

Electronic Supplementary Information (ESI)

Palladium cluster complex $[\text{Pd}_{13}(\mu_4\text{-C}_7\text{H}_7)_6]^{2+}$ ($\text{C}_7\text{H}_7 = \text{Tropylium}$) with *fcc*-close-packed cuboctahedral Pd_{13} core and isomers: Theoretical insight into ligand-control of Pd_{13} core structure

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Table S1. Optimized Pd-Pd distances (in Å) in $[\text{Pd}_{13}(\mu_4\text{-C}_7\text{H}_7)_6]^{2+}$ cubooctahedral Pd_{13} core and their comparisons with experimental values.

Bonds	$[\text{Pd}_{13}(\text{C}_7\text{H}_7)_6]^{2+}$		
	Exp. ^b	This work	Δ ^c
$\text{Pd}^{\text{C}}\text{-Pd}^{\text{T/B}}$ ^a	2.769	2.742	0.027
$\text{Pd}^{\text{C}}\text{-Pd}^{\text{E}}$	2.766	2.738	0.028
$\text{Pd}^{\text{T/B}}\text{-Pd}^{\text{E}}$	2.766	2.740	0.026
$\text{Pd}^{\text{E}}\text{-Pd}^{\text{E}}$	2.766	2.738	0.028
$\text{Pd}^{\text{T/B}}\text{-Pd}^{\text{T/B}}$	2.764	2.746	0.018
	Exp. ^b	This work	Previous work ^d
Longest Pd-Pd	2.737	2.836	2.900
Shortest Pd-Pd	2.631	2.688	2.718
Average	2.716	2.767	2.808

^a Pd^{C} , Pd^{E} , and $\text{Pd}^{\text{T/B}}$ represent Pd atoms at a center, equatorial, and top (or bottom) three-member ring, respectively. Mean value of Pd-Pd distances in each category is presented here because the unique comparison between the optimized geometry and the experimental one is difficult due to various orientations.

^b Ref. S1.

^c This is the absolute value of difference between the optimized Pd-Pd distance and the experimental one.

^d Ref. S2.

Brief discussion: Generally, the DFT-optimized bond-distance of mono-nuclear transition metal complexes with a medium size does not deviate very much from experimental value (less than 0.3 Å to 0.5 Å in many cases). Here, the deviation of the longest Pd-Pd bond seems too large, whereas the average of Pd-Pd distances agree with experimental values. However, the flexibility and large size of this Pd_{13} cluster complex are the obstacle for geometry optimization. Nevertheless, the agreement with the experimental geometry is considerably better than that optimized previously (ref. S2). It is likely to state that the present DFT calculation overestimates Pd-Pd distances but the agreement is not bad, taking the large size and flexibility of the geometry of the Pd_{13} cluster complex.

Table S2. Atomic charge of $[\text{Pd}_{13}(\text{C}_7\text{H}_7)_6]^{2+}$ cuboctahedral and anticuboctahedral Pd_{13} core.

$[\text{Pd}_{13}(\mu_4\text{-C}_7\text{H}_7)_6]^{2+}$ (Cubo; $\mu 4$)				
	Pd^C ^a	Pd^E ^a	Pd^{T/B} ^a	C₇H₇ ^a
NBO^a	-0.517	0.085	0.085	0.249
Hirshfeld^a	-0.064	0.081	0.081	0.181
Mulliken^b	4.42	-0.498	-0.498	0.593
$[\text{Pd}_{13}(\mu_4\text{-C}_7\text{H}_7)_6]^{2+}$ (Anti; $\mu 4$)				
	Pd^C	Pd^E	Pd^{T/B}	C₇H₇
NBO^a	-0.774	0.131	0.099	0.232
Hirshfeld^a	-0.056	0.043	0.097	0.203
Mulliken^b	4.761	-0.472	-0.547	0.559

^a Pd^C, Pd^E, and Pd^{T/B} represent Pd atoms at a center, equatorial, and top (or bottom) three-member ring, respectively.

^b Because the Mulliken population analysis leads to unusual atomic charge when diffuse basis functions are involved, we here employed double-zeta quality basis sets for Pd and other atoms; LANL2DZ was used for Pd and 6-31G(d) were used for other atoms.

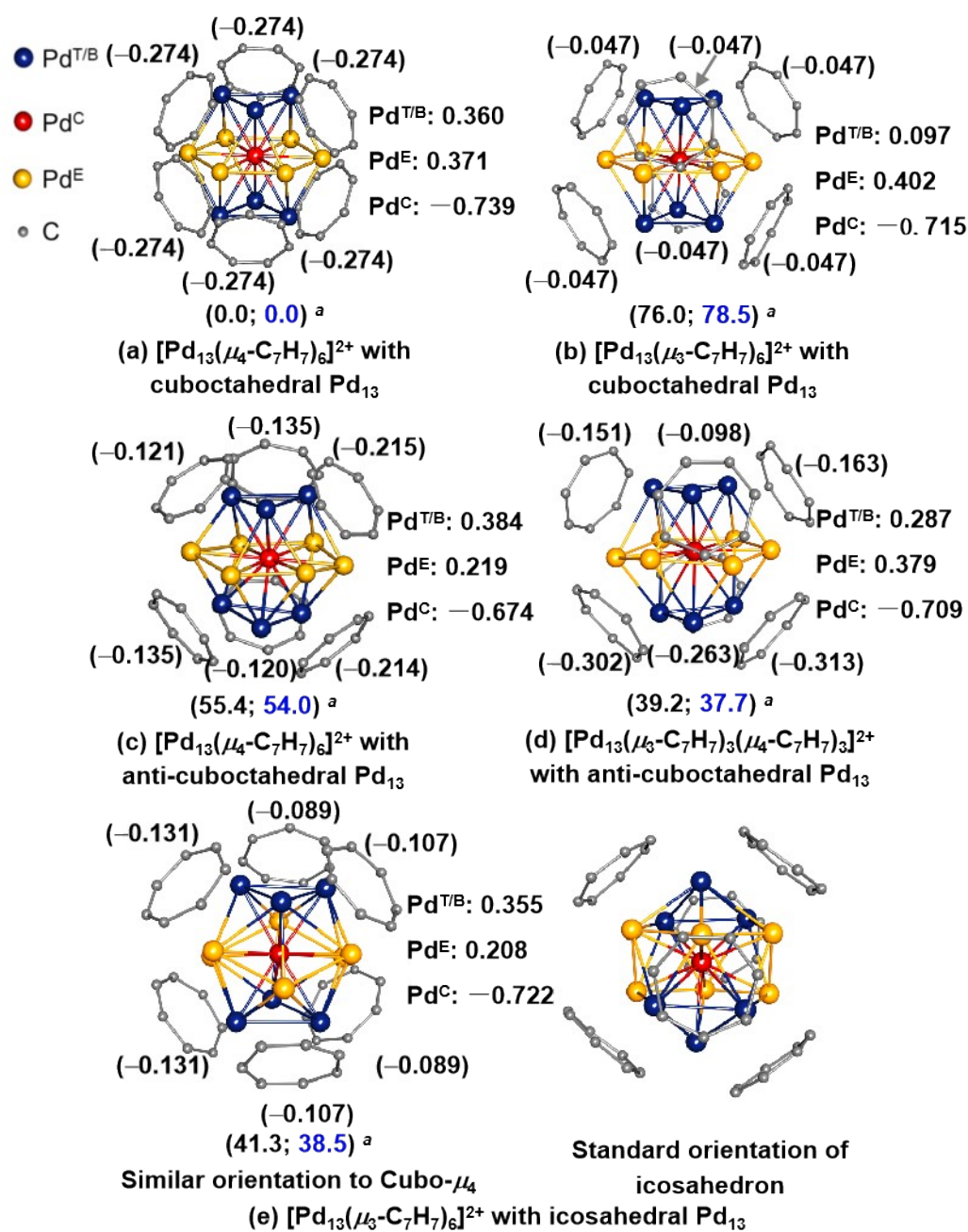
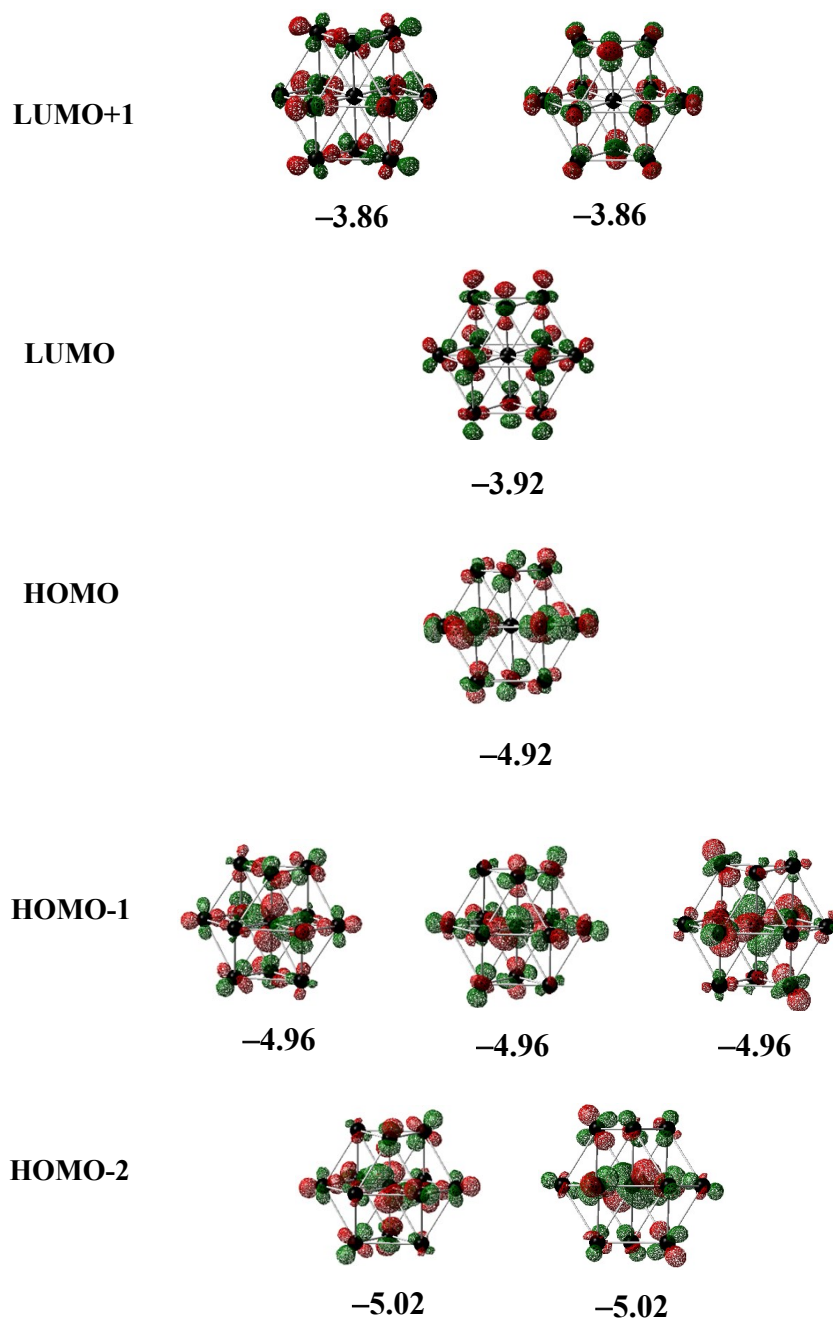
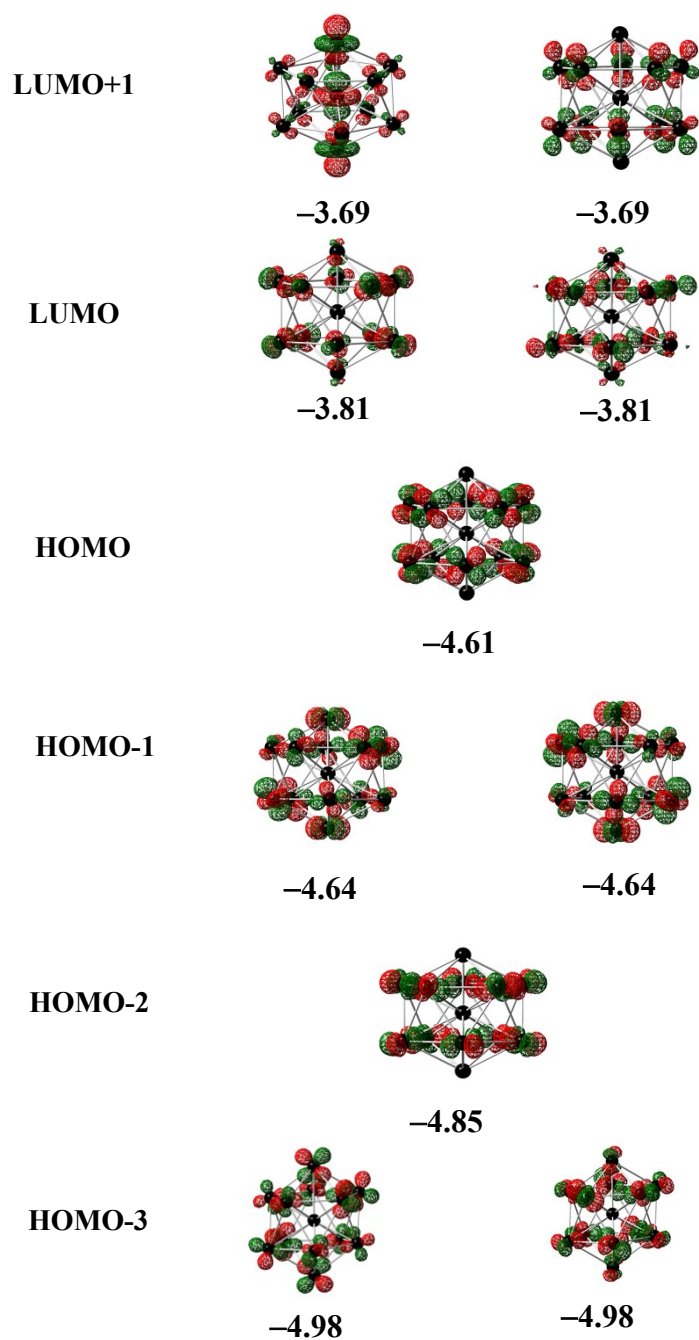


Figure S2. NBO charges (in e unit) of $[\text{Pd}_{13}(\mu_4\text{-C}_7\text{H}_7)_6]^{2+}$ with a cubo-octahedral Pd_{13} core **Cubo- μ_4** (a), $[\text{Pd}_{13}(\mu_3\text{-C}_7\text{H}_7)_6]^{2+}$ with a cubo-octahedral Pd_{13} core **Cubo- μ_3** (b), $[\text{Pd}_{13}(\mu_4\text{-C}_7\text{H}_7)_6]^{2+}$ with an anticubo-octahedral Pd_{13} core **Anti- μ_4** (c), $[\text{Pd}_{13}(\mu_3\text{-C}_7\text{H}_7)_3(\mu_4\text{-C}_7\text{H}_7)_3]^{2+}$ with an anticubo-octahedral Pd_{13} core **Anti- $\mu_3,4$** (d) and $[\text{Pd}_{13}(\mu_3\text{-C}_7\text{H}_7)_6]^{2+}$ with an icosahedral Pd_{13} core **Ih- μ_3** (e). ^aRelative potential energy and Gibbs energy (in kcal/mol) relative to those of $[\text{Pd}_{13}(\mu_4\text{-C}_7\text{H}_7)_6]^{2+}$ with a cubo-octahedral Pd_{13} core.

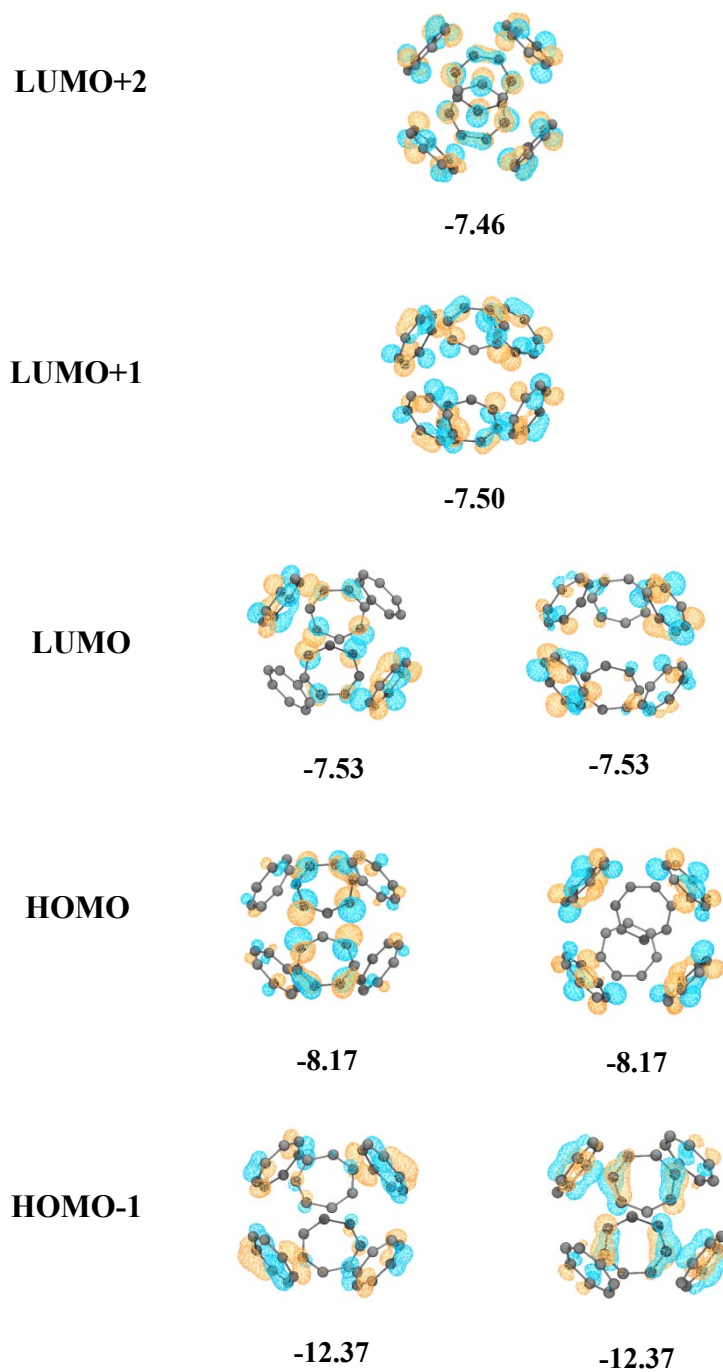


Scheme S1. Frontier orbitals of anticuboctahedral Pd₁₃ cores, where the geometry was optimized. The number below each MO represents the Kohn-Sham orbital energy (in eV). B3PW91-D3(BJ)/BS-II was used here.^{S3}

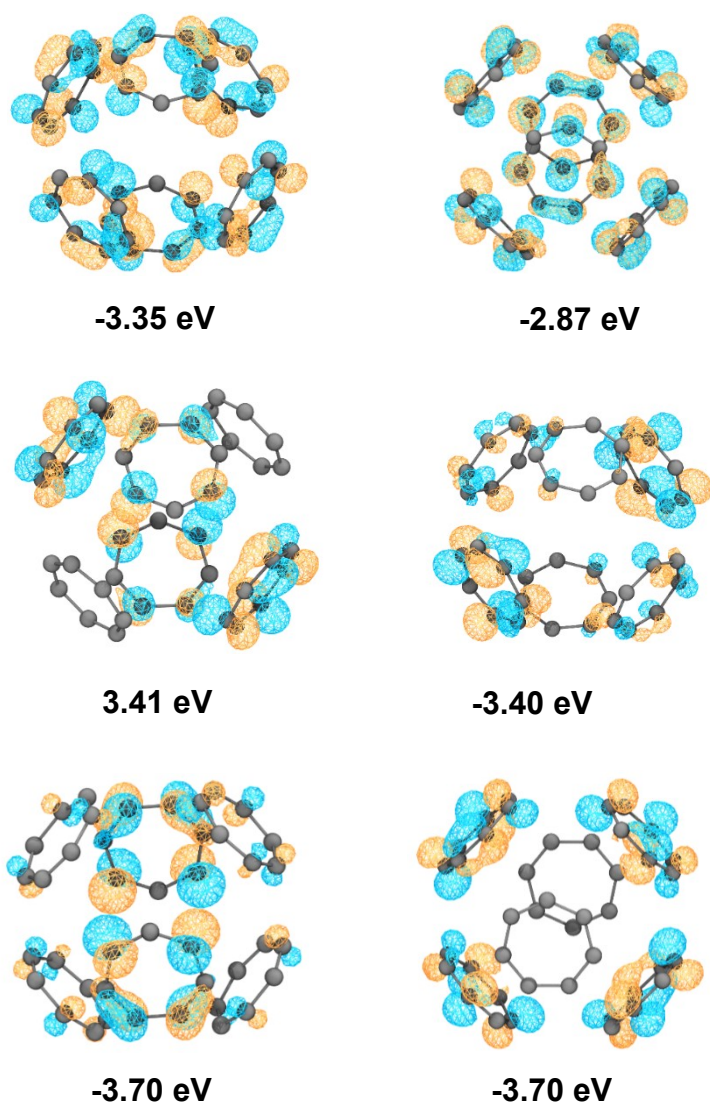


Scheme S2. Frontier orbitals of icosahedral Pd₁₃ cores, where the geometry was optimized. The number below each MO represents the Kohn-Sham orbital energy (in eV). B3PW91-D3(BJ)/BS-II was used here.^{S3}

Discussion: These Schemes 1 and 2 mean that both the neutral Pd₁₃ and the [Pd₁₃]²⁺ clusters can have anticuboctahedral and icosahedral structures at the closed shell singlet state; because both have non-degenerate HOMO, the [Pd₁₃]²⁺ cluster does not induce the Jahn-Teller distortion.



Scheme S3. HOMO and LUMO of $[(C_7H_7)_6]^{2+}$ at the closed shell singlet, where the geometry is taken to be the same as that in $[Pd_{13}(\mu_4-C_7H_7)_6]^{2+}$ with cuboctahedral Pd_{13} core. The number below each MO represents the Kohn-Sham orbital energy (in eV). B3PW91-D3(BJ)/BS-I was used here, because BS-II presents overly diffuse features of MO.



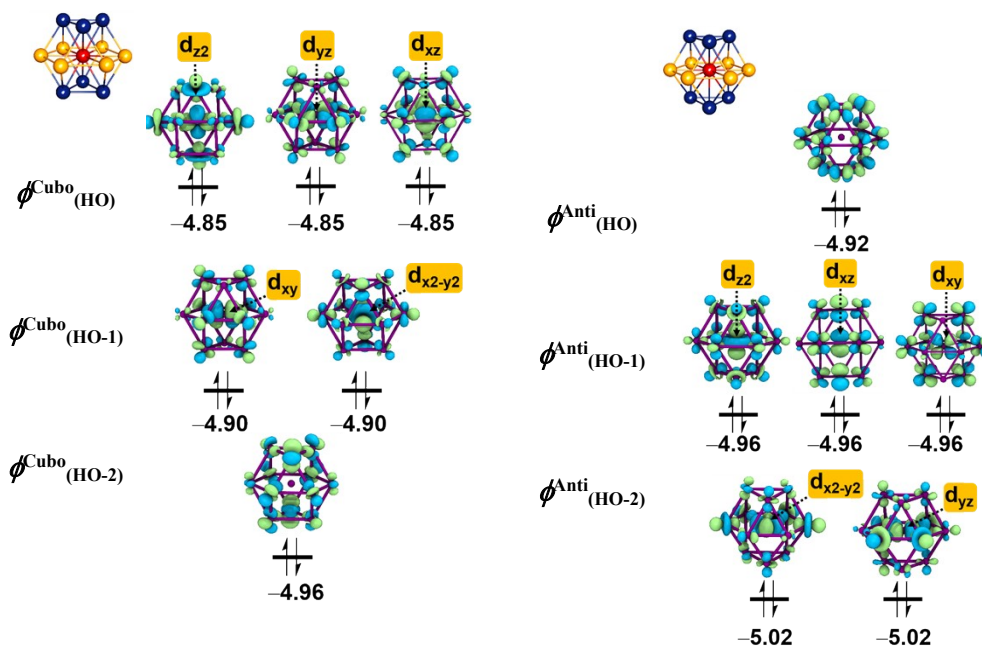
Scheme S4. Six singly occupied molecular orbitals (SOMOs) of $[(C_7H_7)_6]$ in the septet state, where geometries, positions, and orientations of six C_7H_7 molecules are taken to be the same as those in $[Pd_{13}(C_7H_7)]^{2+}$ with a cuboctahedral Pd_{13} core.

B3PW91-D3(BJ)/BS-I was used here, because BS-II presents overly diffuse features of MO.

Discussion: The two MOs at lower energy (-3.70 eV) than the other four MOs become doubly occupied HOMO in the $[(C_7H_7)_6]^{2+}$ at the closed shell singlet, as shown in Scheme S3.

Discussion of closed-shell singlet state of the $[(C_7H_7)_6]^{2+}$ ligand shell

When six neutral C_7H_7 molecules are separated well from each other, there exist six singly occupied molecular orbitals (SOMOs) at the same orbital energy in the septet spin state. When the $[(C_7H_7)_6]^{2+}$ system is calculated at the closed-shell singlet state, two of them are doubly occupied. Such an electronic structure may cause significantly large distortion in the $[(C_7H_7)_6]^{2+}$ moiety due to the Jahn-Teller effect and the six C_7H_7 molecules become non-equivalent. This is not suitable for the nearly octahedral $(C_7H_7)_6$ ligand shell. To check whether the closed shell singlet state of the $[(C_7H_7)_6]^{2+}$ ligand shell is reasonable or not, we calculated the neutral $[(C_7H_7)_6]$ ligand shell at the septet spin state, where the geometry was taken to be the same as that in **Cubo- μ 4**. The neutral $[(C_7H_7)_6]$ system has six SOMOs at the septet state as expected, as shown in Scheme S4 of the Supporting Information. However, their orbital energies are not the same unexpectedly; two of them are calculated at lower energies than the other four SOMOs probably because six C_7H_7 molecules are not completely isolated but interact to some extent with each other in the ligand shell of **Cubo- μ 4**. Consequently, those two MOs at a lower energy become a doubly-degenerated HOMO in the $[(C_7H_7)_6]^{2+}$ system at a closed shell singlet state, as shown in Scheme S3 of the Electronic Supplementary Information (ESI). Thus, there is no unreasonable feature in the $[(C_7H_7)_6]^{2+}$ ligand shell having the closed shell singlet state when the geometry is taken to be the same as that in **Cubo- μ 4**.



Scheme S5. The HOMO, HOMO-1, HOMO-2 of cubo-octahedral and anti-cubo-octahedral Pd₁₃ clusters^{S3}

^a The optimized geometries of cubo-octahedral and anti-cubo-octahedral Pd₁₃ clusters were employed. B3PW91-D3(BJ)/BS-II was used.

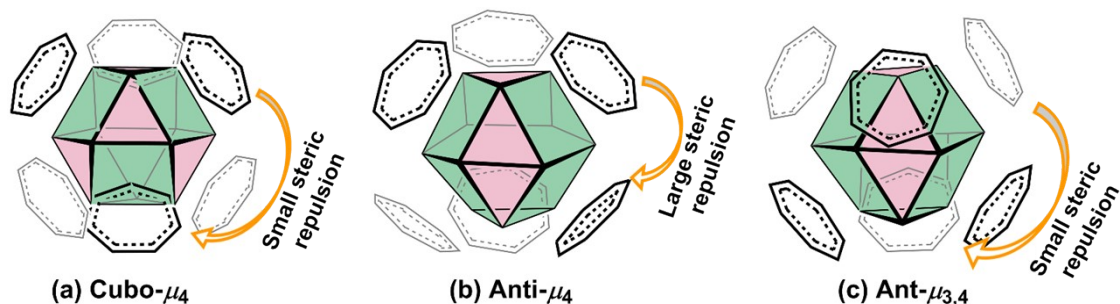
Discussion: The HOMO $\phi^{\text{Anti}}(\text{HO})$ of the anti-cubo-octahedral Pd₁₃ corresponds to the HOMO-2 $\phi^{\text{Cubo}}(\text{HO-2})$ of the cubo-octahedral Pd₁₃, as shown in Scheme S5, but their energies do not differ very much. The HOMO-1 $\phi^{\text{Anti}}(\text{HO-1})$ and HOMO-2 $\phi^{\text{Anti}}(\text{HO-2})$ of the anti-cubo-octahedral Pd₁₃ correspond to the HOMO $\phi^{\text{Cubo}}(\text{HO})$ and HOMO-1 $\phi^{\text{Cubo}}(\text{HO-1})$ of the cubo-octahedral Pd₁₃, respectively. However, the $\phi^{\text{Anti}}(\text{HO-1})$ and $\phi^{\text{Anti}}(\text{HO-2})$ exist at somewhat lower energies than the $\phi^{\text{Cubo}}(\text{HO})$ and $\phi^{\text{Cubo}}(\text{HO-1})$, respectively. As a result, $\phi^{\text{Anti}}(\text{HO})$ becomes the HOMO of the anti-cubo-octahedral Pd₁₃ cluster despite that it resembles $\phi^{\text{Cubo}}(\text{HO-2})$ of the cubo-octahedral Pd₁₃. The orbital energy differences between $\phi^{\text{Cubo}}(\text{HO})/\phi^{\text{Cubo}}(\text{HO-1})$ of the cubo-octahedral Pd₁₃ and $\phi^{\text{Anti}}(\text{HO-1})/\phi^{\text{Anti}}(\text{HO-2})$ of the anti-cubo-octahedral Pd₁₃ may result from the extent of the d orbital participation of the Pd^C atom in these MOs but the details are not clear at the present stage. Further analysis is needed.

9. Comparisons between Anti- $\mu_3,4$ and Anti- μ_4 and between Ih- μ_3 and Cubo- μ_3

9.1 Comparison between Anti- $\mu_3,4$ and Anti- μ_4 . As shown in Figure 2, **Anti- $\mu_3,4$** is more stable than **Anti- μ_4** , which mainly results from the more negative E_{int} value in **Anti- $\mu_3,4$** than in **Anti- μ_4** (Table 2). The larger stability of **Anti- $\mu_3,4$** than that of **Anti- μ_4** is seemingly inconsistent with the conclusion above showing that the μ_4 -coordination bond of C_7H_7 is stronger in **Cubo- μ_4** than the μ_3 -one in **Cubo- μ_3** .

In **Anti- $\mu_3,4$** , the μ_4 - C_7H_7 ligand is less positively charged (average value = +0.156 e) than the μ_3 - C_7H_7 ligand (average value = +0.224 e), suggesting that the CT from the Pd_{13} core to the μ_4 - C_7H_7 ligand is stronger than that to the μ_3 - C_7H_7 ligand like in **Cubo- μ_3** and **Cubo- μ_4** . Consistent with this difference in CT, the $R(Pd^C-X)$ distance (1.976 Å; Table 1) of **Anti- $\mu_3,4$** is considerably shorter in the μ_4 -coordination bond than in the μ_3 -one (2.238 Å). In **Anti- μ_4** , however, the $R(Pd^C-X)$ distance (2.230 Å) is considerably longer than that (1.976 Å) of the μ_4 -coordination of **Anti- $\mu_3,4$** and the averaged charge (+0.203 e) of six μ_4 - C_7H_7 ligands in **Anti- μ_4** is larger than that (+0.156 e) of three μ_4 - C_7H_7 ligands in **Anti- $\mu_3,4$** . These features suggest that the μ_4 -coordination is much stronger and, therefore, the E_{int} is larger in **Anti- $\mu_3,4$** than in **Anti- μ_4** . This is the reason why **Anti- $\mu_3,4$** is more stable than **Anti- μ_4** despite that **Anti- $\mu_3,4$** has three μ_3 - and three μ_4 -coordination bonds of C_7H_7 but **Anti- μ_4** has six μ_4 -coordination bonds.

In the above explanation, we address a new open question why the μ_4 -coordination bond is weaker in **Anti- μ_4** than in **Anti- $\mu_3,4$** . We compared the positions of C_7H_7 ligands between **Anti- μ_4** and **Anti- $\mu_3,4$** . In **Anti- μ_4** , three μ_4 - C_7H_7 ligands in the upper side of the Pd_{13} core are close to three μ_4 - C_7H_7 ligands in the lower side (Figure 1c). Therefore, the steric repulsion between the upper μ_4 - C_7H_7 and the lower μ_4 - C_7H_7 is large (Scheme S6b), and the μ_4 - C_7H_7 ligands are not located at the best position for interacting with the Pd_{13} core to decrease the E_{int} value. In the anticuboctahedral Pd_{13} core, the Pd_3 planes in the upper part of the Pd_{13} core is connected with the Pd_4 planes in the lower part. In **Anti- $\mu_3,4$** , therefore, three μ_3 - C_7H_7 ligands in the upper side are distant from three μ_4 - C_7H_7 ligands in the lower side and the steric repulsion between them is small (Scheme S6c). Therefore, the μ_4 - C_7H_7 ligands are located at the best position to interact with the anticuboctahedral Pd_{13} core in **Anti- $\mu_3,4$** without any interference. Consistent with this understanding, the $R(Pd^C-X)$ distance of the μ_4 - C_7H_7 in **Anti- μ_4** is longer than in **Anti- $\mu_3,4$** , as mentioned above. Thus, it is likely concluded that the μ_4 -coordination bond of the C_7H_7 ligand is weaker in **Anti- μ_4** than in **Anti- $\mu_3,4$** .



Scheme S6. Schematical picture showing the positions of μ_3 - and μ_4 - C_7H_7 ligands.

9.2 Comparison between **Ih- μ_3** and **Cubo- μ_3**

Ih- μ_3 is more stable than **Cubo- μ_3** despite that both have six μ_3 - C_7H_7 ligands. All the C_7H_7 ligands have the moderately smaller positive charges ($+0.211 e \sim +0.214 e$) in **Ih- μ_3** than those ($+0.225 e$) in **Cubo- μ_3** (Figure 2). Also, the $R(Pd^C-X)$ distance (2.202 Å; Table 1) of **Ih- μ_3** is moderately shorter than that (2.242 Å) of **Cubo- μ_3** (Table 1). These features suggest that the μ_3 -coordination bond of C_7H_7 is stronger in **Ih- μ_3** than in **Cubo- μ_3** . One factor is the interaction structure. The Type-A coordination bond of μ_3 - C_7H_7 in **Ih- μ_3** is stronger than in **Cubo- μ_3** because three C=C double bonds interact with three Pd atoms in the Type A μ_3 -coordination bond of **Ih- μ_3** but one C=C double bond coordinates to one Pd atom and three C atoms coordinates to two Pd atoms through a π -allyl type interaction in the μ_3 -coordination bond of **Cubo- μ_3** . The other factor is the HOMO energy of the Pd_{13} core: The HOMO of the icosahedral Pd_{13} cluster exists at a higher energy ($-4.61 eV$; Scheme S2 in the ESI) than that ($-4.85 eV$; Scheme 3) of the cuboctahedral Pd_{13} cluster.²¹ Consequently, the icosahedral Pd_{13} core forms the stronger CT to the μ_3 - $(C_7H_7)_6$ ligand shell than does the cuboctahedral Pd_{13} core. Because of these two factors, the E_{int} value is larger (more negative) in **Ih- μ_3** than in **Cubo- μ_3** . In addition, the icosahedral Pd_{13} core and the $[(C_7H_7)_6]^{2+}$ shell in **Ih- μ_3** are more stable than the cuboctahedral Pd_{13} core and the $[(C_7H_7)_6]^{2+}$ shell in **Cubo- μ_3** , respectively (Table 2). Therefore, **Ih- μ_3** is more stable than **Cubo- μ_3** .

References

- S1. M. Teramoto, K. Iwata, H. Yamaura, K. Kurashima, K. Miyazawa, Y. Kurashige, K. Yamamoto and T. Murahashi, *J. Am. Chem. Soc.*, 2018, **140**, 12682–12686.
- S2. J. Wei, S. Kahlal, J.-F. Halet and J.-Y. Saillard, *J. Cluster Science*, 2019, **30**, 1227-1233).
- S3. Here, we compared HOMO energy using the optimized Pd₁₃ bare cluster because the Pd₁₃ cores in **Cubo-μ4** and **Anti-μ4** distort and the HOMO energy is influenced by the distortion. In other words, the discussion is presented on the intrinsic ability of HOMO of the Pd₁₃ cluster for the CT from the Pd₁₃ core to the [(C₇H₇)₆]²⁺ ligand shell.

Cartesian coordinate

Cubo- μ 4

Pd	7.04727300	8.55225900	35.17618500
Pd	7.41182200	6.11795100	34.09106400
Pd	8.42256900	7.23592900	31.76405100
Pd	8.02436400	9.80280200	32.90126000
Pd	4.85422200	6.76944200	34.93665200
Pd	5.51319200	10.47297400	33.88367500
C	9.17192100	8.67761800	35.57866200
H	9.15635100	9.11552200	36.57579000
C	9.17844700	7.25167500	35.55477000
H	9.02118400	6.77608400	36.51883400
C	9.55490000	6.36762500	34.50213200
H	9.76767300	5.34848000	34.82141900
C	10.10286500	6.70452200	33.21563000
H	10.56866300	5.87025700	32.69672000
C	10.35316400	7.98709700	32.66890000
H	10.99946700	8.00518900	31.79431800
C	10.14245700	9.27863400	33.26774000
H	10.67189100	10.09079900	32.77367800
C	9.61466200	9.58110600	34.54812800
H	9.75639700	10.60465700	34.88545900
C	6.01363100	9.99784400	36.63379000
H	6.84425100	10.58374900	37.01882300
C	5.94452900	8.65748100	37.08997400
H	6.70482100	8.36658100	37.81201600
C	4.84821900	7.73107400	36.98255000
H	4.94144500	6.85905100	37.62581700
C	3.56694600	7.93462800	36.41398200
H	2.81440000	7.19134400	36.66589600
C	3.04176700	9.11579700	35.78215600
H	1.95441400	9.14496900	35.73090300
C	3.67676400	10.38145400	35.62207400
H	3.01439100	11.19477800	35.33900900

C	5.00292400	10.77766000	35.96647100
H	5.14881300	11.85404000	36.04645300
C	7.04574300	3.89759300	33.89708300
H	8.05038200	3.50752400	34.04359800
C	6.38230900	4.32828200	35.07204000
H	6.95118000	4.24215100	35.99463400
C	4.98368300	4.60371300	35.26469900
H	4.66798100	4.58755800	36.30679200
C	3.92354200	4.46112000	34.32249900
H	2.92471000	4.48926400	34.74899700
C	3.98048500	4.09098500	32.94664600
H	3.03638500	3.74274100	32.53002300
C	5.13787000	3.70199100	32.18224900
H	4.90415700	3.21990500	31.23643400
C	6.48876900	3.60860400	32.60162300
H	7.14328000	3.02838500	31.95440200
Pd	4.81738800	7.46000400	30.23626500
Pd	4.45283900	9.89431200	31.32138200
Pd	3.44209000	8.77633100	33.64839300
Pd	3.84029200	6.20945800	32.51118800
Pd	7.01044000	9.24282100	30.47579300
Pd	6.35146500	5.53928700	31.52877400
C	2.69274100	7.33464400	29.83378400
H	2.70831200	6.89674000	28.83665600
C	2.68621500	8.76058700	29.85767600
H	2.84348100	9.23617800	28.89361100
C	2.30976100	9.64463800	30.91031400
H	2.09699000	10.66378300	30.59102600
C	1.76179500	9.30774200	32.19681500
H	1.29599700	10.14200700	32.71572500
C	1.51149300	8.02516600	32.74354500
H	0.86519000	8.00707500	33.61812600
C	1.72220000	6.73362900	32.14470500
H	1.19276500	5.92146400	32.63876700
C	2.24999700	6.43115700	30.86431800
H	2.10826200	5.40760600	30.52698600

C	5.85102900	6.01441500	28.77866100
H	5.02040800	5.42850700	28.39363200
C	5.92012700	7.35477700	28.32247400
H	5.15983400	7.64567500	27.60043400
C	7.01643600	8.28118700	28.42989500
H	6.92320600	9.15320900	27.78662800
C	8.29771100	8.07763500	28.99846100
H	9.05025500	8.82092000	28.74654300
C	8.82289300	6.89646900	29.63028700
H	9.91024600	6.86729900	29.68153900
C	8.18789900	5.63081000	29.79037200
H	8.85027400	4.81748700	30.07343800
C	6.86173900	5.23460100	29.44597900
H	6.71585200	4.15822000	29.36599800
C	4.81891500	12.11467000	31.51536800
H	3.81427600	12.50473900	31.36885500
C	5.48234500	11.68398300	30.34040800
H	4.91347200	11.77011600	29.41781600
C	6.88097000	11.40855000	30.14774600
H	7.19666900	11.42470700	29.10565100
C	7.94111400	11.55114100	31.08994300
H	8.93994400	11.52299700	30.66344200
C	7.88417400	11.92127400	32.46579700
H	8.82827500	12.26951700	32.88241700
C	6.72679100	12.31026700	33.23019700
H	6.96050800	12.79235200	34.17601200
C	5.37589200	12.40365800	32.81082600
H	4.72138400	12.98387700	33.45804800
Pd	5.93233000	8.00613100	32.70622400

Cubo- μ 3

Pd	-0.26152000	-1.50393600	2.27771000
Pd	-1.80323400	-2.14302700	0.03350200
Pd	-2.75729900	0.49021700	-0.03717800
Pd	-1.17344200	0.97539600	2.27766600
Pd	0.95424600	-2.63258400	-0.03654100
Pd	1.42940500	0.52507900	2.27919700
C	0.87366000	-3.64963200	3.25742500
H	1.57104000	-3.48125800	4.07407900
C	1.39651100	-4.25055000	2.11886800
H	2.45105300	-4.51131500	2.15720100
C	0.71873100	-4.65582300	0.93720900
H	1.27410200	-5.34121300	0.30260000
C	-0.63796800	-4.50115100	0.58787400
H	-0.95861400	-5.04632700	-0.29564500
C	-1.69212100	-3.85350800	1.31590500
H	-2.68281800	-4.25550400	1.11813400
C	-1.56599300	-3.24640400	2.62755200
H	-2.51597400	-2.95896800	3.07005400
C	-0.46625700	-3.26026200	3.51986400
H	-0.68092700	-2.95213500	4.54102100
C	3.05148400	1.22565500	3.52300800
H	2.89045100	0.88565900	4.54405500
C	3.58998700	0.26599200	2.63095900
H	3.81437600	-0.70149900	3.07365300
C	4.18018700	0.46046800	1.31967100
H	5.02325200	-0.19693300	1.12181900
C	4.21542300	1.69696400	0.59219100
H	4.84834900	1.69178700	-0.29134200
C	3.67144400	2.95041000	0.94310700
H	3.98916200	3.77427200	0.30981300
C	2.98040900	3.33551100	2.12362800
H	2.67788400	4.37834700	2.16554300
C	2.71999600	2.58042200	3.26106600
H	2.22465800	3.09895200	4.07789100

C	2.49159200	-3.38863900	-1.31670900
H	2.34357600	-4.44765000	-1.11901000
C	3.57818500	-2.80070200	-0.58832500
H	3.88834000	-3.35084900	0.29510700
C	4.39304700	-1.70420700	-0.93622700
H	5.26361900	-1.56706400	-0.30106300
C	4.38373400	-0.91316800	-2.11802600
H	5.13792700	-0.13201500	-2.15481700
C	3.60098700	-1.06503100	-3.25630400
H	3.80429600	-0.37724700	-4.07267300
C	2.59243300	-2.02980900	-3.51872200
H	2.21830500	-2.06061200	-4.53972600
C	2.03068400	-2.97535300	-2.62676200
H	1.30615500	-3.65391500	-3.06978900
Pd	0.26132700	1.50395700	-2.27774400
Pd	1.80322400	2.14304300	-0.03381200
Pd	2.75739900	-0.49013100	0.03677500
Pd	1.17347700	-0.97548000	-2.27772100
Pd	-0.95418800	2.63275200	0.03698000
Pd	-1.42945800	-0.52517800	-2.27941700
C	-0.87400100	3.64981900	-3.25703800
H	-1.57152400	3.48148700	-4.07357800
C	-1.39663900	4.25071400	-2.11838000
H	-2.45118200	4.51150000	-2.15652800
C	-0.71866400	4.65593800	-0.93681000
H	-1.27392200	5.34132600	-0.30210100
C	0.63810400	4.50125600	-0.58772400
H	0.95891900	5.04639200	0.29575900
C	1.69211900	3.85363200	-1.31597600
H	2.68287500	4.25554800	-1.11831800
C	1.56576000	3.24656000	-2.62761800
H	2.51567000	2.95913600	-3.07029100
C	0.46585800	3.26042200	-3.51972300
H	0.68035100	2.95234500	-4.54093400
C	-3.05168800	-1.22591000	-3.52295500
H	-2.89074500	-0.88592200	-4.54401900

C	-3.59019800	-0.26625400	-2.63089700
H	-3.81467500	0.70121500	-3.07360300
C	-4.18027600	-0.46069200	-1.31954700
H	-5.02332400	0.19672400	-1.12163200
C	-4.21543100	-1.69717800	-0.59202500
H	-4.84824200	-1.69197000	0.29159000
C	-3.67155800	-2.95064000	-0.94301800
H	-3.98920700	-3.77451100	-0.30970300
C	-2.98057400	-3.33574400	-2.12357100
H	-2.67806700	-4.37858800	-2.16548800
C	-2.72019700	-2.58067300	-3.26102800
H	-2.22490900	-3.09922200	-4.07787200
C	-2.49132400	3.38884400	1.31740300
H	-2.34322800	4.44787900	1.11989700
C	-3.57790100	2.80112300	0.58887700
H	-3.88807600	3.35151900	-0.29439100
C	-4.39264400	1.70440400	0.93636300
H	-5.26319300	1.56743000	0.30113000
C	-4.38328000	0.91303500	2.11792500
H	-5.13740200	0.13180400	2.15447800
C	-3.60061500	1.06466800	3.25631000
H	-3.80392600	0.37663900	4.07246900
C	-2.59218300	2.02947900	3.51909200
H	-2.21809000	2.06000400	4.54011400
C	-2.03041400	2.97523300	2.62736900
H	-1.30585400	3.65365900	3.07056800
Pd	-0.00000100	0.00001800	-0.00007400

Anti- μ 4

Pd	-1.66035600	2.44814200	-0.21456600
Pd	-2.67997100	-0.10227600	-0.70003500
Pd	-1.53054000	-1.27995100	1.72647100
Pd	-1.19114200	1.33356900	2.19162600
Pd	-0.89426700	1.04806500	-2.44956000
Pd	0.98124300	2.47550100	-0.74291700
C	-4.30391200	1.74281800	0.66432700
H	-4.65656700	2.53659500	0.00990900
C	-4.58977000	0.41094800	0.20363900
H	-5.29185100	0.35464400	-0.62620200
C	-4.26374300	-0.80223000	0.82873600
H	-4.71808600	-1.69175500	0.40205700
C	-3.64419400	-1.02541900	2.12471400
H	-3.85822200	-2.01056600	2.53527900
C	-3.32259400	-0.04798300	3.10231900
H	-3.08845300	-0.43611600	4.09091600
C	-3.36761700	1.35954500	3.00240100
H	-3.23808000	1.89178500	3.94189100
C	-3.85899600	2.13463200	1.91454200
H	-3.89643000	3.20472500	2.09600800
C	0.82818000	2.48594600	-3.68249400
H	0.08078600	3.09900600	-4.18158600
C	1.00834800	1.21254300	-4.18958800
H	0.39257300	0.97291700	-5.05158800
C	1.97127600	0.18587100	-3.83470900
H	2.09168200	-0.58503600	-4.59844600
C	3.14058700	0.38176700	-3.03777400
H	3.91829200	-0.36404700	-3.18361700
C	3.49279500	1.47407600	-2.21018000
H	4.54733200	1.52676800	-1.94888300
C	2.78712500	2.73587100	-2.04727500
H	3.37799700	3.50584200	-1.55611200
C	1.62645800	3.18765900	-2.70332100
H	1.44368000	4.25948000	-2.66495500

C	-3.51084900	-1.47929600	-2.18324900
H	-4.56326300	-1.53295100	-1.91375200
C	-3.16579300	-0.38634900	-3.01278200
H	-3.94572900	0.35811600	-3.15362000
C	-2.00189400	-0.18824000	-3.81756900
H	-2.12946600	0.58200200	-4.58089400
C	-1.04094900	-1.21414200	-4.18025100
H	-0.43215900	-0.97367800	-5.04696200
C	-0.85631100	-2.48789700	-3.67571700
H	-0.11229800	-3.10007100	-4.18085600
C	-1.64623200	-3.19041100	-2.69052500
H	-1.46139300	-4.26189600	-2.65282200
C	-2.80230800	-2.74013800	-2.02528500
H	-3.38846200	-3.51088400	-1.52966900
Pd	1.65598900	-2.44929000	-0.22757500
Pd	1.54501000	1.28046200	1.71483000
Pd	2.67334800	0.10172000	-0.71732300
Pd	0.87320500	-1.04561500	-2.45432100
Pd	1.21135300	-1.33227800	2.18204800
Pd	-0.98837000	-2.47558500	-0.73489100
C	3.39181000	-1.36144000	2.97655100
H	3.26998400	-1.89493800	3.91635400
C	3.34712500	0.04572500	3.07877200
H	3.12048900	0.43253200	4.06962600
C	3.66025900	1.02480700	2.09978200
H	3.87750500	2.00935000	2.51013600
C	4.26916700	0.80356000	0.79852200
H	4.71830500	1.69413700	0.36844900
C	4.59171600	-0.40862400	0.16965200
H	5.28695900	-0.35053700	-0.66579300
C	4.31249400	-1.74136900	0.63179400
H	4.66078400	-2.53396800	-0.02635600
C	3.87600400	-2.13510300	1.88401300
H	3.91490500	-3.20541200	2.06380900
C	0.29386700	-4.50914100	0.24626800
H	0.58596000	-5.13819800	-0.59101300

C	1.34395200	-4.11543300	1.10818600
H	2.28782200	-4.63154300	0.92758000
C	1.21607300	-3.59165400	2.46505400
H	2.09414600	-3.74486200	3.08392200
C	0.07790500	-3.18496900	3.15663000
H	0.20917500	-2.98133600	4.21632800
C	-1.29757400	-3.23251000	2.71891200
H	-2.01977300	-3.11196600	3.52471600
C	-1.80597300	-3.78136700	1.52944900
H	-2.88507500	-3.90851100	1.49437600
C	-1.11412200	-4.31707600	0.39188900
H	-1.72710000	-4.95029000	-0.24686400
C	1.12155000	4.31743400	0.38339900
H	1.73066500	4.94963300	-0.26001000
C	1.82093700	3.78099700	1.51604800
H	2.89987200	3.90744300	1.47354400
C	1.32038600	3.23298100	2.70915500
H	2.04790000	3.11241500	3.51015300
C	-0.05224500	3.18502800	3.15589900
H	-0.17640300	2.98281300	4.21671400
C	-1.19496100	3.59151200	2.47141100
H	-2.06873900	3.74567100	3.09609600
C	-1.33187100	4.11504700	1.11523800
H	-2.27643000	4.63238500	0.94152100
C	-0.28702800	4.50937600	0.24687400
H	-0.58469000	5.13821100	-0.58860500
Pd	-0.00142800	0.00096900	-0.08816500

Anti- μ 3;4

Pd	0.40541900	2.78979900	0.38219900
Pd	0.83770500	1.60137700	-2.04269500
Pd	-1.69607100	0.54709400	-2.27141500
Pd	-2.11971300	1.64314500	0.40513800
Pd	2.58539700	0.93251300	0.05688800
Pd	0.92448900	1.00546600	2.38505100
C	-0.78783100	4.40014200	-0.40795600
H	-0.51203500	5.24489300	0.22154600
C	0.08351500	4.15197300	-1.50441300
H	0.96748700	4.78194200	-1.54467400
C	-0.17427900	3.41988700	-2.71410200
H	0.47684100	3.67954400	-3.54722300
C	-1.37587000	2.75071000	-3.08412800
H	-1.44183500	2.49909600	-4.14036200
C	-2.57533300	2.55833100	-2.33465000
H	-3.44665700	2.28445300	-2.92839800
C	-2.90665200	3.08892000	-1.03350800
H	-3.97394000	3.08893000	-0.82135400
C	-2.11915700	3.91304600	-0.19030600
H	-2.64349300	4.38819600	0.63442700
C	1.43607500	3.80938400	2.33191100
H	0.77830400	4.67349800	2.38333500
C	2.28170400	3.74042300	1.18405800
H	2.30982400	4.64005600	0.57398400
C	3.41265000	2.86685200	0.99317400
H	4.12849200	3.21230300	0.25146500
C	3.78485600	1.73255100	1.73568300
H	4.77057900	1.31870300	1.53286800
C	3.14595100	1.22057500	2.93560100
H	3.60055800	0.31509100	3.32796300
C	2.26883500	1.88451400	3.81583600
H	2.15672900	1.47220700	4.81530700
C	1.42964000	2.99175900	3.48401300
H	0.73611600	3.29560800	4.26402400

C	2.18452300	0.85934300	-3.68964000
H	1.85967500	1.48144100	-4.52036100
C	2.95975700	1.50760100	-2.69544300
H	3.14892300	2.56426700	-2.87415800
C	3.86224500	0.92533800	-1.71082800
H	4.64589300	1.60645800	-1.38384500
C	4.14874100	-0.44004100	-1.50011600
H	5.02016900	-0.64893200	-0.88567000
C	3.51333200	-1.58587500	-2.07107400
H	4.08206200	-2.50898800	-1.97049600
C	2.56967700	-1.61681800	-3.16031000
H	2.44346100	-2.60315700	-3.60051600
C	1.99786800	-0.55320300	-3.89368600
H	1.54450700	-0.82056100	-4.84512800
Pd	-0.43584500	-2.64067500	-0.11666200
Pd	-1.50928900	-0.16581000	2.33471600
Pd	0.70815700	-1.61837500	2.09364200
Pd	2.21847700	-1.71573400	-0.35821700
Pd	-2.68824600	-0.93849000	-0.15401600
Pd	0.44399300	-1.12994600	-2.38705900
C	0.90495400	-4.17806800	0.69630600
H	0.34873400	-4.99577900	0.23818300
C	0.53677300	-3.89769800	2.06523900
H	-0.42956400	-4.29568800	2.36306700
C	1.32549700	-3.38474100	3.12816000
H	0.87238300	-3.45084900	4.11538400
C	2.67477600	-2.91288200	3.09500200
H	3.05645000	-2.54861100	4.04517700
C	3.55272100	-2.89840400	2.02518500
H	4.54682000	-2.50756200	2.22786300
C	3.35523300	-3.36419700	0.69704700
H	4.26643400	-3.46081200	0.11126500
C	2.19762700	-3.93272800	0.10612400
H	2.35036600	-4.42103300	-0.85353600
C	-0.86285500	-3.66182700	-2.07512400
H	-0.19728500	-4.52099000	-2.07702100

C	-1.99823700	-3.77352400	-1.23171900
H	-2.08173200	-4.69726800	-0.66241700
C	-3.15435100	-2.94103100	-1.18209900
H	-3.97133300	-3.34271500	-0.58929100
C	-3.47842000	-1.81344800	-1.99198700
H	-4.52369200	-1.51094400	-1.98032600
C	-2.71587200	-1.22176300	-3.05415100
H	-3.31649900	-0.61629600	-3.73549600
C	-1.49734700	-1.72027500	-3.64279500
H	-1.28668100	-1.33853200	-4.63803300
C	-0.67759800	-2.78855100	-3.21682200
H	0.00758100	-3.17646800	-3.96862800
C	-3.64286000	-1.36081600	3.38060000
H	-3.50537800	-2.12514200	4.14114600
C	-4.24568500	-1.76704700	2.20196800
H	-4.51422000	-2.81824700	2.13967600
C	-4.59499400	-0.99256900	1.06269000
H	-5.26539300	-1.49327400	0.36665200
C	-4.50989900	0.41482900	0.87425900
H	-5.09160300	0.81325300	0.04722600
C	-3.87402700	1.38504300	1.69922000
H	-4.22534400	2.40949300	1.57981100
C	-3.18002900	1.11744500	2.94587000
H	-2.84195800	2.01338000	3.46029000
C	-3.17117900	-0.06731300	3.73209500
H	-2.81934200	0.05151400	4.75455600
Pd	0.00950000	0.01295000	0.04505200

Ih-μ3

Pd	0.00000000	0.00000000	0.04256900
Pd	1.47122600	2.34556800	0.15751000
Pd	-2.71537400	0.12305400	0.11268200
Pd	-0.99877100	2.06961400	-1.42608200
Pd	1.33614800	0.64157900	-2.14918800
Pd	-1.33614800	-0.64157900	-2.14918800
Pd	0.99877100	-2.06961400	-1.42608200
Pd	-1.47122600	-2.34556800	0.15751000
Pd	2.71537400	-0.12305400	0.11268200
Pd	1.15489300	-2.08696700	1.35956200
Pd	-1.27044900	-0.56131200	2.33536000
Pd	1.27044900	0.56131200	2.33536000
Pd	-1.15489300	2.08696700	1.35956200
C	-3.92342900	1.86690400	0.94891400
C	-3.02366600	2.96673700	0.68625500
C	-2.65976700	3.45891900	-0.62261900
C	-3.01631200	2.94967900	-1.89386900
C	-3.80467800	1.81526800	-2.21683400
C	-4.54258000	0.99833400	-1.36484600
C	-4.66926500	1.06951700	0.05024400
H	-5.15544000	0.23752600	-1.84201800
H	-5.44537600	0.45148600	0.49557100
H	-4.20512500	1.76459400	1.99373500
H	-3.02543400	3.72963300	1.46555600
H	-2.21742000	4.45205900	-0.62203400
H	-2.79910500	3.60116100	-2.73697700
H	-3.89811100	1.60729500	-3.27951400
C	-0.20597900	3.97188600	2.15665100
C	-0.69798200	3.23238100	3.25658500
C	0.00000000	2.31604900	4.08918000
C	1.34545500	1.92816000	4.03477200
C	2.32783600	2.25556000	3.03323900
C	2.26345700	3.30543000	2.04034500
C	1.14597400	4.06307400	1.65874600

H	3.23157100	3.65806900	1.69015300
H	1.34753000	4.94555100	1.05844700
H	-0.85717500	4.76800700	1.80289100
H	-1.68980900	3.50587000	3.61051400
H	-0.57563800	1.90877300	4.91603100
H	1.71212800	1.36225300	4.88902500
H	3.34140100	1.93583600	3.26616800
C	2.29558200	3.75478100	-1.39653100
C	0.98534100	4.10673700	-1.77766500
C	0.11916400	3.50028700	-2.72559700
C	0.48484800	2.52175000	-3.72040200
C	1.70909400	1.87971800	-3.92822200
C	2.86687100	1.90989600	-3.07036500
C	3.11804700	2.72953900	-1.95646100
H	3.71322700	1.32355600	-3.41949900
H	4.10745300	2.63572900	-1.51671600
H	2.81712000	4.47046600	-0.76501200
H	0.58240800	4.99273200	-1.29326400
H	-0.73417700	4.11429300	-3.00383900
H	-0.28040500	2.32176600	-4.46665700
H	1.80719000	1.30212300	-4.84359000
C	-0.98534100	-4.10673700	-1.77766500
C	-2.29558200	-3.75478100	-1.39653100
C	-3.11804700	-2.72953900	-1.95646100
C	-2.86687100	-1.90989600	-3.07036500
C	-1.70909400	-1.87971800	-3.92822200
C	-0.48484800	-2.52175000	-3.72040200
C	-0.11916400	-3.50028700	-2.72559700
H	0.28040500	-2.32176600	-4.46665700
H	0.73417700	-4.11429300	-3.00383900
H	-0.58240800	-4.99273200	-1.29326400
H	-2.81712000	-4.47046600	-0.76501200
H	-4.10745300	-2.63572900	-1.51671600
H	-3.71322700	-1.32355600	-3.41949900
H	-1.80719000	-1.30212300	-4.84359000
C	0.20597900	-3.97188600	2.15665100

C	0.69798200	-3.23238100	3.25658500
C	0.00000000	-2.31604900	4.08918000
C	-1.34545500	-1.92816000	4.03477200
C	-2.32783600	-2.25556000	3.03323900
C	-2.26345700	-3.30543000	2.04034500
C	-1.14597400	-4.06307400	1.65874600
H	-3.23157100	-3.65806900	1.69015300
H	-1.34753000	-4.94555100	1.05844700
H	0.85717500	-4.76800700	1.80289100
H	1.68980900	-3.50587000	3.61051400
H	0.57563800	-1.90877300	4.91603100
H	-1.71212800	-1.36225300	4.88902500
H	-3.34140100	-1.93583600	3.26616800
C	3.02366600	-2.96673700	0.68625500
C	2.65976700	-3.45891900	-0.62261900
C	3.01631200	-2.94967900	-1.89386900
C	3.80467800	-1.81526800	-2.21683400
C	4.54258000	-0.99833400	-1.36484600
C	4.66926500	-1.06951700	0.05024400
C	3.92342900	-1.86690400	0.94891400
H	5.44537600	-0.45148600	0.49557100
H	4.20512500	-1.76459400	1.99373500
H	3.02543400	-3.72963300	1.46555600
H	2.21742000	-4.45205900	-0.62203400
H	2.79910500	-3.60116100	-2.73697700
H	3.89811100	-1.60729500	-3.27951400
H	5.15544000	-0.23752600	-1.84201800