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## Supporting Information

### 2 Crystalline Confinement Leads to Broadening of Absorption Spectra through Activated 3 Spin-Forbidden Transitions in Alq<sub>3</sub>-Irppy<sub>2</sub>acac Engineered Crystals

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### 11 Theoretical Background

12 Photoexcitation is a quantum transition described by the Fermi Golden rule as<sup>1-3</sup>

$$\sigma_{\text{ex}}(\omega) = \frac{4\pi^2\omega}{3c} \sum_{v,u} P_{iv}(T) |\langle \Phi_f | \hat{\mu}_{fi} | \Phi_i \rangle|^2 |\langle \Theta_{fu} | \Theta_{iv} \rangle|^2 \delta(\hbar\omega - E_{fi} - E_{fu} + E_{iv}), \quad (1)$$

14 where, the  $\Phi$  and  $\Theta$  are the electronic and vibrational wavefunctions, and indices  $i$  and  $f$   
15 represent the initial and final states, respectively;  $\omega$  is the circular frequency, the  $c$  is the speed of  
16 light in vacuum, the  $E_{fi}$  is the electronic energy difference between the initial and the final states,  
17 the  $E_{iv}$  is the vibrational energy level of the initial state and the  $E_{fu}$  is the vibrational energy level of  
18 the final state. This equation describes both electronic and vibrational transitions. The electronic  
19 transition is characterized by the matrix element  $\langle \Phi_f | \hat{\mu}_{fi} | \Phi_i \rangle$  called TDM, which is spin-allowed if  
20  $\Phi_i$  and  $\Phi_f$  have a same spin state. When the spin states of  $\Phi_i$  and  $\Phi_f$  are different, the value of the  
21 TDM is zero and the transition is spin-forbidden.

22 To allow a spin-forbidden transition, between the pure singlet,  $S$ , and triplet,  $T$ , the states have  
23 to mix with an intermediate state through the SOC as<sup>4-6</sup>

$$|S'\rangle = |S\rangle + \sum_n \sum_{m=-1}^1 \frac{\langle {}^3T_{n,m} | H_{\text{SO}} | S \rangle}{{}^1E_S - {}^3E_n} |{}^3T_{n,m}\rangle, \quad (2)$$

$$|T_m'\rangle = |T_m\rangle + \sum_n^{\{\text{singlets}\}} \frac{\langle {}^1S_n | H_{\text{SO}} | {}^3T_m \rangle}{{}^3E_T - {}^1E_n} |{}^1S_n\rangle, \quad (3)$$

26 where  $S'$  and  $T'$  are the mixed states and  $S_n$  and  $T_n$  are the intermediate states. The index  $n$  is  
 27 for the intermediates,  $m$  is the magnetic spin quantum number ( $m = -1, 0, 1$ ), and  $\hat{H}_{SO}$  is the SOC  
 28 Hamiltonian operator.

29 In this mixing, the TDM and SOC interplay to generate a new spin-forbidden TDM ( $TDM_{SF}$ ).  
 30 There are two processes that result in non-zero  $TDM_{SF}$ . When the intermediate state is a singlet ( $S_n$ ),  
 31 the  $TDM_{SF}$  from the  $S_0$  state to the  $T_k$  state (final state) is expressed as

$$32 \quad TDM_{SF} = \frac{\langle \Phi_{S_n} | \hat{\mu} | \Phi_{S_0} \rangle \langle \Phi_{T_k} | \hat{H}_{SO} | \Phi_{S_n} \rangle}{E_{T_k} - E_{S_n}} \quad (4)$$

33 It involves two parts: the spin-allowed part, naming  $TDM = \langle \Phi_{S_n} | \hat{\mu} | \Phi_{S_0} \rangle$  and the spin-  
 34 forbidden part, naming  $SOC = \langle \Phi_{T_k} | \hat{H}_{SO} | \Phi_{S_n} \rangle$ .

35 The spin-forbidden TDM through a triplet mixed intermediate ( $T_n$ ) is expressed as

$$36 \quad TDM_{SF} = \frac{\langle \Phi_{T_k} | \hat{\mu} | \Phi_{T_n} \rangle \langle \Phi_{T_n} | \hat{H}_{SO} | \Phi_{S_0} \rangle}{E_{T_n} - E_{S_0}} \quad (5)$$

37 The schematic graphic of the spin-forbidden excitation transition is shown in Figure 1(a). To  
 38 sum up, a strong  $TDM_{SF}$  requires both strong spin-allowed TDM and SOC.

39 Calculation Formulas for the Photophysical Parameters:

$$40 \quad k_r = \frac{\Phi_p}{\tau_p} \quad (6)$$

$$41 \quad k_{nr} = \frac{1 - \Phi_p}{\tau_p} \quad (7)$$

42 Where  $\Phi_p$  is the prompt phosphorescence component of  $\Phi_{PL}$ ;  $\tau_p$  is the lifetime of prompt  
 43 phosphorescence;  $k_r$  is the rate constant of radiative transition from  $T_1$  to  $S_0$ ;  $k_{nr}$  is the non-  
 44 radiative decay rate constant from  $T_1$  to  $S_0$ .

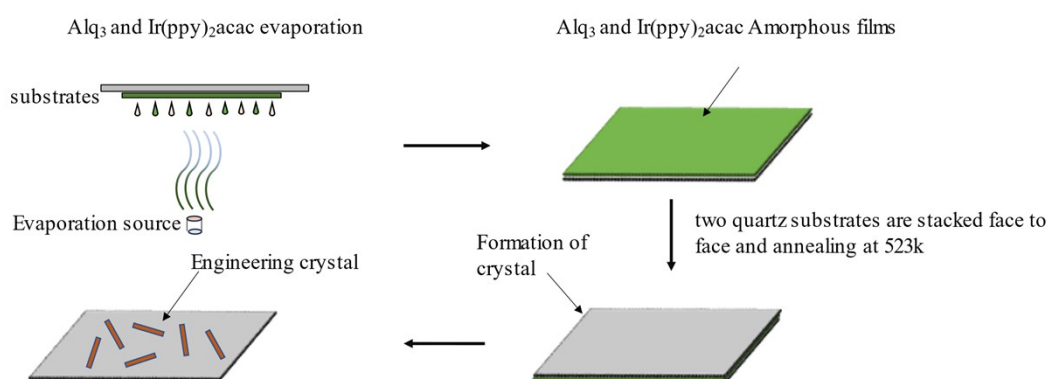
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## 46 Experimental methods:

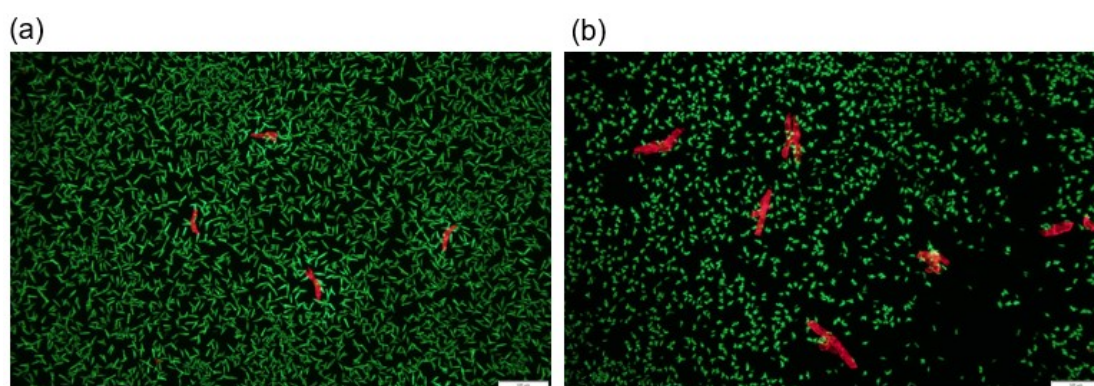
47 (1) Fabrication of engineered crystals

48 (2,4-pentanedionyl) bis (2-phenylpyridine) iridium ( $Irppy_2acac$ , 99%) (CAS:337526-85-9) and  
 49 tris(8-hydroxyquinoline) aluminum ( $Alq_3$ , 98%) (CAS: 2085-33-8) were purchased from Bide

50 Pharmatech Ltd. To take double-film annealing method get high-quality engineering organic crystal  
 51 (fig. S1). Firstly, Irppy<sub>2</sub>acac (90%) and Alq<sub>3</sub> (10%) were mixed and deposited on a quartz substrate  
 52 in a vacuum evaporator. The thickness of the deposited film was 120 nm. Secondly, another quartz  
 53 substrate is horizontally stacked face to face and then heated at 523k for 7 minutes. The temperature  
 54 control precision is ±0.2 K. (Heating at a temperature of 523K, according to TGA data, provided  
 55 in **figure S4**, the sample is safe and has not undergone decomposition). Then the heating was  
 56 stopped and the double substrates were naturally cooled. By comparing the crystal morphology of  
 57 different doping ratios, the optimal crystal morphology was found at the ratio of Irppy<sub>2</sub>acac (90):  
 58 Alq<sub>3</sub> (10), while crystals with other doping ratios are shown in the **figure S2**. The content of Ir and  
 59 Al elements in the sample was determined by inductively coupled plasma mass spectrometry (ICP-  
 60 MS), and the results showed that the content of Alq<sub>3</sub> in the crystal was 4.2%.



61  
 62 **Figure S1.** The process to fabricate engineered crystals.

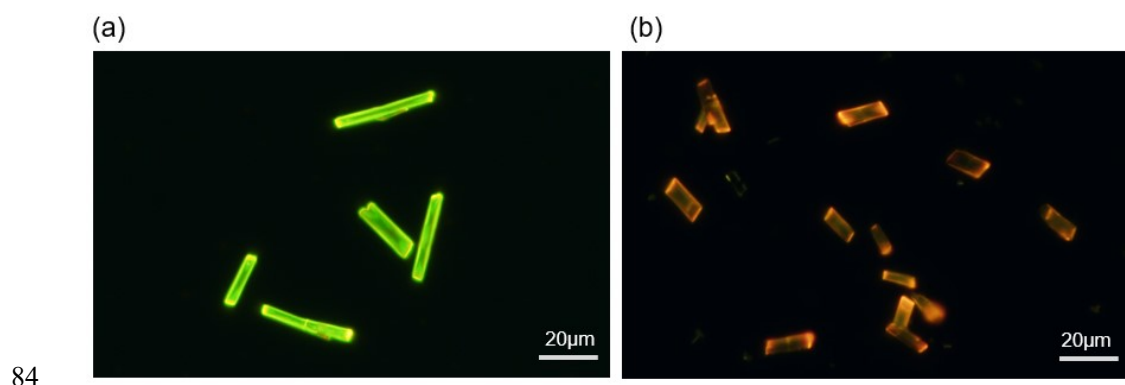


63  
 64 **Figure S2.** (a) Crystal morphology of Irppy<sub>2</sub>acac (30): Alq<sub>3</sub> (70), (b) Crystal morphology of Irppy<sub>2</sub>acac  
 65 (60): Alq<sub>3</sub> (40).

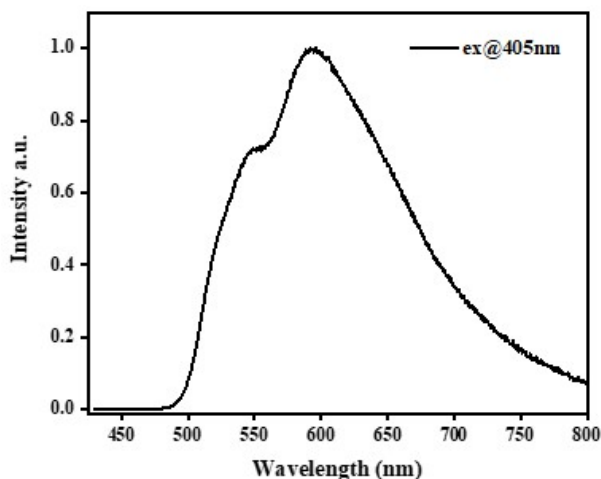
66 Crystals doped with more proportions of host and guest were presented. The images of the  
 67 morphology of some proportions of crystals are show in **Figure S2**. When the ratio between  
 68 Irppy<sub>2</sub>acac and Alq<sub>3</sub> is 3:7 or 6:4 a lot of green crystals can be observed. When the proportion of

69 Irppy<sub>2</sub>acac is increased the size of the red crystals tend to be larger and the green crystals tend to be  
70 smaller. When the ratio is raised to 9:1, as shown in Figure 1 in the main text, all the crystals convert  
71 to red.

72 Two crystallization phases with green and yellow luminescent colors were prepared. As shown  
73 in **Figure S3 (a)** green luminescent crystals were obtained under 523 K and yellow crystals were  
74 obtained when the temperature was increased to 543 K, as shown in **Figure S3 (b)**. The luminescent  
75 spectrum of the yellow crystals excited at 405 nm were measured as shown in **Figure S4**. Two peaks  
76 were observed. One is at 541 nm and the other is at about 592 nm. The spectrum of the green  
77 crystallization phase as shown in **Figure 1(c)** has two peaks. One peak is at 518 nm and the other  
78 peak is at 541 nm. And there is also a shoulder peak at about 585 nm that is close to the peak at 592  
79 nm of the yellow crystallization phase. Therefore, this strong peak at 592 nm should be one of the  
80 vibronic peak of Irppy<sub>2</sub>acac. In some certain crystallization phase, the vibronic coupling can be  
81 significantly enhanced. The thermal gravimetric analyzer (TGA) curves of Irppy<sub>2</sub>acac was also  
82 measured as shown in **Figure S5**. Below 613 K no significant weight loss was observed. The  
83 annealing at 543 K should be safe.



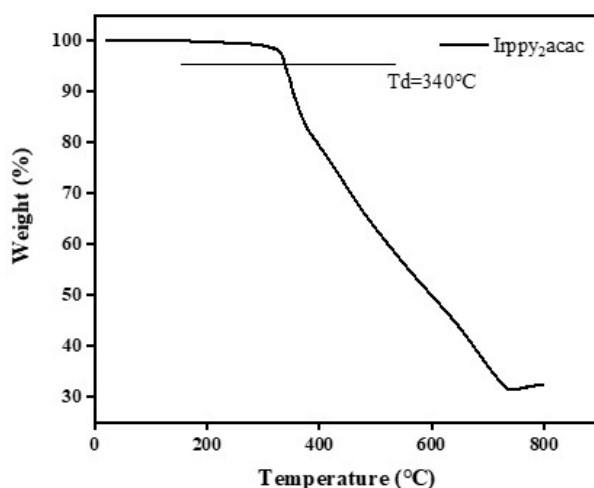
85 **Figure S3.** The different luminescent colors of Irppy<sub>2</sub>acac crystals (a) yellow green (heating  
86 temperature: 250°C); (b) Orange light (heating temperature: 270°C).



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**Figure S4.** The luminescence spectrum of Irppy<sub>2</sub>acac orange single crystals.



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**Figure S5.** The thermal gravimetric analyzer (TGA) curves of Irppy<sub>2</sub>acac.

92 (2) Observation and spectrum measurement

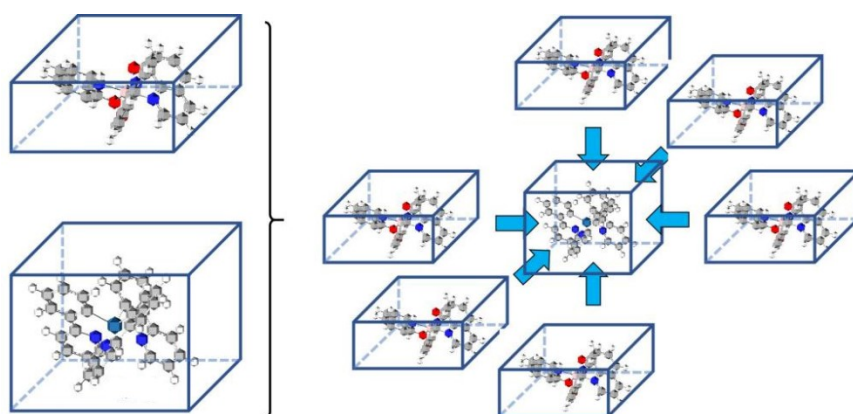
93 The observation of morphology was performed using an Olympus fluorescence microscope  
 94 with UV light source and a Thermo Scientific Prisma E SEM. The photoluminescence spectra  
 95 were measured by a Renishaw microscope. The excitation light sources are 405 nm, 589 nm, and  
 96 632 nm laser. The photoluminescence lifetimes of the long wavelength phosphorescence were  
 97 measured by an Edinburgh spectrophotometer using a  $\mu$ s pulse flash lamp. XRD diffraction spectra  
 98 were measured by a Bruker D8 Advance instrument. The transient absorption spectra were  
 99 measured by a Helios Fire automated femtosecond Transient Absorption Spectrometer with a 390  
 100 nm excitation, and the time window is 8 ns.

101 **Computational details:**

102 Geometry optimizations and single point computations of a ground state were performed by a

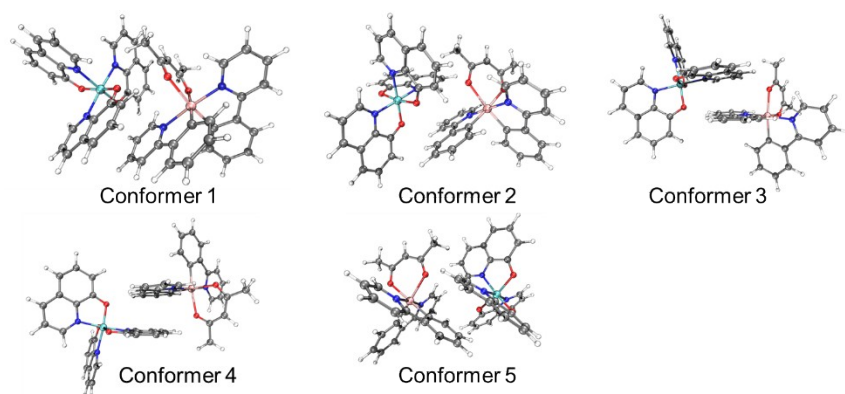
103 DFT method. A hybrid functional M06L<sup>8</sup> was applied with D3 correction<sup>9</sup>, and the basis set was  
104 def2-SVP<sup>10-11</sup> and the auxiliary basis is def2/J. The spin-orbital coupling (SOC) matrix elements  
105 and the transition dipole moment (TDM) were calculated by TDA. For the computation of SOC, the  
106 DKH method was adopted. The basis set was DKH-def2-TZVP<sup>12-13</sup> a version for DKH method<sup>14</sup>  
107 (the auxiliary basis is SARC/J<sup>12-13</sup>). To well describe the SOC matrix elements, no effective core  
108 potential for Ir was adopted. All the methods above were provided by ORCA 5.0.4<sup>15-16</sup> and the grid  
109 size was set as high as “DEFGRID3”<sup>17</sup>.

110 The calculation of transition dipole moments between excited states was performed by  
111 Multiwfn 3.7<sup>18-22</sup> based on the computational results from ORCA. Based on the computations of  
112 SOC and TDM, the spin-forbidden TDMs (TDM<sub>SFS</sub>) can be calculated, and 30 intermediate states  
113 were considered. The conformer searching of the two-molecule system was performed by Crest  
114 module in XTB program<sup>23-27</sup>. The conformer searching depends on the initial geometry sent into the  
115 program. In this work, the initial geometries of a dimer were created by joining the two molecules  
116 in the direction of  $\pm x$ ,  $\pm y$ , and  $\pm z$  (shown in Figure S6). To avoid calculating conformers of extreme  
117 close geometries, e.g., the only difference between two conformers is the rotation of a methyl group,  
118 the minimum energy difference between two conformers was set 0.5 kcal mol<sup>-1</sup>. To cover different  
119 possible conformers in lattice confinements, five conformers with the lowest energies were selected  
120 as shown in Figure S7. The coordinates optimized in ground state were listed in Table S4-S8.  
121 Conformer 1 as the most stable one was studied emphatically. And the excitation spectra of all the  
122 five conformers were calculated and overlapped to create an excitation spectrum for the engineered  
123 crystal.



124

125 **Figure S6.** The generation of initial geometries for conformer searching. The initial geometries of  
126 two-molecule-systems were created by joining the two molecules in the direction of  $\pm x$ ,  $\pm y$ , and  $\pm z$ .  
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129 **Figure S7.** The graph of the five searched conformers with the lowest energies. From Conformer 1  
130 to 5 the energy raises. The coordinates were shown in Table S4 to S8.

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132 **Supplementary data:**

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134 **Table S1.** The luminescence lifetimes and the proportions at different temperatures of the  
 135 engineered crystals excited at 633 nm, 405 nm, and 590 nm.

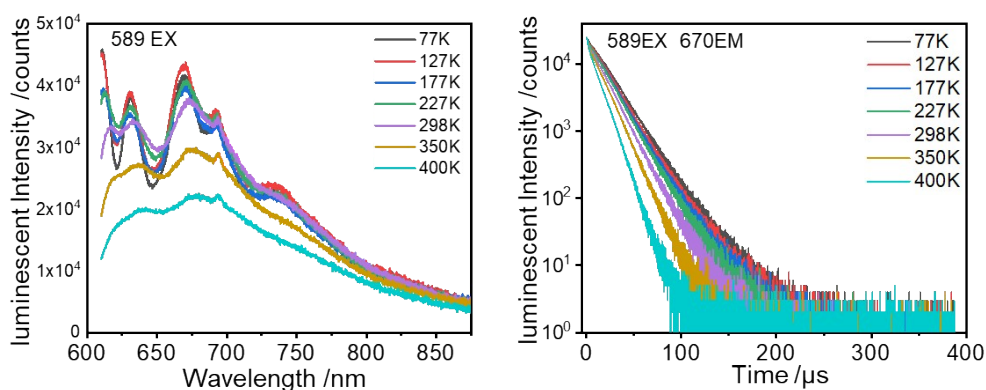
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Ex@633	Temperature /K	Lifetimes	$\chi^2$	Ex@405	Temperature	Lifetimes	$\chi^2$
Em@702		/μs		Em@609	/K	/μs	
	77k	1.27	1.07		77k	12.59	1.06
		12.07				20.34	
	127k	1.27	1.16		127k	10.29	1.05
		11.88				18.89	
	177k	1.31	1.04		177k	9.54	1.09
		11.58				18.09	
	227k	1.40	1.19		227k	9.11	1.12
		12.00				17.56	
	298k	1.44	1.04		298k	6.04	1.12
		12.78				14.98	
	350k	1.33	1.10		350k	3.89	1.05
		12.36				11.67	
	400k	1.25	1.07		400k	3.32	1.11
		10.61				8.59	

Ex@590	Temperature	Lifetimes	$\chi^2$
Em@670	/K	/μs	
	77K	18.3	0.98
		26.9	
	127K	17.3	1.08
		27.2	
	177K	17.5	1.17
		32.4	
	227K	16.7	1.06
		29.5	
	298K	15.5	0.86
		29.6	
	350K	13.2	0.79
		19.0	
	400K	10.6	0.98
		10.0	

137

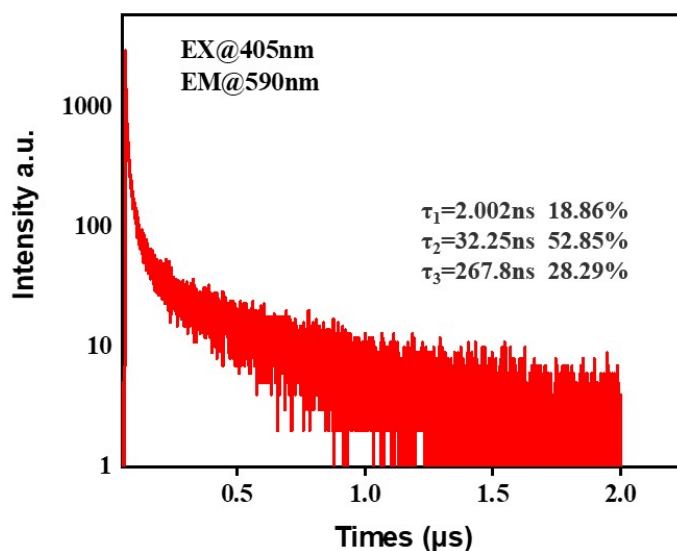




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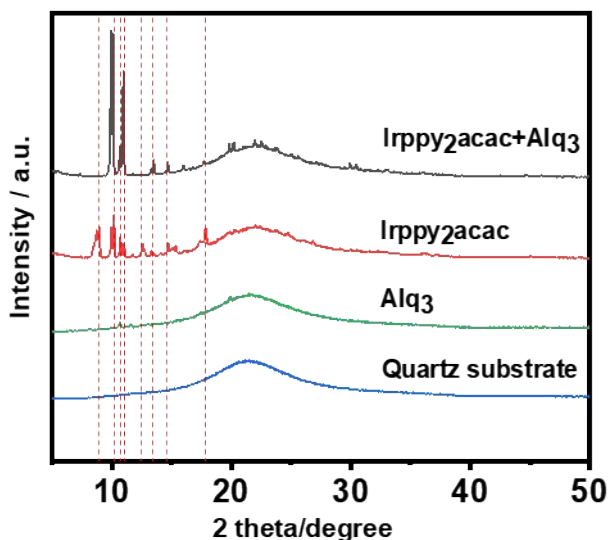
139 **Figure S8.** (a) the spectra excited at 589 nm, (b) the lifetime measurements at 670 nm excited at  
 140 590 nm.

141 For the luminescent spectrum excited at 589 nm, the maximum luminescent intensity is found  
 142 at 127 K. The luminescent intensities below 227 K are close. Higher than 227 K the luminescent  
 143 intensity becomes lower when the temperature increases. This behavior is in between the spectra  
 144 excited at 405 nm and 633 nm. The change of the lifetime with the temperature is similar with that  
 145 excited at 405 nm. The emission at 670 nm contains some component of the phosphorescence of  
 146 Irppy<sub>2</sub>acac as shown in Figure 1 (c) and the excitation at 589 nm also involves some component of  
 147 spin-allowed excitation. These two reasons lead to the results that the luminescent behavior excited  
 148 at 589 nm is to some extent different from the excitation at 633 nm.



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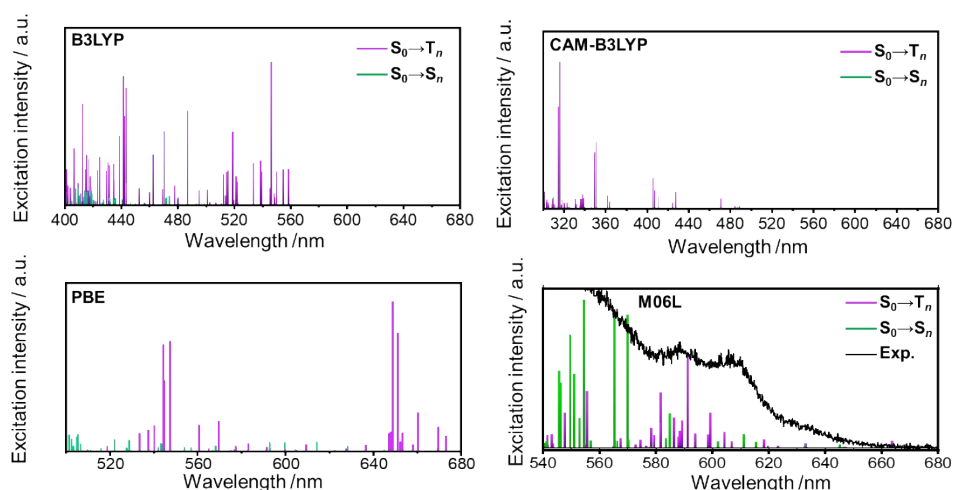
150 **Figure S9.** luminescent lifetimes of the activated emission peak excited at 405 nm about  
 151 Ir(ppy)<sub>2</sub>acac.



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153 **Figure S10.** X-ray diffraction spectra of the engineered crystal and the single-component crystals.

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155

156 **Figure S11.** Calculated excitations using different functionals

157 Four functionals B3LYP (hybrid functional), CAM-B3LYP (range-separated hybrid  
 158 functional), PBE (pure GGA), and M06L (meta-GGA) were tested to calculate the containing spin-  
 159 allowed and spin-forbidden excitations. For B3LYP and cam-B3LYP the excitation energies are  
 160 overestimated. The excitations are shorter than 560 nm. The intensities of the spin-forbidden  
 161 excitations are overestimated. For PBE, the excitation energies are underestimated and the  
 162 wavelength of the strongest excitation extends to over 640 nm, which is away from the experimental  
 163 observation that the maximum absorption is at about 588 nm. The intensities of the spin-forbidden  
 164 excitations are also overestimated. On the whole, M06L is a proper functional to give a best  
 165 performance in reproducing the experimental spectrum. This may be because M06L does not contain

166 the Hartree Fock component as a Meta-GGA functional and thus the low excitation energies are not  
 167 overestimated. But it can approximately describe the dispersion which is difficult for other pure  
 168 GGA functionals like PBE to describe. It also contains the complicated computation of electron  
 169 kinetic energy density which may also help reproduce the excitation intensity.

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s

171 **Table S1.** The original TDM elements between  $S_0$  and  $S_n$  calculated at M06L/def2-svp level.

States		TDM elements between $S_0$ and $S_n$ /debye				
$S_0$	$S_n$	Conformer 1	Conformer 2	Conformer 3	Conformer 4	Conformer 5
0	1	0.44	0.17	0.47	0.52	0.12
0	2	0.39	0.22	0.17	0.26	0.27
0	3	0.68	0.23	0.52	0.36	0.08
0	4	0.07	0.10	0.03	0.18	0.38
0	5	0.35	0.37	0.81	1.57	1.59
0	6	0.11	1.15	1.64	1.08	0.07
0	7	0.43	0.12	0.73	1.17	0.17
0	8	0.36	0.14	0.31	0.34	0.37
0	9	1.43	1.26	0.08	0.62	0.49
0	10	0.40	0.85	0.12	0.09	0.70
0	11	0.42	0.41	1.22	1.23	0.50
0	12	0.18	0.36	0.43	0.47	0.08
0	13	0.93	0.88	0.08	0.20	0.79
0	14	0.90	1.44	0.74	0.75	0.17
0	15	0.62	0.84	0.54	1.32	0.79
0	16	1.04	0.56	1.26	0.45	0.51
0	17	0.29	0.14	1.78	1.16	0.43
0	18	1.23	0.15	0.38	1.78	0.34
0	19	0.50	0.82	0.92	1.10	0.78
0	20	0.17	0.34	0.30	0.16	0.15
0	21	0.70	1.37	0.31	0.47	0.33
0	22	0.73	1.36	1.66	0.09	0.45
0	23	1.23	0.42	0.82	0.36	1.61
0	24	0.68	0.42	0.23	1.21	1.46
0	25	0.37	0.28	0.61	0.50	0.34
0	26	0.90	1.01	0.38	1.99	0.38
0	27	1.35	0.27	2.09	0.83	0.39
0	28	0.98	1.74	0.97	1.09	0.49
0	29	0.79	0.91	0.35	0.28	0.68
0	30	0.73	1.29	0.90	0.18	0.10

172

173 **Table S2.** The original TDM elements between  $T_i$  and  $T_j$  calculated at M06L/def2-svp level.

States		TDM elements between $T_i$ and $T_j$ /debye				
$T_i$	$T_j$	Conformer 1	Conformer 2	Conformer 3	Conformer 4	Conformer 5
1	1	0.00	0.00	0.00	0.00	0.00
1	2	5.30	4.11	2.94	2.84	1.30
1	3	8.61	8.28	2.28	1.08	9.66
1	4	0.34	0.37	3.40	0.38	0.36
1	5	0.00	0.36	2.77	4.43	0.18
1	6	0.46	0.92	0.38	0.18	0.15
1	7	0.62	0.37	0.12	0.11	0.48
1	8	0.44	0.85	0.27	0.36	0.26
1	9	0.33	0.68	0.37	0.36	0.33
1	10	0.82	0.26	0.58	0.36	0.32
1	11	0.31	0.00	1.19	0.17	1.45
1	12	0.90	0.96	0.13	0.24	0.00
1	13	0.72	0.10	0.37	0.12	0.20
1	14	0.24	0.13	1.42	1.37	0.00
1	15	0.31	0.59	0.04	0.01	0.00
1	16	0.07	0.00	0.15	0.05	0.82
1	17	0.00	0.61	0.00	0.00	1.22
1	18	0.00	0.16	0.00	0.00	0.94
1	19	0.08	0.19	0.02	0.00	0.05
1	20	0.18	0.04	0.00	0.00	0.47
1	21	3.23	0.00	0.52	0.05	0.06
1	22	4.56	0.33	5.82	0.02	0.00
1	23	0.05	0.57	0.33	0.00	0.00
1	24	0.85	0.51	0.26	5.82	0.00
1	25	2.53	0.35	0.13	0.00	0.26
1	26	0.83	0.00	0.47	0.55	0.00
1	27	0.55	6.03	0.00	0.00	0.00
1	28	0.08	0.00	0.05	0.09	0.00
1	29	0.11	0.00	0.00	0.11	6.24
1	30	0.39	0.00	0.00	0.27	0.00
2	2	0.00	0.00	0.00	0.00	0.00
2	3	11.03	8.30	0.00	0.00	0.00
2	4	0.00	0.00	2.42	0.34	0.00
2	5	0.34	0.20	1.71	3.00	1.28
2	6	0.12	0.25	0.09	0.66	7.25
2	7	0.18	0.22	1.49	0.42	0.00
2	8	0.82	0.09	0.00	0.56	5.55
2	9	0.63	1.46	0.89	0.67	4.88
2	10	0.54	0.33	1.11	0.29	0.48
2	11	0.22	0.87	0.56	0.30	0.31

2	12	0.19	0.52	0.37	0.19	0.40
2	13	1.48	0.00	1.32	1.12	2.32
2	14	0.87	0.62	0.00	0.00	0.50
2	15	0.68	0.05	0.03	0.02	0.00
2	16	0.18	1.18	0.13	0.14	0.00
2	17	0.00	0.21	0.00	0.35	0.47
2	18	0.43	0.00	1.42	0.97	0.31
2	19	0.29	0.07	0.02	0.93	0.11
2	20	0.16	0.12	0.40	0.00	0.00
2	21	0.16	0.61	0.35	0.07	0.00
2	22	0.11	0.08	0.00	0.04	0.00
2	23	0.03	0.14	0.24	0.00	0.00
2	24	0.55	0.12	0.19	0.00	1.08
2	25	0.00	0.42	0.06	0.00	0.00
2	26	0.00	0.00	0.25	0.27	0.00
2	27	1.29	0.00	0.00	0.00	0.00
2	28	5.36	0.00	0.04	0.15	0.00
2	29	1.08	0.00	0.00	0.19	0.59
2	30	0.53	0.00	0.00	0.47	0.00
3	3	0.00	0.00	0.00	0.00	0.00
3	4	0.00	0.06	0.32	0.02	0.99
3	5	0.00	1.12	0.64	2.66	0.22
3	6	0.40	0.49	2.67	11.51	1.12
3	7	0.31	0.23	4.12	1.11	1.63
3	8	0.85	0.00	0.69	3.48	0.45
3	9	0.18	1.82	16.72	5.81	0.67
3	10	0.20	0.66	16.12	18.09	0.00
3	11	0.10	0.00	0.00	20.15	2.44
3	12	0.30	0.62	1.32	0.00	0.00
3	13	1.47	0.83	15.12	10.51	0.32
3	14	0.44	0.37	0.79	2.36	0.00
3	15	0.39	0.19	0.21	0.17	0.33
3	16	0.89	0.22	0.11	0.03	0.94
3	17	0.00	0.24	6.22	14.39	0.52
3	18	0.63	0.32	0.00	0.32	0.61
3	19	0.84	0.08	0.31	0.16	0.18
3	20	0.28	0.83	0.00	0.14	0.57
3	21	0.02	0.00	0.00	0.13	0.07
3	32	0.00	0.36	4.22	0.33	0.00
3	23	0.19	0.57	0.39	4.90	0.00
3	24	1.28	0.86	5.71	1.05	0.00
3	25	0.00	0.58	1.26	0.05	0.65
3	26	0.00	0.59	0.00	0.00	1.38
3	27	1.35	0.00	0.92	0.09	0.00

3	28	0.19	0.00	0.15	0.14	0.00
3	29	0.26	0.00	0.10	0.25	0.00
3	30	0.89	0.00	0.00	0.07	0.00
4	4	0.00	0.00	0.00	0.00	0.00
4	5	5.25	0.00	12.68	0.43	0.00
4	6	7.55	10.71	18.27	11.24	0.00
4	7	4.63	22.95	0.21	0.00	10.99
4	8	0.00	4.12	0.00	0.00	0.00
4	9	0.87	0.00	1.44	0.02	0.24
4	10	0.84	4.88	0.64	0.12	0.00
4	11	0.23	0.10	4.45	0.15	1.46
4	12	1.71	7.16	0.00	3.29	1.29
4	13	0.00	0.29	0.85	0.07	0.00
4	14	0.58	1.39	0.00	0.00	0.00
4	15	0.74	0.54	0.41	0.64	0.04
4	16	0.18	2.20	1.40	3.31	1.64
4	17	0.43	0.00	1.41	0.08	1.08
4	18	0.00	4.97	0.00	0.03	1.33
4	19	0.20	0.00	0.26	0.02	0.00
4	20	0.00	21.16	0.00	1.21	2.12
4	21	1.45	0.00	1.70	1.00	0.31
4	42	2.05	0.04	0.00	0.57	0.35
4	23	0.10	0.03	1.44	0.03	0.23
4	24	0.00	0.02	0.78	0.00	0.00
4	25	4.70	0.02	0.47	0.73	1.08
4	26	1.90	0.17	1.06	0.07	0.00
4	27	0.00	0.41	0.93	1.17	0.00
4	28	0.00	0.48	0.38	0.32	0.00
4	29	0.09	0.06	0.98	0.18	0.00
4	30	0.00	0.09	0.36	0.78	1.27
5	5	0.00	0.00	0.00	0.00	0.00
5	6	8.29	0.00	18.92	4.29	1.96
5	7	6.82	0.00	0.10	8.38	0.00
5	8	0.00	0.00	0.01	18.84	4.80
5	9	35.13	1.63	0.96	0.90	6.54
5	10	1.28	0.00	0.30	1.10	12.52
5	11	0.59	0.00	5.64	1.49	0.09
5	12	0.45	0.00	0.53	1.71	0.13
5	13	0.00	0.00	0.42	0.74	3.16
5	14	1.37	0.00	0.23	0.48	0.24
5	15	0.79	0.00	0.97	0.22	0.22
5	16	0.43	0.00	2.96	1.31	0.00
5	17	0.80	3.51	1.65	1.26	0.27
5	18	1.31	0.00	0.00	0.41	0.18

5	19	0.89	1.13	0.82	0.21	4.99
5	20	0.00	0.00	0.00	0.24	0.00
5	21	0.32	0.00	0.97	0.65	0.03
5	22	0.23	0.55	0.36	0.26	0.19
5	23	0.07	0.67	0.49	0.17	0.08
5	24	0.00	0.71	0.21	0.34	0.22
5	25	0.43	0.82	0.45	0.05	0.00
5	26	1.33	0.00	0.79	1.02	0.26
5	27	0.27	0.00	0.26	0.06	0.00
5	28	2.42	0.29	1.18	1.31	0.26
5	29	0.62	0.99	0.52	1.09	0.94
5	30	0.00	0.21	0.71	3.78	0.00
6	6	0.00	0.00	0.00	0.00	0.00
6	7	15.56	4.87	0.13	3.42	0.00
6	8	0.00	9.86	0.03	24.42	7.60
6	9	19.04	0.00	0.48	22.43	15.40
6	10	10.07	11.22	0.58	9.39	5.97
6	11	29.55	0.49	0.83	3.55	9.43
6	12	9.36	11.56	2.89	0.37	5.78
6	13	0.00	11.35	0.47	24.19	8.66
6	14	21.00	20.05	1.28	22.10	6.06
6	15	11.17	12.98	1.74	0.23	0.65
6	16	7.68	17.33	1.05	0.36	0.00
6	17	0.62	0.02	1.16	3.54	0.60
6	18	9.88	23.11	0.00	0.47	0.40
6	19	2.78	0.05	3.54	0.45	4.91
6	20	0.00	6.80	0.00	0.23	0.00
6	21	0.30	0.81	0.15	0.14	0.06
6	22	1.70	0.32	1.96	0.58	0.37
6	23	1.42	0.23	0.32	0.40	0.13
6	24	0.00	0.10	0.04	1.48	1.42
6	25	1.23	0.12	0.41	0.09	0.00
6	26	0.76	1.10	0.05	0.23	0.54
6	27	0.05	1.86	0.72	0.09	0.06
6	28	0.32	0.00	0.27	0.36	0.58
6	29	0.10	0.32	0.74	0.26	0.15
6	30	0.00	0.98	0.13	0.82	0.00
7	7	0.00	0.00	0.00	0.00	0.00
7	8	33.48	11.46	9.93	4.88	0.00
7	9	13.86	0.00	4.61	5.89	0.41
7	10	34.43	9.20	10.41	6.38	0.00
7	11	25.67	2.73	0.00	39.16	2.62
7	12	23.52	2.97	5.07	0.00	0.00
7	13	0.71	0.89	9.85	9.16	0.00

7	14	5.18	5.69	0.26	1.25	1.28
7	15	10.78	10.73	0.13	0.05	0.00
7	16	27.41	4.40	0.07	0.01	3.08
7	17	0.55	0.02	3.86	0.47	1.70
7	18	7.67	16.63	0.75	1.31	1.99
7	19	29.00	0.03	0.20	3.09	0.00
7	20	0.37	10.17	1.38	0.07	2.87
7	21	0.83	1.67	0.00	0.03	0.20
7	22	2.21	0.19	0.29	0.29	0.05
7	23	0.47	0.14	0.37	1.12	0.00
7	24	0.03	0.11	3.38	0.34	0.00
7	25	1.16	0.07	1.60	0.14	1.16
7	26	1.26	0.84	0.00	0.00	0.00
7	27	0.09	0.47	0.83	0.10	0.00
7	28	0.30	0.08	0.09	0.15	0.34
7	29	0.16	0.29	0.06	0.35	0.00
7	30	0.05	1.15	0.23	0.22	0.10
8	8	0.00	0.00	0.00	0.00	0.00
8	9	0.00	0.00	0.37	12.27	4.87
8	10	0.00	19.08	2.25	15.06	14.60
8	11	0.00	7.17	0.00	11.16	17.66
8	12	0.00	12.56	10.16	0.00	10.12
8	13	5.87	7.82	2.14	11.29	9.51
8	14	0.00	8.69	2.31	3.95	10.29
8	15	0.00	2.45	0.00	0.06	0.52
8	16	0.00	6.19	0.00	0.01	0.00
8	17	0.21	0.10	0.00	0.98	1.00
8	18	0.00	0.22	0.00	1.19	0.67
8	19	0.00	0.30	0.00	0.68	0.60
8	20	3.11	3.99	2.82	0.13	0.00
8	21	0.00	0.20	0.00	0.02	0.10
8	22	0.00	0.24	0.94	0.55	0.60
8	23	0.00	0.34	0.21	1.84	0.23
8	24	0.28	0.43	0.49	0.65	1.28
8	25	0.00	0.20	0.55	0.11	0.00
8	26	0.17	0.14	0.00	0.00	0.77
8	27	0.63	1.26	0.17	0.11	0.12
8	28	0.20	0.00	0.03	0.18	0.91
8	29	1.02	0.03	0.00	0.21	0.20
8	30	0.44	0.18	0.19	0.14	0.00
9	9	0.00	0.00	0.00	0.00	0.00
9	10	19.52	0.00	5.61	13.25	31.99
9	11	9.14	0.00	0.23	22.14	10.83
9	12	15.07	0.00	16.60	0.00	1.73



9	13	0.00	0.00	22.64	5.24	28.85
9	14	9.50	0.00	0.13	10.86	1.08
9	15	5.00	0.00	8.44	0.35	1.59
9	16	18.62	0.00	0.50	0.07	0.28
9	17	0.12	3.22	3.49	17.31	0.46
9	18	8.23	0.00	0.79	1.96	0.09
9	19	9.85	1.01	1.72	0.95	0.41
9	20	0.00	0.00	1.31	0.29	0.59
9	21	4.48	0.00	0.00	0.12	0.06
9	22	0.23	0.62	0.68	0.81	0.44
9	23	17.16	1.05	13.15	4.66	0.20
9	24	0.00	2.18	15.49	0.57	1.40
9	25	0.48	2.59	13.70	0.14	0.10
9	26	0.11	0.16	0.00	0.00	0.22
9	27	0.25	0.00	15.98	0.13	0.43
9	28	2.67	0.00	3.44	0.17	0.49
9	29	0.54	0.17	0.63	0.21	0.36
9	30	0.00	0.82	0.73	0.08	0.07
10	10	0.00	0.00	0.00	0.00	0.00
10	11	5.45	0.58	0.00	11.40	0.00
10	12	10.26	7.43	35.49	0.00	0.00
10	13	0.00	16.49	15.70	3.71	11.57
10	14	6.54	2.60	1.44	10.19	0.00
10	15	10.82	10.73	0.22	0.95	9.37
10	16	5.90	0.97	0.22	0.28	0.22
10	17	0.06	0.00	11.96	4.53	0.93
10	18	11.40	32.40	0.57	16.76	0.86
10	19	21.12	0.00	0.62	8.56	2.51
10	20	0.00	8.54	0.84	1.80	0.05
10	21	0.84	1.17	0.00	0.29	0.44
10	22	0.77	0.57	1.40	1.69	0.04
10	23	0.65	0.50	0.62	6.07	0.00
10	24	0.00	0.17	13.33	0.57	0.00
10	25	1.06	0.18	2.80	0.25	0.00
10	26	0.25	1.44	0.00	0.00	0.00
10	27	0.19	0.38	1.55	0.51	1.46
10	28	2.09	0.07	0.19	0.97	0.05
10	29	0.59	0.24	0.20	1.93	2.42
10	30	0.00	0.82	0.73	0.48	0.15
11	11	0.00	0.00	0.00	0.00	0.00
11	12	12.09	14.32	0.00	0.00	5.22
11	13	0.00	8.13	0.00	35.28	0.00
11	14	5.77	14.86	0.00	1.35	5.26
11	15	8.50	1.26	0.83	1.22	0.00

11	16	5.91	11.73	3.23	0.34	1.49
11	17	0.02	0.06	0.00	11.47	0.82
11	18	9.10	0.28	0.00	17.92	0.97
11	19	22.22	0.16	0.44	8.80	0.00
11	20	0.00	6.96	0.00	2.14	2.77
11	21	8.72	1.54	2.92	0.33	0.34
11	22	1.24	0.47	0.00	1.53	0.00
11	23	3.53	0.73	1.69	2.99	0.00
11	24	0.00	0.92	1.30	0.27	0.42
11	25	0.69	0.42	0.79	0.12	0.49
11	26	0.21	0.30	0.46	0.00	0.00
11	27	0.07	0.00	1.44	0.78	0.29
11	28	0.81	0.00	0.62	1.23	0.00
11	29	0.17	0.04	0.28	2.28	0.00
11	30	0.00	0.26	0.00	0.61	0.00
12	12	0.00	0.00	0.00	0.00	0.00
12	13	0.00	17.22	34.70	0.00	0.00
12	14	28.48	20.27	0.26	0.00	11.80
12	15	6.37	7.38	1.32	0.60	0.03
12	16	6.82	12.64	0.68	3.92	0.18
12	17	0.01	0.00	0.58	0.00	0.11
12	18	10.79	11.66	1.14	0.00	0.19
12	19	14.54	0.00	2.31	0.00	0.00
12	20	0.00	11.49	0.31	0.00	0.00
12	21	1.89	1.34	0.00	2.84	0.50
12	22	2.43	0.55	0.30	1.77	3.87
12	23	12.94	0.48	0.37	0.16	1.85
12	24	0.00	0.16	0.67	0.00	2.45
12	25	1.42	0.17	0.65	0.12	0.00
12	26	0.45	1.34	0.00	0.25	0.00
12	27	0.08	1.66	0.52	0.55	0.04
12	28	0.84	0.09	0.08	1.41	0.00
12	29	0.19	0.52	0.34	0.81	0.00
12	30	0.00	1.19	0.23	1.98	0.52
13	13	0.00	0.00	0.00	0.00	0.00
13	14	0.00	12.96	0.92	36.39	0.00
13	15	0.00	3.59	0.47	0.65	1.27
13	16	0.00	8.92	0.21	0.18	0.00
13	17	0.00	0.32	15.96	7.53	0.77
13	18	0.00	7.14	1.22	4.53	0.52
13	19	0.00	0.79	0.62	3.42	2.65
13	20	3.58	11.52	1.77	1.12	0.00
13	21	0.00	0.69	0.00	0.19	0.10
13	22	0.00	0.63	0.89	4.19	0.62

13	23	0.00	1.04	1.74	18.35	0.17
13	24	1.95	1.23	13.26	0.10	0.00
13	25	0.10	0.61	4.16	0.20	0.00
13	26	0.34	1.09	0.00	0.00	1.38
13	27	2.68	0.18	1.71	0.23	0.00
13	28	0.34	0.00	0.34	0.61	1.09
13	29	0.48	0.13	0.31	1.31	0.17
13	30	1.14	0.21	0.73	0.34	0.00
14	14	0.00	0.00	0.00	0.00	0.00
14	15	12.27	12.03	0.15	0.05	0.00
14	16	17.65	14.55	0.00	0.00	0.00
14	17	0.12	0.04	0.00	0.73	0.00
14	18	7.36	16.53	2.91	2.11	0.00
14	19	4.74	0.10	0.14	1.71	0.00
14	20	0.00	9.58	0.00	0.05	0.03
14	21	1.47	2.19	0.00	0.00	0.27
14	22	1.50	0.35	0.49	0.04	0.05
14	23	6.57	0.19	0.52	0.00	0.00
14	24	0.00	0.15	0.49	0.46	3.94
14	25	0.38	0.19	0.80	0.37	0.00
14	26	0.14	1.80	0.00	0.00	0.00
14	27	0.11	0.22	1.58	0.49	0.04
14	28	0.53	0.16	1.63	0.82	3.80
14	29	0.67	0.14	3.34	3.92	0.00
14	30	0.00	0.39	0.14	1.14	0.09
15	15	0.00	0.00	0.00	0.00	0.00
15	16	3.54	15.00	1.79	12.26	14.33
15	17	0.02	0.05	5.76	3.05	8.44
15	18	17.14	6.64	0.08	0.49	15.13
15	19	4.57	0.10	5.85	0.25	0.20
15	20	0.00	11.27	0.00	7.30	0.31
15	21	1.24	3.94	1.30	1.35	2.38
15	22	1.66	0.05	0.00	3.86	0.56
15	23	7.82	0.79	0.54	0.75	0.08
15	24	0.00	0.87	1.48	0.00	0.00
15	25	0.49	0.99	4.92	12.27	0.00
15	26	0.27	7.97	0.34	0.11	1.12
15	27	0.10	1.05	8.14	5.51	2.33
15	28	0.73	0.04	1.39	13.65	0.00
15	29	0.15	0.23	2.75	9.11	0.00
15	30	0.00	0.26	6.02	13.60	0.96
16	16	0.00	0.00	0.00	0.00	0.00
16	17	0.10	0.00	20.17	0.22	20.68
16	18	29.84	13.21	0.00	0.00	6.26

16	19	17.47	0.00	4.97	0.00	0.00
16	20	0.00	17.37	0.00	26.02	7.85
16	21	0.88	3.04	5.05	0.88	6.97
16	22	0.72	0.21	0.00	0.53	1.71
16	23	0.16	0.18	3.78	0.07	0.41
16	24	0.00	0.06	2.77	0.00	0.98
16	25	0.11	0.06	0.92	0.92	2.68
16	26	0.19	0.48	1.31	0.71	0.00
16	27	0.17	0.00	3.11	0.65	0.92
16	28	0.52	0.09	5.65	1.12	0.00
16	29	0.55	0.07	0.94	1.59	0.00
16	30	0.00	0.32	0.06	4.10	2.79
17	17	0.00	0.00	0.00	0.00	0.00
17	18	0.00	0.00	0.00	19.26	17.91
17	19	0.02	6.68	29.35	7.02	0.11
17	20	0.00	0.00	0.00	1.53	5.23
17	21	0.00	0.00	0.00	0.53	2.94
17	22	0.00	10.60	0.00	39.96	1.01
17	23	0.00	2.86	1.83	11.37	0.24
17	24	0.00	1.19	7.48	0.00	0.54
17	25	1.71	6.01	2.82	0.89	1.48
17	26	5.92	0.10	0.00	0.00	7.95
17	27	8.25	0.00	5.91	0.69	0.54
17	28	5.55	0.46	4.08	1.00	0.05
17	29	8.13	11.54	0.80	1.87	0.37
17	30	0.00	0.06	6.65	0.66	1.52
18	18	0.00	0.00	0.00	0.00	0.00
18	19	20.75	0.00	0.07	20.85	0.07
18	20	0.00	24.66	2.59	0.62	6.42
18	21	2.65	0.00	0.00	0.07	5.54
18	22	1.93	0.17	0.00	10.02	1.80
18	23	1.45	0.10	0.36	7.79	0.43
18	24	0.00	0.09	0.34	0.00	0.64
18	25	0.00	0.09	0.62	0.17	1.74
18	26	0.08	0.75	0.00	0.00	5.34
18	27	0.14	0.16	1.16	0.29	0.97
18	28	1.23	1.04	1.03	0.50	0.03
18	29	0.30	0.05	2.29	1.97	0.25
18	30	0.00	0.33	0.07	0.65	3.04
19	19	0.00	0.00	0.00	0.00	0.00
19	20	0.00	0.00	0.00	0.32	0.00
19	21	0.87	0.04	0.66	0.03	0.05
19	22	0.55	18.16	0.00	5.08	0.32
19	23	3.40	15.27	5.60	3.95	0.99

19	24	0.00	21.30	0.73	0.00	0.00
19	25	0.12	8.45	15.49	0.08	0.00
19	26	0.07	7.06	0.17	0.00	0.65
19	27	0.09	0.00	6.68	0.11	0.00
19	28	0.97	1.35	8.31	0.23	0.52
19	29	0.14	3.16	18.29	1.28	0.25
19	30	0.00	6.93	5.05	0.29	0.00
20	20	0.00	0.00	0.00	0.00	0.00
20	21	0.00	0.27	0.00	0.39	3.54
20	22	0.00	0.33	0.00	7.15	3.86
20	23	0.00	0.19	0.00	0.60	0.00
20	24	5.28	0.38	0.00	0.00	0.00
20	25	0.00	0.40	0.00	15.43	3.80
20	26	0.00	3.20	0.00	0.00	0.00
20	27	6.86	0.08	0.00	5.61	0.31
20	28	0.98	0.12	0.00	16.30	0.17
20	29	1.32	0.52	0.00	9.88	0.00
20	30	1.29	0.89	0.00	13.92	7.65
21	21	0.00	0.00	0.00	0.00	0.00
21	22	30.91	0.00	0.00	18.59	2.60
21	23	13.04	0.95	6.68	39.66	1.52
21	24	0.00	1.06	25.98	0.00	0.00
21	25	1.91	1.01	7.05	14.53	0.47
21	26	0.62	8.05	17.57	2.94	0.31
21	27	0.31	0.00	0.00	4.35	2.45
21	28	2.83	0.00	5.84	5.34	1.70
21	29	0.50	0.14	0.00	1.38	0.00
21	30	0.00	0.03	0.00	20.18	3.60
22	22	0.00	0.00	0.00	0.00	0.00
22	23	27.08	15.14	0.00	7.46	9.04
22	24	0.00	7.93	0.00	0.00	0.00
22	25	2.69	5.83	0.00	7.08	0.00
22	26	0.88	17.09	0.00	1.53	0.00
22	27	0.48	0.00	0.00	19.43	0.43
22	28	4.31	4.18	0.00	10.25	10.89
22	29	0.77	6.49	0.00	6.58	0.00
22	30	0.00	2.03	0.00	22.27	11.36
23	23	0.00	0.00	0.00	0.00	0.00
23	24	0.00	13.24	19.57	0.00	0.00
23	25	0.22	16.92	10.47	6.44	0.00
23	26	0.07	1.91	1.11	0.00	0.00
23	27	0.01	0.00	5.61	0.90	0.00
23	28	0.06	2.56	8.25	7.42	0.00
23	29	0.01	3.60	12.26	2.92	0.00

23	30	0.00	2.86	1.83	13.07	1.02
24	24	0.00	0.00	0.00	0.00	0.00
24	25	0.00	9.63	14.78	0.00	0.00
24	26	0.00	2.39	9.57	0.00	0.00
24	27	4.62	0.00	21.42	0.00	0.00
24	28	0.71	0.00	8.98	0.00	0.00
24	29	0.95	0.41	25.28	0.00	0.00
24	30	8.17	2.76	4.10	0.00	0.00
25	25	0.00	0.00	0.00	0.00	0.00
25	26	18.47	17.43	43.07	0.00	0.00
25	27	0.00	0.00	5.74	7.81	0.00
25	28	0.00	0.00	10.13	11.31	0.00
25	29	17.83	0.27	7.39	6.16	0.00
25	30	0.00	1.26	9.86	32.72	0.00
26	26	0.00	0.00	0.00	0.00	0.00
26	27	10.45	0.00	0.00	0.00	0.00
26	28	7.03	0.00	16.67	0.92	0.20
26	29	10.70	0.67	0.00	1.12	0.00
26	30	0.00	1.03	0.00	2.77	0.91
27	27	0.00	0.00	0.00	0.00	0.00
27	28	2.95	0.00	2.84	19.07	0.00
27	29	14.44	0.00	10.90	12.00	0.00
27	30	5.16	0.00	12.02	5.68	0.85
28	28	0.00	0.00	0.00	0.00	0.00
28	29	11.78	4.47	8.27	15.56	0.00
28	30	12.70	0.97	4.99	14.26	1.55
29	29	0.00	0.00	0.00	0.00	0.00
29	30	17.06	3.22	0.71	10.48	0.00
30	30	0.00	0.00	0.00	0.00	0.00

174 **Table S3.** The original SOC elements between  $T_i$  and  $S_j$  calculated at M06L/def2-svp level.

States		SOC elements between $T_i$ and $S_j$ /cm <sup>-1</sup>				
$T_i$	$S_j$	Conformer 1	Conformer 2	Conformer 3	Conformer 4	Conformer 5
1	0	18.78	14.55	10.23	7.31	13.62
1	1	1.50	0.27	0.31	0.63	0.39
1	2	1.55	0.58	0.71	1.13	7.28
1	3	4.58	0.64	78.38	395.62	1.42
1	4	642.53	675.49	676.77	563.40	2.23
1	5	5.84	70.40	4.31	11.58	9.45
1	6	22.65	5.99	2.10	24.39	671.42
1	7	689.33	5.34	2.23	11.29	0.84
1	8	373.85	86.56	34.68	113.26	1.41
1	9	4.34	22.25	3.12	20.33	9.00

1	10	44.79	84.26	753.12	742.48	41.65
1	11	11.23	138.31	0.98	1.84	3.83
1	12	6.66	727.82	0.51	1.85	6.90
1	13	8.20	76.99	1.28	1.98	62.99
1	14	8.11	9.01	4.26	3.59	743.31
1	15	9.46	17.64	1.52	2.48	2.06
1	16	18.29	1.96	3.93	2.08	8.16
1	17	122.84	2.87	14.48	3.26	0.70
1	18	2.23	0.35	130.61	14.29	9.11
1	19	8.76	2.97	1.76	8.57	0.78
1	20	1.59	1.10	2.12	114.45	0.56
1	21	352.99	2.08	1.11	2.45	2.45
1	22	250.03	43.72	11.36	0.49	5.42
1	23	10.82	138.09	2.20	0.30	9.17
1	24	5.56	3.18	1.90	2.53	36.29
1	25	1.66	1.61	2.13	2.64	7.69
1	26	11.76	5.07	3.04	1.16	5.19
1	27	37.07	0.77	2.64	2.13	90.87
1	28	24.39	3.54	135.52	17.29	76.90
1	29	127.50	1.10	156.05	9.87	8.36
1	30	19.62	39.43	83.79	246.02	0.52
2	0	40.28	33.67	7.30	7.74	1.43
2	1	2.57	0.88	1.00	0.43	6.30
2	2	2.92	1.15	0.09	0.35	0.09
2	3	4.59	0.40	0.62	2.19	0.31
2	4	3.96	1.68	1.33	0.55	0.98
2	5	643.29	3.66	1.63	2.92	0.28
2	6	11.22	10.25	17.52	405.48	36.23
2	7	5.08	676.26	84.59	129.47	2.64
2	8	12.89	35.17	37.02	151.64	2.72
2	9	7.36	46.24	674.98	525.23	0.67
2	10	98.48	32.63	2.17	7.21	3.41
2	11	761.61	10.38	11.80	4.71	2.04
2	12	40.18	7.10	19.69	19.46	3.32
2	13	129.06	16.92	751.82	741.97	3.36
2	14	37.75	35.21	28.85	52.16	41.50
2	15	61.22	25.58	1.38	74.51	0.04
2	16	2.11	16.40	43.25	2.70	0.62
2	17	4.33	751.35	4.67	4.72	0.09
2	18	6.21	1.24	1.53	6.02	0.58
2	19	7.92	2.19	2.51	2.77	1.97
2	20	1.15	9.17	1.04	1.49	0.10
2	21	4.24	4.09	1.00	1.37	3.42
2	22	3.56	16.95	5.56	0.04	0.20

2	23	109.76	4.47	3.29	0.27	1.50
2	24	18.17	4.66	0.80	2.25	2.00
2	25	14.37	9.19	1.97	7.66	0.70
2	26	114.13	8.71	6.00	13.83	0.35
2	27	84.83	2.36	10.36	2.52	4.60
2	28	77.32	28.86	59.41	3.56	4.00
2	29	34.03	15.62	98.53	1.15	1.22
2	30	15.88	15.71	41.76	2.30	0.19
3	0	15.84	38.57	10.16	2.60	34.59
3	1	1.77	1.52	57.45	320.73	0.88
3	2	0.86	1.13	0.97	3.38	0.24
3	3	3.00	0.48	2.54	8.39	0.67
3	4	4.37	0.73	63.69	85.51	8.17
3	5	12.33	1.72	3.45	14.52	10.13
3	6	641.80	5.92	1.78	10.61	0.42
3	7	31.71	8.10	0.36	4.32	3.83
3	8	25.37	5.07	4.02	49.38	3.96
3	9	7.02	76.65	4.57	9.69	14.14
3	10	34.36	32.75	99.20	330.77	5.88
3	11	0.41	657.19	8.26	5.32	4.45
3	12	707.04	110.35	4.74	8.90	669.16
3	13	82.33	69.12	6.39	4.19	33.93
3	14	24.73	9.17	38.90	22.30	2.86
3	15	321.52	57.99	1.13	0.76	1.44
3	16	12.57	5.60	0.39	1.15	26.34
3	17	4.11	11.05	3.03	2.94	5.95
3	18	2.22	7.42	13.04	4.88	11.96
3	19	3.68	187.09	5.54	5.08	14.43
3	20	1.05	725.10	1.81	34.50	6.31
3	21	2.58	20.62	0.67	0.95	729.75
3	32	2.48	2.86	2.35	0.36	36.94
3	23	32.95	8.80	4.93	0.71	124.09
3	24	5.04	4.05	4.29	4.41	51.38
3	25	2.30	9.03	4.60	20.62	1.98
3	26	80.57	36.65	39.12	9.36	8.87
3	27	65.63	2.94	1.01	1.90	17.17
3	28	38.25	25.57	4.98	9.77	2.13
3	29	17.38	17.22	9.26	4.20	22.58
3	30	95.12	48.34	7.42	63.44	4.61
4	0	23.06	8.44	304.22	350.42	390.23
4	1	639.73	35.10	3.76	23.49	7.40
4	2	5.62	0.56	1.19	4.93	0.09
4	3	10.47	3.81	1.40	3.47	9.78
4	4	1.74	18.31	2.60	0.93	4.46



4	5	4.81	1.94	12.15	18.82	77.23
4	6	21.06	3.62	70.42	36.60	1.43
4	7	616.05	1.15	14.55	126.52	34.28
4	8	311.60	3.62	1.39	4.46	46.51
4	9	1.82	1.77	7.93	3.19	131.34
4	10	17.44	4.39	5.42	21.21	17.28
4	11	6.89	19.18	153.21	111.19	12.32
4	12	4.42	38.14	66.78	277.43	31.48
4	13	1.26	3.93	28.49	3.44	4.40
4	14	7.93	0.85	529.50	73.56	15.19
4	15	5.60	1.01	74.88	3.72	104.88
4	16	4.26	5.34	5.51	214.36	543.56
4	17	58.59	1.35	84.14	20.25	33.98
4	18	1.09	1.75	8.03	394.45	310.96
4	19	4.83	10.48	346.55	392.97	27.84
4	20	1.44	16.70	48.01	18.43	26.90
4	21	52.69	16.89	15.46	60.64	3.52
4	42	37.05	1.80	7.42	10.50	220.59
4	23	4.03	8.27	67.34	135.54	125.32
4	24	2.78	2.32	34.14	52.89	16.48
4	25	0.71	2.53	71.04	25.27	59.51
4	26	7.47	1.32	581.93	13.26	45.69
4	27	26.17	1.13	22.77	142.12	16.61
4	28	18.85	1.21	38.83	58.24	21.85
4	29	145.67	1.07	7.59	18.14	31.79
4	30	16.31	1.78	95.40	18.54	9.18
5	0	35.28	319.09	245.45	82.07	0.74
5	1	7.38	2.81	126.73	144.40	3.04
5	2	625.18	6.47	2.90	6.25	1.09
5	3	11.72	3.98	11.08	17.93	1.72
5	4	1.24	1.74	2.17	10.27	0.18
5	5	15.68	0.68	7.59	15.24	0.11
5	6	1.37	135.83	78.92	7.38	8.11
5	7	3.36	8.40	16.94	11.52	0.57
5	8	11.97	0.53	4.50	20.00	0.90
5	9	10.05	69.89	4.50	3.59	0.33
5	10	83.37	0.44	135.50	156.27	0.48
5	11	660.98	9.28	54.93	97.68	0.33
5	12	51.30	5.21	32.40	298.47	10.75
5	13	71.19	5.66	12.84	48.81	1.60
5	14	27.28	0.84	363.52	560.52	7.34
5	15	42.72	2.63	192.03	16.37	0.16
5	16	1.46	182.55	1.94	24.90	0.46
5	17	3.29	7.99	67.98	100.13	1.93

5	18	4.18	66.56	18.94	58.04	0.12
5	19	9.34	273.74	509.18	52.95	0.42
5	20	2.81	55.87	73.28	13.25	3.53
5	21	2.48	574.02	11.88	10.99	12.88
5	22	3.89	59.95	11.33	2.51	0.81
5	23	41.68	5.63	29.81	12.76	1.99
5	24	7.34	72.50	149.28	35.70	1.42
5	25	6.65	74.85	39.40	586.42	0.60
5	26	46.47	11.96	302.98	271.60	0.28
5	27	21.59	28.24	4.77	48.57	1.26
5	28	21.30	30.79	20.79	196.06	0.98
5	29	4.91	22.32	26.53	63.64	0.58
5	30	2.77	95.44	202.30	55.64	1.68
6	0	16.65	5.61	51.04	11.62	2.61
6	1	130.22	39.06	667.54	571.14	13.08
6	2	16.61	6.00	2.20	11.53	1.62
6	3	479.54	4.75	59.78	82.60	3.32
6	4	101.92	25.36	1.83	17.44	0.78
6	5	9.75	3.85	7.83	8.37	0.08
6	6	12.54	1.17	12.26	15.26	6.97
6	7	67.72	7.39	2.38	8.00	3.26
6	8	51.79	4.34	24.18	76.04	4.59
6	9	3.06	4.30	2.60	18.60	0.57
6	10	7.20	5.53	724.56	610.39	1.59
6	11	11.60	24.04	8.27	20.85	2.17
6	12	473.71	49.20	5.50	77.35	43.23
6	13	56.35	6.41	2.65	16.79	2.10
6	14	16.65	1.16	70.20	123.72	11.84
6	15	199.52	2.76	35.71	3.75	0.96
6	16	0.66	1.28	2.97	2.48	1.53
6	17	22.75	13.90	23.06	22.48	0.78
6	18	1.02	3.11	79.69	16.09	2.13
6	19	16.87	6.44	92.58	8.26	1.57
6	20	3.54	26.41	14.14	59.52	1.19
6	21	42.89	3.31	1.82	3.64	48.57
6	22	22.86	3.15	9.61	0.42	6.78
6	23	10.55	11.99	5.49	1.56	8.74
6	24	3.14	0.75	29.07	7.60	4.24
6	25	2.03	1.60	6.66	135.31	8.51
6	26	15.04	1.86	58.60	60.83	1.00
6	27	12.00	1.27	2.01	16.45	1.10
6	28	10.23	1.90	91.43	51.01	1.51
6	29	38.44	0.52	101.93	17.43	2.30
6	30	38.50	2.59	75.75	124.30	0.67

7	0	36.90	1.78	10.10	4.66	267.61
7	1	253.72	79.93	5.06	44.37	2.84
7	2	10.86	22.90	28.19	56.44	0.26
7	3	408.31	4.41	0.65	5.91	5.25
7	4	223.26	8.47	1.47	1.68	1.28
7	5	14.05	1.15	2.98	2.83	10.59
7	6	7.99	0.52	1.34	6.96	3.12
7	7	65.77	9.72	3.17	1.98	21.82
7	8	77.07	9.02	1.75	4.09	23.58
7	9	12.62	1.93	47.54	21.46	87.80
7	10	24.49	8.40	5.79	47.72	10.02
7	11	18.53	20.55	6.10	3.44	11.34
7	12	398.38	83.97	3.27	11.89	16.07
7	13	51.55	9.50	56.28	54.83	4.61
7	14	29.46	1.96	41.98	21.95	4.20
7	15	174.48	1.48	7.18	5.96	73.58
7	16	5.75	0.69	2.83	1.10	110.92
7	17	49.75	33.07	1.84	3.95	20.24
7	18	2.25	3.30	0.58	3.22	187.73
7	19	64.22	4.25	12.44	2.14	31.75
7	20	11.37	15.95	3.05	4.57	26.96
7	21	77.62	0.66	0.27	0.34	75.92
7	22	80.51	3.68	0.69	0.26	525.33
7	23	7.27	12.76	2.45	0.81	237.39
7	24	15.39	1.06	8.86	1.20	28.38
7	25	6.16	0.89	4.55	21.29	125.25
7	26	19.00	1.53	42.24	10.91	115.65
7	27	32.15	1.05	1.47	1.65	33.27
7	28	21.47	2.16	4.65	7.07	31.52
7	29	63.53	1.21	5.80	2.54	6.80
7	30	26.25	3.38	16.03	11.23	14.03
8	0	363.04	14.25	0.78	3.97	6.24
8	1	10.61	672.01	17.64	142.51	45.25
8	2	9.11	6.88	2.50	80.10	1.68
8	3	51.16	2.38	3.36	28.30	7.48
8	4	10.53	2.66	21.91	7.16	0.74
8	5	6.51	19.43	0.88	7.19	0.74
8	6	2.79	0.97	0.12	4.89	13.58
8	7	9.61	5.64	0.17	1.01	2.39
8	8	7.62	70.34	1.56	21.44	4.19
8	9	123.65	24.17	4.62	25.03	2.42
8	10	2.51	78.72	9.30	150.17	2.80
8	11	6.14	133.03	0.17	6.45	3.07
8	12	57.17	693.18	0.32	23.97	5.22

8	13	30.62	76.25	5.76	78.88	4.32
8	14	142.95	9.92	2.04	38.86	47.13
8	15	48.52	18.30	0.23	8.35	1.23
8	16	3.36	4.02	0.66	0.98	4.41
8	17	11.82	9.08	0.60	7.34	0.48
8	18	21.02	0.85	2.85	3.45	3.67
8	19	524.68	2.26	0.21	3.43	1.72
8	20	93.10	0.66	0.18	15.56	0.59
8	21	162.58	6.08	0.11	0.88	6.59
8	22	239.63	32.41	0.33	0.34	4.98
8	23	4.62	123.44	0.14	0.45	2.64
8	24	142.88	2.29	0.40	2.85	4.32
8	25	56.03	0.78	0.37	41.39	33.32
8	26	11.23	2.85	2.36	20.33	2.25
8	27	26.66	0.60	0.24	4.33	4.12
8	28	24.17	2.22	2.20	15.29	4.68
8	29	18.55	1.95	3.14	6.16	2.41
8	30	9.80	26.57	1.34	31.60	1.20
9	0	10.68	348.43	43.25	7.72	4.82
9	1	346.54	5.05	15.52	84.32	98.45
9	2	136.53	12.37	29.66	350.50	5.36
9	3	44.52	5.07	1.30	43.21	4.20
9	4	307.86	3.41	13.27	51.94	0.59
9	5	65.51	2.01	20.58	21.02	2.15
9	6	20.29	81.26	2.73	4.61	5.40
9	7	48.40	14.52	0.88	2.96	2.54
9	8	36.08	1.07	1.33	1.58	4.29
9	9	3.90	131.63	19.95	77.95	3.84
9	10	8.45	3.97	14.49	45.68	5.05
9	11	138.73	48.04	22.43	2.02	4.27
9	12	30.43	22.61	6.58	11.71	6.70
9	13	24.23	16.07	44.62	369.84	9.68
9	14	7.09	8.00	93.05	25.87	107.15
9	15	19.74	6.63	11.12	36.83	1.54
9	16	4.79	30.08	2.06	1.05	4.61
9	17	74.57	26.66	8.33	4.04	1.07
9	18	1.69	3.86	3.53	4.97	4.27
9	19	7.06	37.37	11.45	1.35	1.97
9	20	1.24	35.21	7.88	11.51	0.32
9	21	116.10	33.00	1.79	0.45	8.34
9	22	79.57	11.40	1.85	0.48	9.58
9	23	9.05	1.36	7.12	1.30	7.14
9	24	2.33	7.84	22.25	1.87	21.37
9	25	1.74	234.29	13.50	14.74	105.10

9	26	8.55	21.19	101.53	13.64	5.47
9	27	21.77	24.88	4.05	3.02	8.97
9	28	15.16	133.81	11.95	6.72	6.99
9	29	78.81	93.20	6.54	2.45	5.26
9	30	12.28	432.16	45.02	12.98	1.04
10	0	12.29	1.51	21.54	8.81	19.58
10	1	609.02	43.23	16.25	27.67	662.67
10	2	104.19	26.62	74.03	168.48	33.13
10	3	65.66	8.80	1.97	14.97	2.70
10	4	541.82	20.87	8.15	16.55	1.67
10	5	68.03	2.29	8.43	64.02	3.76
10	6	12.15	0.28	2.33	3.38	2.52
10	7	59.23	6.57	2.90	4.01	0.76
10	8	73.98	2.24	0.80	1.16	1.06
10	9	2.01	1.76	26.81	34.99	10.05
10	10	46.43	3.26	27.27	14.08	34.53
10	11	79.18	18.30	9.20	5.66	4.32
10	12	78.92	31.32	2.23	25.68	8.16
10	13	17.34	4.47	88.42	180.57	62.70
10	14	17.95	1.74	32.34	44.10	721.62
10	15	16.56	3.27	4.44	18.50	3.31
10	16	8.83	0.66	4.47	2.16	5.85
10	17	130.56	32.19	2.77	10.21	0.44
10	18	1.64	1.31	4.05	4.68	7.19
10	19	16.83	5.41	5.18	4.55	0.99
10	20	2.92	20.68	3.83	4.70	0.48
10	21	201.83	1.22	0.83	1.39	5.10
10	22	138.31	2.06	0.39	0.97	7.62
10	23	25.78	10.25	3.17	4.81	6.12
10	24	8.06	1.48	10.75	8.52	22.33
10	25	3.56	1.55	5.27	69.05	18.68
10	26	14.12	0.87	38.05	34.09	5.24
10	27	38.76	2.26	2.20	8.07	55.60
10	28	26.02	1.64	9.67	23.75	46.42
10	29	142.16	0.88	8.47	7.85	5.36
10	30	23.96	3.94	23.41	4.42	0.48
11	0	5.56	23.93	260.05	9.43	39.11
11	1	134.79	6.21	10.90	99.61	0.62
11	2	46.39	677.39	34.54	416.44	0.37
11	3	44.68	14.51	1.90	54.36	0.98
11	4	117.03	5.30	8.01	71.82	0.62
11	5	22.54	1.95	9.46	78.93	6.78
11	6	5.12	3.95	27.94	39.36	1.98
11	7	24.81	1.47	3.75	10.60	4.10

11	8	10.70	6.64	0.55	12.61	5.91
11	9	1.93	31.91	5.27	48.87	8.27
11	10	2.97	27.50	4.16	33.61	1.13
11	11	48.49	7.26	83.01	6.48	3.89
11	12	40.26	6.82	5.90	28.41	4.62
11	13	11.83	15.67	37.05	442.83	0.74
11	14	2.83	35.51	123.96	65.73	2.74
11	15	19.74	20.01	25.47	44.64	6.98
11	16	1.55	16.98	2.42	3.52	6.96
11	17	29.69	721.35	13.49	11.69	0.69
11	18	0.69	1.87	1.22	8.22	9.81
11	19	2.50	3.54	57.75	4.60	10.80
11	20	0.33	15.93	33.57	13.76	2.26
11	21	42.58	8.34	7.32	2.75	8.50
11	22	29.25	16.78	16.79	1.61	121.07
11	23	2.34	4.28	15.96	2.23	56.44
11	24	1.15	3.79	116.31	17.52	119.51
11	25	0.63	7.60	26.05	86.74	629.30
11	26	2.92	7.67	277.66	31.99	36.55
11	27	7.65	2.24	37.39	21.03	4.82
11	28	5.35	27.24	135.99	26.28	2.02
11	29	31.51	14.14	1.82	8.17	23.40
11	30	6.83	10.58	493.47	17.18	3.20
12	0	3.68	2.57	21.04	307.35	15.58
12	1	172.24	84.72	8.46	15.15	0.68
12	2	55.55	26.31	658.80	18.82	3.65
12	3	79.02	12.18	2.15	7.45	0.43
12	4	143.95	72.92	6.58	10.06	5.68
12	5	51.67	6.71	6.14	26.57	6.82
12	6	30.78	0.90	13.04	13.79	0.21
12	7	51.72	12.38	57.79	40.30	0.35
12	8	4.79	0.99	6.76	1.35	1.97
12	9	0.60	1.86	8.19	2.57	0.78
12	10	14.48	2.45	8.06	2.90	0.57
12	11	10.18	17.09	14.53	113.41	44.26
12	12	75.06	29.32	18.59	14.06	0.08
12	13	3.37	5.17	713.61	17.72	1.01
12	14	4.55	1.02	26.42	40.17	0.70
12	15	37.68	2.10	3.12	3.04	3.57
12	16	3.22	0.65	38.13	31.67	20.93
12	17	38.37	29.14	1.91	6.28	2.56
12	18	0.63	3.07	1.20	47.09	12.94
12	19	2.54	4.96	6.05	36.03	10.13
12	20	0.81	19.11	1.73	8.95	1.42

12	21	50.73	0.55	0.77	5.41	2.23
12	22	34.98	4.87	3.80	23.11	6.55
12	23	11.13	17.42	2.98	20.53	10.50
12	24	0.55	0.99	5.73	191.52	3.19
12	25	1.52	0.34	3.26	104.10	27.92
12	26	6.97	0.84	26.62	34.58	1.42
12	27	13.45	2.16	6.07	509.58	0.70
12	28	8.88	0.49	36.29	88.30	0.39
12	29	39.20	0.32	58.29	14.51	15.85
12	30	1.27	3.39	44.02	74.24	3.00
13	0	292.27	21.63	12.74	14.33	0.54
13	1	4.28	128.28	26.87	85.73	39.89
13	2	35.61	15.53	150.59	382.19	2.59
13	3	27.33	664.76	2.87	47.32	32.91
13	4	5.94	123.65	21.01	68.00	2.15
13	5	35.65	14.73	14.16	56.45	0.78
13	6	21.09	3.73	6.05	90.00	36.79
13	7	8.27	4.99	16.63	29.32	0.86
13	8	5.88	9.27	3.45	24.08	0.79
13	9	18.90	49.03	25.69	26.41	0.78
13	10	3.00	10.35	21.79	7.05	1.11
13	11	17.33	9.09	5.75	3.57	0.27
13	12	9.31	5.34	6.07	11.46	15.00
13	13	11.98	10.56	168.97	397.34	1.38
13	14	109.88	7.39	53.71	24.17	10.27
13	15	36.54	12.18	4.13	41.77	0.17
13	16	5.57	9.58	9.43	1.25	0.56
13	17	4.69	16.44	5.48	8.44	5.34
13	18	3.28	5.43	5.39	5.08	0.41
13	19	59.39	172.48	4.20	2.94	0.88
13	20	10.77	685.54	1.96	12.55	1.91
13	21	39.28	29.71	0.25	0.39	33.18
13	22	59.06	12.66	2.84	0.68	1.66
13	23	60.06	23.97	4.78	0.71	5.53
13	24	511.28	6.23	9.75	2.45	3.62
13	25	296.70	9.87	6.12	43.39	1.01
13	26	12.51	37.59	62.30	26.83	0.57
13	27	34.37	2.45	4.36	2.08	4.77
13	28	94.27	26.32	14.60	15.63	4.08
13	29	21.49	14.77	20.65	5.06	1.27
13	30	13.34	39.29	49.37	14.14	0.36
14	0	28.97	2.93	5.98	12.94	11.08
14	1	22.51	161.53	753.56	736.14	1.50
14	2	576.93	36.72	2.56	16.38	1.94

14	3	141.86	60.95	112.96	408.79	1.95
14	4	5.25	154.12	734.52	596.03	1.84
14	5	520.31	13.53	10.47	26.57	2.73
14	6	104.70	1.60	0.46	9.64	1.31
14	7	15.52	15.58	3.78	4.41	29.47
14	8	17.46	8.37	36.35	56.60	34.23
14	9	4.13	2.69	1.84	1.58	6.48
14	10	26.61	2.58	0.65	0.77	0.88
14	11	23.03	9.54	3.79	4.61	2.37
14	12	50.69	3.09	0.23	3.13	0.51
14	13	85.41	3.26	1.58	17.79	0.29
14	14	24.74	2.52	4.21	6.37	0.39
14	15	47.99	1.21	1.00	2.93	1.29
14	16	2.96	0.72	3.22	1.04	3.26
14	17	2.73	32.52	16.14	3.32	0.73
14	18	3.92	0.95	121.46	13.22	6.31
14	19	4.86	17.24	1.29	8.62	27.63
14	20	2.11	64.86	1.09	108.04	1.10
14	21	1.97	1.63	1.08	1.95	5.26
14	22	4.01	9.31	11.50	0.29	18.74
14	23	109.39	25.29	1.64	0.39	18.27
14	24	29.78	3.18	1.68	4.03	3.73
14	25	11.27	0.38	2.14	16.60	6.07
14	26	65.77	4.17	6.12	9.72	2.78
14	27	42.56	0.79	1.66	6.58	1.87
14	28	34.74	3.43	61.24	10.72	1.87
14	29	27.38	2.17	72.11	5.53	14.45
14	30	12.99	6.22	41.69	113.03	3.31
15	0	19.76	12.40	178.03	229.76	27.16
15	1	24.02	720.55	5.12	3.22	1.67
15	2	506.63	6.54	11.95	4.34	2.42
15	3	85.10	105.63	3.61	2.47	666.18
15	4	1.78	691.90	4.07	4.34	40.34
15	5	445.74	60.60	371.96	17.44	86.21
15	6	73.20	7.79	575.87	200.04	0.85
15	7	23.39	2.23	102.92	646.36	5.18
15	8	15.57	37.09	2.59	9.86	4.59
15	9	9.63	6.68	9.50	4.79	20.00
15	10	27.95	6.19	5.02	2.54	5.05
15	11	57.57	3.83	74.90	34.61	1.89
15	12	62.88	2.66	62.38	126.12	1.19
15	13	54.63	10.51	6.66	4.86	5.45
15	14	28.05	2.91	40.94	67.61	2.45
15	15	35.44	3.16	40.09	1.45	8.41



15	16	1.20	1.43	0.84	32.43	22.54
15	17	4.11	6.05	27.26	18.19	7.80
15	18	3.45	1.21	6.28	218.09	15.47
15	19	3.62	30.17	305.15	221.85	13.93
15	20	1.58	106.07	41.44	14.12	9.32
15	21	2.58	3.55	14.96	34.92	716.76
15	22	1.08	37.54	4.08	8.52	6.34
15	23	103.12	112.80	51.94	71.20	102.96
15	24	26.75	2.59	64.21	19.87	55.32
15	25	9.39	1.56	39.13	64.85	11.34
15	26	42.18	7.88	403.85	48.09	11.28
15	27	24.89	0.88	17.24	163.09	18.68
15	28	24.98	2.68	51.31	32.35	0.81
15	29	9.12	2.48	3.08	11.20	19.76
15	30	13.20	20.13	88.08	17.44	6.25
16	0	25.50	1.06	164.90	167.69	130.75
16	1	13.93	16.13	4.19	2.73	72.32
16	2	64.72	58.30	12.53	5.29	3.69
16	3	724.41	6.94	1.38	1.20	50.99
16	4	5.00	15.82	1.97	1.36	42.49
16	5	37.95	1.48	161.49	15.38	462.85
16	6	648.56	1.07	52.38	14.86	72.30
16	7	40.51	53.21	7.21	44.14	31.72
16	8	20.93	2.15	0.79	1.13	64.28
16	9	6.32	5.12	11.54	4.23	111.65
16	10	33.80	0.51	2.08	0.88	15.24
16	11	10.85	7.31	84.27	58.10	12.67
16	12	31.01	1.58	5.56	19.69	2.81
16	13	26.44	0.54	2.58	0.48	4.66
16	14	22.17	0.41	34.49	13.21	5.02
16	15	78.92	0.51	21.01	0.59	14.26
16	16	11.75	0.79	0.87	15.52	86.20
16	17	4.45	30.59	3.77	2.86	15.20
16	18	1.60	0.29	1.37	35.50	109.04
16	19	5.61	1.26	54.13	37.19	3.60
16	20	1.29	6.30	11.33	2.72	20.69
16	21	2.60	1.32	6.54	18.10	30.02
16	22	3.39	1.50	3.61	1.79	208.89
16	23	41.06	2.82	15.21	17.44	119.40
16	24	13.76	2.22	39.19	40.73	7.79
16	25	6.05	0.49	22.18	3.74	60.87
16	26	93.63	1.37	169.86	3.78	30.14
16	27	84.73	3.33	5.35	58.88	10.25
16	28	50.24	3.43	21.88	36.30	13.71

16	29	17.89	1.68	3.39	10.19	25.95
16	30	99.28	0.66	65.75	19.27	5.48
17	0	198.48	136.37	34.44	69.58	109.59
17	1	2.64	1.47	1.32	12.21	616.62
17	2	11.54	2.94	24.27	199.12	32.04
17	3	0.79	7.30	2.58	3.70	31.70
17	4	4.15	2.16	0.52	8.07	24.56
17	5	9.04	3.74	143.56	650.95	252.36
17	6	9.80	274.87	60.66	121.75	605.82
17	7	1.21	2.30	11.82	41.11	11.98
17	8	5.01	0.27	1.48	45.50	15.80
17	9	694.66	36.95	23.79	142.94	53.26
17	10	6.22	0.71	1.12	5.04	22.93
17	11	8.76	3.90	13.36	46.91	3.78
17	12	2.62	4.36	9.76	17.67	3.91
17	13	13.38	2.20	1.13	2.93	5.03
17	14	47.65	1.13	11.99	11.54	1.02
17	15	11.23	1.12	8.58	2.01	13.12
17	16	2.62	15.42	0.92	17.96	38.15
17	17	3.60	4.02	20.92	57.87	8.90
17	18	8.21	12.98	2.32	42.77	31.41
17	19	218.74	39.20	53.20	49.14	1.87
17	20	47.83	12.83	6.21	6.62	10.97
17	21	114.04	143.60	3.23	13.19	20.15
17	22	173.37	20.57	3.03	4.16	134.31
17	23	5.04	4.44	41.87	2.85	71.07
17	24	199.26	20.33	6.73	36.74	31.64
17	25	95.97	40.77	12.61	567.24	26.35
17	26	9.81	4.27	142.85	256.27	25.32
17	27	24.91	20.19	3.21	32.00	75.08
17	28	31.47	9.55	26.12	189.28	64.87
17	29	14.96	12.40	4.60	62.69	14.86
17	30	5.37	55.45	32.94	27.94	5.17
18	0	13.21	13.45	23.54	18.81	180.01
18	1	3.50	7.12	2.76	2.96	413.96
18	2	45.38	5.47	748.80	574.50	21.51
18	3	211.35	42.98	2.33	1.88	52.68
18	4	0.99	8.56	0.99	2.09	42.31
18	5	13.00	0.84	31.52	163.92	463.77
18	6	174.03	35.11	45.49	327.21	407.53
18	7	5.30	4.57	120.81	104.37	16.21
18	8	9.80	1.16	41.24	123.69	33.50
18	9	21.10	8.73	728.40	430.71	69.71
18	10	4.89	0.72	1.37	4.63	10.46

18	11	47.66	12.32	14.70	8.16	5.16
18	12	78.41	4.49	4.92	5.70	6.04
18	13	6.16	2.11	0.73	3.70	3.43
18	14	6.60	0.48	10.67	2.51	0.78
18	15	3.85	0.54	1.65	7.05	20.29
18	16	3.49	2.81	9.75	3.89	74.42
18	17	0.95	1.25	7.19	14.41	15.64
18	18	1.20	2.40	0.55	9.93	62.24
18	19	6.74	2.54	26.54	12.62	3.93
18	20	1.48	37.13	6.58	2.14	19.68
18	21	3.69	20.95	1.78	3.01	32.86
18	22	5.77	2.50	8.65	1.01	226.04
18	23	6.34	1.44	4.99	0.73	127.51
18	24	4.77	6.70	10.16	9.25	26.28
18	25	1.45	4.97	5.84	141.68	50.15
18	26	26.57	1.73	25.91	64.00	33.65
18	27	23.34	1.51	32.62	10.31	49.90
18	28	14.33	0.76	58.66	48.24	37.99
18	29	5.81	0.95	90.95	16.42	21.16
18	30	29.57	2.20	39.60	7.73	7.93
19	0	8.14	289.29	219.02	10.41	0.52
19	1	8.00	1.96	4.96	2.07	0.25
19	2	50.84	5.68	60.15	434.48	0.29
19	3	134.93	10.93	3.21	1.49	15.04
19	4	1.58	9.39	1.22	0.99	1.14
19	5	27.31	10.44	527.86	78.71	0.75
19	6	116.69	655.68	400.57	247.96	1.96
19	7	4.50	5.78	76.53	78.75	0.16
19	8	9.95	0.61	3.71	93.81	0.18
19	9	5.24	72.90	57.36	327.75	0.14
19	10	12.88	4.10	2.11	3.83	0.15
19	11	48.60	14.20	114.85	3.08	0.07
19	12	14.11	4.33	44.64	3.89	12.33
19	13	5.66	7.29	3.25	8.52	0.59
19	14	8.40	2.81	63.03	1.83	2.41
19	15	12.18	1.85	13.92	3.90	0.10
19	16	2.51	37.98	2.24	1.61	0.14
19	17	0.69	5.56	55.52	6.90	1.04
19	18	0.30	39.33	6.77	5.16	0.21
19	19	1.72	166.36	236.20	7.43	0.20
19	20	0.76	23.73	67.21	1.37	3.61
19	21	1.38	359.83	14.56	1.59	12.25
19	22	2.24	47.42	16.37	0.46	1.03
19	23	1.90	4.62	7.76	0.66	2.37

19	24	1.44	49.90	123.72	4.59	1.16
19	25	2.56	74.11	66.13	67.28	0.15
19	26	21.90	9.46	486.49	30.85	0.17
19	27	16.96	28.84	2.43	5.93	0.51
19	28	10.38	21.09	17.11	23.44	0.30
19	29	5.57	26.93	12.21	7.00	0.65
19	30	19.20	56.73	208.42	3.87	1.45
20	0	167.81	3.64	2.13	102.39	80.51
20	1	4.53	6.33	1.25	0.68	5.23
20	2	2.50	5.41	58.67	7.88	0.23
20	3	3.90	60.47	0.22	1.40	3.55
20	4	0.97	6.69	0.90	2.89	2.02
20	5	0.80	1.46	1.08	28.61	24.67
20	6	2.68	4.78	1.69	89.52	3.65
20	7	3.09	4.80	8.40	292.05	25.97
20	8	1.61	0.70	2.97	5.25	26.55
20	9	29.29	7.33	53.92	6.26	103.45
20	10	0.76	2.48	0.71	1.72	11.41
20	11	1.26	44.52	0.62	25.81	7.00
20	12	2.81	8.98	0.36	103.33	3.70
20	13	19.76	4.74	10.23	1.59	0.92
20	14	79.87	0.65	0.75	39.56	0.97
20	15	15.49	3.06	0.09	0.86	13.88
20	16	1.93	1.11	0.09	51.40	30.84
20	17	4.67	1.47	0.40	5.49	1.39
20	18	2.11	3.06	0.11	165.87	36.61
20	19	38.22	10.00	0.22	166.56	4.34
20	20	7.79	35.80	0.13	7.26	2.61
20	21	25.69	1.86	0.16	24.06	5.05
20	22	38.23	0.26	1.17	3.05	15.68
20	23	4.12	0.95	0.33	54.29	6.94
20	24	35.17	0.71	0.41	46.31	5.55
20	25	37.05	1.04	0.25	6.79	23.30
20	26	6.13	3.30	0.77	6.11	16.13
20	27	3.31	3.87	2.73	77.19	2.18
20	28	16.73	2.22	4.51	15.55	3.21
20	29	2.21	0.80	7.14	8.51	6.32
20	30	8.20	3.46	2.52	9.87	1.63
21	0	11.51	35.30	92.87	245.71	419.78
21	1	70.94	4.93	7.78	2.40	4.19
21	2	4.01	748.37	0.62	1.46	0.49
21	3	6.67	19.02	0.54	1.39	26.14
21	4	33.57	4.05	5.86	2.02	24.86
21	5	3.09	5.47	21.11	16.77	248.79

21	6	4.19	12.65	15.53	11.61	5.24
21	7	86.87	716.79	2.87	39.08	154.17
21	8	48.05	35.24	0.40	0.85	155.40
21	9	3.66	96.06	1.09	3.65	592.65
21	10	3.24	26.51	8.52	1.12	65.59
21	11	2.49	21.14	57.60	290.20	49.06
21	12	6.88	5.48	4.30	30.72	24.46
21	13	3.11	7.19	1.70	1.82	2.58
21	14	12.59	19.12	16.71	31.76	2.53
21	15	3.69	11.50	12.43	2.64	31.74
21	16	0.87	9.04	0.58	36.66	235.54
21	17	0.52	4.93	11.60	1.46	8.58
21	18	0.40	1.70	1.48	28.70	245.34
21	19	2.27	6.72	10.58	29.42	22.49
21	20	0.69	6.74	16.44	3.19	5.19
21	21	18.44	9.54	11.63	16.91	6.79
21	22	12.77	5.35	3.69	2.96	120.24
21	23	0.76	2.32	16.28	9.91	68.24
21	24	4.65	1.86	23.16	38.11	16.51
21	25	2.93	9.49	4.22	12.35	42.33
21	26	1.49	4.32	31.59	5.95	86.56
21	27	2.63	2.57	2.49	123.37	21.76
21	28	1.56	18.80	5.47	30.33	28.23
21	29	9.44	9.34	1.14	6.09	2.60
21	30	1.95	32.80	17.33	21.31	3.44
22	0	16.89	127.72	6.54	246.00	75.21
22	1	98.91	3.53	131.41	5.45	0.32
22	2	5.63	31.88	0.78	7.29	0.05
22	3	2.81	56.10	13.49	3.84	3.83
22	4	48.45	3.29	82.72	4.51	2.21
22	5	4.57	3.27	4.72	82.45	19.61
22	6	2.62	54.87	2.14	35.17	0.70
22	7	116.88	41.53	0.64	106.52	28.55
22	8	64.56	2.75	7.28	2.29	31.10
22	9	4.93	255.78	0.81	8.02	102.86
22	10	4.87	13.02	123.70	4.59	16.15
22	11	3.38	63.84	5.11	488.11	8.58
22	12	1.63	17.36	1.50	44.21	4.63
22	13	4.57	14.11	0.74	5.06	0.78
22	14	15.77	5.78	2.99	91.65	0.79
22	15	3.20	9.44	0.54	4.99	7.14
22	16	1.98	36.50	0.33	44.88	15.88
22	17	1.08	7.73	0.84	4.38	0.32
22	18	0.33	5.89	0.40	60.25	27.79

22	19	2.76	22.61	1.37	55.95	1.71
22	20	0.67	7.90	1.92	4.78	1.38
22	21	25.68	48.13	1.26	48.93	3.10
22	22	17.89	8.41	1.40	7.34	11.33
22	23	1.74	1.71	1.35	24.27	11.16
22	24	6.45	6.37	1.10	40.67	9.61
22	25	3.82	21.29	0.21	33.24	3.45
22	26	1.70	42.66	4.81	14.73	14.75
22	27	3.51	6.50	0.59	190.97	27.31
22	28	2.40	24.93	13.35	56.13	27.16
22	29	13.53	10.76	15.33	10.39	3.21
22	30	2.02	107.91	8.89	40.01	1.24
23	0	2.12	206.47	209.64	92.48	21.35
23	1	13.80	4.75	2.63	1.78	0.17
23	2	3.44	45.04	4.57	1.26	0.05
23	3	13.52	79.71	0.75	0.91	0.91
23	4	2.07	4.79	1.54	1.36	4.11
23	5	1.35	3.27	97.37	59.95	41.06
23	6	9.18	51.83	45.63	6.10	0.24
23	7	8.64	60.79	8.70	16.20	7.74
23	8	4.77	6.12	0.74	0.65	10.62
23	9	0.32	396.13	3.68	2.58	27.16
23	10	0.87	15.48	1.11	1.35	4.60
23	11	5.09	92.55	241.05	183.27	2.30
23	12	6.94	26.27	27.26	28.45	0.25
23	13	1.86	21.20	3.68	3.95	0.82
23	14	1.40	10.12	98.83	46.44	1.08
23	15	2.61	6.82	44.41	2.08	6.24
23	16	0.68	56.00	0.65	17.05	24.54
23	17	1.14	12.28	8.05	2.17	1.80
23	18	0.43	8.33	1.37	13.68	10.65
23	19	0.15	23.71	44.47	11.61	1.08
23	20	0.09	16.29	11.13	1.77	0.42
23	21	0.24	55.82	4.96	13.47	0.35
23	22	0.34	7.53	4.18	2.23	3.36
23	23	0.46	2.71	11.04	9.29	2.69
23	24	0.32	8.91	19.82	12.82	6.48
23	25	0.70	31.32	3.21	25.08	1.39
23	26	1.38	31.79	45.99	11.22	1.74
23	27	1.16	12.31	5.68	74.23	2.03
23	28	0.79	25.36	25.89	24.17	1.41
23	29	3.73	19.22	1.15	5.71	8.39
23	30	1.24	172.67	121.00	15.48	44.31
24	0	19.77	176.52	113.46	7.34	11.58

24	1	1.47	6.92	5.69	115.91	0.32
24	2	0.65	46.43	2.96	0.70	1.90
24	3	0.45	88.82	1.13	43.13	4.55
24	4	0.45	8.80	3.35	59.95	1.99
24	5	1.12	3.51	71.40	4.82	4.01
24	6	0.63	59.42	80.86	4.01	0.48
24	7	1.35	61.46	15.47	7.47	10.56
24	8	0.66	5.99	0.95	16.00	8.73
24	9	4.90	457.38	4.31	3.19	6.27
24	10	0.17	18.96	4.73	111.03	0.34
24	11	1.45	107.31	77.54	12.43	31.12
24	12	0.50	29.97	24.28	1.40	4.39
24	13	1.11	25.12	2.33	0.70	0.88
24	14	5.69	10.39	59.44	3.17	0.47
24	15	1.39	10.00	22.66	0.32	8.72
24	16	0.57	46.66	0.65	1.77	3.06
24	17	0.37	14.10	10.02	0.47	0.58
24	18	0.12	6.90	2.02	3.05	3.34
24	19	3.39	28.56	56.60	3.07	1.08
24	20	0.73	27.53	24.75	0.48	0.07
24	21	1.55	44.66	14.26	1.67	0.19
24	22	1.45	9.91	3.66	0.27	1.80
24	23	1.32	1.46	9.41	1.23	3.10
24	24	11.92	4.06	13.24	1.48	1.17
24	25	9.07	18.51	6.58	4.27	0.55
24	26	2.59	5.04	28.06	1.79	1.74
24	27	5.05	9.62	4.36	5.64	0.75
24	28	6.81	26.90	16.73	3.17	0.60
24	29	2.91	16.14	0.78	1.05	7.48
24	30	1.34	193.78	60.46	15.66	2.30
25	0	132.31	163.07	169.72	161.96	193.64
25	1	412.19	5.91	2.76	1.79	3.65
25	2	6.07	18.22	5.94	1.20	0.13
25	3	6.18	89.33	1.75	0.94	46.63
25	4	62.86	6.94	3.05	1.32	4.47
25	5	3.29	2.23	208.71	84.14	57.29
25	6	6.09	53.46	145.43	136.70	3.07
25	7	269.91	21.43	26.12	447.16	27.75
25	8	130.89	2.60	1.58	6.46	30.20
25	9	55.56	154.45	6.72	5.54	84.06
25	10	7.41	5.23	2.58	3.13	9.25
25	11	7.65	90.69	283.81	61.57	6.73
25	12	7.51	17.52	29.91	169.92	44.89
25	13	46.53	12.14	6.73	2.49	3.27

25	14	194.58	3.71	191.31	6.91	1.87
25	15	39.07	6.05	19.44	2.62	31.79
25	16	6.24	18.48	1.47	131.61	32.82
25	17	35.42	4.72	24.82	20.94	3.78
25	18	1.77	5.17	2.97	223.78	49.51
25	19	48.23	16.96	116.06	220.51	2.46
25	20	8.07	13.23	38.05	8.14	3.21
25	21	20.84	25.74	20.46	45.23	4.43
25	22	31.11	2.86	5.54	4.80	33.94
25	23	6.67	1.57	19.97	82.90	13.63
25	24	49.62	2.40	11.67	14.38	3.26
25	25	44.28	3.08	8.94	12.34	5.09
25	26	10.32	13.68	48.46	3.99	46.72
25	27	17.57	4.15	4.94	6.35	5.75
25	28	10.56	10.67	34.61	8.18	10.15
25	29	109.08	4.87	1.08	9.49	2.48
25	30	6.78	85.99	162.20	3.90	2.42
26	0	371.26	62.64	56.59	27.90	32.15
26	1	138.25	2.84	0.53	0.53	0.28
26	2	14.08	6.37	0.78	0.26	2.55
26	3	15.31	734.77	0.38	0.21	743.99
26	4	16.26	2.10	0.83	0.28	39.84
26	5	14.72	8.87	34.09	5.69	29.05
26	6	9.03	24.87	51.44	2.10	0.34
26	7	93.04	9.84	9.51	6.49	13.09
26	8	46.12	8.55	0.34	0.28	13.11
26	9	170.95	186.93	1.21	0.44	36.22
26	10	4.88	37.19	0.69	0.65	4.92
26	11	24.08	680.69	13.02	6.36	5.89
26	12	19.26	113.36	7.09	4.64	730.10
26	13	142.46	66.87	1.39	0.37	37.16
26	14	591.05	6.34	29.64	5.38	2.63
26	15	119.59	47.21	16.22	0.35	7.91
26	16	13.99	7.81	0.72	2.51	23.71
26	17	29.51	2.09	7.00	1.35	2.30
26	18	4.41	3.05	1.01	5.59	9.84
26	19	147.94	6.57	35.88	4.76	2.26
26	20	26.72	4.72	10.48	0.29	0.48
26	21	64.86	14.03	5.07	4.12	2.05
26	22	96.19	4.26	1.29	0.19	7.44
26	23	11.43	2.35	4.83	1.80	1.45
26	24	156.16	2.40	6.49	4.70	7.84
26	25	132.60	3.34	2.94	4.04	1.94
26	26	16.27	10.66	6.13	1.69	4.16



26	27	15.82	3.11	1.00	3.89	3.07
26	28	68.78	10.49	3.27	3.49	2.08
26	29	25.04	4.34	0.28	0.91	2.11
26	30	25.04	46.43	17.23	1.01	1.77
27	0	123.67	5.77	245.90	172.42	25.79
27	1	3.56	144.63	4.45	1.22	1.51
27	2	13.96	1.61	7.97	1.20	0.27
27	3	6.54	12.34	3.35	0.61	10.30
27	4	1.14	129.58	4.15	2.70	5.57
27	5	6.93	11.08	315.69	113.44	61.99
27	6	4.06	1.38	324.75	130.86	1.63
27	7	1.83	1.36	57.48	428.56	453.91
27	8	1.29	19.93	2.55	6.92	442.12
27	9	169.56	6.11	7.76	5.63	215.97
27	10	3.24	15.01	4.15	3.37	19.79
27	11	17.17	21.29	314.82	139.93	27.52
27	12	5.95	121.56	68.04	173.24	10.91
27	13	14.16	12.65	11.86	4.74	4.48
27	14	44.94	1.40	297.15	23.44	2.51
27	15	5.25	2.79	50.33	2.68	30.71
27	16	1.53	0.64	3.11	117.11	34.59
27	17	3.81	1.26	53.80	13.41	2.60
27	18	4.58	0.38	4.78	205.02	40.68
27	19	122.39	1.43	253.06	197.93	11.07
27	20	16.02	1.38	48.77	7.90	0.95
27	21	26.66	1.75	18.79	42.89	2.27
27	22	41.83	1.46	5.55	5.04	9.05
27	23	3.61	0.28	35.42	77.62	9.26
27	24	18.11	0.74	38.84	11.55	6.67
27	25	4.16	1.13	20.60	19.18	7.10
27	26	9.78	0.67	41.50	7.68	5.60
27	27	3.60	0.44	4.40	30.95	2.15
27	28	6.04	1.03	36.12	9.67	0.72
27	29	3.59	0.70	1.45	8.77	13.28
27	30	2.97	3.78	200.27	7.87	1.68
28	0	37.92	6.43	213.06	164.52	22.28
28	1	5.87	0.05	2.54	3.23	11.56
28	2	121.88	0.18	4.26	2.75	0.54
28	3	7.29	0.08	0.48	0.46	10.36
28	4	3.88	0.32	2.00	2.58	4.42
28	5	56.20	0.38	228.50	151.82	40.90
28	6	3.06	19.89	625.42	63.33	7.84
28	7	5.31	0.27	116.72	215.56	33.82
28	8	4.19	1.32	0.73	3.20	37.27

28	9	106.27	2.55	4.44	5.77	107.81
28	10	21.02	2.10	2.16	1.15	12.88
28	11	171.91	0.43	247.93	309.28	11.12
28	12	12.86	0.95	86.03	124.60	9.02
28	13	28.11	4.59	6.10	9.85	2.06
28	14	24.04	3.65	273.24	132.96	10.95
28	15	10.89	1.28	206.51	6.50	17.65
28	16	0.85	3.58	3.11	103.48	9.97
28	17	1.09	0.16	69.53	19.21	46.60
28	18	2.86	2.82	3.87	95.71	24.41
28	19	78.39	8.06	448.27	97.67	4.89
28	20	11.93	1.67	74.74	1.84	4.54
28	21	18.49	16.69	37.03	29.14	11.57
28	22	32.03	4.22	3.40	4.24	78.10
28	23	7.45	1.24	28.66	36.19	35.12
28	24	9.78	1.89	65.34	30.03	3.54
28	25	2.65	1.23	30.68	12.94	16.75
28	26	7.28	0.94	15.46	6.08	12.32
28	27	8.07	1.79	12.87	51.06	4.54
28	28	5.86	1.91	24.44	14.06	4.26
28	29	2.79	1.61	1.32	8.33	2.68
28	30	2.38	14.34	103.30	14.43	2.48
29	0	241.08	184.77	221.34	107.88	7.37
29	1	2.72	2.55	7.73	14.12	124.20
29	2	24.95	9.41	5.85	11.65	5.86
29	3	5.85	25.72	4.34	7.82	1.59
29	4	1.27	7.70	5.22	11.18	1.89
29	5	5.39	8.26	585.98	687.86	18.58
29	6	2.33	606.20	39.84	49.73	77.61
29	7	3.51	4.34	3.01	109.42	5.93
29	8	4.68	1.78	2.90	4.85	6.37
29	9	682.54	70.22	5.37	8.59	2.85
29	10	6.77	2.84	3.98	4.28	7.00
29	11	31.95	24.08	476.10	147.94	0.59
29	12	13.61	4.80	44.53	296.61	2.36
29	13	51.71	9.15	27.65	46.99	10.49
29	14	211.71	2.33	516.56	569.52	123.57
29	15	44.02	5.35	26.65	15.79	2.89
29	16	3.41	144.20	2.57	30.55	14.18
29	17	12.37	2.52	37.43	81.42	4.25
29	18	19.00	58.81	7.82	68.09	8.49
29	19	469.31	205.38	39.20	63.27	0.72
29	20	74.18	41.88	10.49	10.42	0.28
29	21	114.05	385.14	11.29	27.05	1.01

29	22	172.20	32.79	7.53	4.79	2.26
29	23	9.75	2.59	24.91	14.68	0.74
29	24	74.62	40.51	36.68	29.10	0.60
29	25	49.61	28.11	13.55	27.67	2.42
29	26	2.53	6.40	46.78	14.50	3.99
29	27	8.56	36.99	11.43	19.97	0.43
29	28	11.09	4.61	26.19	30.35	0.98
29	29	2.43	4.44	1.86	10.07	0.51
29	30	6.24	19.25	72.41	7.02	0.30
30	0	42.82	36.40	36.84	164.60	284.64
30	1	1.24	1.09	0.30	4.44	2.21
30	2	5.01	1.75	0.47	3.81	0.79
30	3	6.79	2.11	0.21	2.36	36.71
30	4	1.52	2.21	0.39	3.82	65.82
30	5	1.13	1.48	7.78	203.88	682.68
30	6	3.03	120.04	67.52	55.68	2.06
30	7	2.28	2.93	12.65	163.73	33.95
30	8	0.81	0.66	1.51	3.33	82.91
30	9	31.97	48.37	0.64	4.01	262.25
30	10	1.06	2.20	0.52	3.77	31.67
30	11	7.05	5.83	34.94	146.50	25.53
30	12	7.55	2.47	10.49	54.43	46.18
30	13	1.90	2.14	0.34	15.20	4.59
30	14	6.68	1.00	12.95	190.38	17.51
30	15	4.13	1.04	24.57	4.42	99.16
30	16	0.63	32.33	0.42	40.19	486.44
30	17	0.96	2.58	9.23	24.90	28.31
30	18	1.62	51.95	0.50	94.62	260.92
30	19	22.18	44.69	58.14	92.49	20.73
30	20	4.14	7.63	8.35	4.24	11.80
30	21	8.34	79.33	4.15	31.69	4.07
30	22	10.95	7.31	0.80	2.43	43.83
30	23	4.19	0.55	6.62	30.27	20.82
30	24	9.94	8.23	9.81	23.52	4.70
30	25	4.89	14.78	4.12	23.20	8.08
30	26	3.06	3.29	1.37	11.78	51.38
30	27	3.98	7.40	3.24	28.75	12.72
30	28	2.48	6.21	7.52	17.16	16.44
30	29	2.13	4.98	0.91	4.94	3.19
30	30	2.73	22.18	13.53	6.06	7.56

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**Table S4.** Coordinate of Conformer 1.

Element	X /Bohr	Y /Bohr	Z /Bohr
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Ir	-2.608552	-0.038520	0.161827
O	-1.992924	-2.042790	-0.520425
O	-1.030308	-0.293498	1.703937
C	-1.154334	-2.814900	0.030110
C	-0.301769	-1.315987	1.878102
C	-0.331071	-2.512174	1.133658
C	-1.068922	-4.187428	-0.575927
C	0.672464	-1.219440	3.020157
H	0.190841	-0.758041	3.893448
H	1.099384	-2.188965	3.308332
H	1.492060	-0.553801	2.711055
H	-0.337765	-4.839230	-0.082036
H	-2.059412	-4.662865	-0.534666
H	-0.809115	-4.106961	-1.640896
H	0.349919	-3.302046	1.460234
C	-2.420323	2.842436	-0.063431
C	-2.653326	4.202320	0.190395
C	-3.531245	4.585907	1.196883
C	-4.181500	3.602707	1.950704
C	-3.959689	2.249714	1.699391
H	-3.712424	5.644647	1.395572
H	-2.134412	4.967770	-0.395115
H	-4.875955	3.899009	2.742411
H	-4.494235	1.506183	2.299361
C	-1.482568	2.363049	-1.065670
C	-0.750906	3.156199	-1.955834
C	0.143478	2.563331	-2.837049
H	-0.888583	4.238365	-1.944032
C	-0.446127	0.434394	-1.917546
C	0.313250	1.179416	-2.810058
H	0.720013	3.178901	-3.532557
H	-0.363152	-0.653451	-1.853789
H	1.035724	0.671341	-3.447372
C	-5.202856	-1.256788	0.702034
C	-6.213423	-1.938742	1.393689
C	-6.012290	-2.340342	2.705956
C	-4.794951	-2.052881	3.328038
C	-3.824040	-1.380057	2.601722
H	-6.800260	-2.870610	3.245399
H	-7.158541	-2.144589	0.889521
H	-4.595841	-2.343482	4.360008
H	-2.853081	-1.113796	3.026901
C	-5.285155	-0.747762	-0.656858
C	-6.419876	-0.899734	-1.469399

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C	-6.440876	-0.361595	-2.749006
H	-7.294731	-1.442235	-1.098892
C	-4.190642	0.489633	-2.418893
C	-5.321966	0.335830	-3.217399
H	-7.323395	-0.479702	-3.381457
H	-3.334248	1.040241	-2.820265
H	-5.334988	0.765928	-4.223118
C	-4.134645	-0.051917	-1.124875
C	-3.074391	1.830976	0.693274
N	-4.016391	-0.999332	1.329162
N	-1.330916	1.004408	-1.085830
Al	3.337926	-0.151932	-0.511591
N	4.536705	0.331080	1.147274
N	3.143629	-2.092224	0.129314
C	4.209202	0.770668	2.354102
C	5.840437	0.109482	0.847084
N	3.358341	1.802135	-1.094533
O	1.854598	0.403858	0.482037
O	2.482474	-0.787674	-2.031062
O	5.028721	-0.487681	-1.260382
C	3.456437	-2.688149	1.268759
C	2.654159	-2.830340	-0.899591
C	5.182006	1.007347	3.343519
H	3.145554	0.940609	2.544213
C	6.897998	0.314588	1.778549
C	6.068850	-0.349031	-0.494195
C	4.137284	2.432537	-1.960222
C	2.446380	2.497649	-0.370241
C	1.638329	1.692491	0.500227
C	2.308505	-2.075549	-2.072108
C	3.311622	-4.076119	1.445777
H	3.839543	-2.053333	2.073833
C	2.498660	-4.243684	-0.830284
C	6.513590	0.777964	3.059506
H	4.869392	1.370121	4.323705
C	8.220991	0.046617	1.358630
C	7.394673	-0.599098	-0.858381
C	4.045431	3.824070	-2.147514
H	4.853826	1.813793	-2.508028
C	2.302928	3.910417	-0.461007
C	0.710947	2.357281	1.305278
C	1.806300	-2.792722	-3.160690
C	2.846240	-4.848407	0.400508
H	3.580021	-4.524602	2.403478

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C	2.011814	-4.923633	-1.970212
C	8.437442	-0.399664	0.064806
H	7.604050	-0.952081	-1.869853
C	3.144818	4.556323	-1.396329
H	4.698359	4.311091	-2.873283
C	1.346221	4.537227	0.371815
C	0.588339	3.758585	1.231526
H	0.056610	1.768051	1.952612
C	1.679116	-4.192663	-3.097789
H	1.530742	-2.249741	-4.067325
H	9.461333	-0.607024	-0.258269
H	-0.155923	4.241849	1.870939
H	1.294144	-4.717945	-3.976236
H	1.888854	-6.008202	-1.942015
H	1.217488	5.621184	0.328675
H	9.050924	0.194055	2.052366
H	7.280985	0.954804	3.818477
H	2.740272	-5.931142	0.514729
H	3.076274	5.641201	-1.519183

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**Table S5.** Coordinate of Conformer 2.

Element	X /Bohr	Y /Bohr	Z /Bohr
Ir	-2.600548	-0.718232	-0.563740
O	-3.123146	-1.846751	-2.402740
O	-0.672877	-1.816528	-0.699985
C	-2.433897	-2.737125	-2.978763
C	-0.363525	-2.729887	-1.524591
C	-1.146815	-3.183089	-2.604602
C	-3.080564	-3.362783	-4.184474
C	0.971211	-3.380081	-1.283483
H	1.757486	-2.614221	-1.199377
H	0.958462	-3.895693	-0.310298
H	1.253931	-4.099507	-2.061187
H	-3.325986	-2.583159	-4.919648
H	-2.458706	-4.124707	-4.668637
H	-4.038580	-3.815918	-3.893350
H	-0.712937	-3.977302	-3.215433
C	-1.167590	1.489986	0.620520
C	-0.589782	2.314828	1.595337
C	-0.798486	2.056702	2.942361
C	-1.607276	0.978219	3.316608
C	-2.190456	0.154983	2.353010
H	-0.336372	2.695979	3.699159

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H	0.036845	3.157480	1.297134
H	-1.787945	0.778583	4.377101
H	-2.824673	-0.676123	2.679331
C	-1.018617	1.721289	-0.804255
C	-0.265709	2.741180	-1.405305
C	-0.266533	2.880548	-2.784861
H	0.341816	3.393765	-0.777589
C	-1.724944	0.988156	-2.925052
C	-1.025827	2.001502	-3.562986
H	0.325481	3.668971	-3.255559
H	-2.322661	0.253150	-3.471451
H	-1.063242	2.084535	-4.649953
C	-4.846824	-2.051016	0.732669
C	-5.558847	-3.026824	1.443552
C	-4.900333	-4.133684	1.958579
C	-3.522424	-4.260596	1.763688
C	-2.862442	-3.274701	1.046201
H	-5.454237	-4.892357	2.515831
H	-6.631764	-2.899401	1.593842
H	-2.963385	-5.109176	2.159515
H	-1.784853	-3.306205	0.862894
C	-5.383671	-0.815223	0.189936
C	-6.734737	-0.446617	0.290187
C	-7.164309	0.775529	-0.208745
H	-7.455667	-1.119112	0.764649
C	-4.894908	1.281043	-0.909783
C	-6.236662	1.639296	-0.803006
H	-8.215096	1.062581	-0.130955
H	-4.191927	1.980788	-1.372170
H	-6.569075	2.608735	-1.185384
C	-4.434053	0.043973	-0.433073
C	-1.970060	0.374208	0.984467
N	-3.502976	-2.209674	0.538169
N	-1.710000	0.840691	-1.590207
Al	3.309094	1.171778	0.482778
N	4.158628	-0.715084	0.756079
N	3.969173	1.794384	2.323515
C	5.369782	-1.166410	0.466205
C	3.243520	-1.547219	1.314506
N	2.762356	0.660639	-1.400828
O	4.883895	1.741683	-0.366246
O	2.475920	2.840045	0.445237
O	1.828388	0.306654	1.228906
C	4.722604	1.206846	3.239506

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C	3.437187	3.017950	2.576569
C	5.739530	-2.499537	0.729138
H	6.062426	-0.446231	0.020531
C	3.522023	-2.910054	1.623454
C	1.967208	-0.940540	1.564771
C	1.678869	0.032132	-1.830853
C	3.726157	1.039696	-2.276059
C	4.875856	1.648482	-1.666551
C	2.636176	3.561852	1.514541
C	4.992176	1.812615	4.479822
H	5.130249	0.222525	2.991629
C	3.650898	3.712623	3.801270
C	4.825311	-3.361102	1.303658
H	6.746100	-2.835787	0.475931
C	2.495005	-3.682922	2.210896
C	0.982659	-1.748805	2.140690
C	1.499176	-0.257429	-3.197767
H	0.943361	-0.263504	-1.075120
C	3.619722	0.821622	-3.676698
C	5.895959	2.062682	-2.525889
C	2.076409	4.824138	1.729886
C	4.459199	3.054536	4.758137
H	5.617512	1.291047	5.205680
C	3.052484	4.982559	3.969856
C	1.263716	-3.089576	2.453659
H	-0.003849	-1.317443	2.323436
C	2.453651	0.144280	-4.110880
H	0.604340	-0.799827	-3.511999
C	4.680670	1.263272	-4.502090
C	5.779451	1.869784	-3.916050
H	6.783516	2.535993	-2.101425
C	2.288656	5.507201	2.941034
H	1.469239	5.272419	0.940197
H	0.468869	-3.690497	2.906930
H	6.599243	2.209103	-4.555428
H	1.828323	6.490570	3.069517
H	3.201849	5.531855	4.901328
H	4.626278	1.114514	-5.582409
H	2.680105	-4.728625	2.465213
H	5.100443	-4.398358	1.515585
H	4.655963	3.540707	5.717784
H	2.324356	-0.067291	-5.176411

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**Table S6.** Coordinate of Conformer 3.



Element	X /Bohr	Y /Bohr	Z /Bohr
Ir	-3.051723	0.166898	1.433172
O	-2.026977	-1.532295	2.450417
O	-4.792949	-1.208605	1.367379
C	-2.539462	-2.646547	2.777871
C	-4.883334	-2.362620	1.874647
C	-3.858249	-3.073706	2.536774
C	-1.615050	-3.576082	3.519424
C	-6.230375	-3.018027	1.735549
H	-6.511186	-3.065985	0.674033
H	-6.272169	-4.026975	2.161976
H	-6.992667	-2.393469	2.222313
H	-2.078721	-4.533512	3.783760
H	-0.714696	-3.774391	2.919753
H	-1.263226	-3.090130	4.441871
H	-4.118821	-4.067923	2.904579
C	-4.914482	2.360089	1.678182
C	-5.779575	3.402532	1.309216
C	-5.894824	3.776709	-0.022701
C	-5.137397	3.108190	-0.990799
C	-4.277965	2.071165	-0.634369
H	-6.567029	4.587990	-0.310057
H	-6.366542	3.926996	2.068971
H	-5.219297	3.404169	-2.040620
H	-3.698858	1.573045	-1.417811
C	-4.715805	1.934056	3.053092
C	-5.319137	2.492160	4.188547
C	-5.010797	2.008893	5.451443
H	-6.027064	3.312908	4.066286
C	-3.524321	0.442534	4.423519
C	-4.091344	0.964519	5.576104
H	-5.478952	2.445332	6.336471
H	-2.790682	-0.367109	4.449349
H	-3.813954	0.557999	6.549121
C	-1.155962	0.258712	-0.792386
C	-0.556619	-0.052031	-2.021539
C	-1.086081	-1.062164	-2.810883
C	-2.217965	-1.754910	-2.370722
C	-2.779206	-1.402212	-1.152382
H	-0.606744	-1.317106	-3.759155
H	0.351272	0.475230	-2.325859
H	-2.665689	-2.556507	-2.959393
H	-3.672956	-1.892004	-0.758068
C	-0.679744	1.242946	0.163151

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C	0.473394	2.017693	-0.040903
C	0.901174	2.894535	0.948533
H	1.057532	1.910174	-0.960050
C	-0.974942	2.236111	2.344049
C	0.177143	2.995723	2.140239
H	1.807764	3.485720	0.799242
H	-1.526955	2.352192	3.282522
H	0.518495	3.679296	2.923688
C	-1.433998	1.341483	1.365502
C	-4.149087	1.662295	0.701999
N	-2.265921	-0.427091	-0.384281
N	-3.828311	0.905231	3.200865
Al	3.718204	-0.271165	-1.884116
N	2.542404	-0.812209	-0.320344
N	5.083111	-1.783734	-1.480872
C	1.528673	-1.664089	-0.251563
C	2.872908	-0.070129	0.768966
N	4.917044	0.486253	-3.381394
O	2.519203	1.018590	-2.497130
O	2.993944	-1.600203	-3.000942
O	4.553981	0.806355	-0.606709
C	6.130599	-1.796821	-0.669932
C	4.765128	-2.893050	-2.192196
C	0.755901	-1.796047	0.913049
H	1.311999	-2.236636	-1.157823
C	2.162310	-0.148063	1.997264
C	3.985027	0.813351	0.571501
C	6.137178	0.169342	-3.784312
C	4.243208	1.484046	-4.010346
C	2.930018	1.748130	-3.494030
C	3.604476	-2.747802	-3.025713
C	6.935194	-2.943469	-0.530125
H	6.333620	-0.869330	-0.126558
C	5.514928	-4.100602	-2.119891
C	1.058852	-1.032845	2.024545
H	-0.107269	-2.463451	0.914834
C	2.589654	0.672143	3.067947
C	4.373351	1.592523	1.661803
C	6.764624	0.835068	-4.852782
H	6.644063	-0.638472	-3.248288
C	4.791430	2.218126	-5.100913
C	2.198006	2.767778	-4.105227
C	3.230311	-3.861585	-3.782271
C	6.630706	-4.083096	-1.248708

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H	7.790434	-2.917972	0.146755
C	5.093077	-5.196905	-2.906747
C	3.675341	1.509566	2.883329
H	5.219076	2.273927	1.547871
C	6.096381	1.850658	-5.505679
H	7.771335	0.538976	-5.150900
C	4.008161	3.239267	-5.686420
C	2.743205	3.490353	-5.181872
H	1.197042	2.995722	-3.732054
C	3.973281	-5.054562	-3.710572
H	2.353199	-3.788303	-4.428291
H	4.003555	2.143833	3.711683
H	2.139101	4.281422	-5.634281
H	3.648049	-5.903765	-4.317913
H	5.648275	-6.136229	-2.871982
H	4.408100	3.815050	-6.523268
H	2.048984	0.644183	4.016188
H	0.426909	-1.099642	2.914267
H	7.247355	-4.980994	-1.149739
H	6.567846	2.379670	-6.338648

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**Table S7.** Coordinate of Conformer 4.

Element	X /Bohr	Y /Bohr	Z /Bohr
Ir	-3.408430	0.065717	0.552295
O	-4.772809	-1.198217	1.758097
O	-2.531222	0.624138	2.527146
C	-4.832920	-1.283892	3.017789
C	-2.951182	0.272330	3.672771
C	-4.007032	-0.615361	3.948204
C	-5.902275	-2.201622	3.544281
C	-2.230805	0.903153	4.835647
H	-2.619087	0.588086	5.811134
H	-1.156620	0.668966	4.794049
H	-2.305046	1.998661	4.765182
H	-6.883394	-1.871475	3.175308
H	-5.751870	-3.213153	3.140980
H	-5.931919	-2.256507	4.638623
H	-4.216730	-0.803316	5.002853
C	-0.933750	0.305436	-0.934002
C	0.082135	0.862514	-1.724813
C	0.011116	2.199514	-2.099845
C	-1.076234	2.976798	-1.687186

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C	-2.096093	2.422757	-0.913039
H	0.805090	2.635531	-2.711227
H	0.949127	0.265140	-2.024438
H	-1.135518	4.028303	-1.984569
H	-2.944756	3.051909	-0.625371
C	-0.922354	-1.075481	-0.481295
C	0.075506	-2.019983	-0.755462
C	-0.058840	-3.318041	-0.286254
H	0.959796	-1.715539	-1.321253
C	-2.135341	-2.685249	0.723260
C	-1.186860	-3.662717	0.462143
H	0.720278	-4.055200	-0.493850
H	-3.042426	-2.890374	1.297124
H	-1.332379	-4.672823	0.847083
C	-5.784060	1.516553	-0.319412
C	-6.820952	2.459736	-0.319042
C	-6.875331	3.437315	0.663147
C	-5.882685	3.471752	1.645741
C	-4.880532	2.514187	1.606495
H	-7.682048	4.173606	0.662556
H	-7.578649	2.418036	-1.102517
H	-5.880444	4.227154	2.432116
H	-4.068613	2.489363	2.337627
C	-5.580653	0.468512	-1.305396
C	-6.433765	0.261239	-2.401216
C	-6.154167	-0.734998	-3.326724
H	-7.322436	0.885273	-2.534245
C	-4.160469	-1.328805	-2.074522
C	-5.011178	-1.524780	-3.160029
H	-6.817610	-0.895288	-4.179212
H	-3.273945	-1.962787	-1.978710
H	-4.781312	-2.304512	-3.891882
C	-4.425299	-0.340083	-1.113869
C	-2.056017	1.076543	-0.518496
N	-4.833306	1.561654	0.661914
N	-2.008275	-1.428409	0.269621
Al	4.280861	-0.216600	-0.629620
N	5.771683	0.367240	0.675264
N	5.599927	-1.344097	-1.738589
C	6.211970	-0.213258	1.781872
C	6.327524	1.529594	0.253651
N	2.816982	0.681304	0.446528
O	3.901959	-1.684391	0.471047
O	3.075407	-0.703927	-1.973306

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O	4.778877	1.379654	-1.488857
C	6.880954	-1.629564	-1.567694
C	4.939554	-1.846459	-2.814160
C	7.258214	0.348882	2.537315
H	5.722435	-1.149826	2.063884
C	7.390550	2.175002	0.945950
C	5.753174	2.053219	-0.954066
C	2.332130	1.913267	0.396154
C	2.207235	-0.252165	1.224882
C	2.839911	-1.540477	1.220069
C	3.555322	-1.473533	-2.906351
C	7.589339	-2.442183	-2.471268
H	7.372700	-1.205867	-0.687021
C	5.566600	-2.682295	-3.782385
C	7.842550	1.529915	2.122400
H	7.595795	-0.156613	3.443302
C	7.897845	3.381920	0.412367
C	6.292310	3.248092	-1.439044
C	1.183236	2.282784	1.116803
H	2.855974	2.615240	-0.259241
C	1.027893	0.020218	1.966975
C	2.269750	-2.529520	2.021592
C	2.833495	-1.974488	-3.992260
C	6.936498	-2.964544	-3.568966
H	8.645184	-2.648930	-2.291103
C	4.791185	-3.160533	-4.863383
C	7.343884	3.884908	-0.754287
H	5.883215	3.676110	-2.356081
C	0.518224	1.340169	1.875304
H	0.800478	3.300433	1.025996
C	0.456453	-1.035729	2.718423
C	1.089960	-2.265896	2.744052
H	2.734995	-3.517389	2.050375
C	3.456345	-2.800979	-4.944696
H	1.779728	-1.705957	-4.094518
H	7.737627	4.818512	-1.165527
H	0.648377	-3.075218	3.332830
H	2.857352	-3.169601	-5.781859
H	5.251677	-3.801923	-5.617295
H	-0.479399	-0.870794	3.253608
H	8.712765	3.900921	0.920595
H	8.657629	1.976901	2.698734
H	7.469793	-3.599576	-4.281936
H	-0.418266	1.583219	2.385109

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**Table S8.** Coordinate of Conformer 5.

Element	X /Bohr	Y /Bohr	Z /Bohr
Ir	-2.452346	-1.232898	0.365948
O	-3.253220	-2.265146	2.160152
O	-0.500183	-2.095811	1.018078
C	-2.596938	-2.868358	3.057155
C	-0.273413	-2.778138	2.064111
C	-1.208526	-3.129202	3.056085
C	-3.412246	-3.359820	4.221999
C	1.149326	-3.243079	2.225447
H	1.840940	-2.390891	2.147687
H	1.333954	-3.760809	3.174264
H	1.411861	-3.924308	1.402391
H	-2.822565	-3.898846	4.972499
H	-3.910626	-2.507470	4.704907
H	-4.217057	-4.015549	3.861109
H	-0.822391	-3.696197	3.905353
C	-1.779223	-1.266938	-2.452120
C	-1.296575	-0.828203	-3.695859
C	-0.688093	0.414589	-3.809200
C	-0.552098	1.220087	-2.672984
C	-1.060561	0.806178	-1.442592
H	-0.285120	0.744305	-4.769607
H	-1.366809	-1.475373	-4.576273
H	-0.031502	2.180146	-2.740232
H	-0.928360	1.463264	-0.578918
C	-2.314818	-2.594114	-2.217266
C	-2.453175	-3.614798	-3.168816
C	-2.954067	-4.853502	-2.796228
H	-2.163772	-3.419059	-4.202565
C	-3.160006	-4.033254	-0.558591
C	-3.314938	-5.072279	-1.463989
H	-3.064662	-5.647367	-3.538229
H	-3.435000	-4.131985	0.494545
H	-3.712732	-6.030089	-1.127655
C	-3.371066	1.316479	1.462665
C	-3.341313	2.521618	2.179025
C	-2.230322	2.854950	2.938945
C	-1.147613	1.973274	2.985354
C	-1.227272	0.791999	2.263351
H	-2.201287	3.797798	3.489635
H	-4.197927	3.194577	2.123490
H	-0.251451	2.202144	3.564689

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H	-0.417814	0.058071	2.247436
C	-4.450419	0.846851	0.612868
C	-5.656994	1.542817	0.439047
C	-6.639730	1.037774	-0.401252
H	-5.829893	2.485865	0.965851
C	-5.216887	-0.864176	-0.908322
C	-6.411837	-0.167165	-1.074974
H	-7.579067	1.577797	-0.536894
H	-5.073452	-1.800789	-1.455774
H	-7.180119	-0.566486	-1.743305
C	-4.208839	-0.385250	-0.056951
C	-1.691671	-0.436755	-1.300894
N	-2.297654	0.473332	1.521345
N	-2.676938	-2.834184	-0.919014
Al	3.304516	1.538954	-0.699290
N	3.188829	0.783665	1.258026
N	2.467286	-0.141514	-1.469199
C	2.211533	0.859992	2.147388
C	4.321913	0.107704	1.561849
N	3.959504	3.334717	0.023883
O	1.613930	2.317100	-0.409592
O	3.545626	2.010919	-2.471109
O	5.015127	0.760482	-0.569964
C	1.819706	-1.149937	-0.909079
C	2.665542	-0.133527	-2.813927
C	2.314002	0.247624	3.411388
H	1.324871	1.418724	1.835860
C	4.524601	-0.546817	2.810812
C	5.306883	0.113308	0.516797
C	5.191648	3.782313	0.203902
C	2.899500	4.116590	0.346683
C	1.619306	3.517628	0.089489
C	3.254175	1.072197	-3.326327
C	1.374084	-2.253007	-1.658730
H	1.613335	-1.091833	0.162585
C	2.272508	-1.210891	-3.653093
C	3.460618	-0.449407	3.739515
H	1.483944	0.323051	4.117601
C	5.746792	-1.226295	3.016669
C	6.497633	-0.573181	0.772766
C	5.434580	5.063814	0.732486
H	6.000320	3.104038	-0.082602
C	3.040817	5.420721	0.895358
C	0.490342	4.273964	0.416248

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C	3.439260	1.144454	-4.709065
C	1.622220	-2.293706	-3.014230
H	0.812808	-3.039768	-1.151589
C	2.500166	-1.094918	-5.043771
C	6.693883	-1.224574	2.004285
H	7.269049	-0.594548	0.000758
C	4.369954	5.873067	1.077761
H	6.463208	5.402311	0.864694
C	1.866082	6.145211	1.207333
C	0.632547	5.563018	0.964627
H	-0.501503	3.854016	0.231173
C	3.069821	0.067749	-5.536630
H	3.873948	2.051413	-5.133353
H	7.638766	-1.750956	2.165510
H	-0.274217	6.126889	1.202205
H	3.235310	0.161905	-6.613669
H	2.209604	-1.910069	-5.710201
H	1.943828	7.149677	1.627976
H	5.931032	-1.739278	3.962569
H	3.553997	-0.933595	4.715866
H	1.279871	-3.142303	-3.615331
H	4.544439	6.869917	1.492597

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