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Packing polymorphism of dicarbonyl-[2-(phenylamino)pent-3-en-4-onato]rhodium(I)



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ABSTRACT

Depending on the crystallization conditions, the interaction between the rhodium metal centres of the separate $[Rh(CH_3COCHCN(Ph)CH_3)(CO)_2]$ molecular units, as described by the interplanar separation and lateral shift of two of the units, leads to packing polymorphism of $[Rh(CH_3COCHCN(Ph)CH_3)(CO)_2]$, which means the same molecule crystallises in different fashions, resulting in different polymorphs (α and β), with a difference in crystal packing. Six different sets of solid state single crystal data of $[Rh(CH_3COCHCN(Ph)CH_3)(CO)_2]$, show that this complex is polymorphic, forming dinuclear units that either stack in wire-like chains with weak metallophilic rhodium-rhodium interactions (β -polymorph), or with packing of the dinuclear units that does not result in rhodium-rhodium chains (α -polymorph). A DFT study on the inter-molecular interactions in different dinuclear $[Rh(CH_3COCHCN(Ph)CH_3)(CO)_2]_2$ models, using different DFT methods, provides an understanding on a molecular level of the rhodium-rhodium and other inter-molecular interactions between the two separate $[Rh(CH_3COCHCN(Ph)CH_3)(CO)_2]$ molecules in the dinuclear unit.

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1. Introduction

Polymorphism [1] has been defined by McCrone in 1965 as "a solid crystalline phase of a given compound resulting from the possibility of at least two different arrangements of the molecules of that compound in the solid state" [2]. A few years later, Rosenstein and Lamy (1969) simplified the definition to "when a substance can exist in more than one crystalline state it is said to exhibit polymorphism" [3]. Polymorphs are also described as "isomers at the individual crystal level in which the same molecule or salt crystallises in different fashions, which frequently, but not necessarily, result in differences in the space group and the cell dimensions", sometimes leading to different arrangements (packing) of the molecules in the solid state [4]. When polymorphism exists as a result of differences in crystal packing, it is called packing polymorphism. It has been observed that the molecules of different $[Rh(\beta-diketonato)(CO)_2]$ complexes in the solid state, either pack in separate dinuclear [Rh(β diketonato)($(CO)_2$)₂ dimeric units [5–7], while others of these complexes pack in dinuclear units which form extended linear wirelike chains of rhodium atoms [8-11]. The extended metal-metal interactions along the linear chains in the latter case, have been described as resulting from the overlap of the filled d_{z^2} orbitals and the empty p_z orbitals that lie along the direction of the metal-metal axis [12]. Similarly to the latter, it has been shown recently that crystals of the complex [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] (dicarbonyl-[2-(phenylamino)pent-3-en-4-onato]rhodium(I)) [13] (see Fig. 1), in the solid state also pack in linear chains of consecutive unit cells along the intermolecular rhodium-metal axis in a wire-like structure (called the β-polymorph), while a related molecule of dicarbonyl-(4-((2,6-dichlorophenyl)imino)pent-2-en-2-olato)-rhodium did not show any inter-molecular rhodium-rhodium interaction in the solid state at all [13]. In this contribution we present an additional packing polymorphism (the α-polymorph) of the same [Rh(CH₃COCHCN(Ph) CH₃)(CO)₂] molecule (as in Fig. 1), complemented by a DFT study on the inter-molecular rhodium-rhodium interactions in four different dinuclear [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ models.

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Fig. 1. Structure of [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂].

2. Experimental

2.1. Synthesis

Reagents were obtained from Sigma-Aldrich. Solid reagents employed in preparations were used directly without further purification. Solvents were distilled prior to use. Di- μ -chloro-tetra-carbonyldirhodium(I) was obtained from Sigma-Aldrich.

2.1.1. [CH₃COCHCN(HPh)CH₃]

The ligand [CH₃COCHCN(HPh)CH₃] was synthesised using published methods [14–17] with slight modifications. Acetylacetone (5 g, 5.2 ml) was placed in a round bottom flask outfitted with a condenser for refluxing. Aniline (6.9763 g, 7 ml) and concentrated hydrochloric acid (5 g, 4.3 ml) was added to the flask whilst stirring and the mixture was brought to reflux for 5 h. Diethyl ether was added to the mixture and then placed in the fridge (-5° C). The yellow crystalline precipitate was filtered and washed with cold diethyl ether and recrystallised.

 $\label{eq:Yield} Yield = 50\%. \ ^1H \ NMR; \ 12.493 \ ppm \ (s, O-H); \ 7.388 \ ppm-7.114 \ ppm \ (m, C_6H_5-N); \ 5.208 \ ppm \ (s, C-H); \ 2.122 \ ppm \ (s, CH_3-CN); \ 2.014 \ ppm \ (s, CH_3-CO).$

2.1.2. [Rh(CH3COCHCN(Ph)CH3)(CO)2]

The complex [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] was synthesised using published methods with slight modifications [18]. The metal, di- μ -chloro-tetracarbonyldirhodium(I) (0.05 g, 0.1286 mmol), was dissolved in methanol (5 ml). The ligand, [CH₃COCHCNHPhCH₃] (0.2572 mmol), also dissolved in methanol (2 ml) was added dropwise during 10 min whilst stirring. The mixture was left to stir for 1 h. The mixture was then extracted with n-hexane until the n-hexane solvent became clear. The solvents were combined and evaporated under reduced pressure. The precipitated solid was collected and recrystallised, either from n-hexane at -5 °C, or from DCM at room temperature (25 °C).

Yield = 78%. ¹H NMR: 7.388 ppm-7.061 ppm (m, C₆H₅-N); 5.298 ppm (s, C-H); 2.195 ppm (s, CH₃-CN); 2.141 ppm (s, CH₃-CO). ν_{CO} : 2059 cm⁻¹; 1998 cm⁻¹. UV/Vis (CHCl₃): $\lambda_{\text{max}} = 329$ nm ($\varepsilon = 5102 \text{ mol}^{-1} \text{ dm}^{+3} \text{ cm}^{-1}$), 265 nm ($\varepsilon = 6497 \text{ mol}^{-1} \text{ dm}^{+3} \text{ cm}^{-1}$).

2.2. Crystal structure analysis

The solid state crystal data of two crystals, obtained under different crystallisation conditions, collected at three different temperatures, for two different crystalline forms of [Rh(CH₃COCHCN(Ph) CH₃)(CO)₂], is presented here. The first crystalline form, labelled the α -polymorph, was obtained from a solution of n-hexane at -5 °C. The second crystalline form, labelled the β -polymorph, was crystallised from a solution of DCM at room temperature. Data for both crystals was collected, both at RT and at 150 K for each of these two

crystalline forms (α - and β -polymorphs), on a Bruker D8 Venture kappa geometry diffractometer, with duo Iµs sources, a Photon 100 CMOS detector and APEX II [19] control software, using Quazar multi-layer optics, and monochromated Mo- $K\alpha$ radiation, by means of a combination of φ and ω scans. An additional set of data for the α -polymorph was collected at 100 K, on a Bruker APEX-II CCD.

Data reduction was performed using SAINT+ [19] and the intensities were corrected for absorption, using SADABS [19]. The five crystal structures were solved by intrinsic phasing, using SHELXTS, and refined by full-matrix least squares, using both SHELXTL+ [20] and SHELXL-2014+ [20]. In the structure refinement, all hydrogen atoms were added in the calculated positions and treated as riding on the atom to which they are attached. All non-hydrogen atoms were refined with anisotropic displacement parameters; all isotropic displacement parameters for hydrogen atoms were calculated as (X \times Ueq) of the atom to which they are attached, where X = 1.5 for the methyl hydrogens, and X = 1.2 for all other hydrogens. Crystal data, data collections, structure solutions and refinement details for all five crystal structures, are available in the CIFs (with CCDC deposit numbers 1548178, 1558071, 1571132, 1571133 and 1571134).

2.3. Theoretical approach

Density functional theory (DFT) calculations of this study were performed with the hybrid B3LYP [21,22] functional, as implemented in the Gaussian 09 program package [23]. Geometries of the neutral complexes were optimised in the gas phase, using the 6-311G(d,p) basis set on all atoms except rhodium, for which the def2tzvpp [24] (valens electrons) and SDD (core electrons) basis set was used. These B3LYP optimised gas phase structures were used to further conduct a natural bond orbital (NBO) analysis (using the NBO 3.1 module [25] in Gaussian 09), a fragment analysis, as well as an electronic density analysis (using Bader's quantum theory of atoms in molecules (QTAIM) [26–28], as implemented in ADF2013 [29–31]), at the same level of theory.

The single molecular unit of complex [Rh(CH₃COCHCN(Ph) CH₃)(CO)₂] was optimised by calculation, as well as the [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ dimeric unit. Further additional single point calculations were done on the coordinates of such a dinuclear unit which were obtained from the solid state crystal structure data of this study (α -polymorph at 150 K), as well as on two additional dinuclear models, calculated on the coordinates of a dinuclear unit obtained from the solid state crystal structure data of a previous study (β -polymorph at 100 K), as described in Section 3.2.2. The coordinates of the DFT calculations are given in the Supporting Information.

3. Results and discussion

3.1. X-ray structure

In this section, crystallographic results are presented of the solid state crystal data of two polymorphs of complex [Rh(CH₃COCHCN(Ph) CH₃)(CO)₂]; see Table 1 and Table 2 (column 2–6). Data for each crystalline structure was repeatedly collected at decreasing temperatures, to determine whether temperature would transform the structure of each polymorph. The two structures were labelled the α -polymorph (with space group P 2_1 /n, data collected at 298 K, 150 K and 100 K), as well as the β -polymorph (with space group I2/a, data collected at 298 K and 150 K). These experimental results were then compared to a previously published structure [13] and packing of the same complex (β -polymorph with space group I2/a, data collected at 100 K); see Table 2 (last column of the β -polymorphs). Crystals suitable for X-ray diffraction were obtained by slow evaporation from

Table 1 Crystal data and structure refinement of both the α -and β -polymorphs of complex [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂], collected for two crystals at three different temperatures (298 K, 150 K and 100 K).

Polymorph	α	α	α	β	β
Empirical formula	C ₁₃ H ₁₂ N O ₃ Rh	C ₁₃ H ₁₂ N O ₃ Rh	C ₁₃ H ₁₂ N O ₃ Rh	C ₁₃ H ₁₂ N O ₃ Rh	C ₁₃ H ₁₂ N O ₃ Rh
Formula weight	333.15	333.15	333.15	333.15	333.15
Temperature	298(2) K	150(2) K	100(2) K	298(2) K	150(2) K
Wavelength	0.71073 Å				
Crystal system	Monoclinic	Monoclinic	Monoclinic	Monoclinic	Monoclinic
Space group	P 2 ₁ /n	P 2 ₁ /n	P 2 ₁ /n	I 2/a	I 2/a
Unit cell dimensions	a = 8.7171(4) Å	a = 8.5599(4) Å	a = 8.5127(5) Å	a = 13.6872(10) Å	a = 13.4204(14) Å
	b = 12.0141(5) Å	b = 11.9808(6) Å	b = 11.9666(7) Å	b = 9.3364(16) Å	b = 9.3030(8) Å
	c = 13.1833(6) Å	c = 13.1030(7) Å	c = 13.0725(8) Å	c = 21.350(3) Å	c = 21.226(3) Å
	$lpha=90^\circ$	$\alpha=90^{\circ}$	$\alpha=90^{\circ}$	$lpha=90^\circ$	$lpha=90^\circ$
	$\beta = 105.6850(10)^{\circ}$	$\beta = 105.775(2)^{\circ}$	$\beta = 105.874(2)^{\circ}$	$\beta = 95.000(4)^{\circ}$	$\beta = 95.817(4)^{\circ}$.
	$\gamma=90^{\circ}$	$\gamma=90^{\circ}$	$\gamma=90^{\circ}$	$\gamma=90^\circ$	$\gamma=90^{\circ}$
Volume	1329.25(10) Å ³	1293.16(11) Å ³	1280.89(13) Å ³	2717.9(6) Å ³	2636.4(5) Å ³
Z	4	4	4	8	8
Density (calculated)	1.665 Mg/m ³	1.711 Mg/m ³	1.728 Mg/m ³	1.628 Mg/m ³	1.679 Mg/m^3
Absorption coefficient	1.283 mm ⁻¹	1.319 mm ⁻¹	1.332 mm ⁻¹	1.255 mm ⁻¹	1.294 mm ⁻¹
F(000)	664	664	664	1328	1328
Crystal size	$0.185 \times 0.158 \times 0.148 \text{ mm}^3$	$0.185 \times 0.158 \times 0.148 \text{ mm}^3$	$0.185 \times 0.158 \times 0.148 \text{ mm}^3$	$0.480 \times 0.120 \times 0.110 \text{ mm}^3$	$0.480 \times 0.120 \times 0.110 \text{ mm}^3$
Theta range for data collection	2.520 to 28.387°	2.345 to 25.339°	3.014 to 25.349°	2.382 to 26.408°.	2.392 to 28.536°
Index ranges	$-11 \le h \le 11$,	$-10 \le h \le 10$,	$-10 \le h \le 10$,	$-17 \le h \le 17, -11 \le k \le 11,$	$-17 \le h \le 17, -12 \le k \le 12, -28 \le l \le 2$
•	$-16 \le k < =16, -17 \le l < =17$	$-14 \le k < =14, -15 \le l < =15$	$-14 \le k <= 14, -15 \le l <= 15$	-26 ≤ l<=26	
Reflections collected	48660	14021	49469	33261	41844
Independent reflections	3347 [R(int) = 0.0335]	2370 [$R(int) = 0.0252$]	2350 [$R(int) = 0.0182$]	2789 [R(int) = 0.0863]	3342 [R(int) = 0.0632]
Completeness to theta $= 25.242^{\circ}$	100%	99.9%	99.9%	100.0%	100.0%
Refinement method	Full-matrix least-squares on F ²				
Data/restraints/parameters	3347/0/165	2370/0/163	2350/0/165	2789/0/165	3342/0/165
Goodness-of-fit on F ²	1.041	1.043	1.134	1.013	1.040
Final R indices [I > 2sigma(I)]	R1 = 0.0214, $wR2 = 0.0591$	R1 = 0.0177, $wR2 = 0.0440$	R1 = 0.0146, $wR2 = 0.0375$	R1 = 0.0344, $wR2 = 0.0582$	R1 = 0.0232, $wR2 = 0.0449$
R indices (all data)	R1 = 0.0318, $wR2 = 0.0668$	R1 = 0.0.021, $wR2 = 0.0456$	R1 = 0.0152, $wR2 = 0.0379$	R1 = 0.0714, $wR2 = 0.0662$	R1 = 0.0376, $wR2 = 0.0479$
Extinction coefficient	n/a	n/a	n/a	n/a	n/a
Largest diff. peak and hole	0.348 and -0.470 e.Å ⁻³	0.499 and -0.183 e.Å ⁻³	0.260 and -0.494 e.Å ⁻³	0.472 and -0.463 e.Å ⁻³	0.410 and -0.439 e.Å ⁻³

Table 2 Selected geometrical data (bond lengths in Å and bond angles in degrees) of complex [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂], obtained for both its α - and β -polymorphs, as well as the optimized DFT calculated data. Data for the two crystals from this study (columns 2 - 6) is compared to the corresponding crystallographic data of a previously published β -polymorph, collected at 100 K [13] (column 7). Atom numbering is shown in Fig. 2.

Polymorph α-polymor		norph		β-polymorph		difference min and	DFT	max difference	
Temperature	298 K	150 K	100 K	298 K	150 K	100 K	max experimental parameter	optimised molecule	DFT and experimental parameter
Rh(1)-C(1)	1.841(2)	1.847(2)	1.8439(17)	1.827(4)	1.844(2)	1.841(3)	0.020	1.855	0.028
Rh(1)-C(2)	1.866(2)	1.870(2)	1.8709(17)	1.857(4)	1.868(2)	1.866(2)	0.014	1.878	0.021
Rh(1)-O(3)	2.0187(14)	2.0239(13)	2.0251(11)	2.022(2)	2.0312(13)	2.032(2)	0.013	2.040	0.021
Rh(1)-N(1)	2.0501(17)	2.0540(16)	2.0557(13)	2.054(3)	2.0596(15)	2.058(2)	0.010	2.092	0.042
O(3)-C(3)	1.284(3)	1.292(2)	1.291(2)	1.282(4)	1.289(2)	1.294(2)	0.012	1.280	0.014
N(1)-C(5)	1.314(2)	1.321(2)	1.320(2)	1.318(4)	1.320(2)	1.318(3)	0.007	1.330	0.016
N(1)-C(6)	1.448(3)	1.443(2)	1.444(2)	1.437(4)	1.441(2)	1.448(3)	0.011	1.439	0.009
C(4)-C(5)	1.419(3)	1.417(3)	1.418(2)	1.395(5)	1.408(3)	1.410(3)	0.024	1.412	0.017
C(3)-C(4)	1.357(3)	1.372(3)	1.374(2)	1.376(5)	1.380(3)	1.381(3)	0.024	1.390	0.033
C(1)-Rh(1)-C(2)	86.79(10)	87.03(9)	86.98(7)	86.54(16)	86.60(9)	86.5(1)	0.53	89.0	2.54
C(2)-Rh(1)-O(3)	89.52(8)	89.57(7)	89.72(6)	89.33(13)	89.66(7)	89.75(8)	0.42	94.3	4.98
C(1)-Rh(1)-N(1)	93.06(8)	92.67(7)	92.67(6)	93.72(13)	93.26(8)	93.13(9)	1.05	89.4	4.28
O(3)-Rh(1)-N(1)	90.62(6)	90.72(6)	90.63(5)	90.41(10)	90.47(6)	90.55(6)	0.31	89.4	1.28
N(1)-C(5)-C(4)	123.07(19)	123.76(18)	123.68(14)	123.8(3)	124.38(18)	123.9(2)	1.31	124.6	1.51
C(3)-C(4)-C(5)	128.09(19)	127.50(18)	127.42(14)	128.3(3)	127.57(19)	127.8(2)	0.88	127.2	1.12
O(3)-C(3)-C(4)	126.38(19)	126.33(18)	126.37(14)	125.4(3)	125.66(18)	125.7(2)	0.98	125.8	0.60
Rh·····Rh	3.5560(3)	3.4816(3)	3.4577(3)	3.4204(5);	3.3272(5);	3.3085(2);		1.855	
				3.4945(5)	3.4425(5)	3.4358(2)			

either an n-hexane solution at -5 °C (to yield the α -polymorph), or from a DCM solution at RT (to yield the β -polymorph). Both polymorphs exhibit the same experimental properties, such as colour or UV/vis, IR and 1 H NMR spectra as provided in the experimental section. The different polymorphs seemed to result from different crystallisation conditions (solvent and temperature) and did not transform into one another by cooling, since it was found by crystallography that the space group of a specific polymorph stayed the same upon cooling from 298 K to 150 K or 100 K; see Table 1.

3.1.1. Structure

The crystal data and structure refinement of the two $[Rh(CH_3COCHCN(Ph)CH_3)(CO)_2]$ crystals of this study are summarised in Table 1, while Table 2 compares selected geometrical data of the two crystals from this study (obtained for both the α -and β -polymorphs, from data collections at different temperatures) to the corresponding data of a previously published β -polymorph, collected at 100 K [13]. A molecular diagram of the α -polymorph at 150 K, showing the atom labelling used for all six structures, is illustrated in Fig. 2.

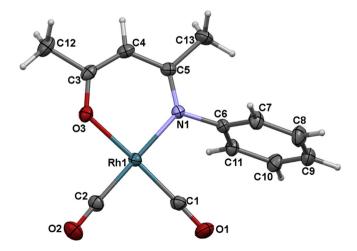


Fig. 2. A perspective drawing of the molecular structure of [Rh(CH₃COCHCN(Ph) CH₃)(CO)₂] (the α -polymorph at 150 K), showing the atom numbering scheme. Atomic displacement parameters (ADPs) are shown at the 50% probability level.

The α-polymorph of [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] crystallised in the P2₁/n space group, with four molecules per unit cell. The four bond angles around the rhodium atom deviate slightly (<3°) from the normally expected 90° bond angle for a square planar complex, as can be seen from the bond angles of the α -polymorph at 150 K (see Table 2, second column of the α -polymorphs): namely, bond angle C(1)-Rh(1)-C(2) of 87.03(9)°, bond angle C(2)-Rh(1)-O(3) of 89.57(7)°, bond angle C(1)-Rh(1)-N(1) of 92.67(7)° and bond angle O(3)-Rh(1)-N(1) of 90.72(6)°. The atoms around rhodium (N1, O3, C1 and C2) as well as rhodium itself all lie in the same plane, with C1 having the maximum deviation from this plane of 0.031(2) Å. Similarly, the atoms of the backbone of the bidentate ligand (O3, C3, C4, C5 and N1) as well as Rh itself, all lie in another plane (slanted at 0.49° with respect to the plane through Rh, N1, O3, C1 and C2), with O3 having the maximum deviation from that plane of 0.021(1) Å. The angle between the latter plane and the plane through the phenyl ring is 81.70°. The separate molecules form dimers, which are packed about a centre of symmetry.

The β -polymorph of [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] crystallised in the I2/a space group, with eight molecules per unit cell. The structure of the β -polymorph is very similar to that of the α -polymorph of [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂], as can be seen from the structure overlay of the two polymorphs, as illustrated in Fig. 3 (top left).

The metal-ligand distances in transition metal complexes are important in determining the structure and reactivity of the metal complex. Therefore in Table 2 the bond lengths and bond angles around the rhodium atom, as well as selected geometrical parameters involving the backbone of the bidentate ligand of the α - and β-polymorph obtained crystallographically in this study, are compared to each other and also to the corresponding parameters of the previously published structure of the β -polymorph, collected at 100 K [13], of complex [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]. The maximum difference between the selected bond lengths and bond angles of both polymorphs obtained at different temperatures, is only 0.024 Å and 1.3° respectively. The main difference between the geometries of the α - and β -polymorphs lies in the orientation of the phenyl and methyl groups on the ligand, as is clear from the overlay of the different structures, as presented in Fig. 3 (top left). However, the orientation of the phenyl and methyl groups on the ligand did

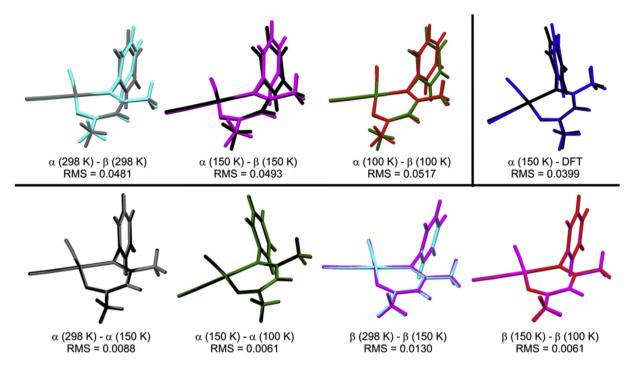


Fig. 3. Overlay of the structures of the α -polymorph (298 K, grey), α -polymorph (150 K, black), α -polymorph (100 K, green), β -polymorph (298 K, cyan), β -polymorph (150 K, magenta) from this study and β -polymorph (100 K, red) from a previous study [13], of [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] with each other. The root means square (RMS) overlay values, when using the Rh atom, the atoms of the backbone of the bidentate ligand, as well as the carbons of the two carbonyl groups of each structure in the overlay, are indicated. Also shown (top right) is the overlay of the experimentally obtained structure of the α -polymorph (150 K, black) of [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] with the theoretical mono nuclear DFT optimised molecule (blue). (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

not change with temperature for a specific polymorph, see Fig. 3 (bottom).

3.1.2. Packing

In Fig. 4, packing diagrams are compared of two crystals in this study, namely of the α-polymorph (collected at 150 K, with space group $P2_1/n$) and the β -polymorph (collected at 150 K, with space group I2/a). Crystals of both polymorphs (α and β) form dinuclear [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ units in the solid state, with differing inter-molecular rhodium-rhodium distances which decrease slightly upon temperature lowering, namely decreasing from 3.5560(3) (298 K) to 3.4816(3) (150 K) to 3.4577(3) (100 K) for the α -polymorph; and decreasing from 3.4204(5) (298 K) to 3.3272(5)(150 K) to 3.3085(2)(100 K) for the β -polymorph (values given in Å, Table 2). This rhodium-rhodium distance is larger in a dinuclear unit of the α -polymorph at the lowest temperature of 100 K, than in a dimer of the β -polymorph at room temperature (298 K). Consequently, the dinuclear [Rh(CH₃COCHCN(Ph) CH₃)(CO)₂|₂ units of the β-polymorphs with shorter intermolecular rhodium-rhodium distances, rather form extended linear chains of metal atoms along the a-axis, regardless of temperature; while packing in the solid state of the α -polymorph does not form any linear chains of rhodium atoms, but only pack in pairs of dinuclear [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ units. In the case of the α -polymorph, the formation of linear wire-like chains of intermolecular rhodium-rhodium atoms is further impeded, due to the larger lateral shift (of ca. 0.9 Å) between the two separate molecules in each dinuclear [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ unit, relative to a smaller lateral shift of only ca. 0.3–0.4 Å between the molecules of the β -polymorph dimer. The difference in crystal packing of the α and β dimers show that these are therefore packing polymorphs, since dinuclear [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ can exist in more than one crystalline state. Packing polymorphism is a result of differences in crystal packing for the same molecule, also leading to differences in the space group and the cell dimensions of the different polymorphs.

The molecular pair-wise packing observed for the α -polymorph (150 K) is further stabilised by parallel ring1 ... ring1 interaction (ring1 defined by Rh1,03, C3, C4, C5, N1) at a distance of 4.465(1) Å, as well as ring1 ... ring2 interaction (ring2 defined by C6, C7, C8, C9, C10, C11) at a distance of 4.607(1) Å between adjacent molecules in each dimeric unit. Some additional interaction of the type X-Y ... ring-centroid (perpendicular) was observed: for C9-H9 ... ring1 the distance is 2.79 Å and for C1-O1 ... ring2 the distance is 3.47 Å.

From the differences found in the solid state packing of all six crystal structures we can therefore conclude that the interaction of the rhodium atoms between two adjacent [Rh(CH₃COCHCN(Ph) CH₃)(CO)₂] molecules can lead to either crystallising as extended linear chains of metal atoms (β -polymorph), or otherwise under differing crystallising conditions (such as solvent and temperature), they do not form infinite metal-metal chains, but crystallise as dinuclear units in the solid state (α -polymorph) instead. It has previously been found that the solid state packing of other related $[M(\beta-diketonato)(CO)_2]$ complexes either displayed extended linear chains of metal atoms [8-10,32-35] in some instances, or otherwise they instead formed dinuclear [M(β-diketonato)(CO)₂]₂ units for other complexes [5–7]. However, to our knowledge, no case of polymorphism has as yet been reported for these related $[M(\beta-diketonato)(CO)_2]$ complexes, as was found in this study for complex [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂].

3.2. Computational chemistry

3.2.1. Structure of mono nuclear [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]

The overlay of the theoretical DFT optimised single molecular unit of $[Rh(CH_3COCHCN(Ph)CH_3)(CO)_2]$, onto the coordinates of the

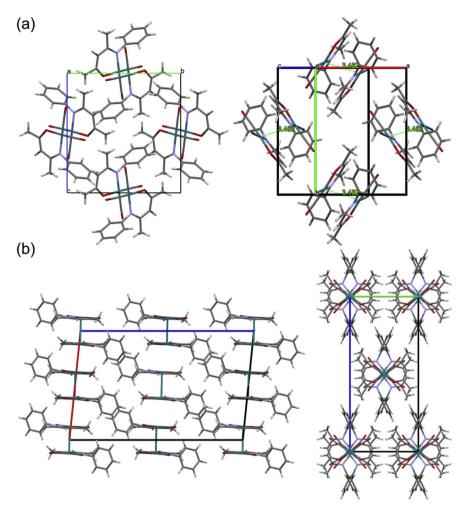


Fig. 4. Packing diagrams of [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] of (a) the α-polymorph (150 K, space group P2₁/n) along the a (top left) and c^* (top right) axis, and of (b) the β-polymorph (150 K, space group I2/a) along the b (bottom left) and a (bottom right) axis.

experimentally obtained solid state data of the α-polymorph (150 K) of this study, is presented in Fig. 3 (top right). The maximum difference between the bond lengths and bond angles around rhodium of any experimental structure and the theoretical DFT optimised bond lengths and bond angles around rhodium, is 0.042 Å and 4.98° respectively (See Table 2 last column). This is comparable with the maximum difference between the selected bond lengths and bond angles of all six crystal structures obtained at different temperatures for both polymorphs reported in this study, namely 0.024 Å and 1.3° respectively (see Table 2, column 8). The slightly overestimated DFT calculated metal-ligand bond lengths are generally observed for gas phase optimisations [36], and therefore considered insignificant. Similar slightly overestimated rhodium-ligand bond lengths for gas phase optimisations, relative to experimentally obtained solid state crystal data, have previously been obtained for related rhodium-β-diketonato complexes [37–39].

3.2.2. Structure of the [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ dimer

Both the α - and β -polymorphs of [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] thus pack in dimeric units in the solid state. An understanding on a molecular level of the rhodium-rhodium and other inter-molecular interactions between the two separate [Rh(CH₃COCHCN(Ph) CH₃)(CO)₂] molecules in the dinuclear unit is presented here. In this section the computational chemistry results, calculated on the coordinates of four models of the [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂

dimeric unit, are described. The four theoretical models will be referred to hereafter as the **DFT optimised dimer 1**, **DFT single point** 2, DFT single point 3 and DFT single point 4 respectively. The first model, DFT optimised dimer 1, is the gas phase optimised dinuclear [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ structure, based on the theoretical coordinates obtained by DFT calculations for a dimeric molecular unit in this study. The second, third and fourth models, involve single point calculations based on crystallographically obtained coordinates, obtained in this and a previous study [13] for dimer [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂, and were chosen so that the lateral shift between adjacent rhodium atoms in the dimeric unit decreases consistently, in going from the first model (DFT optimised dimer 1, lateral shift = 1.590 Å) to the fourth model (**DFT single point 4**, lateral shift = 0 Å), where in the latter case, the two rhodium atoms of adjacent molecular units are aligned directly "on top of each other" (with inclination angle $\alpha = 90^{\circ}$, see Fig. 5), as shown in Fig. 6. The second model (DFT single point 2) was calculated, based on the coordinates of the solid state crystal structure data of dimeric unit of the α -polymorph (at 150 K, with the exact experimental lateral shift of 0.925 Å between adjacent rhodium atoms), while the third model (**DFT single point 3**) was based on the coordinates of the dimeric unit of the β-polymorph (at 100 K, from a previous study [13], with the exact experimental lateral shift of 0.310 Å between adjacent rhodium atoms). The fourth model (DFT single point 4) was conducted on the coordinates of the same previously published βpolymorph crystal structure (at 100 K, as used for **DFT single point**

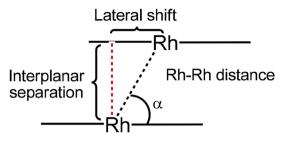


Fig. 5. Description of inter-molecular distances between two adjacent molecular units of the dimer $[Rh(CH_3COCHCN(Ph)CH_3)(CO)_2]_2$. The plane through each of the two molecules is defined by Rh1-O3-N1-C2-C1.

3), however with the two molecular units being aligned directly above each other, with the rhodium-rhodium axis near perpendicular to the square plane through the molecules, i.e. with the lateral shift between the two separate molecular units reduced to near 0 Å (inclination $\alpha = 90^{\circ}$, see Fig. 5). The DFT calculated electronic energies of DFT single point 2 (α-polymorph at 150 K with lateral shift = 0.925 Å), DFT single point 3 (β -polymorph at 100 K with lateral shift = 0.310 Å) and **DFT single point 4** (with no lateral shift) are very close to each other (namely 8.42 eV \approx 8.28 eV \approx 8.30 eV respectively), but all of them are ca. 8.3 eV higher than the **DFT** optimised dimer 1, see Table 3. The interactions between the two separate mono nuclear units [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] in each model, can collectively be described by the rhodium-rhodium distances, the interplanar separation between the two units, the lateral shift between the two individual units, as well as the inclination angle a, between the Rh-Rh vector and the plane through one molecular unit, as defined in Fig. 5 [4] and as summarised in Table 3. The four models are visualised in Fig. 6, indicating the lateral shift between adjacent rhodium atoms in red. It is clear that the lateral shift decreases in going from the **DFT optimised dimer 1** (1.590 Å) to **DFT single point 2** (0.925 Å) to **DFT single point 3** (0.310 Å) to **DFT single point 4** (0 Å), where in the latter, the two rhodium atoms of adjacent molecular units are aligned directly "on top of each other" (inclination angle $\alpha = 90^\circ$, see Fig. 5). It is expected that in the parallel aligned dimers of **DFT single point 4**, wire-like linear metal-metal stacking of the [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ units will be favoured, as has also been observed in the three experimental crystal structures of the β-polymorph of [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ (listed in Table 2), as well as in the solid state structure of a related complex from literature, namely [Rh(acetylacetono)(CO)₂] [12].

3.2.2.1. Molecular orbitals. The character of the highest occupied molecular orbital (HOMO) of the DFT optimised molecule [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] (mono nuclear unit), is distributed over the backbone of the 2-(phenylamino)pent-3-en-4-onato ligand, while the HOMO-1 is of mainly d_{z^2} character on rhodium (Fig. 7 left). The order of these two orbitals is opposite to what has previously been found for complexes [Rh(β -diketonato)(CO)₂] [11,40] and [Ir(acetylacetonato)(CO)₂] [41], where the HOMO was of mainly d_{z^2} character on the metal, instead of on the ligand. Interaction between these two MOs (ligand HOMO and d_{z^2} HOMO-1) from each separate mono nuclear unit in dinuclear [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂, leads to the top four occupied molecular orbitals of the dinuclear [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ pair, namely two bonding MOs (one ligand based and the other rhodium d_{z^2} based) and two antibonding MOs (also one ligand based and the other rhodium d_{z^2}

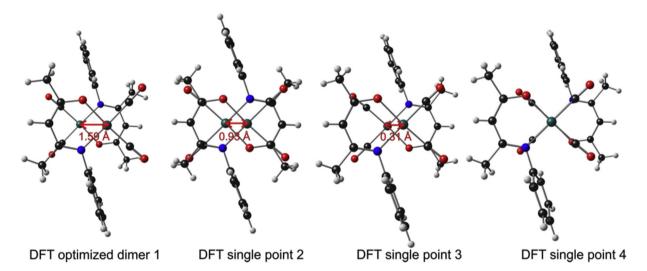


Fig. 6. Visualisation of the lateral shift between two adjacent molecular units (indicated in red), in each of the four different DFT models of the dimer [Rh(CH₃COCHCN(Ph) CH₃)(CO)₂]₂, namely 1.590 Å > 0.925 Å > 0.310 Å) > 0 Å, as seen directly from top. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

Table 3

DFT calculated energies and inter-molecular distances for four different [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ dimer models, with the latter three models based on coordinates obtained from crystallographic data, while the first model is based on theoretically calculated coordinates for a dimeric molecular unit of the complex.

Complex	Relative electronic energy (eV)	Rh-Rh distance (Å)	Angle α (°)	Interplanar separation (Å)	Lateral shift (Å)
DFT optimised dimer 1	0.00	4.474	69.8	4.182	1.590
DFT single point 2 (α-polymorph at 150 K)	8.42	3.482	73.9	3.357	0.925
DFT single point 3 (β-polymorph at 100 K)	8.28	3.309	77.6	3.294	0.310
DFT single point 4 (with no lateral shift)	8.30	3.309	90.0	3.309	0.000

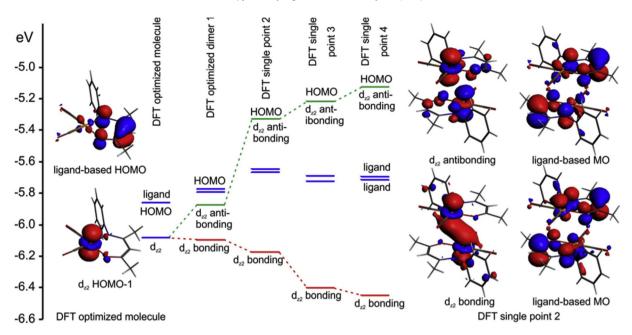


Fig. 7. Visualisation of the relative energies of the top four occupied molecular orbitals of the single **DFT optimised molecule** [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] (mono nuclear unit) (left), together with the four dimolecular [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ models calculated in this study. Visualisation of selected MOs of DFT calculations for **DFT single point 2** (α-polymorph at 150 K), as representative of the top four MOs of the dimolecular models, are shown on the right.

based), see Fig. 7 (right) for the top four MOs of **DFT single point 2** as representative example. Since the $d_{\rm Z^2}$ MOs from each [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] unit are already are occupied, overlapping of these d_{72} orbitals do not form a formal bond between the rhodium centres, but lead to weak intermolecular interactions between the rhodium centres from adjacent molecules. The d_{72} based bonding orbitals of rhodium stabilise the interaction, or weak metallophilic bond, between the two rhodium centres of a dinuclear $[Rh(CH_3COCHCN(Ph)CH_3)(CO)_2]_2$ unit, while the d_{7^2} based antibonding orbitals destabilise the rhodium-rhodium bond. A balance between the bonding and antibonding intermolecular orbitals of mainly d_{z2} type, determine the strength of the weak intermolecular interactions between adjacent [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] molecules. The energy of the d_{z^2} based bonding orbitals of **DFT single** point 3 and DFT single point 4 is more than 0.2 eV lower (more negative i.e. more stable) than that of the DFT optimised dimer 1 and DFT single point 2 models (Fig. 7), therefore contributing to the Rh-Rh wire formation observed for both **DFT single point 3** and **DFT** single point 4 in the solid state. On the other hand, the energies of the d_{7^2} based antibonding orbitals, increases with decrease in the lateral shift between the two separate molecular units in dinuclear $[Rh(CH_3COCHCN(Ph)CH_3)(CO)_2]_2$, see Fig. 7.

3.2.2.2. Natural bond orbitals (NBO). The inter-molecular interactions between the two molecular units of dimeric [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ were evaluated by the natural bond orbitals (NBO) analysis method of Weinhold, since an NBO analysis provides information about interactions between filled Lewis type donor NBOs (occupation numbers near 2) and empty non-Lewis acceptor NBOs (occupation numbers near 0). NBO types important for the inter-molecular interactions between the two single molecular units of dimeric [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂, are the following types: the 1-centre core (CR), the 1-centre non-bonded (lone pair, LP) and the 2-centre bond (BD) NBOs [42,43]. Selected results obtained from the NBO analysis of the four dimeric models used in this study, are summarised in Table 4, and selected NBOs are visualised in Fig. 8, while elected donor — acceptor NBO

interactions between the two rhodium centres in bimolecular [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ are visualised in Fig. 9.

In the **DFT single point 3** model (based on the coordinates of the β-polymorph at 100 K) where the two molecular units exhibit a large overlap (with a smaller lateral shift of 0.310 Å), the NBO calculations generally produced larger donor-acceptor interaction energies E(2) between a filled (bonding or lone pair) Lewis type NBO on rhodium of the one molecular unit (which acts as a donor) and an empty (antibonding or Rydberg) non-Lewis NBO on rhodium of the other molecular unit (which acts as an acceptor), than the calculated E(2) for the **DFT optimised dimer 1** (with a larger lateral shift of 1.590 Å) or **DFT single point 2** (with lateral shift of 0.925 Å), see Table 4. The donor NBOs on rhodium, involved in donor-acceptor interactions, are a CR NBO of mainly s character, a LP NBO of mainly d_{z^2} character, as well as two BD NBOs, see Fig. 8. The acceptor NBOs on rhodium, involved in donor-acceptor interactions, are LP* NBOs of mainly p_z , p_V or p_x character respectively, see Fig. 8. Selected donor-acceptor interactions between the two molecular units of **DFT single point 3** (βpolymorph at 100 K) are shown in Fig. 9. The slightly lateral shift (of 0.310 Å) between the two units of **DFT single point 3**, enables interaction between the filled donor LP on Rh of mainly d_{z^2} character and the three empty acceptor LP*s on Rh, of mainly p_z , p_V and p_X character respectively. This slightly lateral shift also favours donoracceptor interactions between the CR on Rh of mainly s character, and the two 2-centre bond NBOs, namely BD(Rh-C_{trans N}) and BD(Rh- $C_{trans\ O}$), with the acceptor LP* on Rh of mainly p_z character. The intermolecular donor-acceptor interaction energies, E(2), between the two rhodium atoms of the single molecular units of dimeric [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂, generally decrease in going from DFT single point 3 to DFT single point 2 to the DFT optimised dimer **1** gas phase model (for example 23.9 kJ $\text{mol}^{-1} > 11.4 \text{ kJ } \text{mol}^{-1} >$ 0.7 kJ mol⁻¹, for the LP(Rh mainly d_{z^2}) donor \rightarrow LP*(Rh mainly p_z) acceptor interaction, also see Table 4 for the other indications), as the lateral shift between the two single molecular units increases (namely 0.310 Å < 0.925 Å < 1.590 Å). Similarly, for **DFT single point 4**, where the lateral shift approaches 0 Å, the interaction energies E(2) of the donor-acceptor interactions involving the acceptor LP*(Rh

Table 4 Selected second order perturbation theory donor-acceptor interaction energies, E(2) in $kJ \cdot mol^{-1}$, with the calculated natural bond orbital (NBO) occupations, for each of the four different theoretically calculated DFT models of the dimer $[Rh(CH_3COCHCN(Ph)CH_3)(CO)_2]_2$, with lateral shift between adjacent molecules decreasing from model 1 to model 4.

Complex	DFT optimised dimer 1	9 1	DFT single point 3 (coordinates of β polymorph at 100 K)	DFT single point 4 (no lateral shift)
E(2)/kJ·mol ⁻¹				
$LP(Rh \text{ mainly } d_{z^2}) \rightarrow LP^*(Rh \text{ mainly } p_z)$	0.7	11.4	23.9	30.5
$LP(Rh \text{ mainly } d_{z^2}) \rightarrow LP^*(Rh \text{ mainly } p_x)$	0.8	4.8	8.5	3.5
$LP(Rh \text{ mainly } d_{z^2}) \rightarrow LP^*(Rh \text{ mainly } p_y)$	0.2	9.7	6.6	2.5
$BD(Rh-C_{trans\ N}) \rightarrow LP^*(Rh\ mainly\ p_z)$	0.3	15.6	15.6	22.9
$BD(Rh-C_{trans\ O}) \rightarrow LP^*(Rh\ mainly\ p_z)$	0.3	7.7	15.4	22.8
$CR(Rh \text{ mainly s}) \rightarrow LP^*(Rh \text{ mainly } p_z)$	0.1	8.1	15.2	13.8
Occupancy				
LP(Rh mainly d_{z^2})/e ⁻	1.972	1.963	1.957	1.955
CR(Rh mainly s)/e ⁻	1.989	1.989	1.988	1.988
$LP^*(Rh mainly p_z)/e^-$	0.061	0.078	0.088	0.090
$LP^*(Rh mainly p_x)/e^-$	0.182	0.183	0.184	0.188
$LP^*(Rh mainly p_y)/e^-$	0.166	0.168	0.167	0.171
BD(Rh C _{trans N})/e ⁻	1.960	1.955	1.951	1.949
BD(Rh-C _{trans O})/e ⁻	1.960	1.953	1.949	1.948

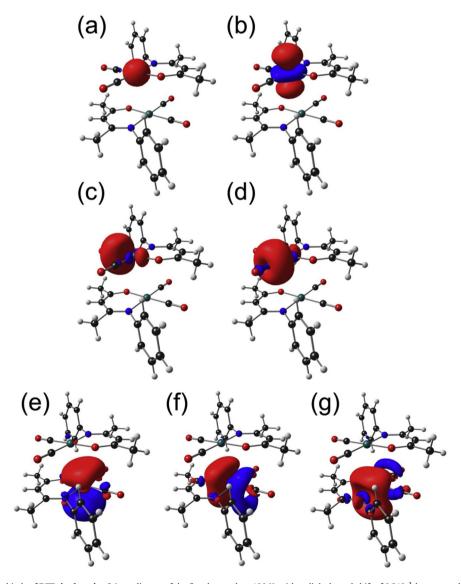


Fig. 8. Selected natural bond orbitals of DFT single point 3 (coordinates of the β-polymorph at 100 K, with a slight lateral shift of 0.310 Å between adjacent rhodium atoms). Donor NBOs: (a) LP on Rh is of mainly s character, (b) LP on Rh is of mainly d_{z^2} character, (c) BD (Rh-C_{trans O}), (d) BD (Rh-C_{trans N}). Acceptor NBOs: (e) LP* on Rh is of mainly p_z character, (f) LP* on Rh is of mainly p_y character and (g) LP* on Rh is of mainly p_z character. The NBO plots utilise a contour of 0.05 e/ų. Colour code of atoms (online version): Rh (green), C (black), N (blue), O (red), H (white). (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

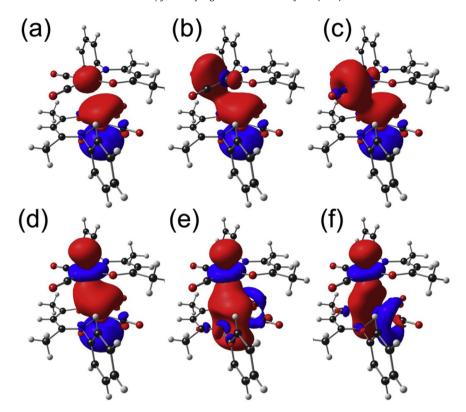


Fig. 9. Visualisation of selected donor — acceptor NBO interactions between the two rhodium centres in bimolecular [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ of **DFT single point 3** (coordinates of the β-polymorph at 100 K, with a slight lateral shift of 0.310 Å between adjacent rhodium atoms). (a) LP(Rh of mainly s character) — LP*(Rh of mainly p_s character), (b) LP(Rh of mainly p_s character) — LP*(Rh of mainly p_s character) — LP*(Rh of mainly p_s character) — LP*(Rh of mainly p_s character). (c) LP (Rh of mainly p_s character) — LP*(Rh of mainly p_s character). (b) LP (Rh of mainly p_s character) — LP*(Rh of mainly p_s character). (c) LP (Rh of mainly p_s character) — LP*(Rh of mainly p_s character). The NBO plots utilise a contour of 0.05 e/Å³. Colour code of atoms (online version): Rh (green), C (black), N (blue), O (red), H (white). (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

of mainly p_z character), generally increases relative to the other three models who have a larger lateral shift between the two separate molecules (for example 30.4 kJ mol⁻¹ > 23.9 kJ mol⁻¹ > 11.4 kJ mol⁻¹ > 0.7 kJ mol⁻¹, for the LP(Rh mainly d_{z^2}) donor \rightarrow LP*(Rh mainly p_z) acceptor interaction of DFT models **4**, **3**, **2** and **1** respectively, also see Table 4 for the other indications). However, as is also expected for a ca. 0 Å lateral shift, the donor-acceptor interactions involving the acceptors LP*(Rh of mainly p_x character) and LP* (Rh of mainly p_y character), both decrease dramatically relative to

E(2) for **DFT single point 3** and **DFT single point 2**. E(2) of LP(Rh mainly d_{z^2}) donor \rightarrow LP*(Rh mainly p_x) acceptor decreases from 8.5 eV to 3.5 eV, while E(2) of LP(Rh mainly d_{z^2}) donor \rightarrow LP*(Rh mainly p_y) acceptor decreases from 6.6 eV to 2.5 eV (when going from **DFT single point 3** to **DFT single point 4**) see Table 4.

In summary, these identified donor-acceptor interactions thus support the experimental observation that the [Rh(CH₃COCHCN(Ph) CH₃)(CO)₂] molecules are packed in pairs in the solid state, either as separate dinuclear [Rh(β -diketonato)(CO)₂]₂ dimeric units or in

Table 5
Selected topological parameters of the inter-molecular bond-paths for the four dimolecular [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ models of this study, indicating that Rh-Rh bond paths are only present for models 2 to 4. The critical point (CP) numbers are shown in Fig. 10.

Bond/Ring critical point	Bond type	Atoms involved	inter-atomic distance	BP length	Electron density	Laplacian of electron density	
			/Å	/Å	$\rho/e \ a_0^{-3}$	$\nabla^2 \rho / e \ a_0^{-5}$	
DFT optimised dimer 1							
CP# 68/88	hydrogen bond	O-H	2.6314	2.6420	0.0062	0.0228	
DFT single point 2							
CP# 48	rhodium-rhodium	Rh-Rh	3.4811	3.4844	0.0113	0.0264	
CP# 115/75	hydrogen bond	O-H	2.9315	3.0967	0.0034	0.0153	
CP# 130/61	hydrogen bond	О-Н	2.7761	2.8373	0.0046	0.0196	
CP# 113/73	carbon-hydrogen	C-H	2.9478	3.0076	0.0033	0.0139	
CP# 89/90	hydrogen bond	O-H	2.5858	2.6171	0.0071	0.0281	
DFT single point 3							
CP# 90	rhodium-rhodium	Rh-Rh	3.3085	3.3109	0.0160	0.0367	
CP# 93/77	hydrogen bond	О-Н	2.7644	2.8699	0.0048	0.0213	
CP# 92/78	hydrogen bond	О-Н	2.9263	3.2006	0.0040	0.0181	
CP# 9785	hydrogen bond	О-Н	2.9244	3.1336	0.0041	0.0174	
CP# 129/87	hydrogen bond	O-H	2.8607	2.9078	0.0041	0.0159	
DFT single point 4							
CP# 87	rhodium-rhodium	Rh-Rh	3.3085	3.3097	0.1720	0.0383	
CP# 76/99	hydrogen bond	O-H	2.4708	2.5016	0.0087	0.0382	

dinuclear units that form extended linear wire-like chains of rhodium, as was indeed found for the α - and β -polymorphs of this study.

3.2.2.3. OTAIM. The interaction between the two [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] mono nuclear units in dinuclear [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ was further analysed, using Bader's quantum theory of atoms in molecules (QTAIM) theory with selected topological parameters of the inter-molecular bond-paths for the four dimolecular [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ models of this study listed in Table 5. The topological analysis of the charge density $\rho(r)$ distribution, as well as the properties of the critical points (CPs) determined by QTAIM, provides a universal indicator of bonding between atoms [44]. For the **DFT optimised dimer 1**, the **DFT single point 2**, the **DFT single point 3** and the **DFT single point** 4, a total amount of two, nine, nine and three inter-molecular bonds respectively, were identified via bond critical points, see Fig. 10. Due to symmetry, half of the bond paths are identical to the other half, except for the rhodium-rhodium bond path.

No rhodium-rhodium bond path was identified for the **DFT optimised dimer 1** with the largest lateral shift of 1.590 Å, indicating no formation of the linear metal-metal chains. However, the **DFT optimised dimer 1** is stabilised by two inter-molecular hydrogen bonds, thereby favouring the formation of the observed dinuclear units in the solid state, as observed for the α -polymorphs of this study.

For **DFT single point 2**, **DFT single point 3** and **DFT single point 4**, in addition to inter-molecular hydrogen bonds, rhodium-rhodium bond paths was identified. The electron density (ρ) and the Laplacian of electron density $(\nabla^2 \rho)$ at the rhodium-rhodium bond critical point, increased in value in going from **DFT single point 2** to **DFT single point 3** and to **DFT single point 4**,

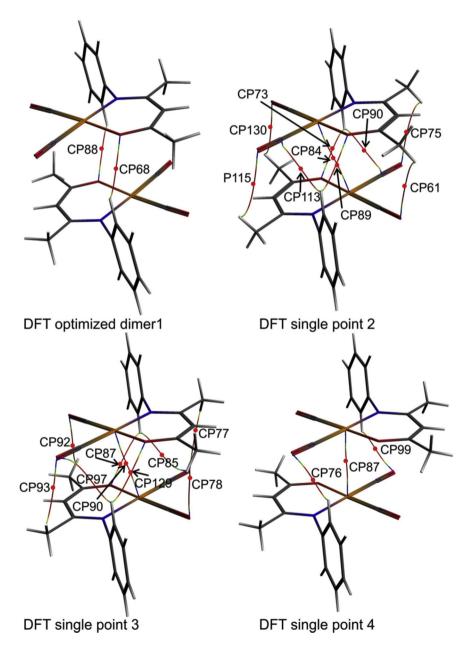


Fig. 10. Schematic representation of the inter-molecular bond-paths (BP), indicated by red bond critical points (CP), which are present in the DFT optimised geometry of all four dimolecular [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ models of this study. The colour range of the bond-paths decreases in frequency, according to the value of the electron density: from blue (high density) to green to red (low density). (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

corresponding to decreasing lateral shift: For example, electron density (ρ) at the rhodium-rhodium bond critical point increased in the order 0.0113 e.a $_{0}^{-3}$ < 0.0160 e.a $_{0}^{-3}$ < 0.1720 e.a $_{0}^{-3}$ from **model 2** to **model 4**, while the Laplacian of electron density ($\nabla^{2}\rho$) increased accordingly, 0.0264 e.a $_{0}^{-5}$ < 0.0367 e.a $_{0}^{-5}$ < 0.0383 e.a $_{0}^{-5}$, indicating a rhodium-rhodium bond path in the latter two models with much higher ρ and $\nabla^{2}\rho$.

To conclude, both NBO and QTAIM calculations shed light on intermolecular interactions leading to the formation of dimeric $[Rh(CH_3COCHCN(Ph)CH_3)(CO)_2]_2$ units as was found for both the α -and β -polymorphs of this study. The rhodium-rhodium bond paths of the QTAIM results suggested the formation of wire-like rhodium-rhodium chains as the lateral shift between the two molecular units decreases as was found for the β -polymorphs of this study.

4. Conclusion

The geometry of the solid state structure of two [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] crystals presented in this contribution, compares well with the previously published structure [13] for the same complex, as well as with the DFT calculated optimised single molecular unit [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] from this study. Different crystallization conditions (solvent and temperature) led to different polymorphs (α and β), with a difference in crystal packing of [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]. The solid state crystallographic data of both the α - and β -polymorph [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] crystal structures, show that this complex forms dimeric units that either stacked in wire-like linear metal-metal chains (β-polymorph) or the dimeric units do not stack onto each other (α-polymorph). Additional theoretical QTAIM computations supported the experimental crystallographic findings of dimer formation, since it was found that the DFT optimised dimer of [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂]₂ is stabilised by two intermolecular hydrogen bonds, thereby favouring the formation of the observed dinuclear units in the solid state. QTAIM results further suggested the formation of wire-like rhodium-rhodium chains as the lateral shift between the two molecular units decreases as was found for the β-polymorphs of this study. An NBO analysis of the DFT optimised dinuclear [Rh(CH₃COCHCN(Ph) CH₃)(CO)₂]₂ unit, showed that various donor-acceptor interactions between the two separate [Rh(CH₃COCHCN(Ph)CH₃)(CO)₂] molecules do favour the formation of dimeric [Rh(CH₃COCHCN(Ph) $CH_3)(CO)_2|_2$ pairs.

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Appendix A. Supplementary data

Supplementary data related to this article can be found at https://doi.org/10.1016/j.jorganchem.2017.09.039.

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