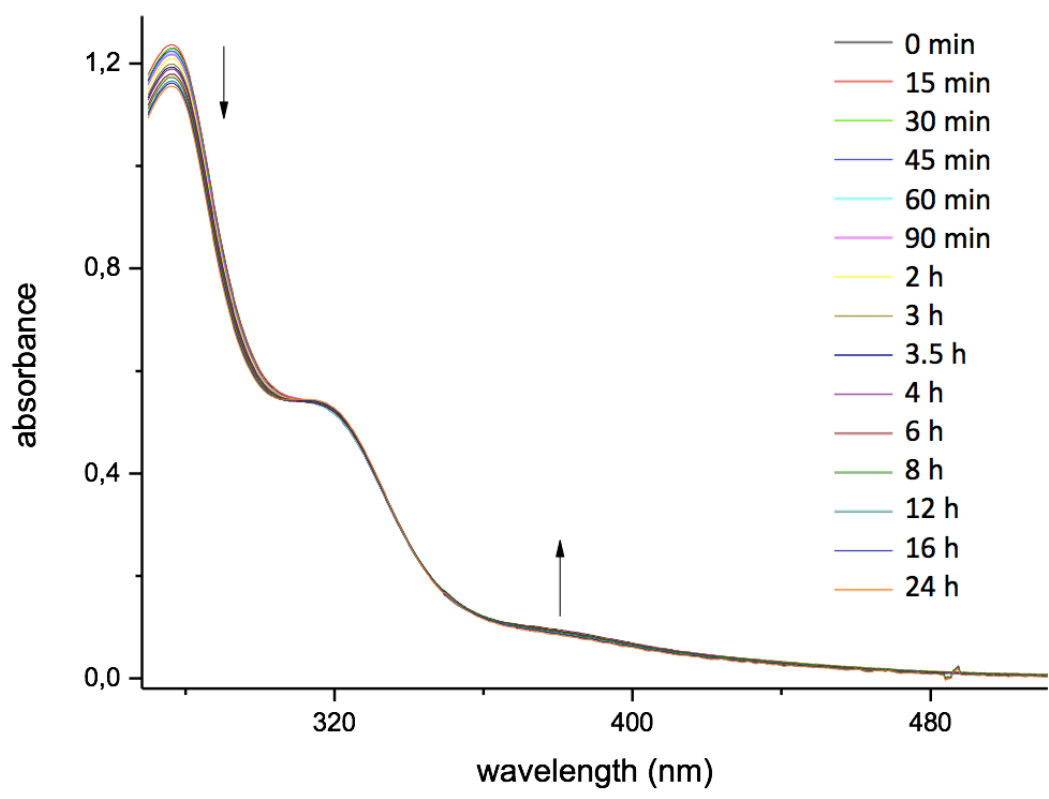


Figure S12. Electronic spectra of IT03 (50 μM) recorded in DMSO at 25°C for 24 h.

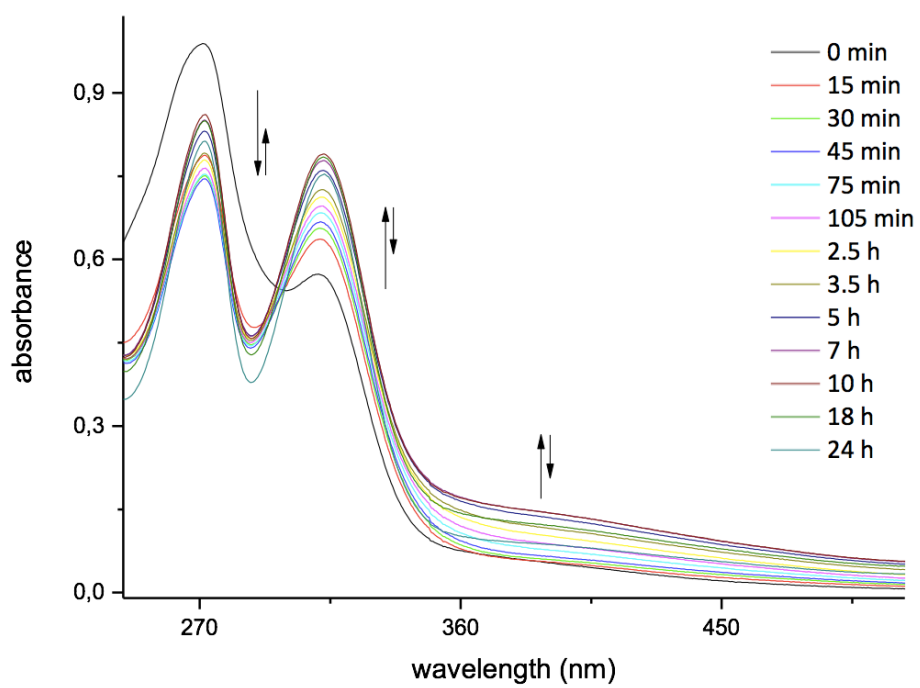


Figure S13. Electronic spectra of IT03 (50 μM) recorded in deionized water at 25°C for 24h.

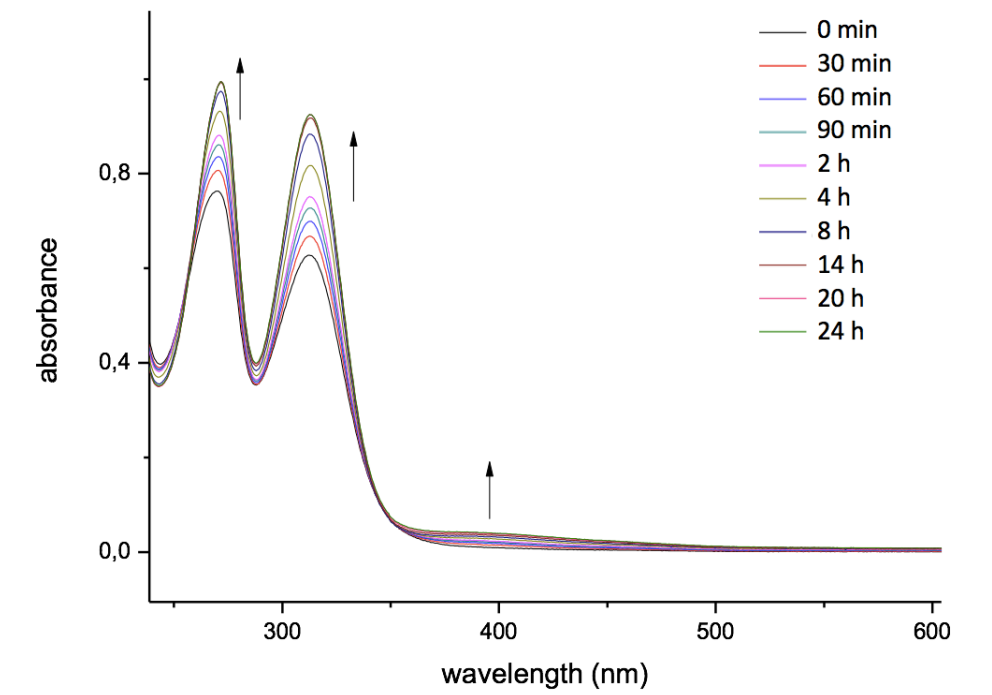


Figure S14. Electronic spectra of IT03 (50 μM) recorded in saline solution (NaCl 0.9% w/v) at 25°C for 24 h.

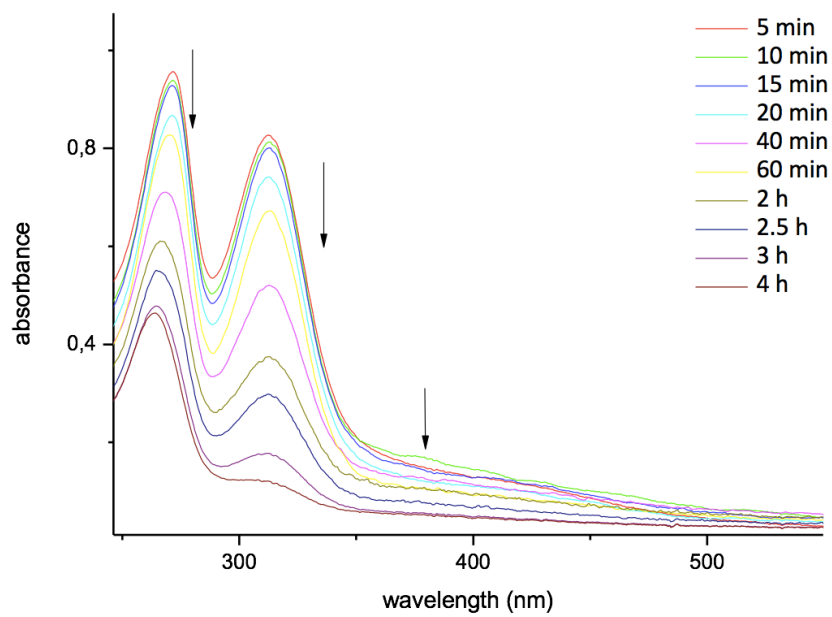


Figure S15. Electronic spectra of IT03 (50 μM) recorded in PBS buffer pH=7.4 at 25°C over 4 h.

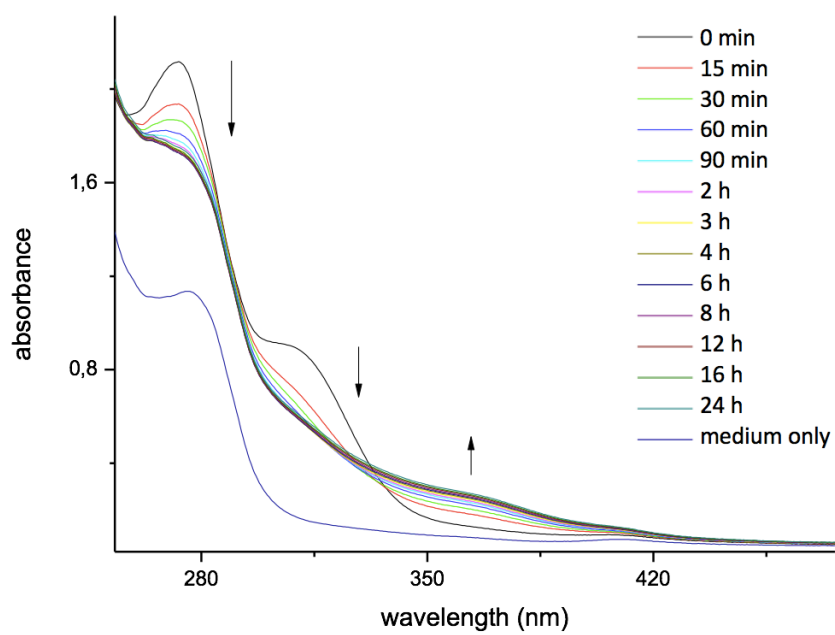


Figure S16. Electronic spectra of IT03 (30 μM) in PBS added of 20% v/v complete medium (RPMI-1640 medium supplemented with 10% fetal calf serum) at 25°C for 24 hours.

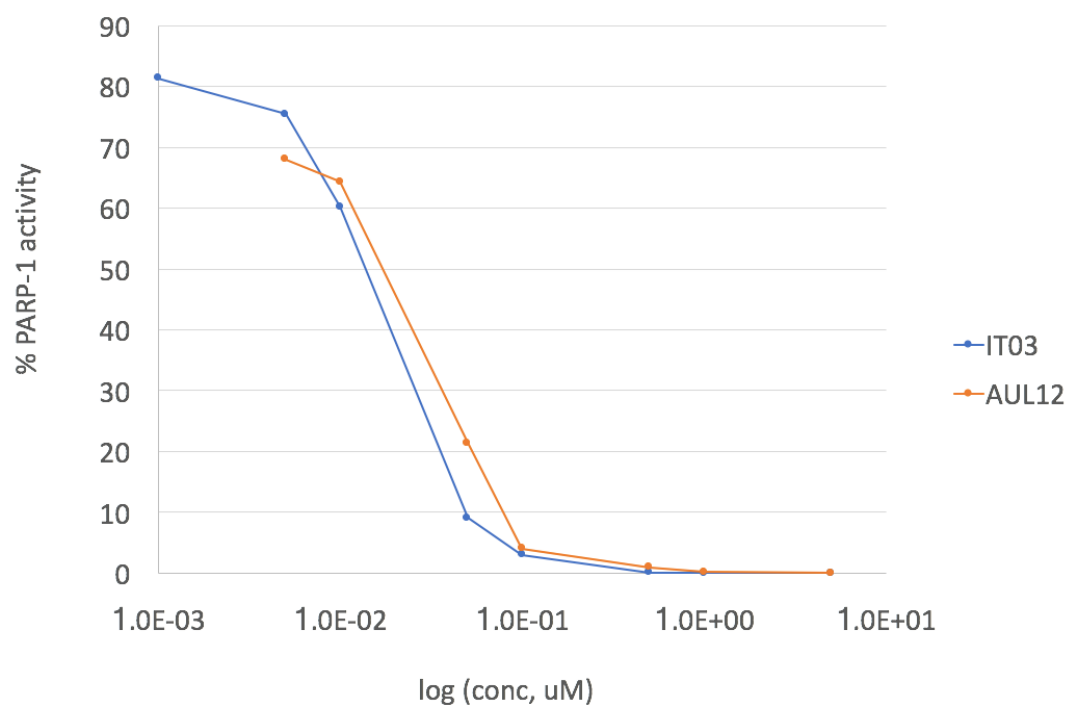


Figure S17. Inhibition of PARP-1 activity by test compounds at different concentrations (from 10 to 0.01 μM). For the assay each compound was incubated for 1 h with 0.5 U of purified protein.

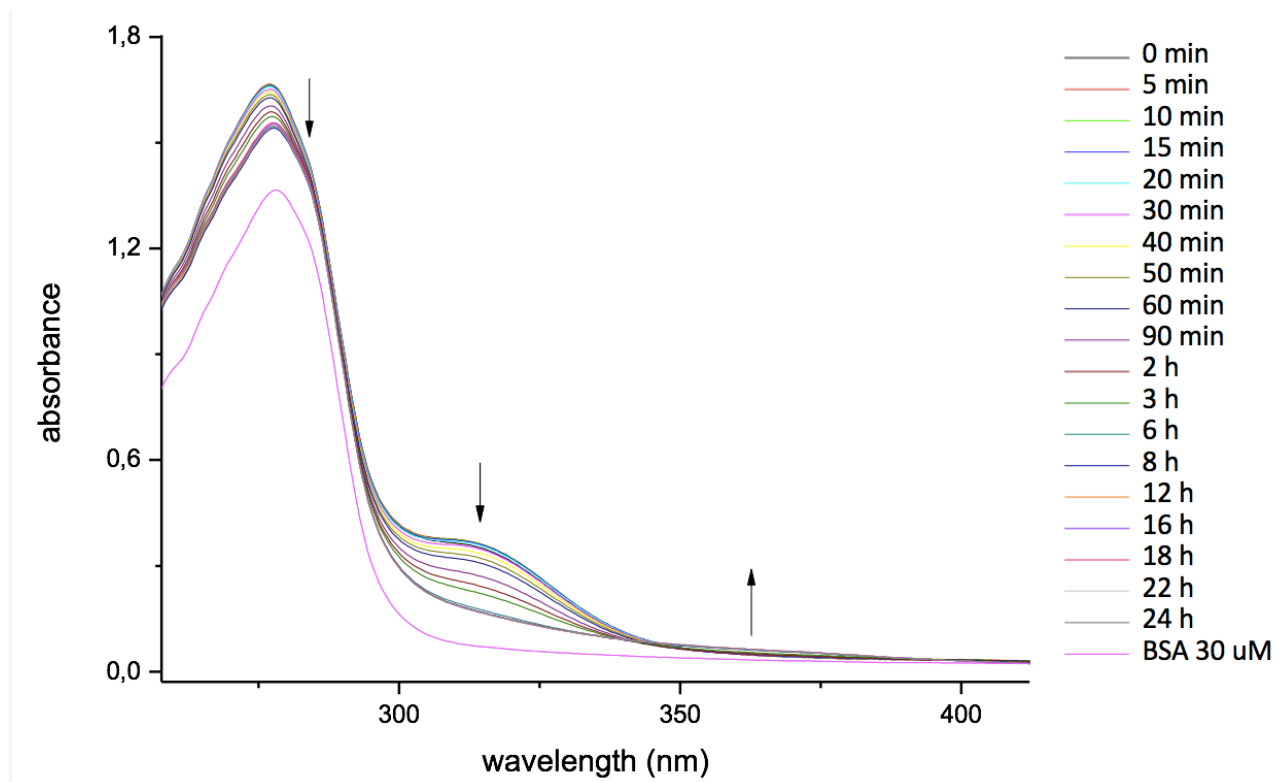


Figure S18. Electronic spectrum of BSA and IT03 (equimolar, 30 μM) in PBS at 25°C over 24 h.

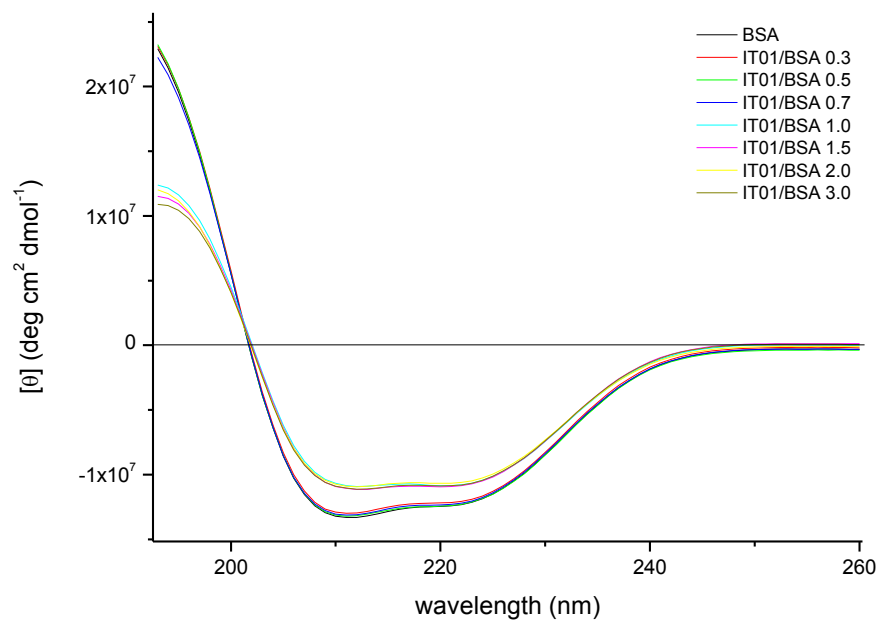


Figure S19. CD spectra of BSA incubated with IT03 at different molar ratios (from 0 to 3) in PBS at 25°C.

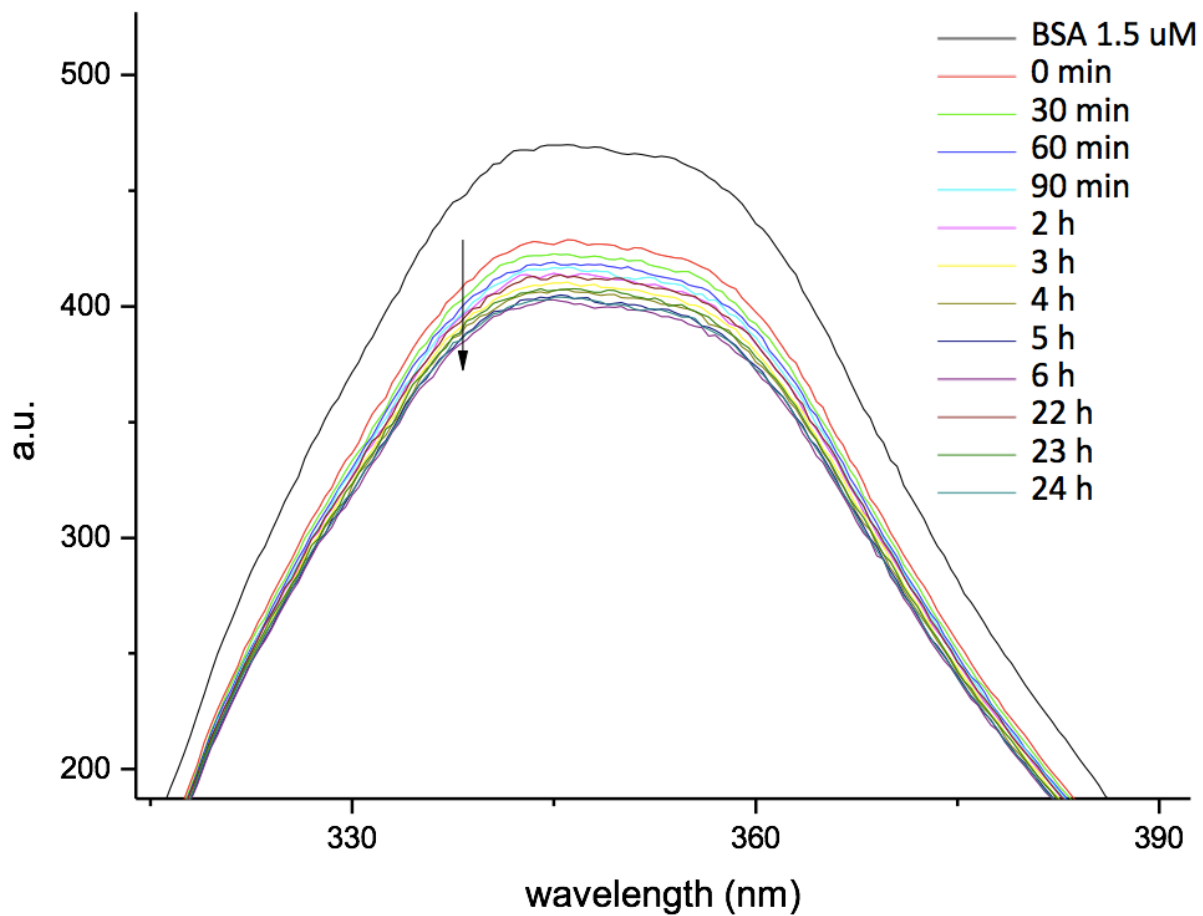


Figure S20. Fluorescence spectra of BSA (15 μM) in presence of IT03 (15 μM) in PBS at 25°C over 24h.

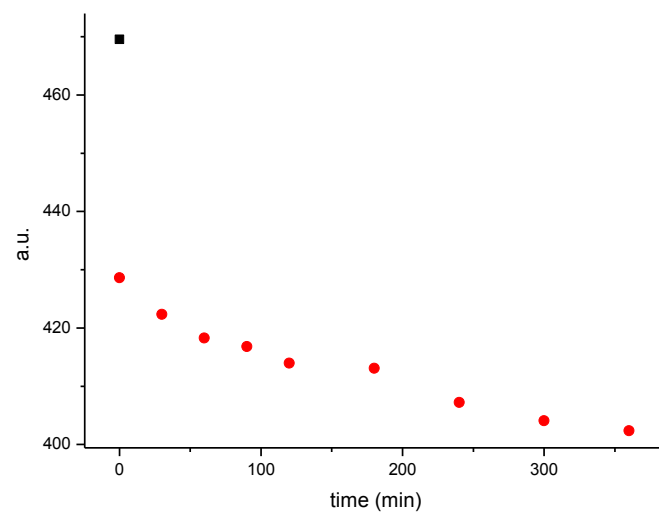


Figure S21. Intensity fluorescence at the λ_{em} maximum (349 nm) of BSA alone (■) and when incubated with IT03 as a function of time (●, 0-6 h).

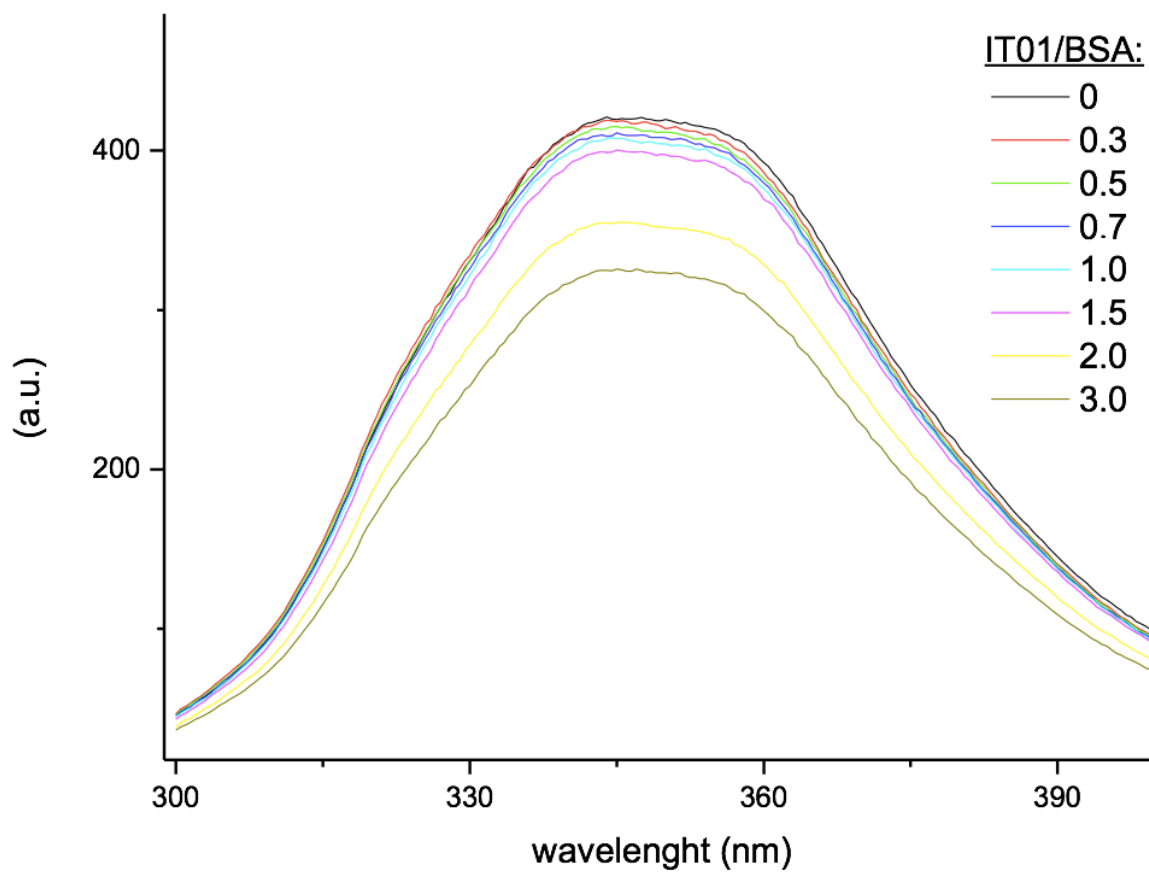


Figure S22. Fluorescence spectra of BSA (15 μM) in presence of IT03 at different molar ratios (from 0 to 3) soon after mixing, in PBS at 25°C.

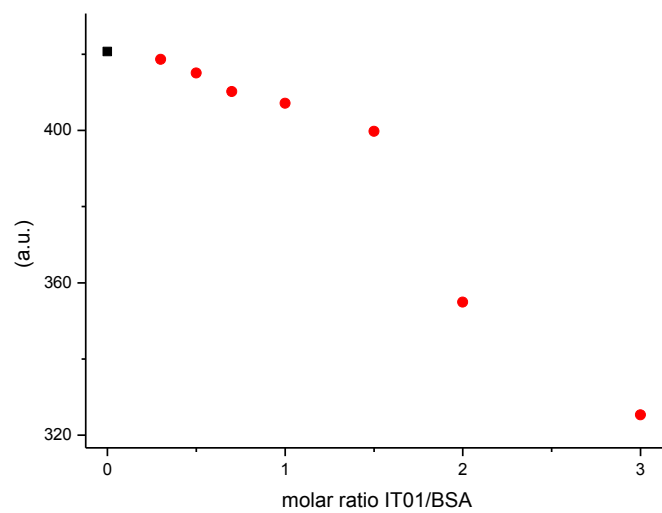


Figure S23. Intensity fluorescence at the λ_{em} maximum (349 nm) of BSA alone (■) and when incubated with IT03 at different IT03/BSA molar ratios (From 0.3 to 3; ●).