

Supporting Information

The important role of the hydroxyl group on the
conformational adaptability in
bis(L-threoninato)copper(II) compared to
bis(L-*allo*-threoninato)copper(II): Quantum Chemical
study

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Assignment of chair / boat / twisted boat / envelope / other twist conformations in the Oo coordination mode

The procedure for the assignment of the chelate-ring conformations in the Oo coordination mode has been as follows:

1. Definition of the torsion angles (atom numbering is defined in Figure 5):

$$\begin{aligned}\tau_1 &= \text{Cu} - \text{O}_9 - \text{C}_7 - \text{C}_5 = \text{Cu} - \text{O}_h - \text{C}^\beta - \text{C}^\alpha \\ \tau_2 &= \text{O}_9 - \text{C}_7 - \text{C}_5 - \text{C}_{15} = \text{O}_h - \text{C}^\beta - \text{C}^\alpha - \text{C}' \\ \tau_3 &= \text{O}_9 - \text{Cu} - \text{O}_{17} - \text{C}_5 = \text{O}_h - \text{Cu} - \text{O} - \text{C}^\alpha \\ \tau_4 &= \text{O}_9 - \text{Cu} - \text{O}_{17} - \text{C}_{15} = \text{O}_h - \text{Cu} - \text{O} - \text{C}' \\ \tau_5 &= \text{Cu} - \text{O}_{17} - \text{C}_{15} - \text{C}_5 = \text{Cu} - \text{O} - \text{C}' - \text{C}^\alpha \\ \tau_6 &= \text{O}_{17} - \text{C}_{15} - \text{C}_7 - \text{O}_9 = \text{O} - \text{C}' - \text{C}^\beta - \text{O}_h \\ \tau_7 &= \text{C}_7 - \text{C}_5 - \text{C}_{15} - \text{O}_{17} = \text{C}^\beta - \text{C}^\alpha - \text{C}' - \text{O} \\ \tau_8 &= \text{O}_{17} - \text{Cu} - \text{O}_9 - \text{C}_7 = \text{O} - \text{Cu} - \text{O}_h - \text{C}^\beta \\ \tau_9 &= \text{Cu} - \text{O}_9 - \text{C}_5 - \text{C}_{15} = \text{Cu} - \text{O}_h - \text{C}^\alpha - \text{C}' \\ \tau_{10} &= \text{H}_{10} - \text{O}_9 - \text{Cu} - \text{O}_{17} = \text{H}_h - \text{O}_h - \text{Cu} - \text{O}\end{aligned}$$

2. The applicable criterions used for the assignment chair / boat / twisted boat / envelope / other twist forms:

chair 1 (ch 1):

$$\begin{aligned}|\tau_1 + \tau_2 + \tau_5 + \tau_7| &< 15^\circ \text{ and} \\ \min(\tau_1, \tau_2, \tau_5, \tau_7) &\geq 45^\circ \text{ and} \\ \max(\tau_1, \tau_2, \tau_5, \tau_7) &\leq 90^\circ \text{ and } \tau_1 > 0\end{aligned}$$

chair 2 (ch2):

$$\begin{aligned}|\tau_1 + \tau_2 + \tau_5 + \tau_7| &< 15^\circ \text{ and} \\ \min(\tau_1, \tau_2, \tau_5, \tau_7) &\geq 45^\circ \text{ and} \\ \max(\tau_1, \tau_2, \tau_5, \tau_7) &\leq 90^\circ \text{ and } \tau_1 \leq 0\end{aligned}$$

boat 1 (bt1):

$$\begin{aligned}|\tau_1| &\leq 20^\circ \text{ and} \\ |\tau_5| &\leq 20^\circ \text{ and} \\ |\tau_6| &\leq 10^\circ \text{ and } \tau_2 < 0\end{aligned}$$

boat 2 (bt2):

$$\begin{aligned}|\tau_1| &\leq 20^\circ \text{ and} \\ |\tau_5| &\leq 20^\circ \text{ and} \\ |\tau_6| &\leq 10^\circ \text{ and } \tau_2 \geq 0\end{aligned}$$

boat 3 (bt3):

$|\tau_8| \leq 22^\circ$ and
 $|\tau_7| \leq 22^\circ$ and
 $|\tau_9| \leq 20^\circ$ and $\tau_1 < 0$

boat 4 (bt4):

$|\tau_8| \leq 22^\circ$ and
 $|\tau_7| \leq 22^\circ$ and
 $|\tau_9| \leq 20^\circ$ and $\tau_1 \geq 0$

twisted boat 1 (tb1):

$|\tau_1| \leq 29^\circ$ and
 $|\tau_5| \leq 29^\circ$ and
 $|\tau_6| \leq 40^\circ$ and $\tau_2 < 0$

twisted boat 2 (tb2):

$|\tau_1| \leq 29^\circ$ and
 $|\tau_5| \leq 29^\circ$ and
 $|\tau_6| \leq 40^\circ$ and $\tau_2 \geq 0$

envelope 4 down (e4d): $|\tau_4| \leq 40^\circ$ and $|\tau_8| \leq 40^\circ$ and $|\tau_6| \leq 20^\circ$ and $\tau_2 > 0$

envelope 4 up (e4u): $|\tau_4| \leq 40^\circ$ and $|\tau_8| \leq 40^\circ$ and $|\tau_6| \leq 20^\circ$ and $\tau_2 \leq 0$

envelope 3 up (e3u): $|\tau_5| \leq 33^\circ$ and $|\tau_4| \leq 33^\circ$ and $|\tau_9| \leq 20^\circ$ and $\tau_1 > 0$

envelope 3 down (e3d): $|\tau_5| \leq 33^\circ$ and $|\tau_4| \leq 33^\circ$ and $|\tau_9| \leq 20^\circ$ and $\tau_1 \leq 0$

twist 1 (tw1): $|\tau_8 - \tau_4| \leq 22^\circ$ and $\tau_8 * \tau_4 > 0^\circ$ and $|\tau_3| \leq 15^\circ$

twist2 (tw2): $|\tau_8 - \tau_5| \leq 22^\circ$ and $\tau_8 * \tau_5 > 0^\circ$ and $|\tau_4| \leq 15^\circ$

3. For all of the above, the sign minus (m) was used to define the value $\tau_0 > 0$, whereas the sign plus (p) defined the values with $\tau_0 \leq 0$.

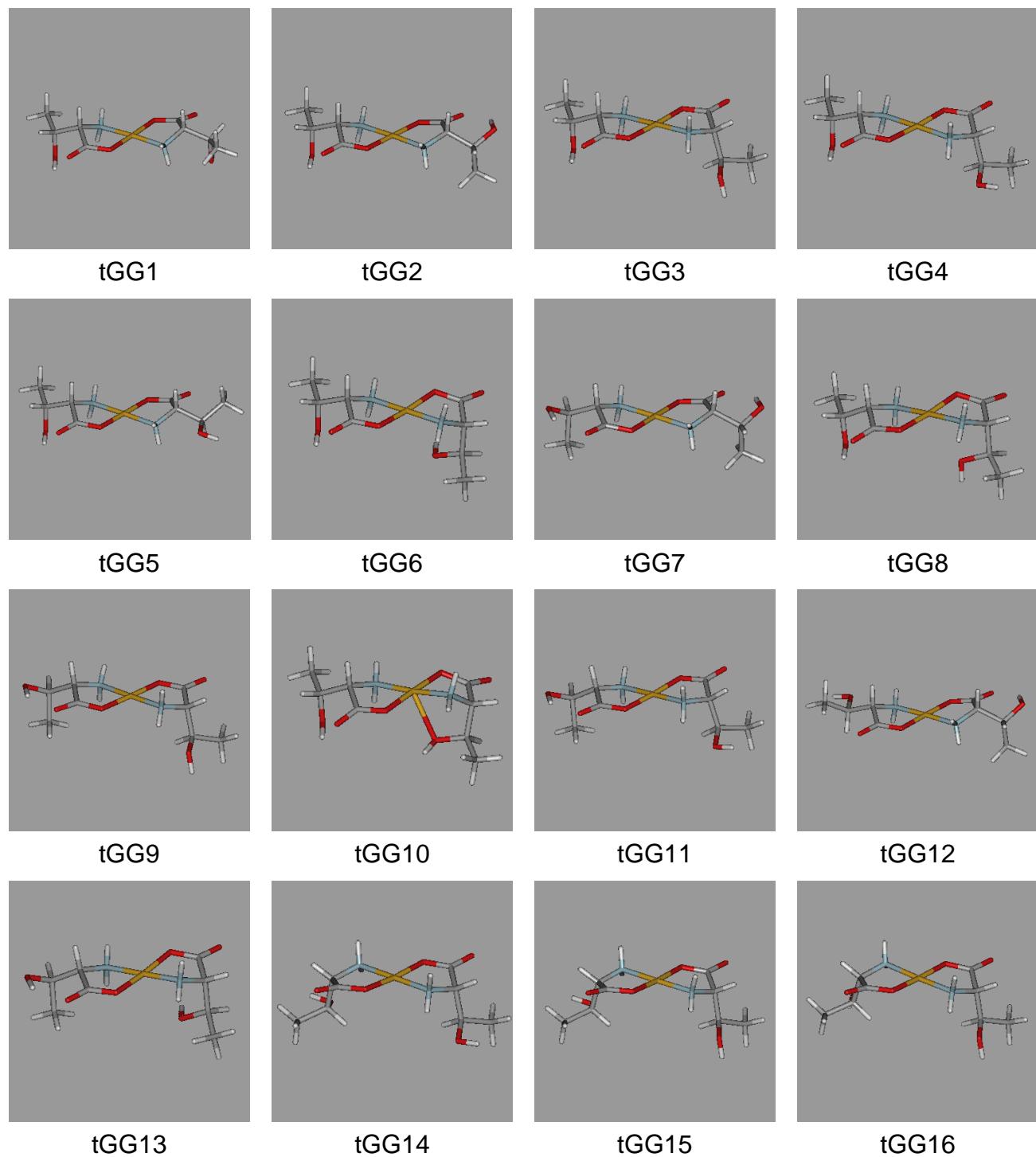


Figure S1. Illustrations and names of 196 Cu(L-Thr)₂ and 267 Cu(L-*a*Thr)₂ conformers computed using B3LYP/BS0 in the gas phase.

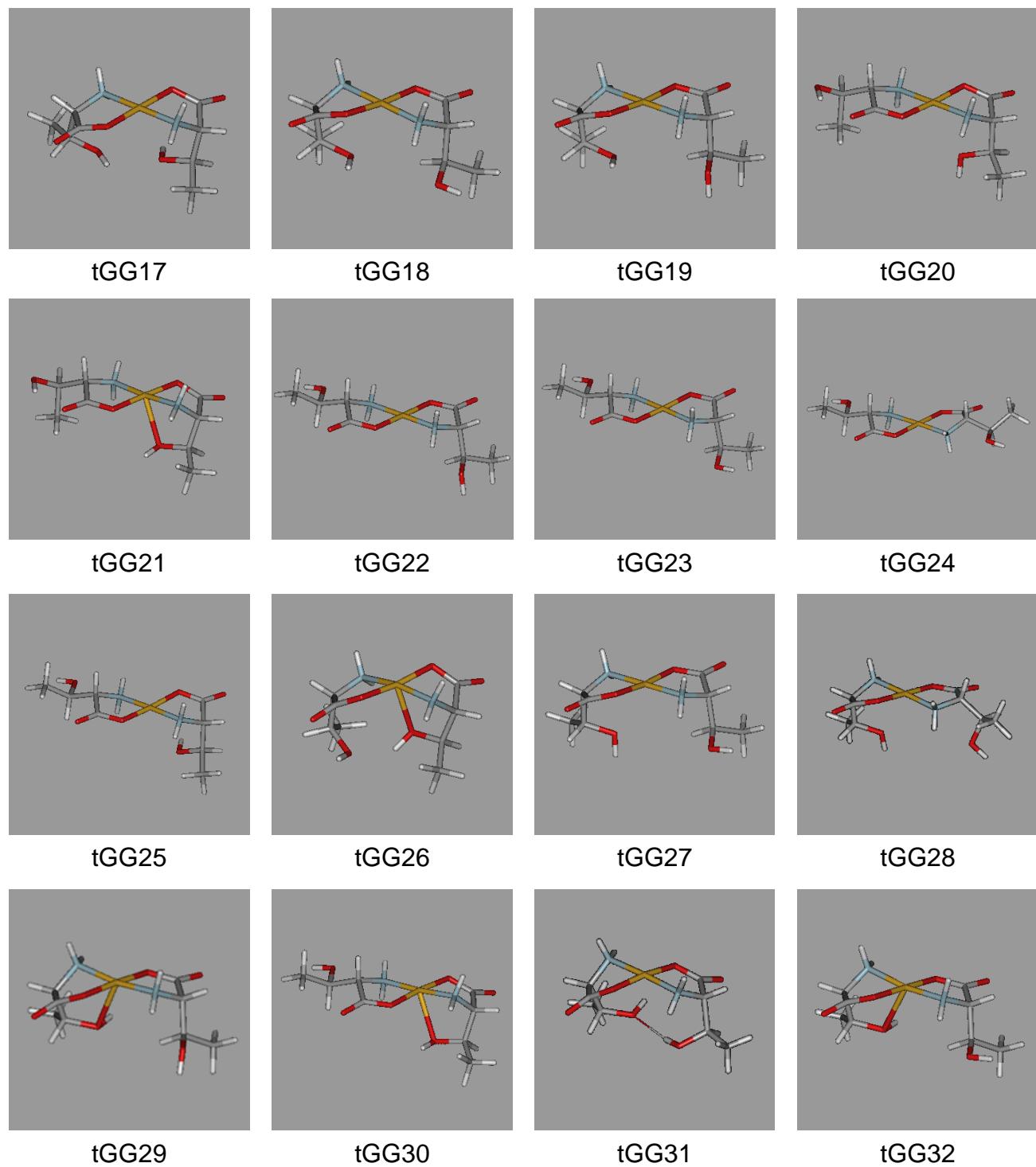


Figure S1. continued

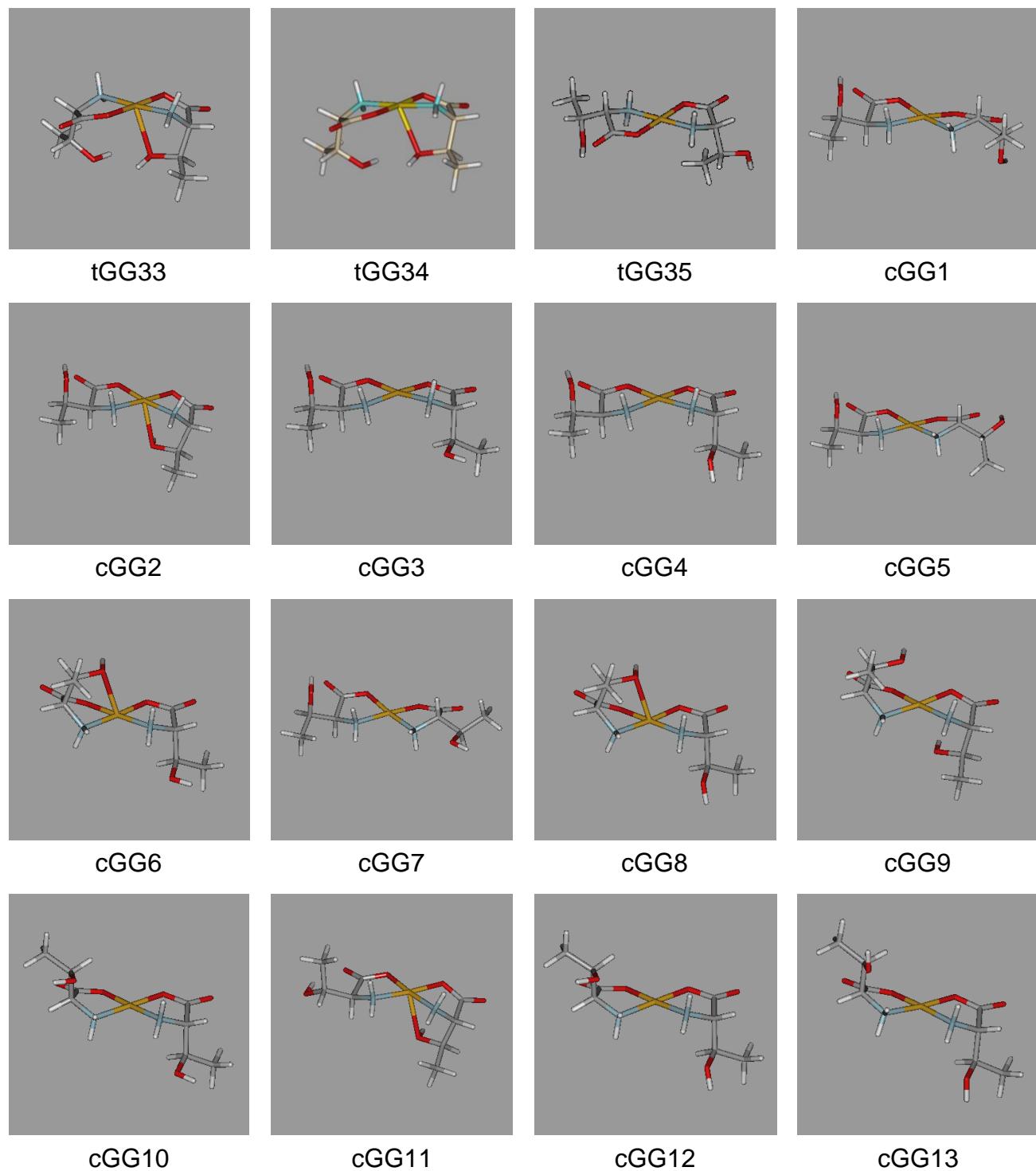


Figure S1. continued

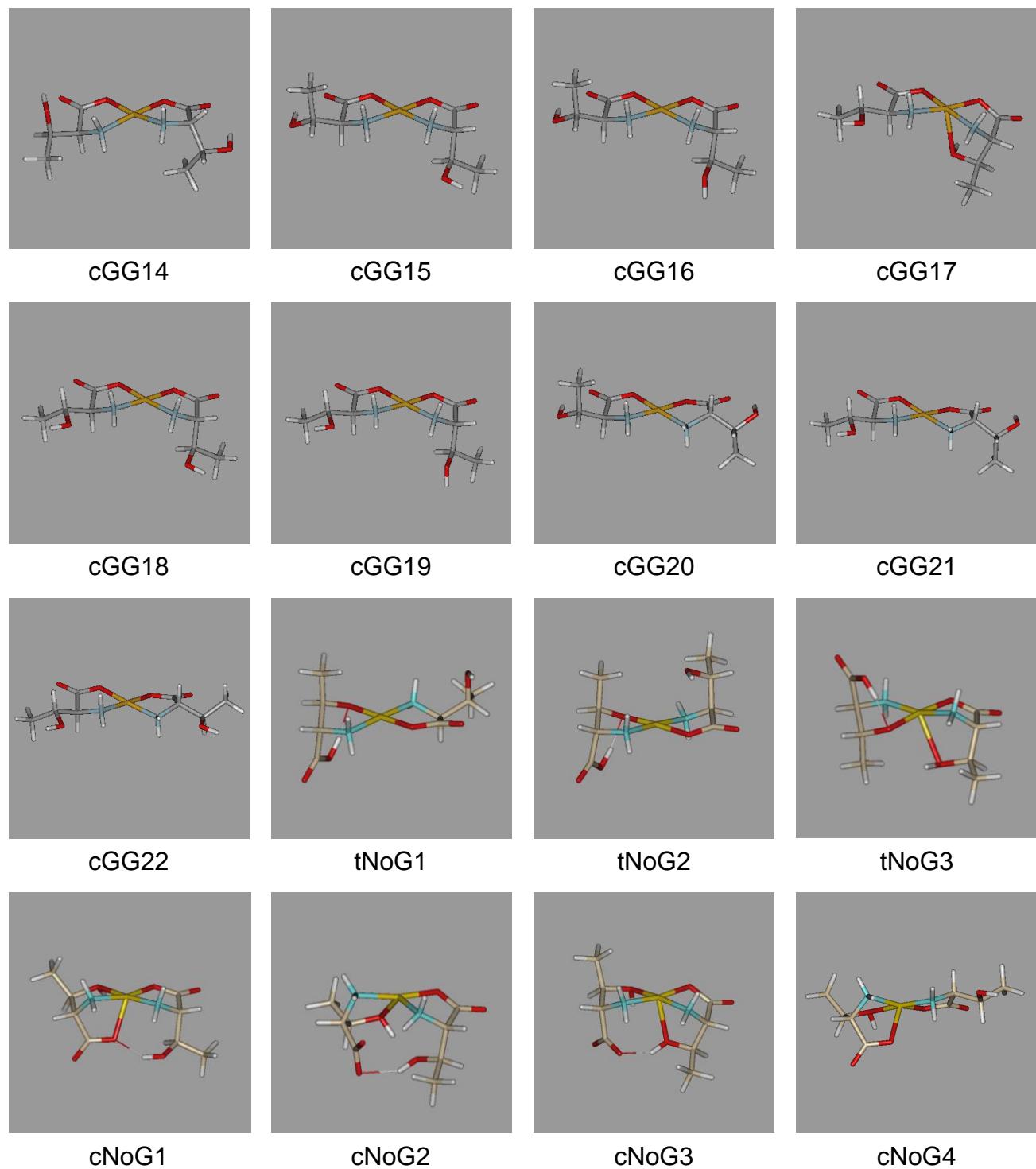


Figure S1. continued

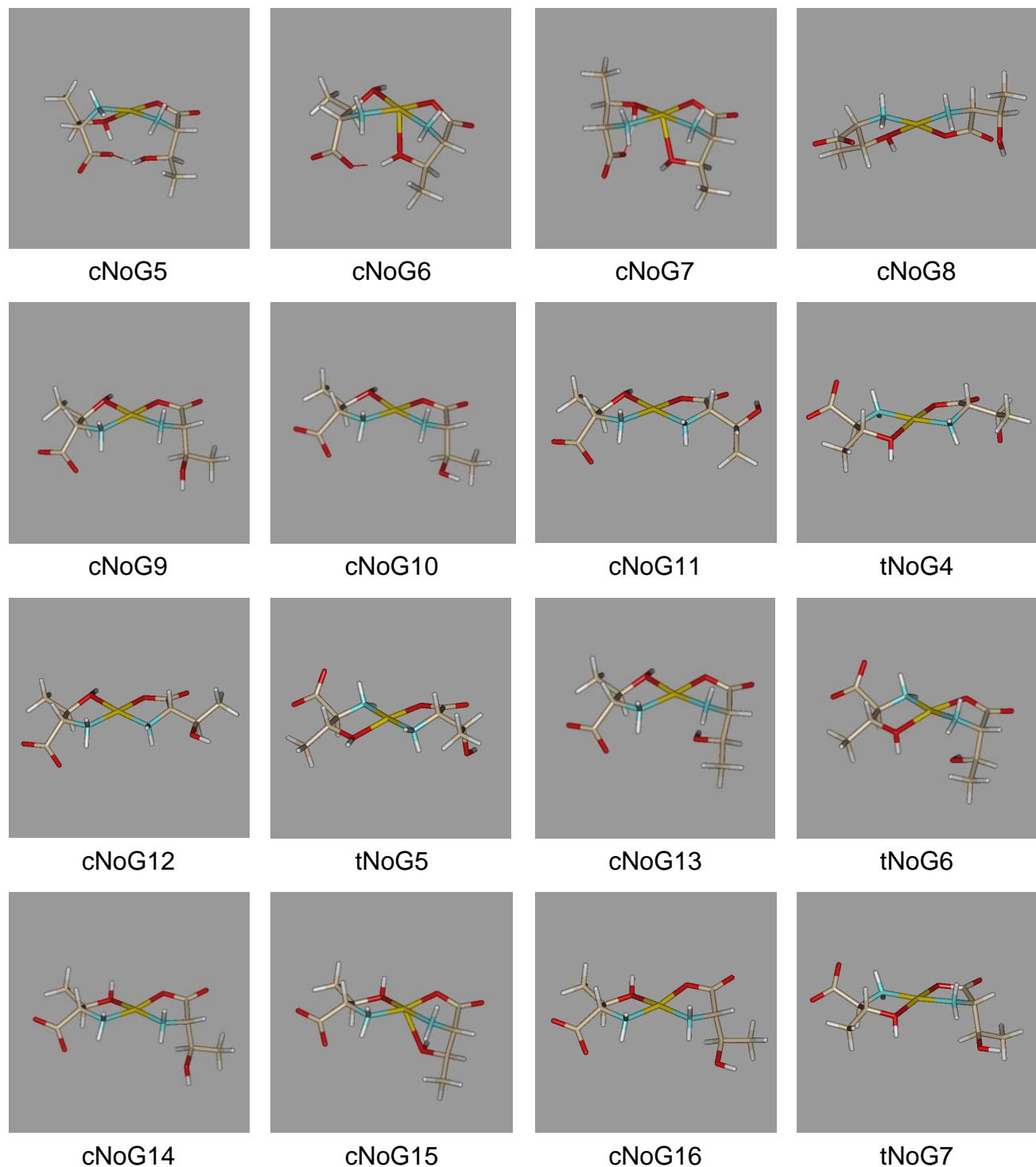


Figure S1. continued

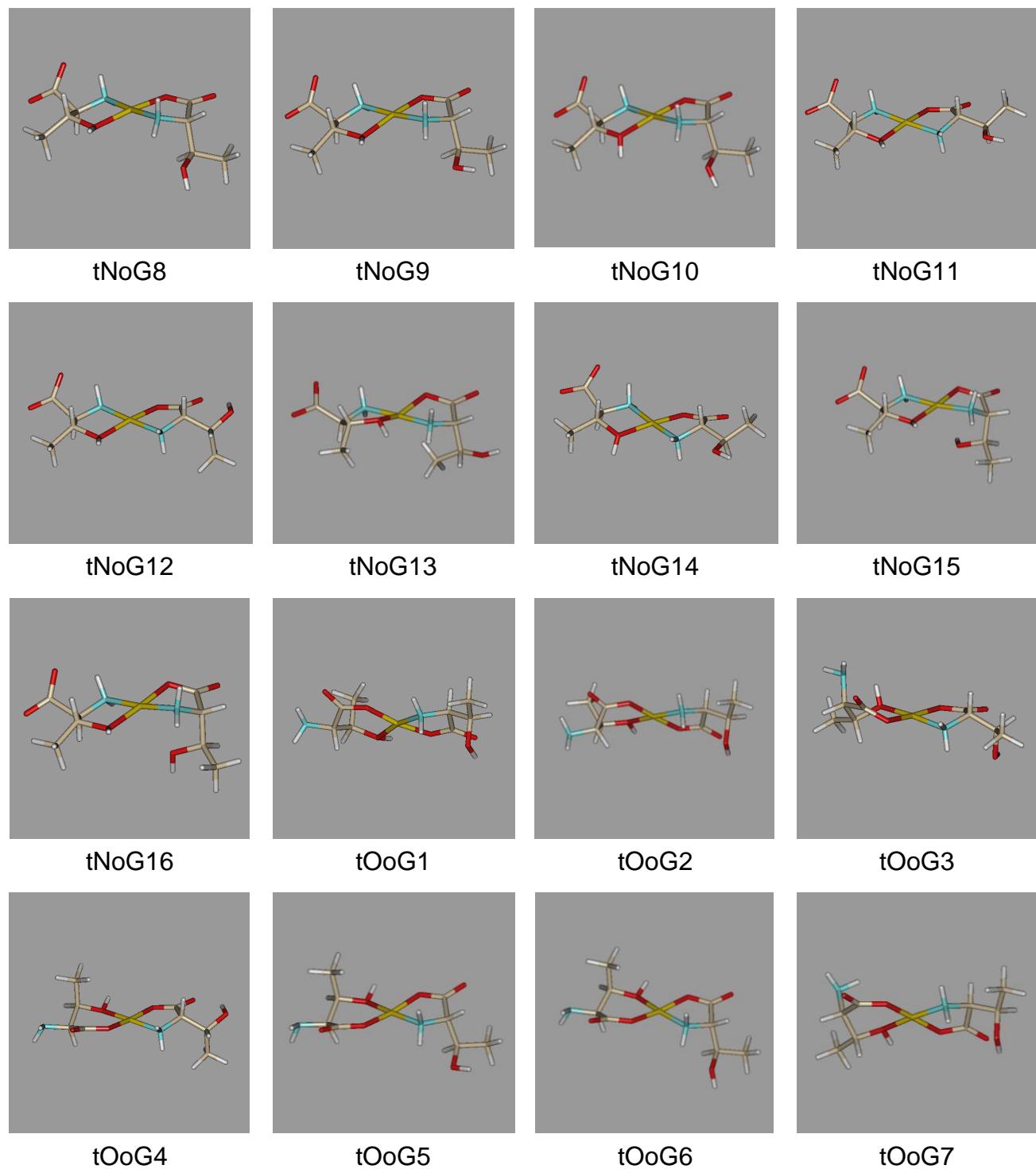


Figure S1. continued

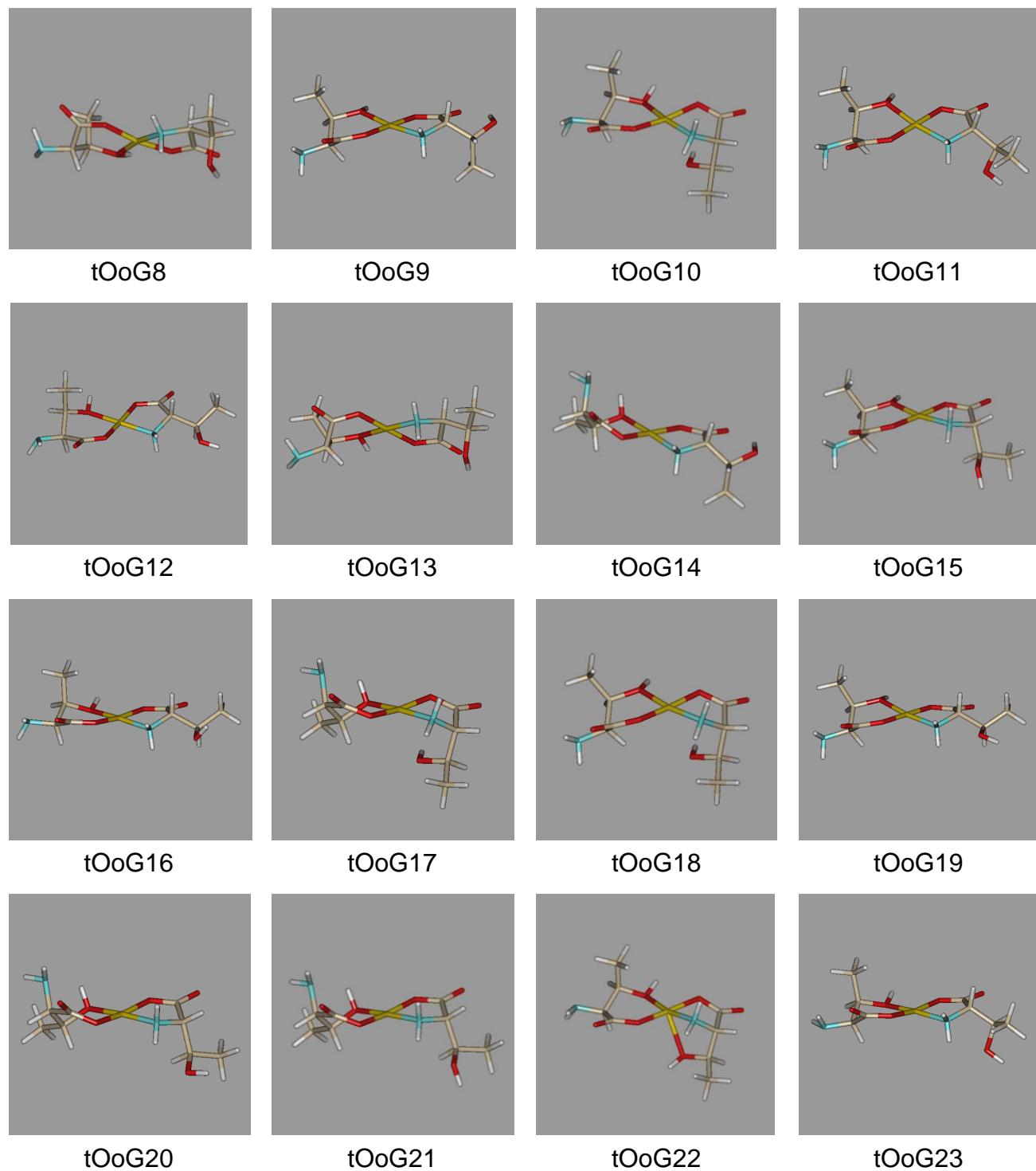


Figure S1. continued

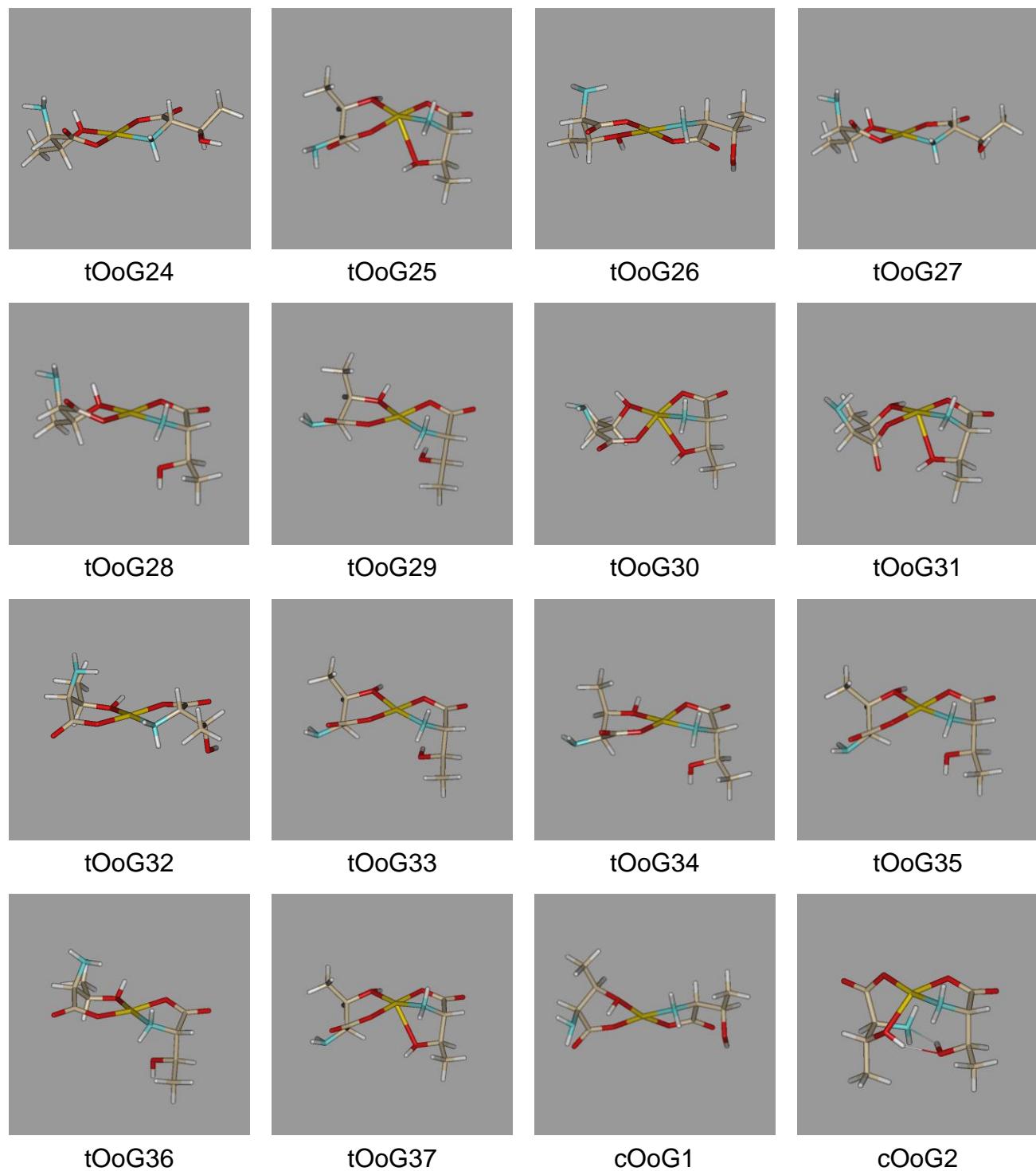


Figure S1. continued

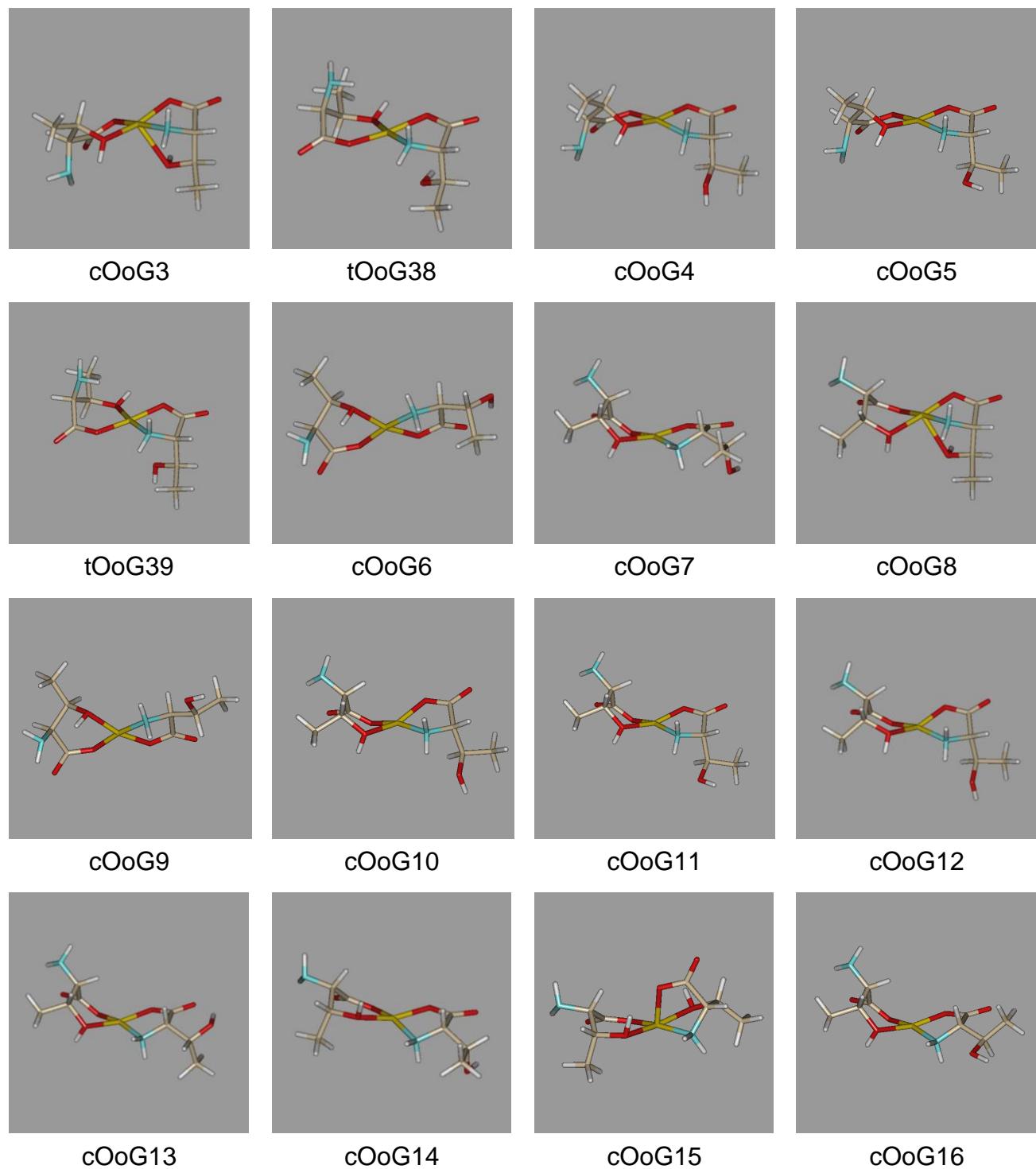


Figure S1. continued

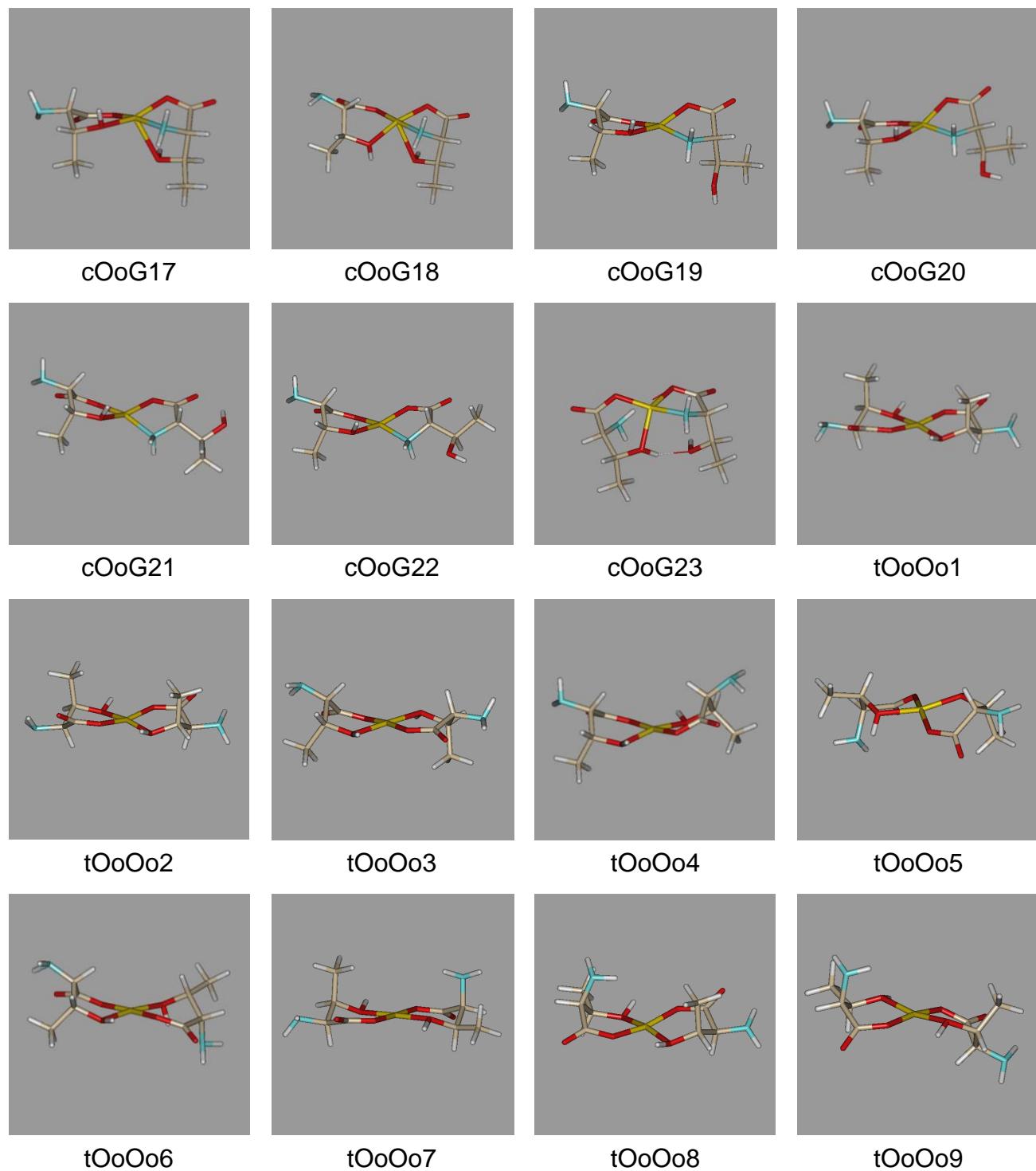


Figure S1. continued

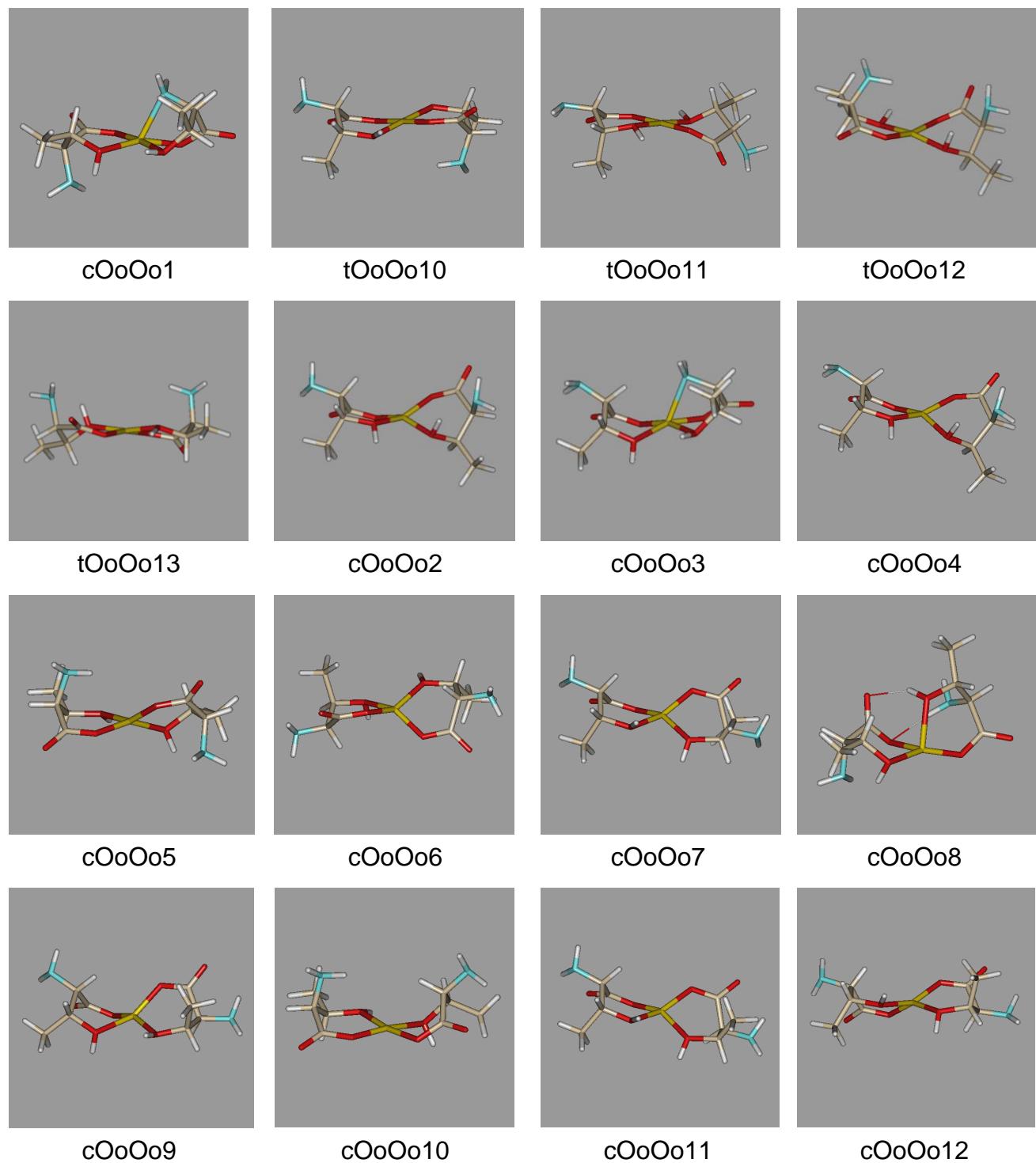


Figure S1. continued

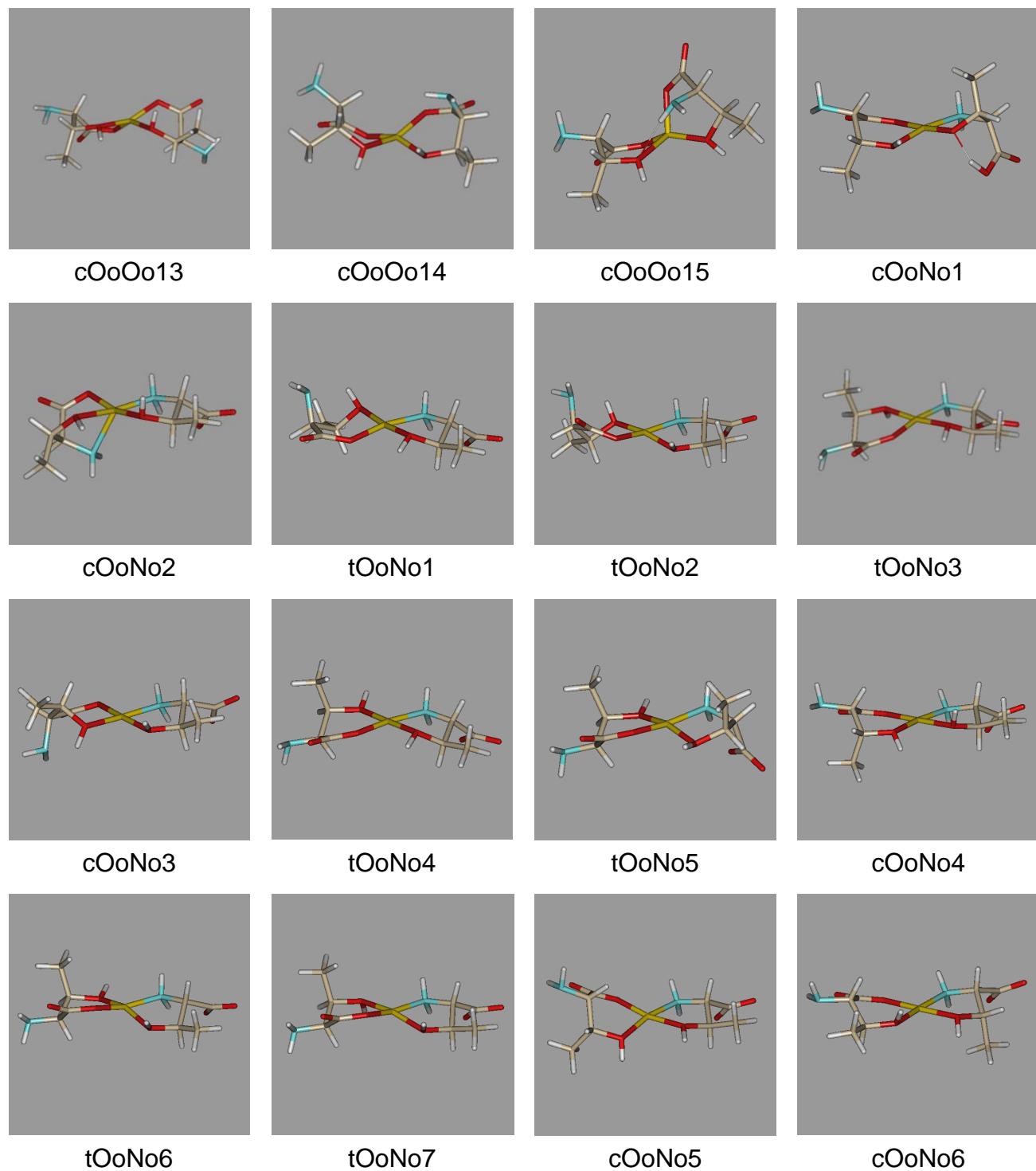


Figure S1. continued

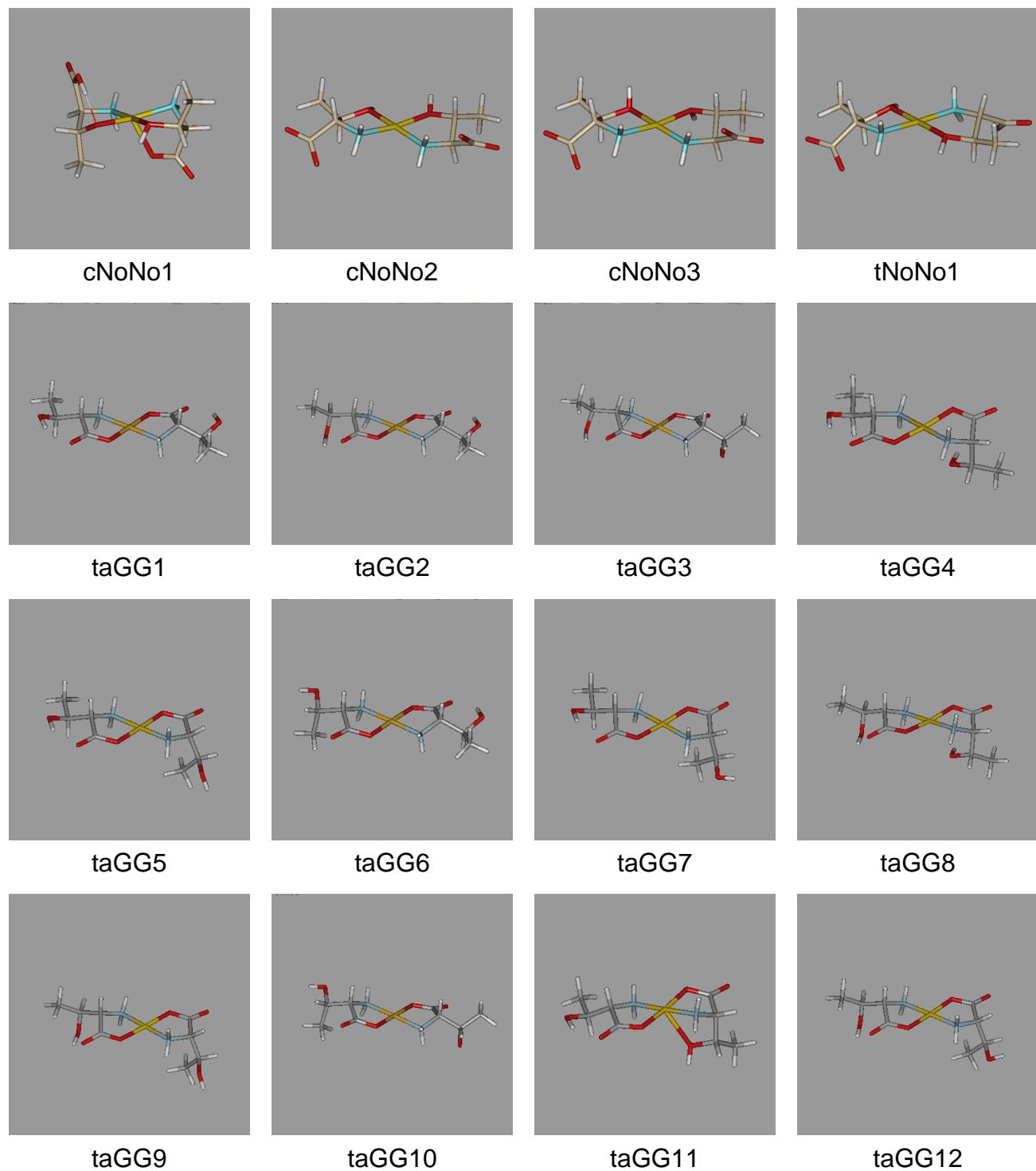


Figure S1. continued

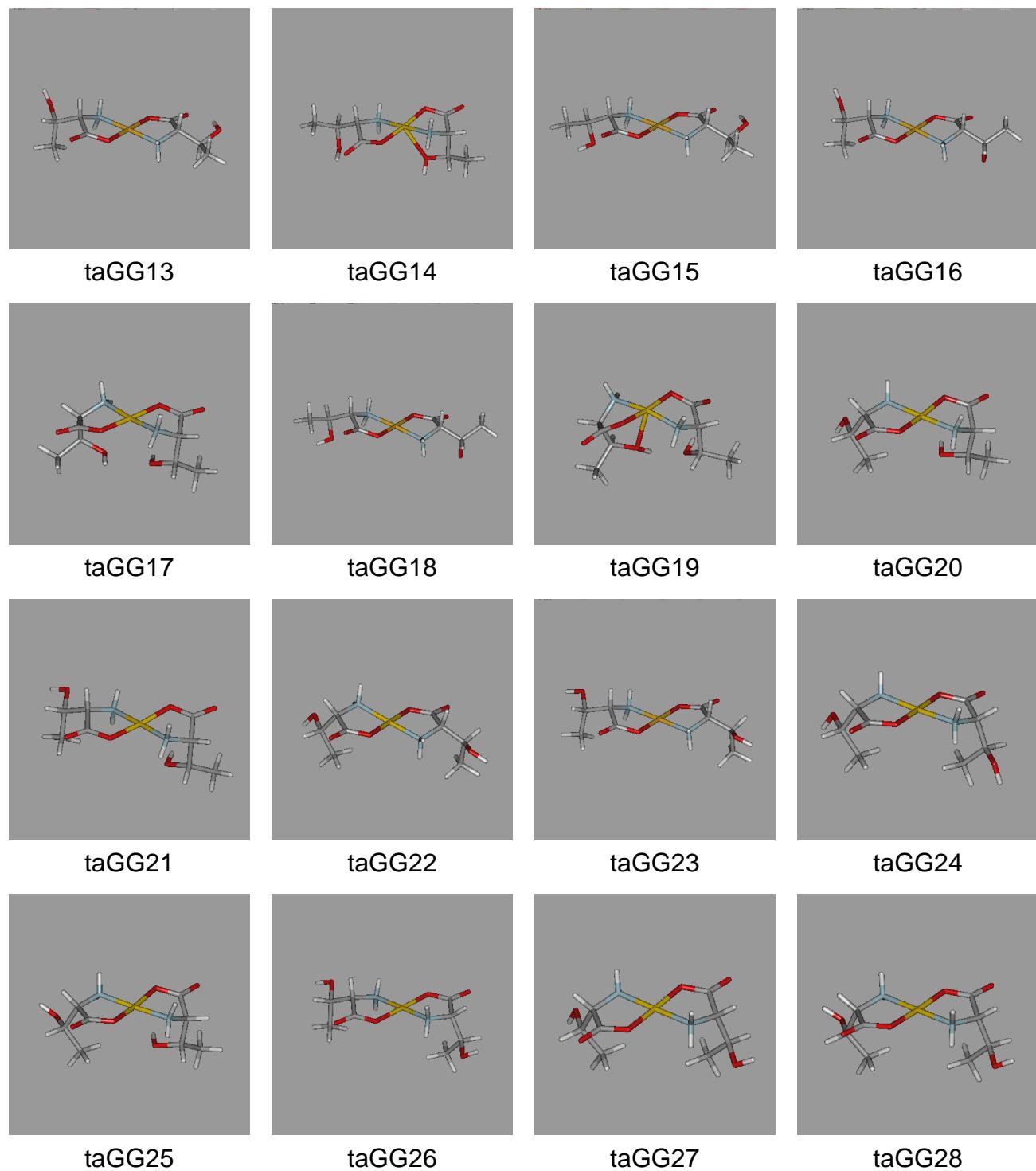


Figure S1. continued

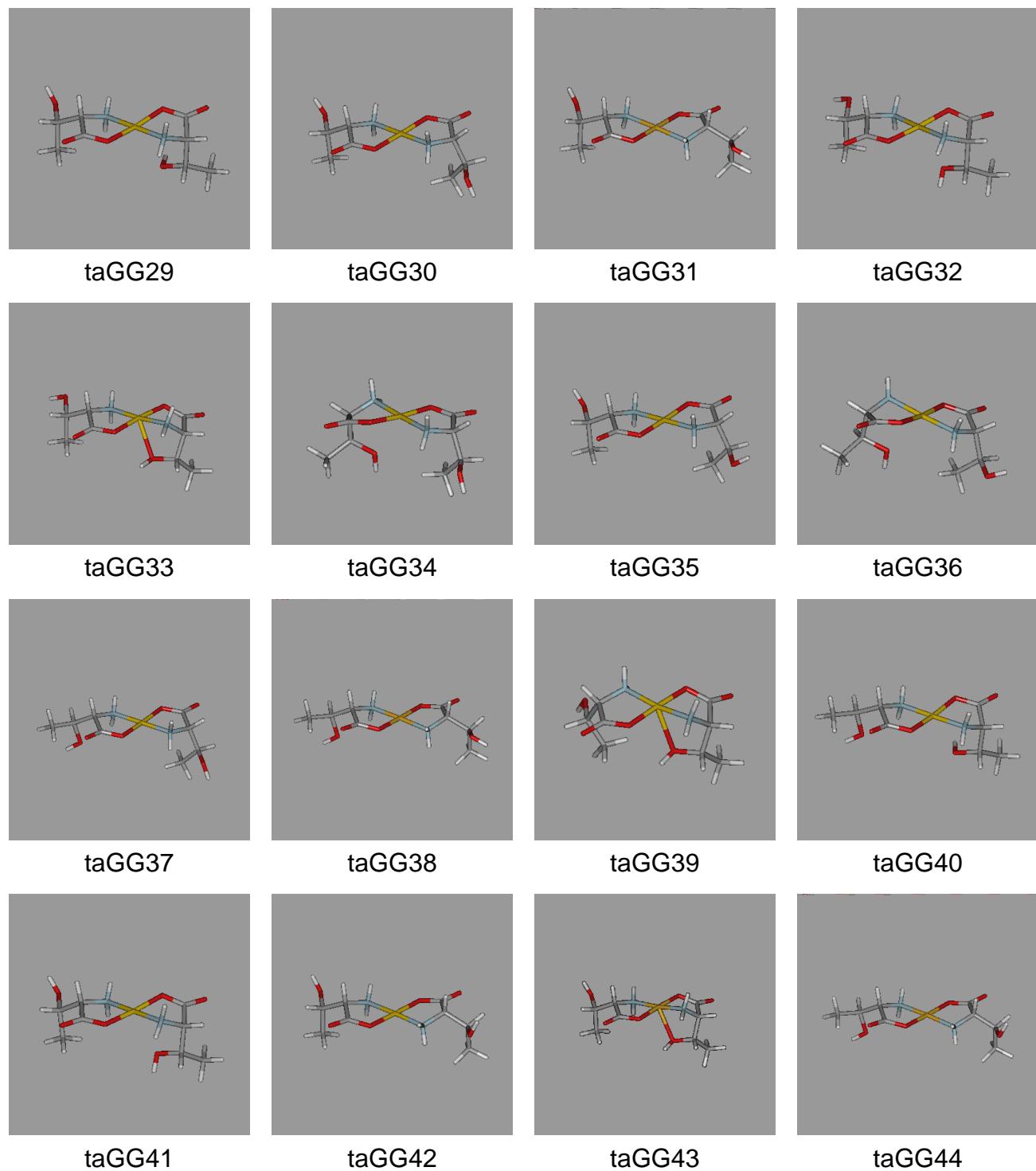


Figure S1. continued

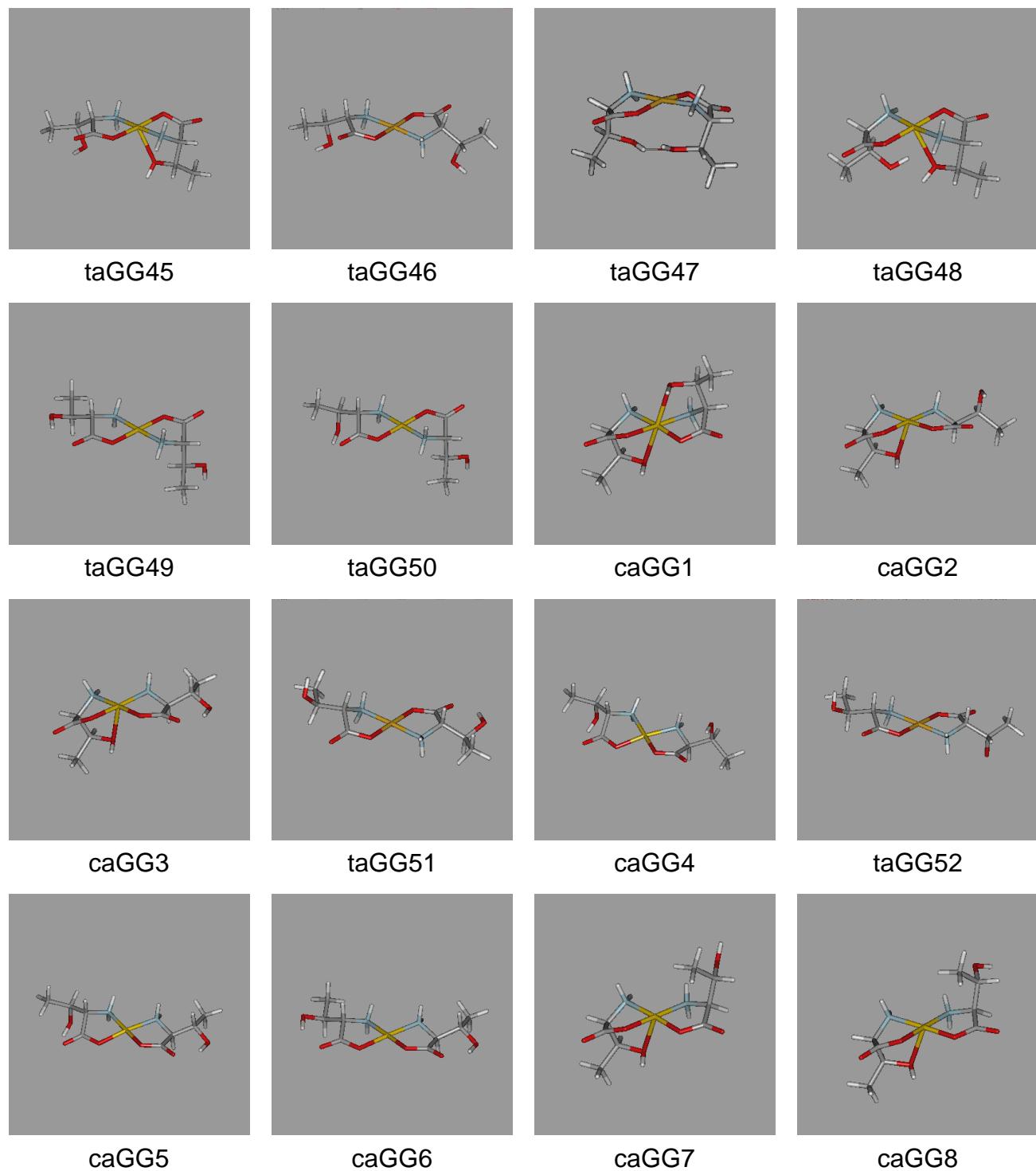


Figure S1. continued

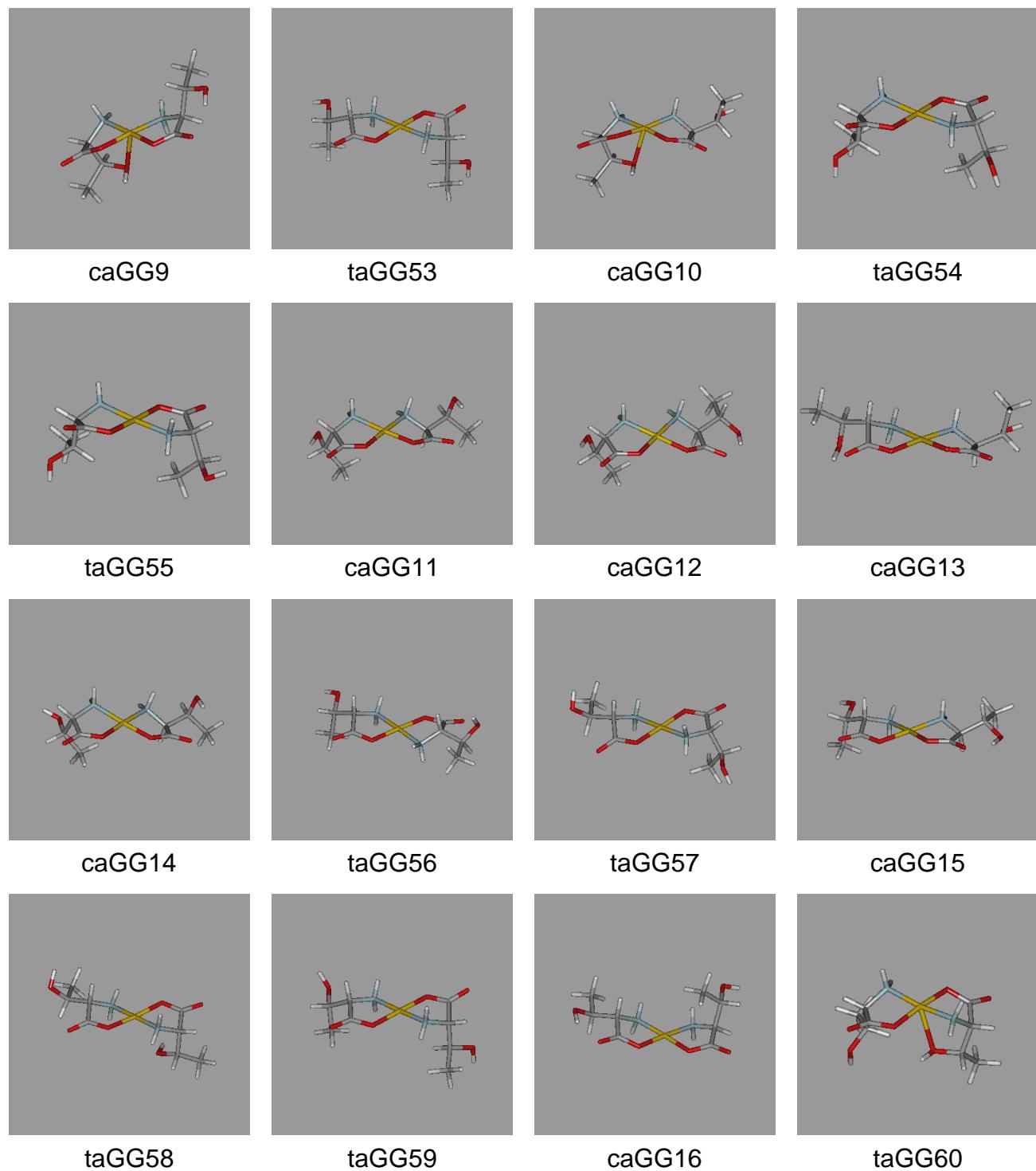


Figure S1. continued

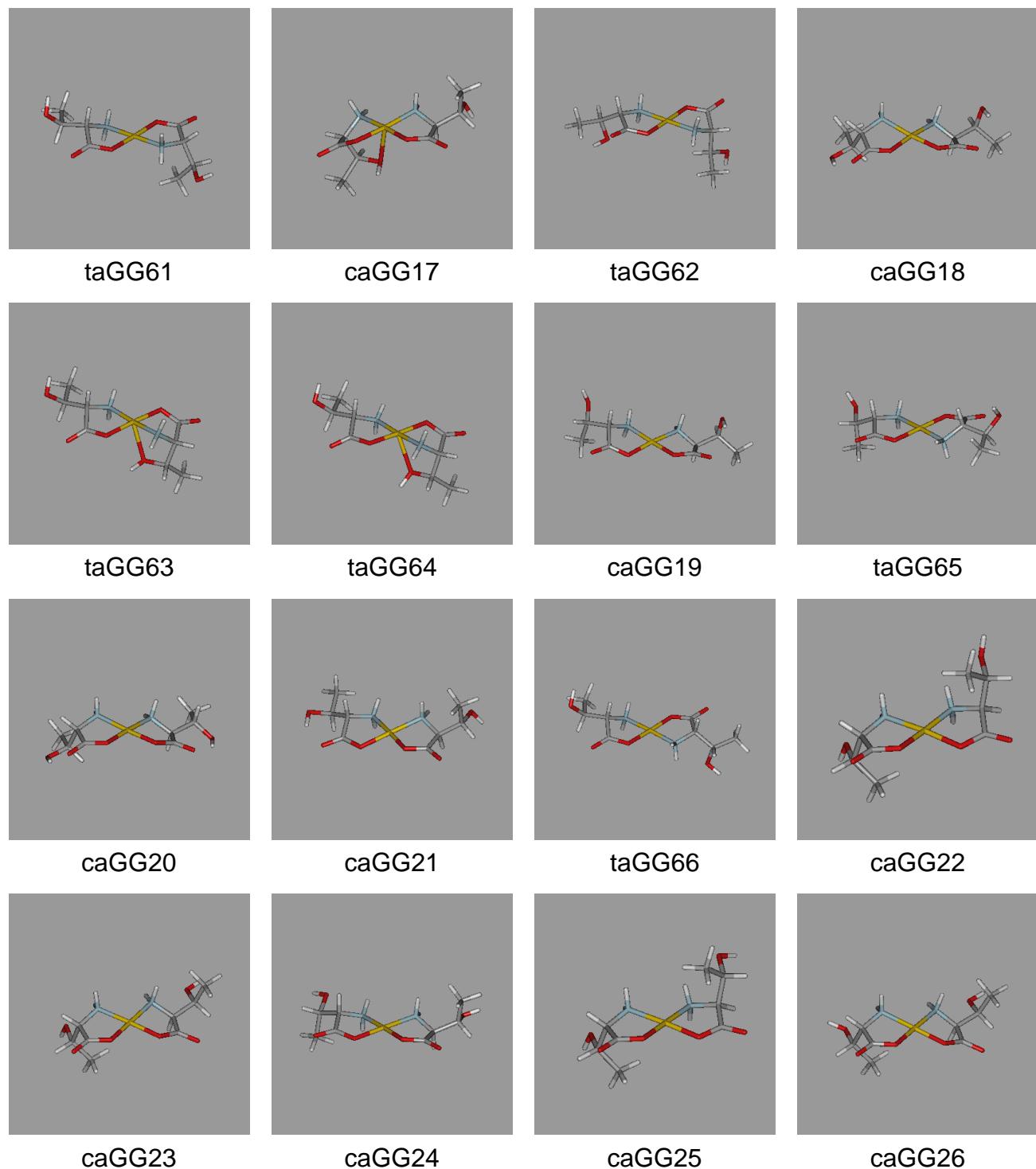


Figure S1. continued

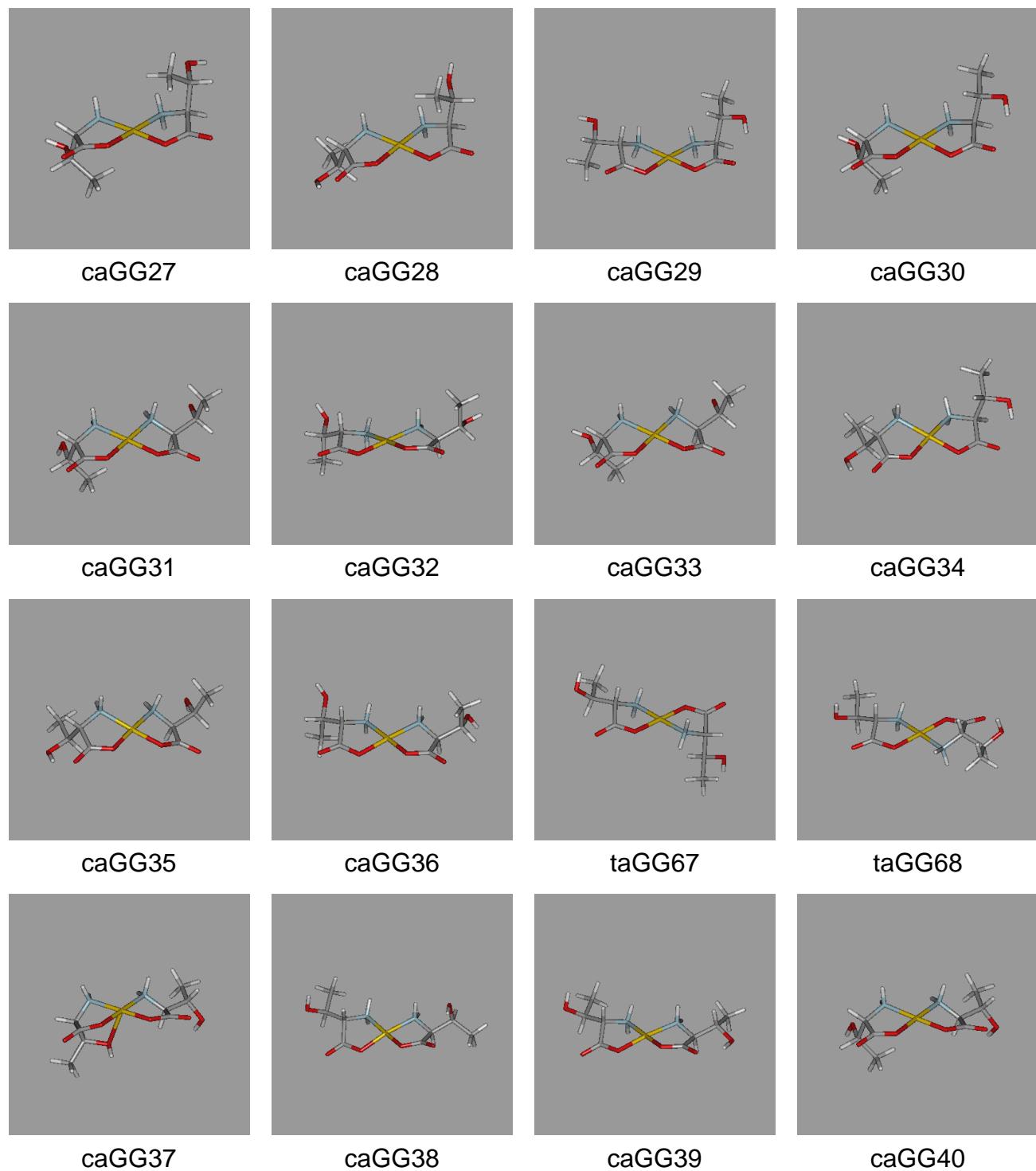


Figure S1. continued

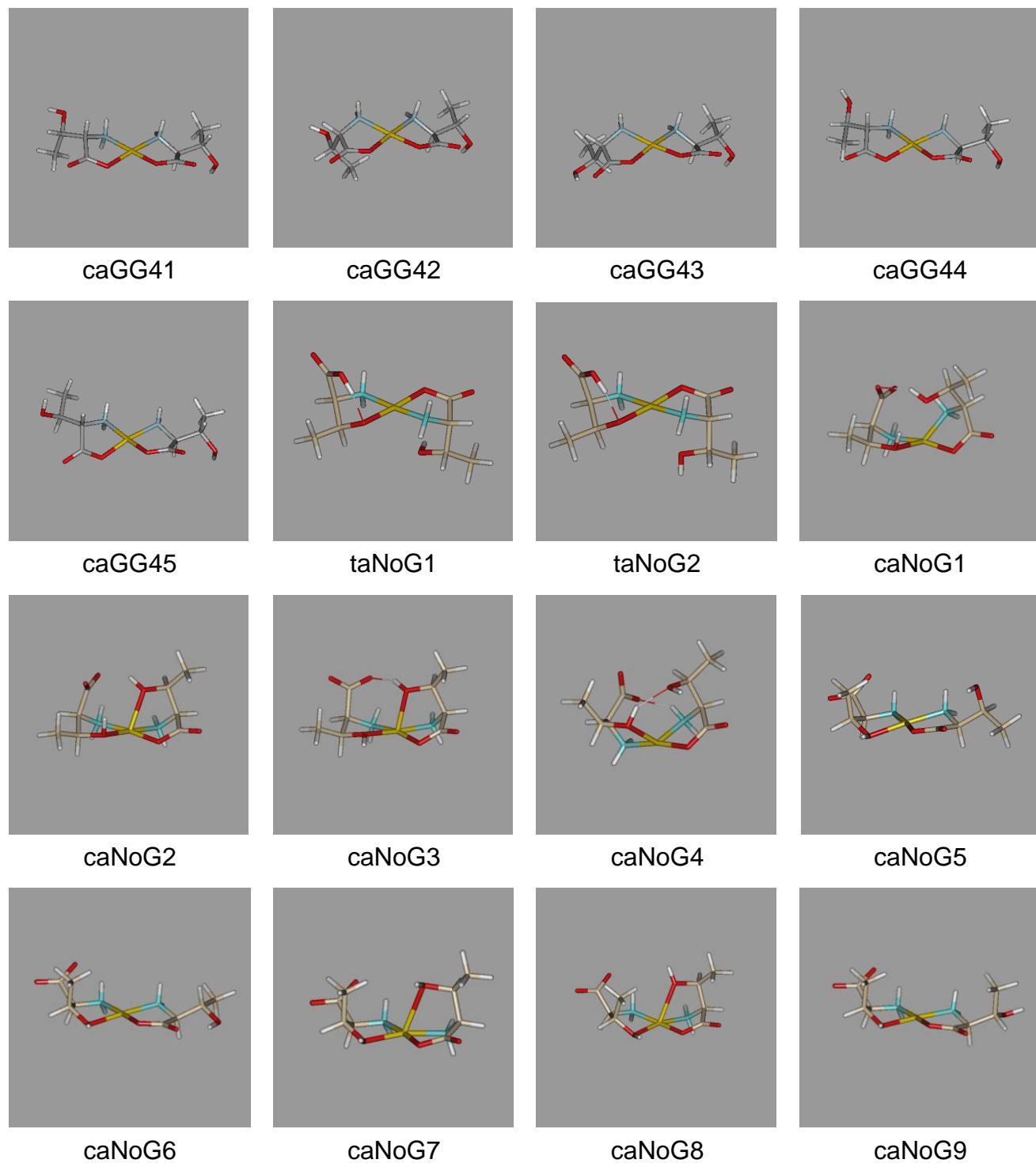


Figure S1. continued

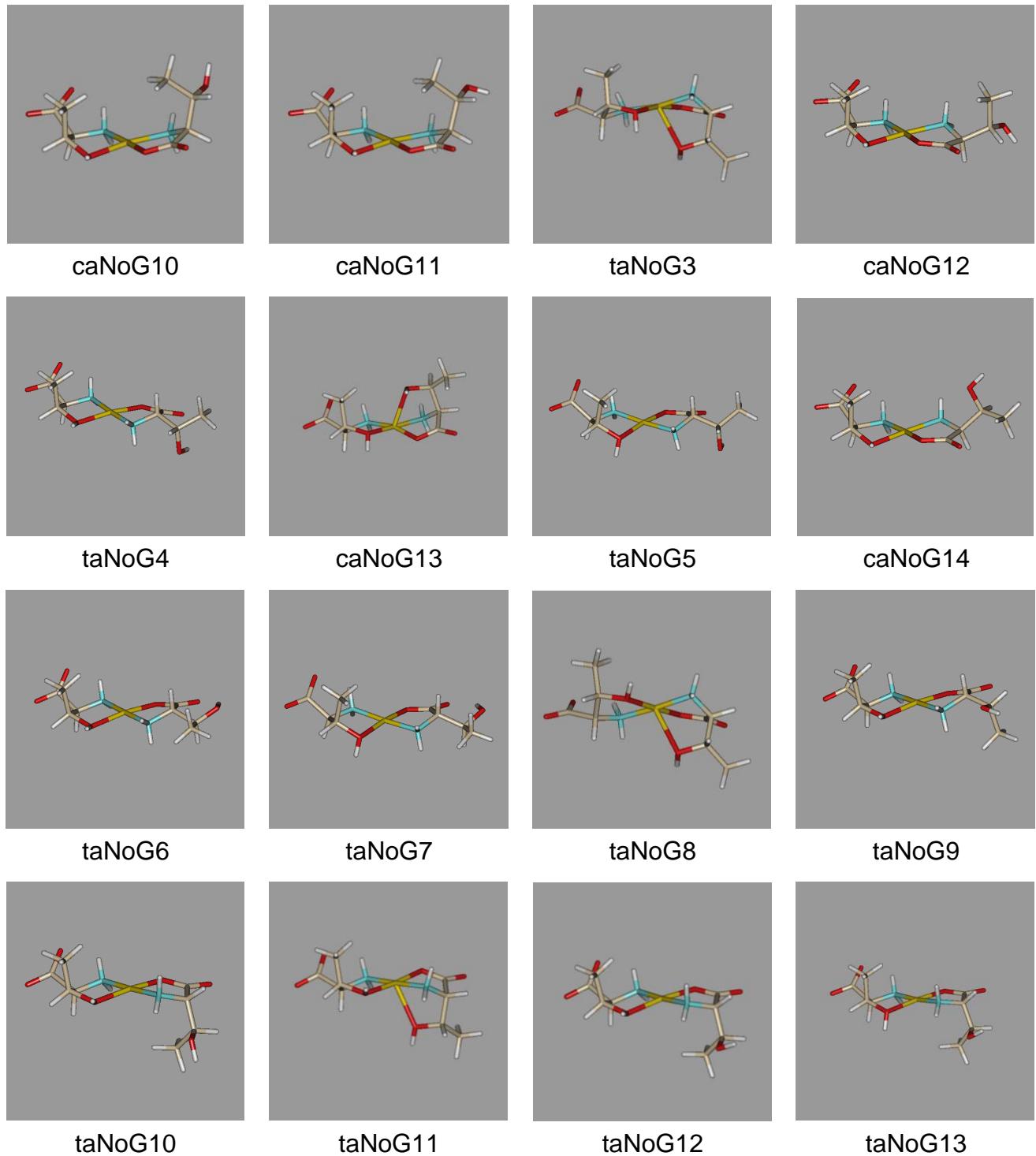


Figure S1. continued

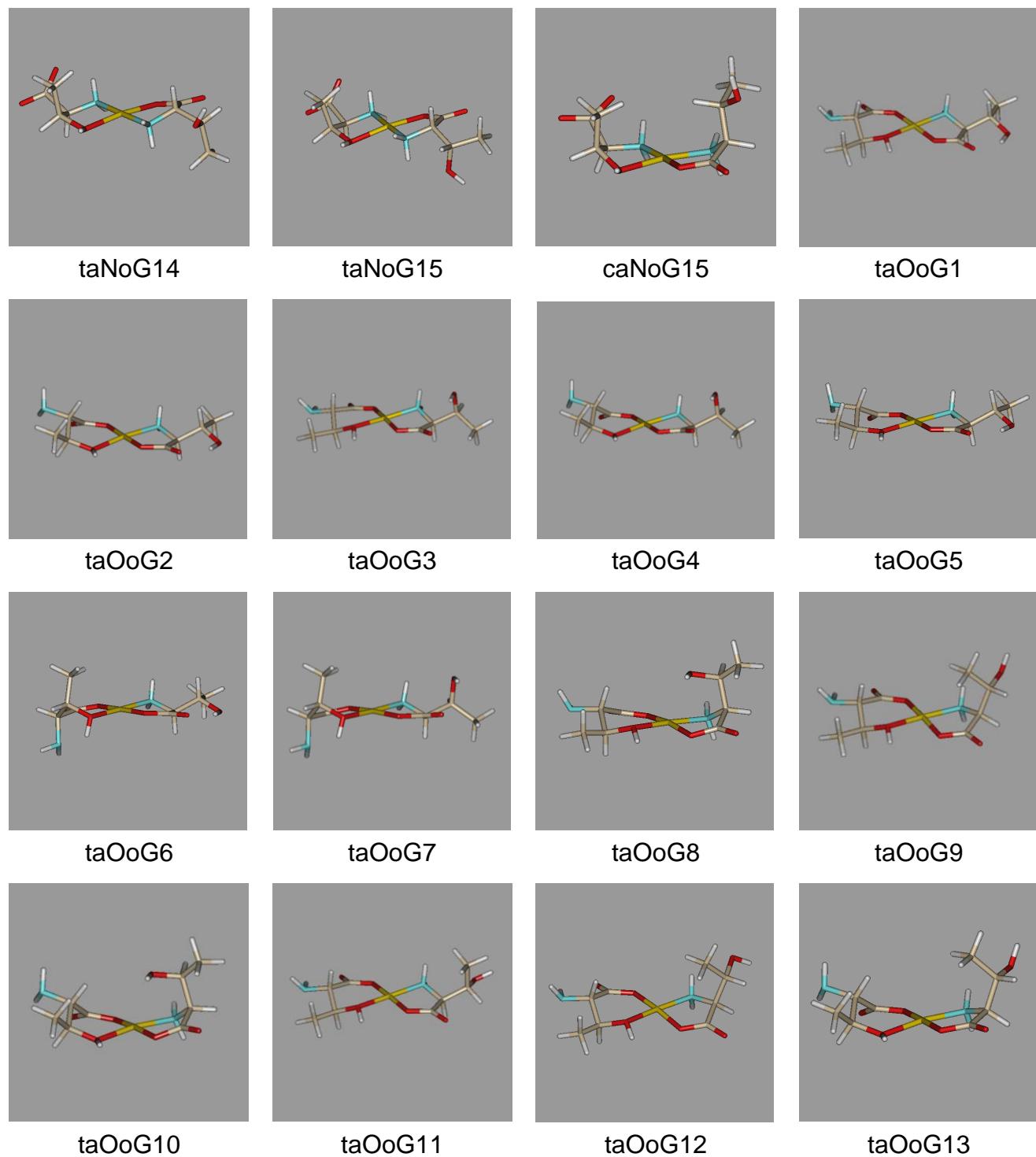


Figure S1. continued

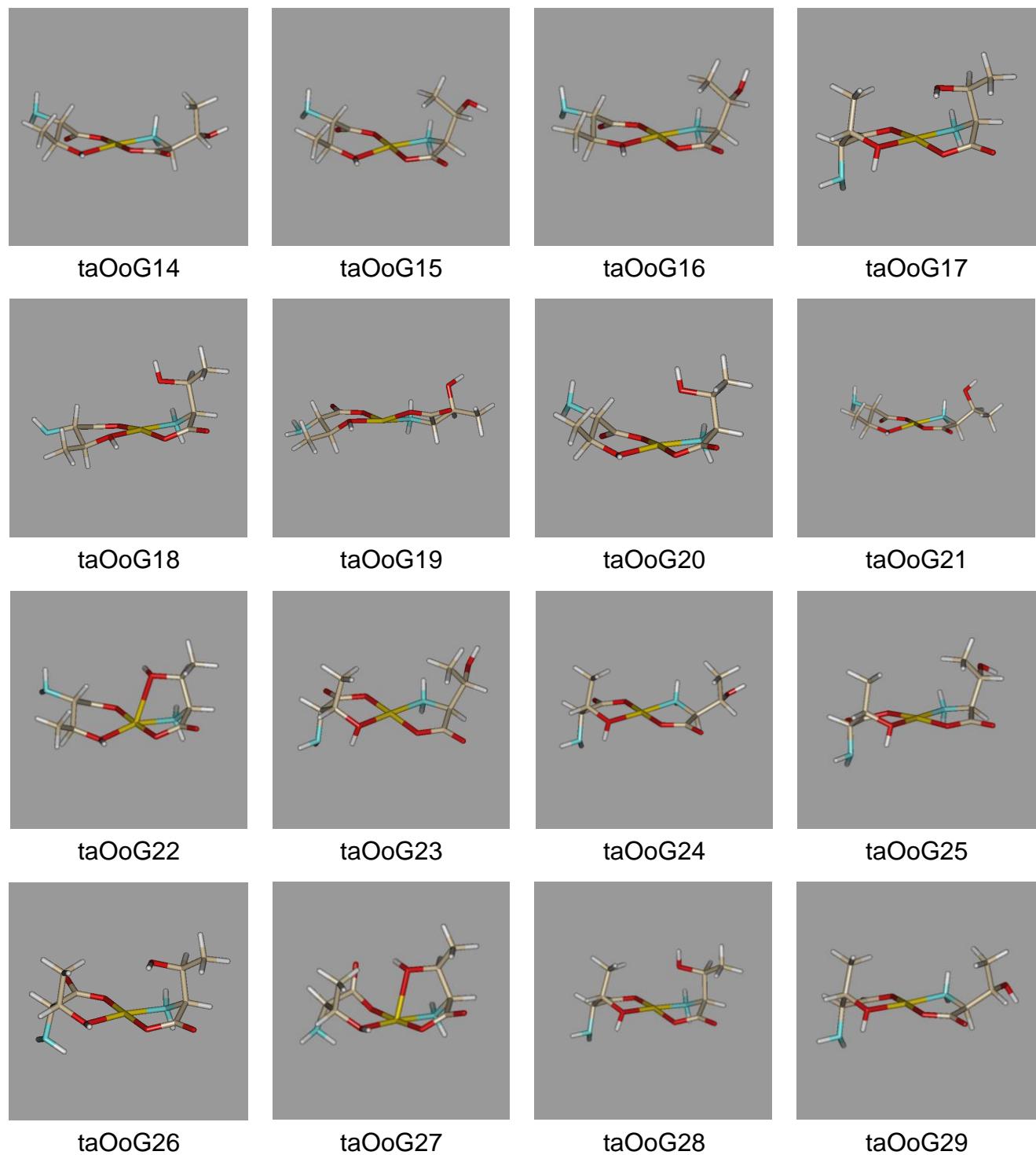


Figure S1. continued

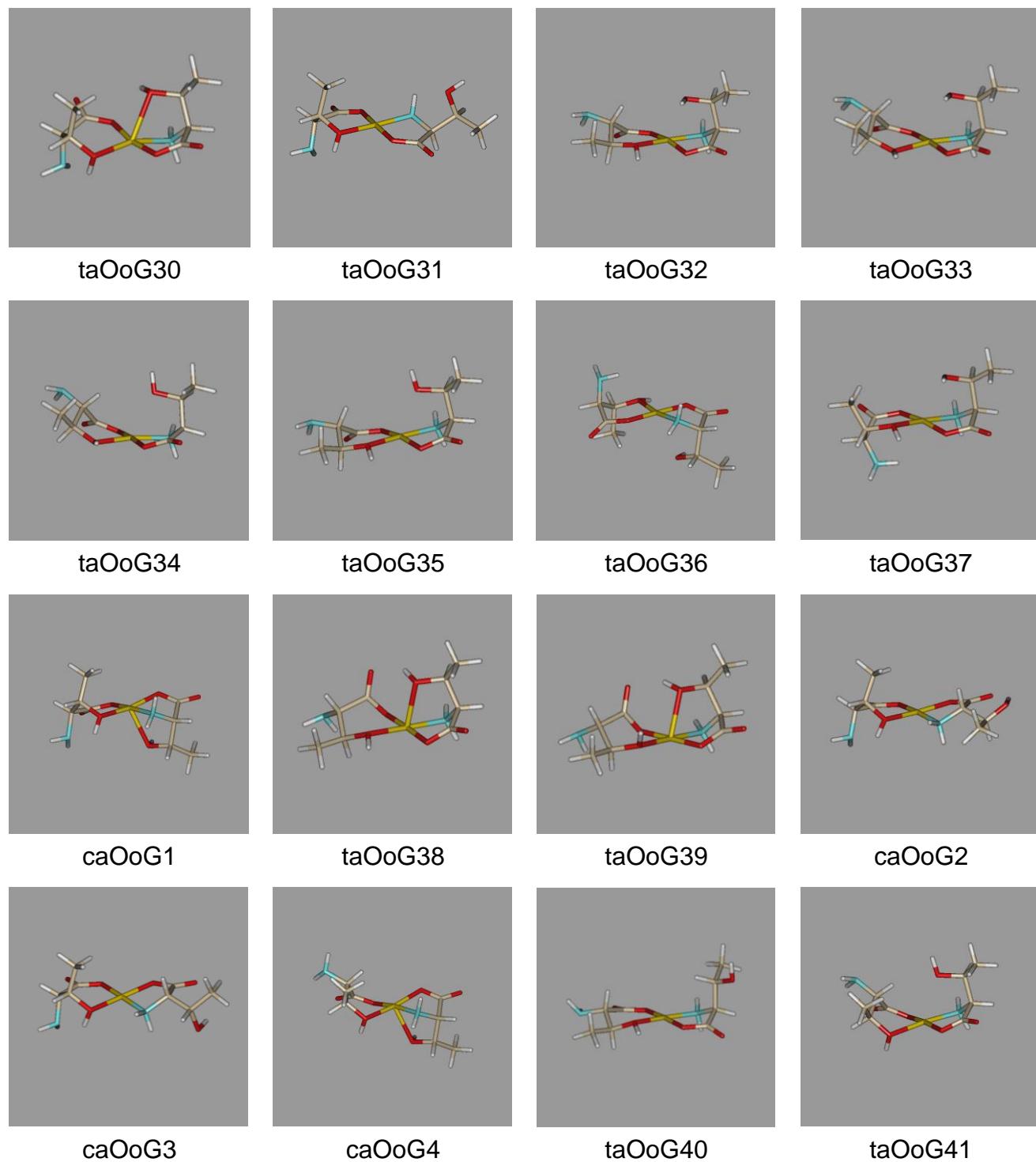


Figure S1. continued

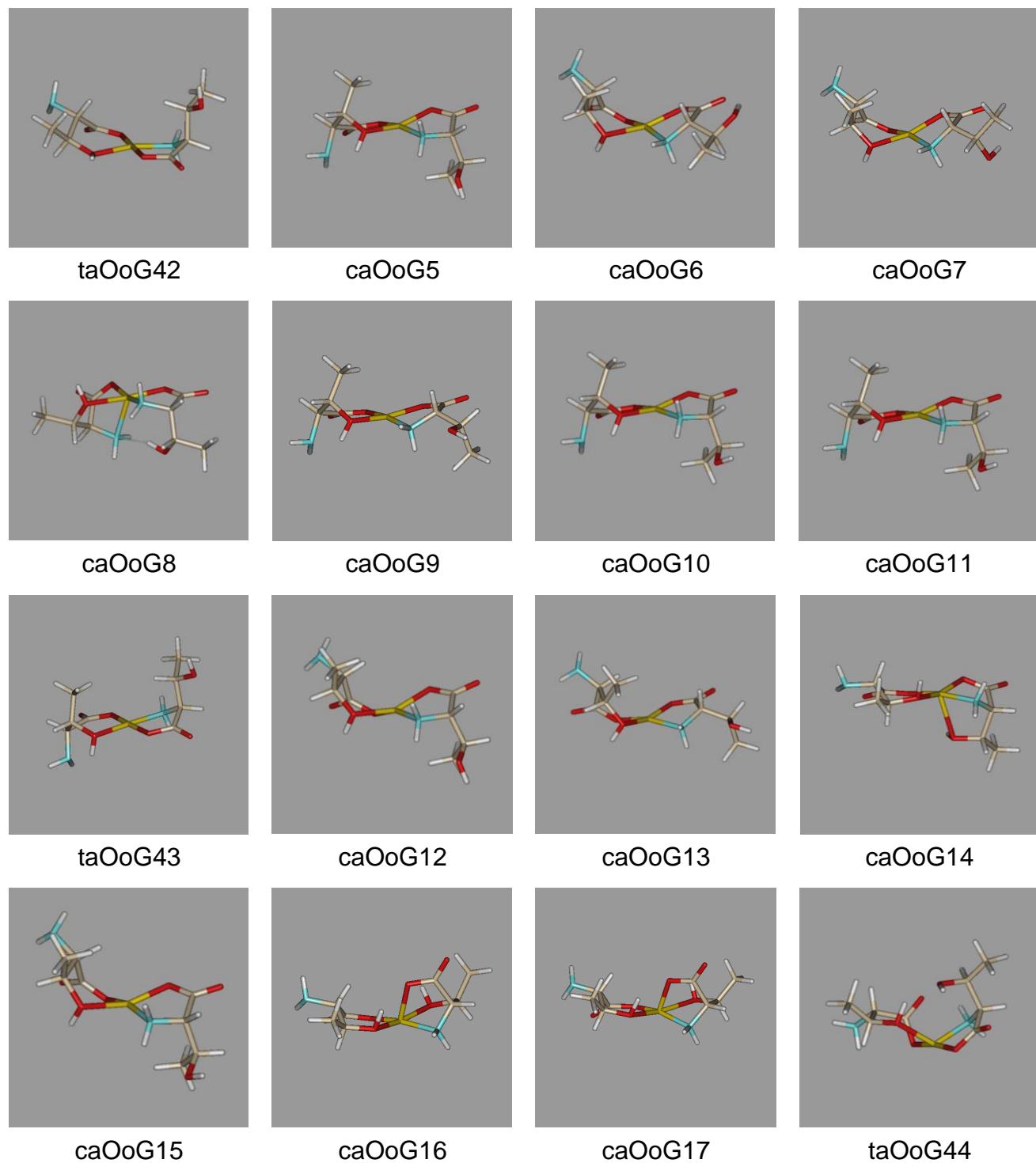


Figure S1. continued

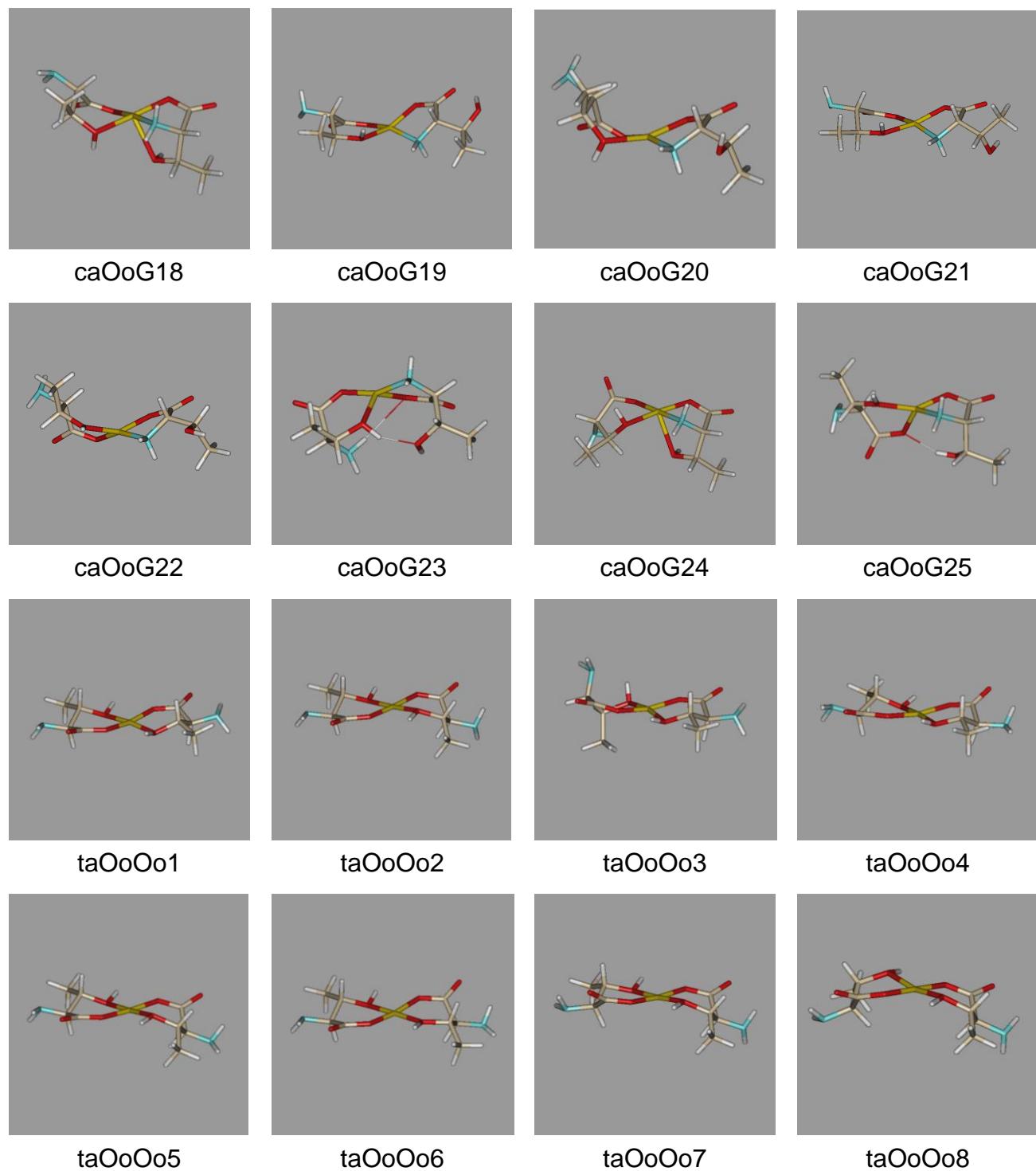


Figure S1. continued

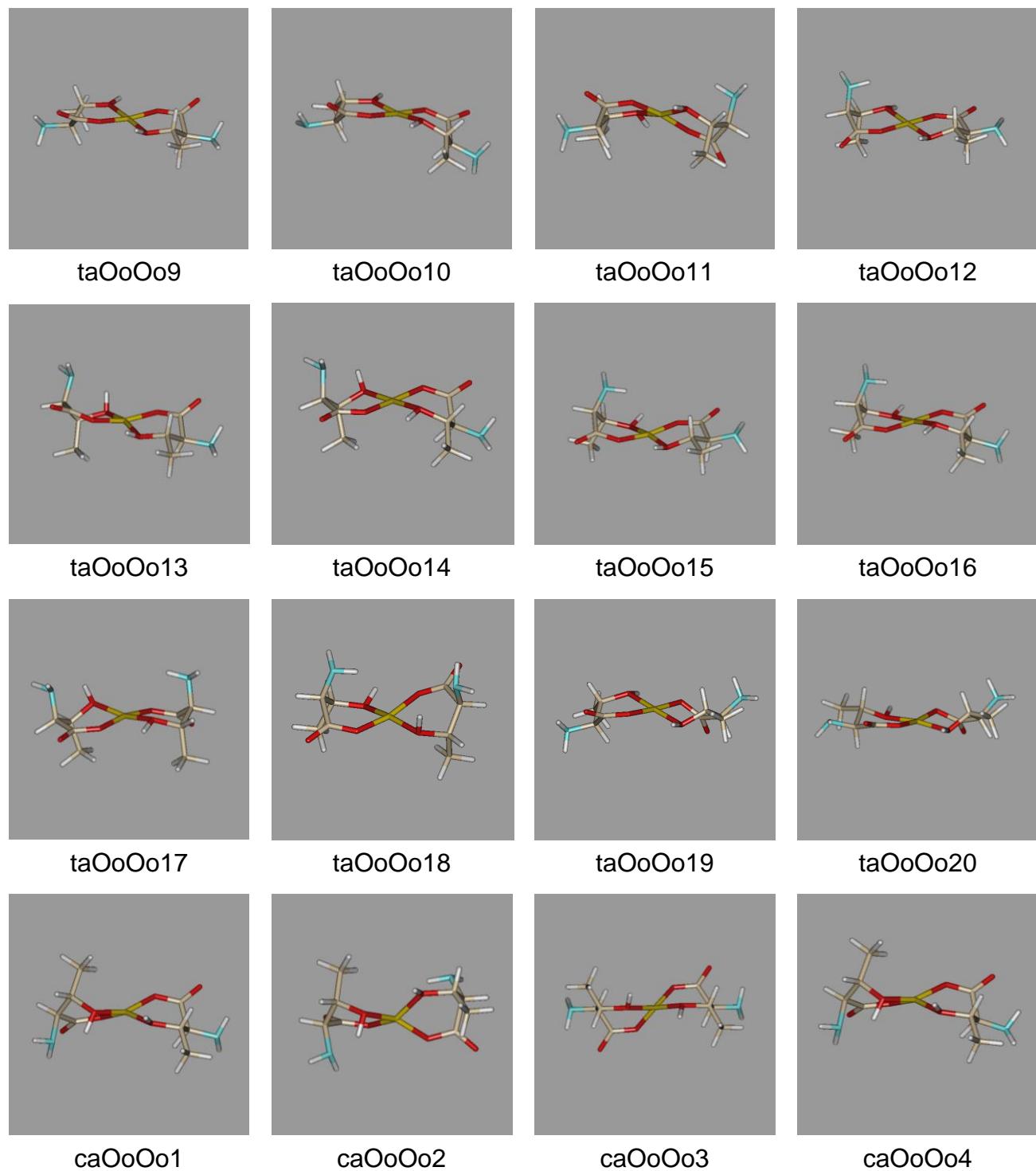


Figure S1. continued

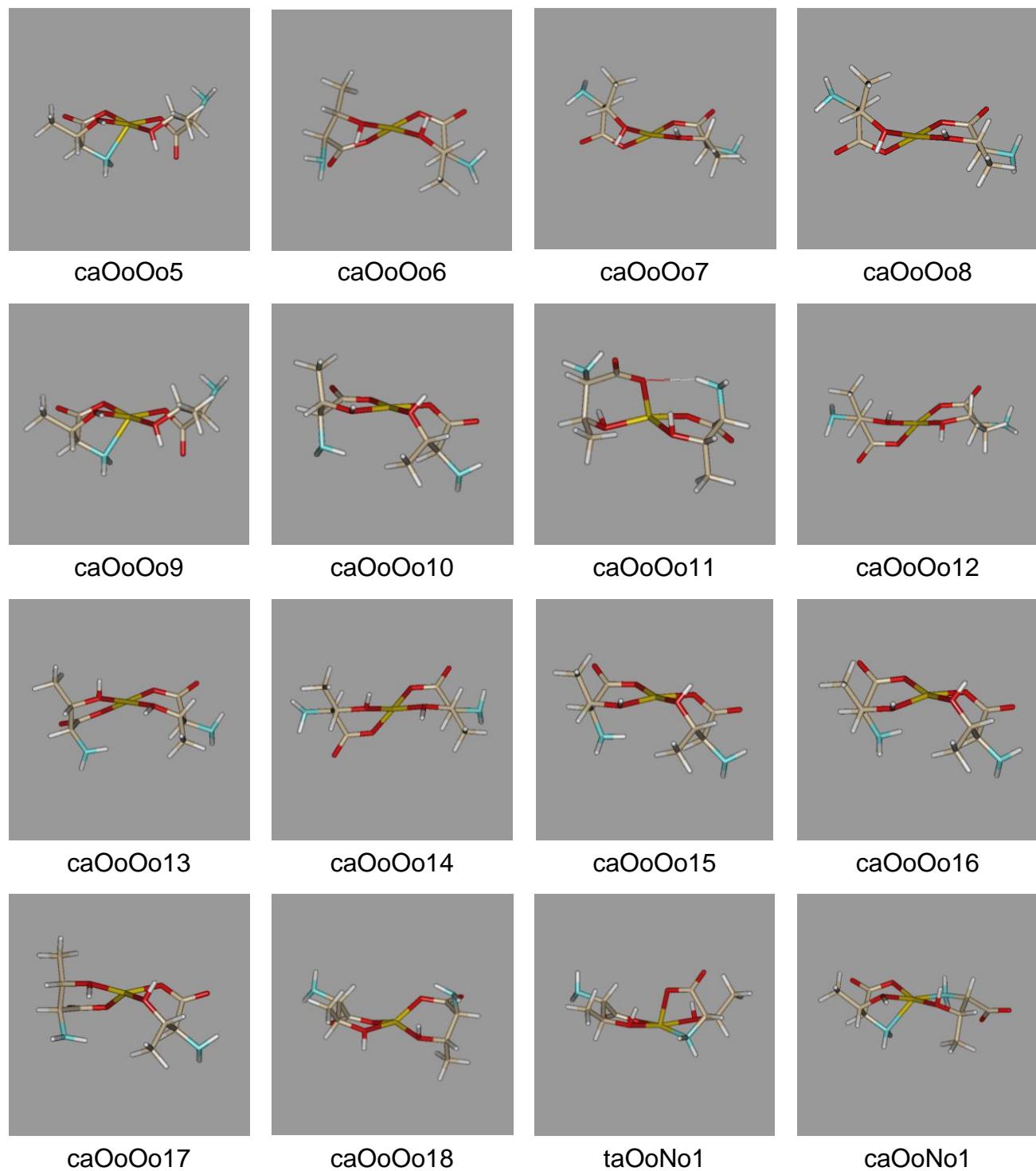


Figure S1. continued

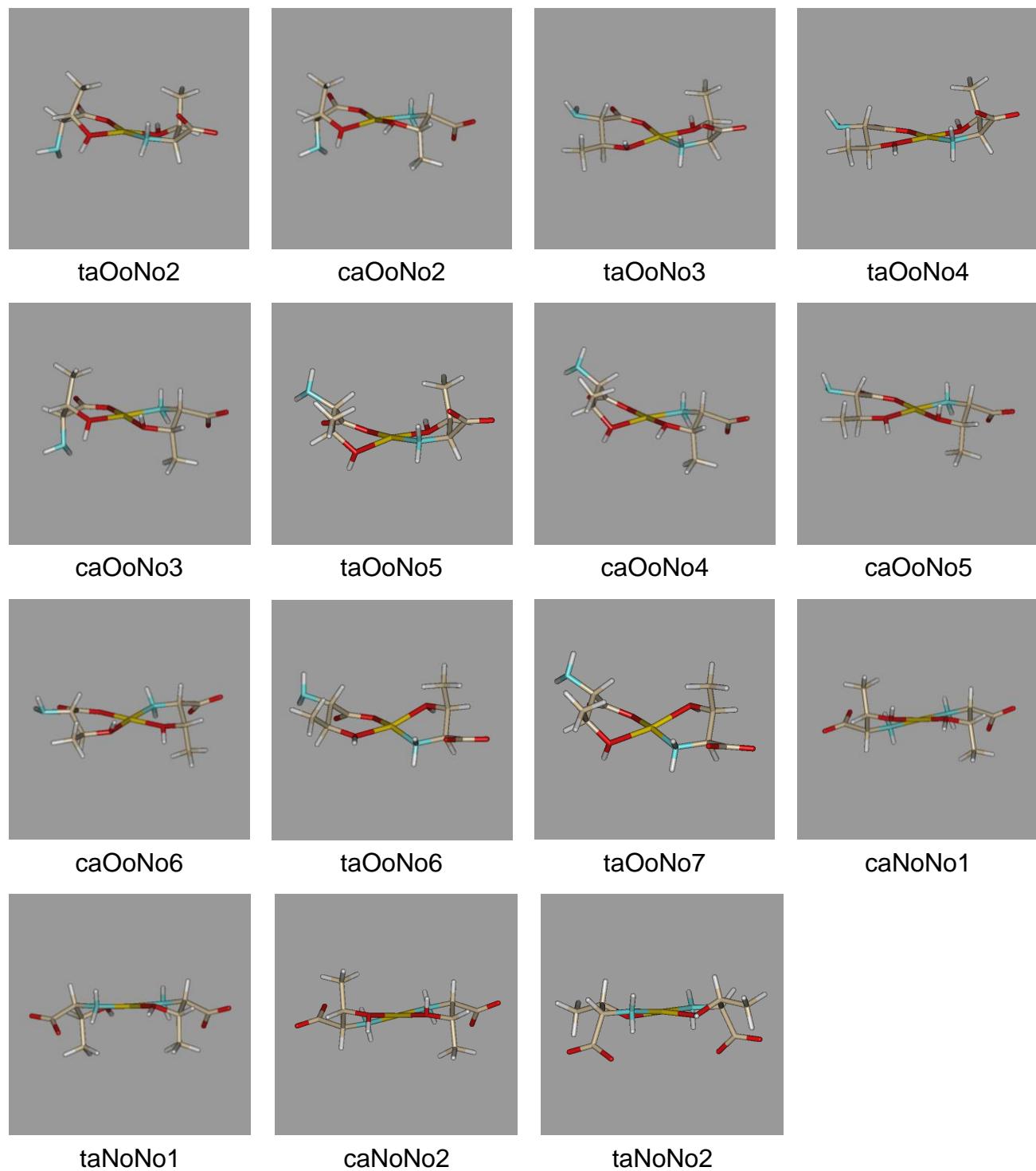


Figure S1. continued

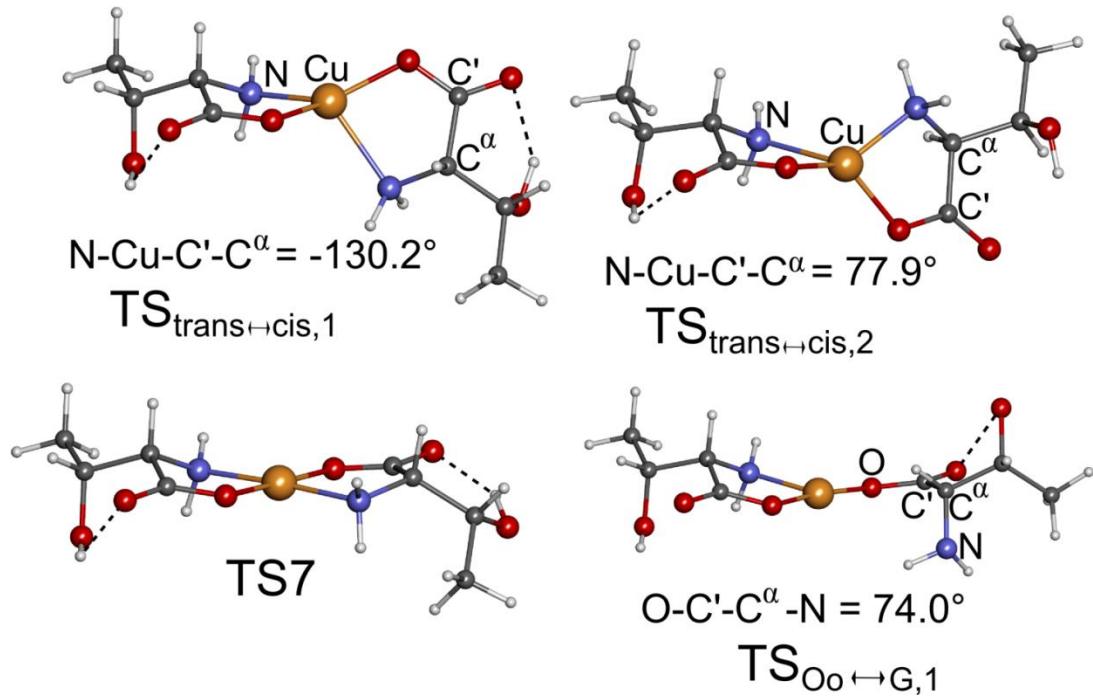


Figure S2. Illustrations of the B3LYP/BS0 TS structures calculated for Cu(L-Thr)₂. The hydrogen bonds are denoted by the broken lines.

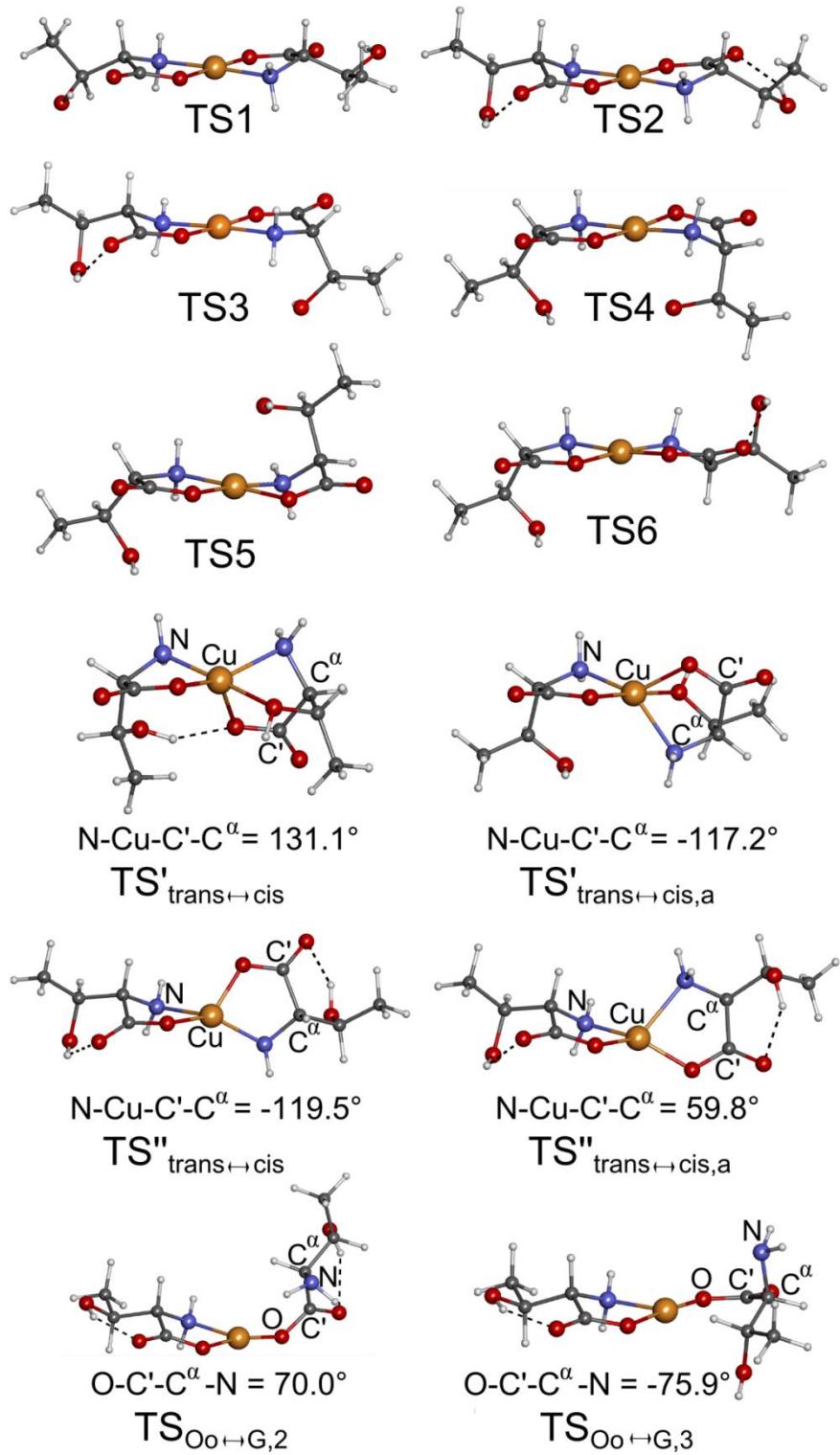


Figure S3. Illustrations of the B3LYP/BS0 TS structures calculated for Cu(L-*a*Thr)₂. The hydrogen bonds are denoted by the broken lines.

Table S1. Characteristic torsion angles ($^{\circ}$) and B3LYP/BS0 relative electronic energy ΔV (kJ mol $^{-1}$) of 57 Cu(L-Thr) $_2$ conformers in the G-G coordination mode^a

name	conformer	ΔV	$\varphi_1^{[b]}$	φ_2	φ_3	φ_4	φ_5	φ_6
tGG1	te1-te1	0.0	-154.4	60.8	74.0	-154.5	60.7	74.1
tGG2	te1-te8	7.0	-153.8	60.2	74.9	-148.5	-169.0	-52.6
tGG3	ta6-te1	10.5	-95.4	-47.5	179.2	-154.6	61.2	73.3
tGG4	ta5-te1	10.9	-95.6	-44.4	-89.4	-154.6	60.9	73.4
tGG5	te1-te6	13.8	-154.7	61.0	73.6	-157.2	-41.2	-170.4
tGG6	ta1-te1	14.4	-78.7	59.5	77.3	-154.7	60.5	73.3
tGG7	te8-te8	15.0	-150.2	-169.3	-53.3	-150.2	-169.3	-53.3
tGG8	ta3-te1	17.5	-85.4	49.4	175.5	-155.2	62.3	72.1
tGG9	ta6-te8	18.2	-95.4	-47.3	-179.9	-148.8	-169.4	-52.2
tGG10	ta2-te1	18.3	-73.2	51.6	-109.3	-156.4	62.8	72.3
tGG11	ta5-te8	18.4	-95.5	-44.1	-89.8	-149.5	-169.2	-52.4
tGG12	te6-te8	20.8	-157.8	-40.8	-170.1	-150.9	-169.4	-52.8
tGG13	ta1-te8	21.6	-81.3	60.8	77.2	-152.3	-169.6	-52.8
tGG14	ta5-ta5	22.0	-95.3	-44.6	-89.1	-95.4	-44.9	-89.0
tGG15	ta5-ta6	22.1	-95.4	-44.5	-88.9	-95.3	-47.6	-179.1
tGG16	ta6-ta6	22.5	-96.3	-47.3	-178.2	-96.3	-47.3	-178.2
tGG17	ta1-ta1	23.1	-76.7	57.1	82.9	-101.0	43.0	68.0
tGG18	ta1-ta5	23.6	-76.2	59.1	80.0	-89.0	-45.2	-89.7
tGG19	ta1-ta6	24.0	-78.7	59.9	74.3	-91.3	-47.7	-179.8
tGG20	ta3-te8	24.1	-86.2	49.9	176.8	-149.6	-169.4	-51.9
tGG21	ta2-te8	24.3	-73.3	51.6	-109.3	-152.2	-169.5	-53.3
tGG22	ta6-te6	25.4	-95.1	-48.0	179.4	-157.4	-41.1	-169.9
tGG23	ta5-te6	25.6	-95.4	-44.8	-88.7	-157.2	-41.1	-170.0
tGG24	te6-te6	28.0	-158.1	-41.2	-170.1	-158.1	-41.2	-170.2
tGG25	ta1-te6	28.6	-80.6	60.5	74.5	-157.9	-40.7	-169.6
tGG26	ta1-ta3	28.6	-100.1	84.2	69.5	-70.3	60.6	-139.4
tGG27	ta3-ta5	29.5	-88.5	49.1	166.3	-94.4	-45.8	-87.1
tGG28	ta3-te3	30.2	-80.0	52.0	-176.2	-145.9	46.1	163.0
tGG29	ta2-ta6	30.7	-74.7	50.5	-99.0	-93.1	-49.6	170.9
tGG30	ta2-te6	31.9	-73.5	51.6	-102.5	-158.4	-42.1	-171.0

tGG31	ta1-ta2	32.0	-106.0	30.9	66.7	-75.5	48.8	-88.3
tGG32	ta2-ta5	32.4	-74.2	50.0	-102.2	-95.6	-46.9	-87.4
tGG33	ta2-ta3	39.3	-96.8	63.6	-77.4	-73.8	56.8	-121.0
tGG34	ta2-ta2	39.4	-74.0	56.3	-118.9	-96.6	64.1	-77.0
tGG35	ta9-te1	43.7	-96.1	-156.2	-160.8	-153.5	60.8	74.3
cGG1	ce1-ce1	43.8	-156.6	59.7	73.5	-156.6	59.7	73.5
cGG2	ca1-ce1	49.5	-72.5	55.4	82.5	-158.3	60.5	72.5
cGG3	ca5-ce1	51.0	-87.7	-44.5	-95.2	-157.0	60.3	73.0
cGG4	ca6-ce1	51.9	-87.7	-45.3	-168.0	-157.1	60.3	73.0
cGG5	ce1-ce8	54.6	-155.4	58.9	74.6	-156.8	-170.6	-53.0
cGG6	ca1-ca5	55.3	-72.0	54.5	83.6	-90.3	-46.0	-93.5
cGG7	ce1-ce6	55.6	-156.9	60.5	73.0	-164.0	-42.0	-167.6
cGG8	ca1-ca6	55.8	-71.9	54.0	84.0	-90.4	-46.9	-172.4
cGG9	ca1-ca1	55.9	-73.2	54.5	80.1	-73.2	54.5	80.2
cGG10	ca5-ca5	58.3	-88.5	-45.1	-94.4	-88.4	-45.0	-94.3
cGG11	ca1-ce8	58.7	-71.9	54.8	84.9	-156.9	-170.5	-52.3
cGG12	ca5-ca6	59.0	-88.7	-44.9	-94.2	-88.7	-45.8	-170.3
cGG13	ca6-ca6	59.5	-87.8	-47.0	-173.3	-87.8	-47.0	-173.3
cGG14	ce1-ca8	59.8	-156.8	59.2	74.3	-108.7	-169.6	-44.9
cGG15	ca5-ce8	60.8	-87.4	-44.8	-96.1	-155.5	-170.4	-52.6
cGG16	ca6-ce8	61.1	-87.3	-46.1	-170.7	-156.1	-170.4	-52.7
cGG17	ca1-ce6	61.3	-72.1	55.0	81.8	-165.6	-41.7	-168.4
cGG18	ca5-ce6	63.5	-89.0	-45.2	-94.1	-162.9	-41.6	-167.9
cGG19	ca6-ce6	63.8	-88.9	-46.3	-173.3	-163.2	-41.9	-168.7
cGG20	ce8-ce8	65.0	-156.0	-170.3	-53.3	-156.0	-170.3	-53.3
cGG21	ce6-ce8	65.2	-164.1	-42.2	-168.0	-155.4	-170.6	-52.7
cGG22	ce6-ce6	67.2	-163.8	-42.1	-168.2	-163.9	-42.1	-168.2

^a The conformers are depicted in Figure S1. The electronic energy of the most stable conformer (te1-te1) is used as the reference value ($V_0 = -1071.76257129$ a.u.). The torsion angles $\varphi_1, \varphi_2, \varphi_3, \varphi_4, \varphi_5$, and φ_6 are: Cu–N11–C12–C13, N11–C12–C13–O13, C12–C13–O13–HO13, Cu–N21–C22–C23, N21–C22–C23–O23, C22–C23–O23–HO23. The atom labels are defined in Figure 5.

Table S2. Characteristic torsion angles ($^{\circ}$) and B3LYP/BS0 relative electronic energy ΔV (kJ mol $^{-1}$) of 32 Cu(L-Thr) $_2$ conformers in the No-G coordination mode^a

name	conformer	ΔV	φ_{11}	φ_{12}	φ_{13}	φ_4	φ_5	φ_6
tNoG1	tap-te1	41.3	86.5	-87.9	-71.9	-154.6	61.1	73.3
tNoG2	tap-ta1	56.1	86.5	-87.7	-71.9	-81.0	61.7	74.8
tNoG3	tap-ta2	61.4	83.5	-89.8	-74.4	-73.8	52.0	-92.5
cNoG1	cap-ca4	62.8	55.9	-26.6	-146.2	-81.7	-25.8	97.3
cNoG2	cap-ca2-A	75.0	70.2	28.6	-124.4	-84.7	55.7	-116.9
cNoG3	cap-ca3-A	82.8	75.6	20.7	-111.9	-69.6	54.8	-123.8
cNoG4	cap-ce6	84.1	54.4	-26.2	-140.7	-155.0	-44.1	-173.1
cNoG5	cap-ca2-B	85.2	69.7	3.9	-120.1	-78.2	49.0	-93.4
cNoG6	cap-ca3-B	89.7	102.4	-12.7	-165.7	-70.2	53.8	-123.7
cNoG7	cap-ca3-C	91.7	75.7	-99.2	-87.9	-70.4	38.5	143.7
cNoG8	cep-ce1	106.9	138.3	-11.0	-156.2	-153.9	58.8	76.7
cNoG9	cep-ca6	113.2	135.3	-10.3	-156.8	-90.9	-47.1	175.9
cNoG10	cep-ca5	115.5	139.7	-12.0	-155.6	-93.6	-44.1	-95.6
cNoG11	cep-ce8	116.1	135.4	-9.8	-157.5	-151.7	-168.9	-54.5
tNoG4	tem-te1	116.7	145.8	-17.1	104.7	-155.1	58.2	76.3
cNoG12	cep-ce6	117.0	137.2	-10.9	-156.3	-160.4	-42.2	-170.4
tNoG5	tep-te1	117.1	148.5	-18.2	-155.6	-154.7	58.6	76.1
cNoG13	cep-ca1	118.4	140.4	-11.4	-157.4	-74.1	58.2	90.9
tNoG6	tem-ta1	122.1	146.7	-17.2	102.3	-75.5	57.3	88.4
cNoG14	cem-ca6	123.0	151.6	-16.4	130.7	-91.2	-46.8	-179.6
cNoG15	cem-ca1	123.1	151.3	-16.0	119.6	-71.2	56.1	88.7
cNoG16	cem-ca5	124.1	153.4	-16.8	131.1	-91.4	-44.2	-94.9
tNoG7	tem-ta5	124.5	148.3	-18.1	109.2	-90.9	-43.1	-95.5
tNoG8	tep-ta6	124.7	149.1	-18.5	-154.0	-92.0	-46.3	-175.1
tNoG9	tep-ta5	124.8	148.6	-18.3	-155.6	-91.3	-44.6	-95.6
tNoG10	tem-ta6	125.4	148.7	-18.2	109.5	-92.1	-44.3	-168.0
tNoG11	tep-te6	127.6	149.1	-18.3	-153.4	-162.3	-41.5	-168.2
tNoG12	tep-te8	128.4	149.7	-19.0	-153.9	-154.0	-169.0	-54.3
tNoG13	tem-te8	128.7	146.7	-17.9	105.2	-154.7	-169.5	-54.4
tNoG14	tem-te6	129.0	147.5	-18.0	107.4	-162.3	-40.9	-168.1

tNoG15	tep-ta1	129.8	149.9	-19.1	-153.2	-75.7	58.2	86.4
tNoG16	tep-ta3	132.6	150.2	-19.2	-153.1	-79.0	48.1	159.2

^a The conformers are depicted in Figure S1. The reference energy value V_0 is given in Table S1. The φ_4 , φ_5 , and φ_6 torsions are defined in Table S1. The torsion angles φ_{11} , φ_{12} , and φ_{13} are: Cu–N11–C12–C11, N11–C12–C11–O11, HO13–O13–Cu–N11. The atom labels are defined in Figure 5.

Table S3. Characteristic torsion angles ($^{\circ}$) and B3LYP/BS0 relative electronic energy ΔV (kJ mol $^{-1}$) of 62 Cu(L-Thr) $_2$ conformers in the Oo-G coordination mode^a

name	conformer	ΔV	φ_7	φ_8	φ_9	φ_{10}	φ_4	φ_5	φ_6
tOoG1	te4dm-te1-A	49.4	-51.7	71.3	-8.7	150.7	-154.2	59.9	74.9
tOoG2	tbt2p-te1-A	53.5	-15.1	61.9	-9.0	-171.7	-154.4	60.1	74.9
tOoG3	ttw1m-te1	56.6	72.6	-66.3	-22.1	86.3	-154.8	60.4	73.9
tOoG4	te4dm-te8	57.2	-52.7	71.5	-8.1	151.1	-150.3	-169.2	-53.3
tOoG5	te4dm-ta5	60.1	-51.4	71.4	-8.9	149.1	-97.2	-43.5	-89.2
tOoG6	te4dm-ta6	60.4	-52.9	71.5	-9.0	150.6	-96.7	-47.4	-179.8
tOoG7	te3up-te1	60.7	62.3	-67.2	3.7	-160.1	-153.6	60.3	74.8
tOoG8	te4dm-te1-B	61.4	-45.0	73.5	-12.8	148.0	-154.4	59.9	75.1
tOoG9	tbt2p-te8	61.8	-17.0	63.2	-7.4	-167.0	-151.0	-169.4	-53.4
tOoG10	te4dm-ta1-A	63.1	-49.5	72.4	-6.8	145.6	-77.4	57.3	79.9
tOoG11	tbt2p-te3	63.3	-14.1	62.2	-8.0	-167.2	-145.4	45.5	165.1
tOoG12	te4dm-te6	63.8	-51.5	71.4	-8.4	151.0	-156.0	-40.6	-170.3
tOoG13	tbt2p-te1-B	63.8	-14.1	66.0	-7.2	-162.8	-154.3	60.3	74.7
tOoG14	ttw1m-te8	64.5	72.5	-66.8	-20.9	85.7	-150.4	-169.9	-52.7
tOoG15	tbt2p-ta6	65.1	-14.6	61.9	-8.7	-173.4	-96.9	-46.6	-178.7
tOoG16	te4dm-te5	66.7	-51.5	71.4	-8.5	150.9	-156.6	-40.1	-103.9
tOoG17	ttw1m-ta1	67.7	74.2	-67.3	-19.9	83.0	-73.1	58.3	77.7
tOoG18	tbt2p-ta1-A	67.7	-11.1	61.9	-6.8	-177.3	-77.5	60.3	79.0
tOoG19	tbt2p-te6	67.8	-14.6	62.5	-7.8	-166.0	-157.7	-40.9	-170.1
tOoG20	ttw1m-ta5	67.9	72.1	-65.9	-22.7	87.3	-96.3	-44.4	-88.5
tOoG21	ttw1m-ta6	68.3	71.6	-66.0	-23.2	88.2	-96.5	-47.2	-178.0
tOoG22	te3dm-ta2	70.4	-57.6	71.3	-7.7	155.2	-73.3	51.6	-114.0
tOoG23	te4dm-te3	71.2	-44.9	73.4	-13.0	147.6	-145.0	46.0	172.7
tOoG24	ttw1m-te6	71.7	71.8	-66.2	-22.6	87.5	-157.3	-40.8	-169.9

tOoG25	ttb2p-ta2	72.4	-23.6	67.2	-4.3	-149.5	-74.6	51.6	-100.1
tOoG26	te4up-te1	74.4	47.8	-69.0	18.6	-156.4	-153.5	60.3	74.9
tOoG27	ttw1m-te5	74.4	71.8	-65.9	-23.3	87.9	-157.0	-40.4	-100.6
tOoG28	ttw1m-ta3	75.2	73.4	-66.4	-21.6	85.2	-89.2	48.0	168.2
tOoG29	te4dm-ta1-B	75.2	-43.1	74.3	-11.3	142.0	-76.7	57.1	80.9
tOoG30	ttw1m-ta2	75.3	74.6	-68.1	-17.4	81.1	-72.6	50.5	-99.1
tOoG31	tch1p-ta2	75.7	51.9	-60.5	44.8	-166.1	-71.4	56.8	-119.1
tOoG32	te4um-te1	77.3	17.3	-61.2	16.2	161.0	-152.7	59.3	75.7
tOoG33	tbt2p-ta1-B	79.0	-13.3	65.7	-6.9	-167.6	-85.7	59.7	78.5
tOoG34	te4dm-ta3	79.1	-45.1	73.6	-12.4	146.9	-85.9	48.7	171.2
tOoG35	tbt2p-ta3	80.1	-14.4	68.0	-3.7	-159.2	-84.8	51.5	178.0
tOoG36	te4um-ta3-A	82.0	41.4	-72.0	4.0	127.3	-83.7	48.4	170.6
tOoG37	tbt2p-ta2	82.0	-19.2	68.4	-6.6	-153.3	-74.7	52.7	-112.7
cOoG1	ctw1m-ce1	86.5	78.8	-67.3	-17.8	72.8	-156.2	59.3	74.1
cOoG2	cbt1m-ca1	86.7	13.6	-69.8	8.3	141.8	-86.8	45.3	65.6
cOoG3	ctw1m-ca1	90.9	78.6	-65.8	-21.3	74.5	-70.8	51.6	87.2
tOoG38	ttb1m-ta1	91.7	22.4	-65.2	12.6	147.0	-78.1	57.4	79.8
cOoG4	ctw1m-ca6	94.2	78.4	-65.6	-21.8	75.4	-89.0	-47.4	-178.0
cOoG5	ctw1m-ca5	94.6	78.1	-66.5	-20.4	75.2	-89.4	-46.0	-93.1
tOoG39	te4um-ta3-B	94.7	14.2	-60.6	16.7	158.7	-81.5	50.2	-179.5
cOoG6	ctw1m-ce8	95.4	79.3	-67.1	-17.3	71.5	-154.5	-170.2	-52.8
cOoG7	ctb2m-ce1	95.9	10.8	55.6	-3.0	60.7	-154.9	58.6	75.1
cOoG8	cbt2m-ca1-A	98.5	8.3	57.0	-3.1	63.2	-71.0	51.8	88.7
cOoG9	ctw1m-ce6	99.1	79.5	-66.4	-18.7	72.5	-161.8	-41.8	-169.6
cOoG10	ctb2m-ca6	103.1	11.2	55.6	-3.2	60.3	-90.0	-47.2	-177.5
cOoG11	ctb2m-ca5	103.4	11.0	55.7	-3.1	60.5	-90.3	-45.8	-92.8
cOoG12	cbt2m-ca6	104.1	-9.4	66.4	-2.9	94.3	-89.7	-47.3	-177.7
cOoG13	ctb2m-ce8	105.3	12.2	54.8	-3.1	58.7	-152.8	-169.4	-53.0
cOoG14	ce4dp-ce1	105.7	-40.8	72.6	-6.4	-120.4	-153.9	58.6	75.2
cOoG15	ce4dp-ca1-A	107.9	-34.3	71.8	-7.4	-114.2	-65.9	47.0	107.3
cOoG16	ctb2m-ce6	108.3	12.8	54.4	-3.4	58.5	-160.8	-41.2	-169.2
cOoG17	ce4dp-ca1-B	108.4	-45.4	72.7	-8.0	-114.0	-70.9	50.3	87.1
cOoG18	cbt2m-ca1-B	110.7	-10.1	68.9	-1.9	81.1	-72.0	52.9	87.5

cOoG19	ce4dp-ca6	111.7	-33.9	72.0	-5.3	-131.4	-86.8	-48.3	179.3
cOoG20	ce4dp-ca5	112.9	-36.9	72.3	-5.7	-127.2	-87.2	-46.7	-93.3
cOoG21	ce4dp-ce8	113.6	-38.0	72.4	-6.2	-124.2	-154.2	-169.6	-53.1
cOoG22	ce4dp-ce6	116.0	-33.5	71.5	-4.8	-132.8	-161.3	-42.5	-170.2
cOoG23	ce4um-ca1	118.3	20.4	-66.5	17.3	133.9	-92.2	71.0	73.5

^a The conformers are depicted in Figure S1. The reference energy value V_0 is given in Table S1. The φ_4 , φ_5 , and φ_6 torsions are defined in Table S1. The torsion angles φ_7 , φ_8 , φ_9 , and φ_{10} are: Cu–O13–C13–C12, O13–C13–C12–C11, Cu–C12–C11–O11, HO13–O13–Cu–O11. The atom labels are defined in Figure 5.

Table S4. Characteristic torsion angles ($^{\circ}$) and B3LYP/BS0 relative electronic energy ΔV (kJ mol $^{-1}$) of 28 Cu(L-Thr) $_2$ conformers in the Oo-Oo coordination mode^a

name	conformer	ΔV	φ_7	φ_8	φ_9	φ_{10}	φ_{14}	φ_{15}	φ_{16}	φ_{17}
tOoOo1	te4dm-te4dm-A	95.5	-54.7	71.9	-8.4	149.4	-54.1	71.8	-8.3	148.7
tOoOo2	te4dm-te4dm-B	107.7	-52.9	71.6	-8.8	147.8	-47.5	73.8	-13.2	146.0
tOoOo3	tbt2p-te3dm	112.4	-9.8	63.8	-7.3	-170.0	-57.4	71.3	-7.5	153.3
tOoOo4	tbt2p-tbt2p	117.3	-19.4	65.5	-3.4	-156.4	-15.8	67.5	-4.7	-157.6
tOoOo5	tbt4m-te4dm	117.3	68.3	-66.9	-23.3	92.8	-45.3	73.6	-12.9	142.2
tOoOo6	tbt2p-te3um	122.1	-7.6	63.7	-6.1	-167.2	71.0	-69.9	-14.9	84.4
tOoOo7	te3dm-te4up	122.9	-58.6	70.4	-8.0	157.3	54.8	-70.0	14.3	-160.5
tOoOo8	tbt1m-te4dm	123.3	19.2	-64.0	10.5	149.8	-53.4	71.4	-9.4	150.9
tOoOo9	te4up-ttb2p	124.0	-20.9	66.3	-3.5	-149.7	48.4	-69.2	20.1	-155.0
cOoOo1	ce3um-ctb1m	126.8	64.5	-80.7	14.5	73.9	-8.9	-62.2	7.8	177.0
tOoOo10	tbt2p-te4um	127.5	-13.8	62.9	-6.2	-160.9	9.8	-59.5	15.7	160.8
tOoOo11	te4dm-ttw1p	130.2	72.9	-47.4	-26.8	-164.0	-51.5	73.8	-12.5	150.7
tOoOo12	ttb1m-ttw1m	133.1	68.9	-65.6	-25.0	93.2	18.1	-63.7	10.8	147.5
tOoOo13	te3um-te4up	133.7	74.5	-68.3	-15.3	79.8	49.3	-68.5	18.9	-156.0
cOoOo2	ctb2m-ctw1m-A	137.7	-22.3	69.2	-5.1	118.0	79.3	-66.2	-20.1	73.0
cOoOo3	cbt2m-ctb1m	150.7	-7.5	-62.9	8.3	177.4	-0.4	64.3	-1.2	69.0
cOoOo4	ctb2m-ctw1m-B	151.4	-25.5	73.1	-5.6	119.2	79.2	-67.0	-18.4	72.3
cOoOo5	ce3um-ctb1p	154.2	63.9	-81.1	10.9	76.3	25.4	-66.3	14.3	-141.2
cOoOo6	ce4dp-ctb2m	155.2	-22.1	68.4	-7.1	133.1	-37.0	73.6	-7.0	-123.0
cOoOo7	cbt2p-ctb2p-A	155.8	-18.7	68.2	-3.5	-153.3	-23.7	69.9	-3.7	-145.8
cOoOo8	cch1m-ce4um	156.4	64.2	-71.4	46.1	59.7	32.9	-71.1	15.9	119.6

cOoOo9	cbt2m-ctb2m	158.3	9.3	57.3	-2.9	62.5	-25.0	73.0	-7.0	119.2
cOoOo10	ce4up-ctb2m	166.3	13.1	54.1	-3.3	58.0	42.2	-67.5	12.4	-155.2
cOoOo11	cbt2p-ctb2p-B	169.1	-15.3	66.5	-4.2	-159.2	-26.7	73.4	-4.9	-140.4
cOoOo12	cbt2p-ce4dp	169.1	-10.7	64.4	-5.0	-167.8	-29.4	74.3	-5.6	-135.8
cOoOo13	ctb2p-ctb2p	169.2	-21.6	68.9	-3.7	-149.7	-25.6	73.3	-3.7	-143.2
cOoOo14	cbt2m-ce4um	180.7	6.2	58.4	-4.1	68.5	42.9	-71.9	13.5	116.9
cOoOo15	cbt2m-ce4up	188.2	-9.7	66.3	-7.3	110.3	37.0	-73.4	8.8	-132.4

^a The conformers are depicted in Figure S1. The reference energy value V_0 is given in Table S1. The φ_7 , φ_8 , φ_9 , and φ_{10} torsions are defined in Table S2. The torsion angles φ_{14} , φ_{15} , φ_{16} , and φ_{17} are: Cu–O23–C23–C22, O23–C23–C22–C21, Cu–C22–C21–O21, HO23–O23–Cu–O21. The atom labels are defined in Figure 5.

Table S5. Characteristic torsion angles ($^{\circ}$) and B3LYP/BS0 relative electronic energy ΔV (kJ mol $^{-1}$) of 13 Cu(L-Thr) $_2$ conformers in the Oo-No coordination mode^a

name	conformer	ΔV	φ_7	φ_8	φ_9	φ_{10}	φ_{18}	φ_{19}	φ_{20}
cOoNo1	ctb2p-cap	96.2	-21.4	66.1	-4.6	-152.5	87.9	-87.3	-71.2
cOoNo2	ctb1m-cem	151.2	-4.4	-65.7	7.8	174.1	138.2	-15.4	93.5
tOoNo1	te3um-tep	151.6	76.4	-70.6	-11.4	72.4	140.3	-13.4	-154.5
tOoNo2	ttw1m-tem	159.9	76.8	-68.0	-18.5	75.5	155.5	-18.1	131.5
tOoNo3	te4dp-tep	162.3	-43.2	71.8	-3.7	-124.3	134.2	-8.6	-157.4
cOoNo3	ctw1m-cem	162.7	74.3	-70.1	-17.2	78.7	155.5	-20.6	128.7
tOoNo4	te4dm-tep	163.4	-43.8	71.9	-4.8	146.9	136.5	-10.1	-157.7
tOoNo5	te4dm-tam	169.5	-37.6	69.8	-6.3	151.8	94.2	12.8	150.1
cOoNo4	ce4dm-cem	169.8	-39.6	72.6	-5.8	135.7	149.0	-19.6	108.7
tOoNo6	te4dm-tem	170.0	-37.8	71.3	-6.7	143.5	152.9	-16.5	124.1
tOoNo7	te4dp-tem	171.6	-34.1	70.6	-5.4	-137.0	151.8	-16.2	125.8
cOoNo5	ctb2m-cep	173.0	-24.8	71.5	-2.9	113.1	148.4	-19.0	-161.1
cOoNo6	ce4dp-cep	173.5	-48.6	72.9	-6.3	-112.6	152.4	-21.4	-144.5

^a The conformers are depicted in Figure S1. The reference energy value V_0 is given in Table S1. [b] The φ_7 , φ_8 , φ_9 , and φ_{10} torsions are defined in Table S2. The torsion angles φ_{18} , φ_{19} , and φ_{20} are: Cu–N21–C22–C21, N21–C22–C21–O21, HO23–O23–Cu–N21. The atom labels are defined in Figure 5.

Table S6. Characteristic torsion angles ($^{\circ}$) and B3LYP/BS0 relative electronic energy ΔV (kJ mol $^{-1}$) of 4 Cu(L-Thr) $_2$ conformers in the No-No coordination mode^a

name	conformer	ΔV	$\varphi_{11}^{[b]}$	φ_{12}	φ_{13}	φ_{18}	φ_{19}	φ_{20}
cNoNo1	cap-cap	99.7	53.9	-25.5	-137.2	86.6	-90.0	-72.6
cNoNo2	cep-cep	211.7	148.4	-18.4	-150.3	148.5	-18.2	-150.1
cNoNo3	cem-cem	212.8	146.6	-17.2	109.5	147.1	-17.2	110.3
tNoNo1	tep-tep	212.8	146.2	-15.5	-153.7	146.1	-15.4	-153.3

^a The conformers are depicted in Figure S1. The reference energy value V_0 is given in Table S1. The φ_{11} , φ_{12} , and φ_{13} torsions are defined in Table S3. The φ_{18} , φ_{19} , and φ_{20} torsions are defined in Table S5.

Table S7. Characteristic torsion angles ($^{\circ}$) and B3LYP/BS0 relative electronic energy ΔV (kJ mol $^{-1}$) of 113 Cu(L-*a*Thr) $_2$ conformers in the G-G coordination mode^a

name	conformer	ΔV	φ_1	φ_2	φ_3	φ_4	φ_5	φ_6
taGG1	te8-te8	7.4	-153.8	-175.5	-51.0	-154.1	-175.5	-51.1
taGG2	te1-te8	8.6	-158.8	68.3	71.0	-153.9	-175.7	-50.7
taGG3	te1-te1	10.5	-159.4	68.8	70.4	-159.3	68.9	70.4
taGG4	ta1-te8	16.5	-77.8	59.9	83.7	-154.6	-176.0	-50.1
taGG5	te8-ta6	16.9	-153.7	-176.0	-50.4	-106.4	-55.3	178.0
taGG6	te6-te8	17.0	-154.9	-57.0	-176.4	-154.4	-175.9	-50.6
taGG7	te8-ta5	18.1	-154.0	-175.9	-50.5	-105.5	-55.5	-74.5
taGG8	ta1-te1	18.4	-77.0	58.8	84.8	-160.0	68.8	69.1
taGG9	te1-ta6	18.4	-158.7	69.1	69.8	-104.6	-55.9	177.3
taGG10	te1-te6	18.6	-158.7	68.9	69.8	-153.8	-56.9	-176.9
taGG11	ta3-te8	18.9	-78.1	57.0	-165.0	-153.2	-176.2	-49.6
taGG12	te1-ta5	19.6	-158.7	69.0	70.0	-104.0	-56.5	-74.0
taGG13	te5-te8	20.7	-147.5	-57.6	-83.5	-153.4	-175.8	-50.4
taGG14	ta3-te1	21.1	-77.9	56.3	-168.1	-158.3	70.4	68.5
taGG15	te3-te8	21.7	-144.6	54.9	173.7	-153.1	-175.7	-50.1
taGG16	te1-te5	22.3	-158.3	68.9	70.0	-147.9	-57.8	-83.0
taGG17	ta1-ta1	23.6	-80.5	52.3	84.1	-90.2	48.3	75.8
taGG18	te1-te3	23.8	-159.4	69.4	69.5	-145.2	54.8	173.1
taGG19	ta1-ta3	25.5	-99.2	93.5	65.5	-71.3	62.8	-143.8
taGG20	ta1-ta6	27.0	-78.8	59.9	78.9	-107.7	-53.1	177.4

taGG21	ta1-te6	27.4	-78.9	59.8	81.0	-154.0	-56.6	-176.3
taGG22	te6-ta6	27.7	-154.5	-57.1	-176.4	-106.0	-55.6	177.8
taGG23	te6-te6	27.7	-153.5	-57.1	-176.7	-153.9	-56.7	-176.6
taGG24	ta6-ta6	27.9	-107.1	-55.6	178.5	-107.2	-55.5	178.7
taGG25	ta1-ta5	28.2	-80.3	59.1	81.6	-107.4	-54.4	-74.0
taGG26	te6-ta5	28.6	-153.1	-57.2	-176.3	-104.3	-56.1	-73.8
taGG27	ta5-ta6	28.6	-105.3	-56.0	-73.9	-106.8	-55.8	178.2
taGG28	ta5-ta5	29.5	-105.1	-55.8	-74.3	-105.2	-55.5	-74.0
taGG29	te5-ta1	30.8	-151.6	-57.9	-82.2	-78.6	59.6	82.0
taGG30	te5-ta6	31.1	-148.1	-58.0	-81.7	-106.2	-55.5	177.8
taGG31	te5-te6	31.2	-148.6	-57.7	-82.5	-151.4	-56.9	-176.7
taGG32	ta3-te6	31.3	-79.2	56.7	-167.7	-153.7	-57.2	-176.4
taGG33	te6-ta2	31.6	-153.7	-57.9	-177.5	-73.9	54.3	-108.1
taGG34	ta3-ta6	32.0	-85.0	54.1	178.1	-108.0	-55.7	178.3
taGG35	te5-ta5	32.1	-149.1	-57.7	-82.4	-104.3	-55.8	-74.0
taGG36	ta3-ta5	32.4	-84.0	53.7	176.8	-108.5	-56.1	-72.6
taGG37	te3-ta6	32.6	-145.8	55.3	172.7	-106.0	-56.0	177.4
taGG38	te3-te6	32.8	-145.1	55.2	172.9	-151.4	-56.8	-177.3
taGG39	ta2-ta6	33.6	-74.6	54.8	-112.8	-113.5	-54.2	177.5
taGG40	ta1-te3	33.8	-79.1	59.0	81.8	-149.1	54.6	168.9
taGG41	ta3-te5	34.4	-80.3	55.5	-174.3	-146.7	-58.1	-80.3
taGG42	te5-te5	34.7	-149.0	-57.8	-82.8	-148.9	-57.7	-82.8
taGG43	ta2-te5	35.2	-73.9	55.0	-116.2	-153.5	-58.6	-83.0
taGG44	te3-te5	36.2	-145.3	55.1	173.0	-149.3	-57.9	-82.1
taGG45	ta3-te3	36.3	-76.3	58.3	-156.8	-144.9	56.1	173.9
taGG46	te3-te3	38.3	-145.8	55.4	172.6	-145.9	55.2	173.3
taGG47	ta1-ta2	38.4	-98.8	31.7	73.2	-76.8	50.5	-84.5
taGG48	ta3-ta2	39.8	-74.7	58.3	-132.1	-95.7	67.0	-70.3
taGG49	te8-ta9	49.0	-153.7	-176.0	-50.4	-82.8	177.9	-174.4
taGG50	te1-ta9	50.2	-158.9	69.2	69.9	-81.9	-179.7	-170.6
caGG1	ca1-ca1	51.1	-73.8	54.1	85.3	-73.7	54.4	84.7
caGG2	ca1-ce1	51.7	-72.9	56.0	85.9	-163.0	68.2	68.2
caGG3	ca1-ce8	52.0	-72.3	55.0	88.3	-159.0	-176.0	-50.5

taGG51	te7-te8	52.0	-165.0	-159.4	78.2	-153.7	-175.8	-50.6
caGG4	ce1-ce1	53.2	-160.8	67.1	69.7	-161.5	67.3	69.4
taGG52	te1-te7	54.0	-158.9	69.1	70.0	-165.1	-158.7	77.5
caGG5	ce8-ce1	55.2	-157.1	-176.4	-50.8	-161.0	66.7	70.2
caGG6	ce8-ce8	56.9	-157.7	-175.8	-51.4	-157.0	-175.7	-51.3
caGG7	ca1-ca6	57.8	-73.0	53.7	88.3	-99.3	-54.6	-179.8
caGG8	ca1-ca5	58.4	-72.8	54.2	88.3	-99.1	-55.5	-77.4
caGG9	ca1-ca8	59.1	-72.7	54.6	89.0	-100.5	-173.8	-41.0
taGG53	te6-ta9	59.5	-151.7	-56.6	-176.6	-81.7	-178.4	-170.0
caGG10	ca1-ce6	59.9	-72.3	54.6	87.1	-162.1	-56.0	-175.8
taGG54	ta6-ta9	60.0	-108.1	-55.7	178.7	-82.9	-179.6	-170.9
taGG55	ta5-ta9	60.6	-106.7	-55.8	-73.7	-83.0	-178.6	-168.0
caGG11	ce1-ca6	61.6	-161.3	67.6	69.2	-99.0	-53.7	-178.1
caGG12	ce8-ca6	61.8	-157.2	-175.9	-50.7	-98.0	-53.7	-179.2
caGG13	ce1-ce6	62.0	-161.0	67.5	69.1	-159.6	-55.8	-174.5
caGG14	ce1-ca5	62.2	-161.3	67.5	69.4	-97.6	-54.0	-78.7
taGG56	te6-te7	62.4	-153.8	-57.1	-176.9	-165.5	-159.0	76.7
taGG57	te7-ta6	62.5	-165.6	-159.2	76.9	-107.0	-55.6	177.6
caGG15	ce6-ce8	62.5	-159.1	-55.6	-175.3	-156.6	-175.8	-50.6
taGG58	te7-ta1	62.6	-166.0	-158.6	75.9	-78.0	60.4	81.3
taGG59	te5-ta9	62.9	-148.9	-57.5	-82.1	-82.0	-178.4	-169.7
caGG16	ce8-ca5	62.9	-156.7	-176.0	-50.6	-97.3	-54.3	-79.6
taGG60	ta2-ta9	63.4	-75.2	54.7	-101.0	-83.3	166.3	179.4
taGG61	te7-ta5	63.6	-165.5	-159.4	77.2	-105.7	-56.1	-73.9
caGG17	ca1-ce5	63.9	-72.5	55.1	86.4	-161.8	-58.0	-87.8
taGG62	te3-ta9	64.1	-144.9	55.3	173.4	-81.8	-179.4	-170.1
caGG18	ce1-ca8	64.4	-161.4	66.7	70.1	-99.0	-173.8	-41.2
taGG63	te7-ta2	65.0	-164.7	-159.9	76.3	-73.7	55.1	-112.7
taGG64	te7-ta3	65.2	-164.6	-159.1	75.3	-75.5	57.9	-145.9
caGG19	ce1-ce5	66.0	-161.1	67.4	69.3	-159.3	-57.4	-91.6
taGG65	te5-te7	66.2	-147.0	-57.8	-82.1	-165.3	-158.7	77.2
caGG20	ce8-ca8	66.2	-157.3	-176.0	-51.2	-98.0	-173.3	-41.7
caGG21	ce5-ce8	66.9	-159.0	-57.2	-95.1	-156.8	-175.9	-50.7

taGG66	te3-te7	67.4	-145.1	55.4	174.0	-165.5	-158.1	76.8
caGG22	ca6-ca6	69.0	-97.0	-54.2	-179.6	-98.7	-54.1	-179.8
caGG23	ce6-ca6	69.4	-160.4	-56.3	-175.9	-97.2	-54.0	179.7
caGG24	ce6-ce6	69.7	-159.0	-56.2	-175.5	-160.6	-55.8	-175.5
caGG25	ca5-ca6	69.9	-97.6	-54.7	-78.1	-98.5	-54.0	-179.0
caGG26	ce6-ca5	70.6	-160.4	-56.0	-175.3	-96.7	-55.1	-78.4
caGG27	ca5-ca5	70.7	-96.3	-54.4	-78.4	-97.6	-54.5	-78.2
caGG28	ca6-ca8	71.5	-98.5	-53.4	-178.4	-98.6	-173.5	-41.4
caGG29	ce6-ca8	72.2	-161.9	-56.0	-174.9	-98.0	-173.6	-41.3
caGG30	ca5-ca8	72.3	-97.7	-54.2	-79.3	-98.0	-173.5	-41.3
caGG31	ce5-ca6	73.8	-158.8	-57.8	-91.7	-97.6	-53.8	-179.3
caGG32	ce5-ce6	74.2	-158.4	-57.8	-91.6	-159.2	-55.8	-175.1
caGG33	ce5-ca5	74.6	-159.1	-57.6	-90.4	-96.7	-54.8	-78.4
caGG34	ca8-ca8	75.4	-100.7	-173.3	-41.8	-99.5	-173.6	-41.6
caGG35	ce5-ca8	76.4	-161.0	-57.3	-93.3	-97.9	-173.7	-41.1
caGG36	ce5-ce5	78.3	-158.7	-57.7	-91.8	-159.6	-57.6	-91.5
taGG67	te7-ta9	94.0	-165.5	-160.3	77.0	-82.4	178.6	-174.1
taGG68	te7-te7	97.3	-165.5	-158.8	78.1	-165.4	-158.8	77.8
caGG37	ce7-ca1	100.0	-170.1	-158.4	70.7	-72.6	55.7	86.4
caGG38	ce7-ce1	102.5	-168.7	-160.0	72.2	-161.4	67.1	69.3
caGG39	ce7-ce8	104.5	-168.2	-159.7	73.2	-158.0	-176.1	-51.0
caGG40	ce7-ca6	110.8	-167.7	-158.3	73.6	-97.5	-53.2	-177.8
caGG41	ce6-ce7	111.0	-161.3	-55.6	-175.0	-168.2	-159.5	72.2
caGG42	ce7-ca5	111.3	-168.1	-159.1	73.0	-97.0	-53.8	-79.2
caGG43	ce7-ca8	113.7	-168.4	-160.8	73.1	-97.2	-173.9	-40.8
caGG44	ce5-ce7	115.1	-160.2	-57.3	-92.5	-168.4	-159.7	72.2
caGG45	ce7-ce7	152.4	-168.8	-160.0	71.3	-169.0	-160.1	71.6

^a The conformers are depicted in Figure S1. The reference energy value V_0 is given in Table S1. The φ_1 , φ_2 , φ_3 , φ_4 , φ_5 , and φ_6 torsions are defined in Table S1.

Table S8. Characteristic torsion angles ($^{\circ}$) and B3LYP/BS0 relative electronic energy ΔV (kJ mol $^{-1}$) of 30 Cu(L-*a*Thr) $_2$ conformers in the No-G coordination mode^a

name	conformer	ΔV	φ_{11}	φ_{12}	φ_{13}	φ_4	φ_5	φ_6
taNoG1	tap-ta1	54.1	86.6	-83.6	-72.0	-76.9	60.9	82.2
taNoG2	tap-ta3	59.2	86.9	-82.9	-71.7	-82.8	55.7	175.3
caNoG1	cap-ca3-A	81.4	68.0	31.1	-122.4	-81.9	58.8	-125.4
caNoG2	cap-ca3-B	85.6	70.7	36.0	-102.1	-69.9	59.6	-137.8
caNoG3	cam-ca3	94.8	80.5	4.7	149.8	-69.4	60.0	-127.2
caNoG4	cap-ca1	104.8	70.4	29.9	-118.3	-81.9	84.5	73.8
caNoG5	cep-ce1	119.0	142.5	-10.4	-154.8	-158.3	66.7	73.2
caNoG6	cep-ce8	119.4	140.0	-9.7	-156.4	-154.7	-174.6	-52.5
caNoG7	cep-ca1	122.6	145.7	-11.0	-155.6	-74.2	58.5	92.9
caNoG8	cep-ca3	123.0	134.3	-10.7	-156.4	-77.8	54.4	-169.9
caNoG9	cep-ce6	124.4	141.3	-10.2	-155.5	-157.5	-56.2	-176.8
caNoG10	cep-ca6	124.5	141.8	-10.6	-155.2	-103.0	-53.7	177.5
caNoG11	cep-ca5	127.4	143.6	-11.0	-154.8	-103.0	-53.9	-79.3
taNoG3	tem-ta1	127.5	149.5	-13.7	94.4	-74.6	57.3	92.8
caNoG12	cep-ce5	129.3	141.9	-10.4	-155.3	-156.8	-57.3	-96.0
taNoG4	tep-te1	130.0	151.7	-14.8	-152.2	-159.3	66.8	71.9
caNoG13	cem-ca1	130.4	155.9	-13.7	113.3	-72.0	55.8	92.1
taNoG5	tem-te1	130.4	147.5	-13.5	94.0	-159.5	66.4	72.2
caNoG14	cep-ce3	130.5	141.6	-9.8	-155.3	-145.8	54.1	172.0
taNoG6	tep-te8	131.9	152.8	-15.1	-150.6	-155.8	-174.8	-52.5
taNoG7	tem-te8	133.1	148.2	-13.8	94.1	-155.6	-175.5	-52.3
taNoG8	tep-ta1	134.5	154.6	-15.0	-148.2	-74.9	57.5	92.7
taNoG9	tep-te6	135.9	153.2	-15.0	-149.5	-158.1	-55.4	-175.1
taNoG10	tep-ta6	136.0	153.6	-15.0	-149.6	-101.4	-53.0	179.8
taNoG11	tep-ta3	137.4	155.5	-15.0	-146.9	-77.9	50.9	164.2
taNoG12	tep-ta5	137.5	152.7	-14.9	-150.9	-99.6	-53.9	-80.2
taNoG13	tem-ta5	139.2	148.0	-13.6	94.7	-102.3	-53.5	-78.2
taNoG14	tep-te5	140.4	152.6	-14.9	-150.2	-157.2	-56.6	-97.7
taNoG15	tep-te3	143.7	152.3	-14.5	-151.0	-147.8	53.7	162.8
caNoG15	cep-ca9	158.7	138.7	-9.4	-156.8	-81.4	-176.8	-167.9

^a The conformers are depicted in Figure S1. The reference energy value V_0 is given in Table S1. The φ_4 , φ_5 , and φ_6 torsions are defined in Table S1. The torsion angles φ_{11} , φ_{12} , and φ_{13} are: Cu–N11–C12–C11, N11–C12–C11–O11, HO13–O13–Cu–N11. The atom labels are defined in Figure 5.

Table S9. Characteristic torsion angles ($^{\circ}$) and B3LYP/BS0 relative electronic energy ΔV (kJ mol $^{-1}$) of 69 Cu(L-*a*Thr) $_2$ conformers in the Oo-G coordination mode^a

name	conformer	ΔV	φ_7	φ_8	φ_9	φ_{10}	φ_4	φ_5	φ_6
taOoG1	te4dm-te8	55.6	-53.1	67.7	-6.6	155.8	-153.5	-175.8	-50.7
taOoG2	ttb2p-te8	56.8	-1.8	53.6	-14.0	-171.6	-153.7	-176.0	-50.9
taOoG3	te4dm-te1	57.1	-52.4	68.0	-7.0	155.5	-159.4	68.5	71.1
taOoG4	ttb2p-te1	57.3	1.2	52.4	-14.5	-172.3	-158.9	68.5	71.0
taOoG5	ttb2m-te8	57.5	-20.3	58.8	-13.4	172.3	-153.6	-175.8	-50.7
taOoG6	te3um-te8	62.2	67.3	-69.5	-3.0	79.3	-154.7	-176.5	-50.4
taOoG7	te3um-te1	63.5	68.5	-67.5	-6.9	79.6	-159.2	68.4	70.1
taOoG8	te3dm-ta1	64.0	-51.6	70.0	-4.2	149.0	-75.6	57.3	87.0
taOoG9	te3dm-ta6	65.1	-54.9	68.0	-6.6	155.2	-105.5	-56.2	177.9
taOoG10	ttb2p-ta1-A	65.4	1.3	54.1	-11.4	-176.6	-77.5	59.5	84.4
taOoG11	te3dm-te6	65.8	-55.1	68.3	-6.2	155.5	-153.2	-56.7	-176.5
taOoG12	te4dm-ta5	65.9	-53.8	68.0	-7.3	154.7	-104.7	-56.1	-74.1
taOoG13	ttb2p-ta6	66.4	1.5	51.8	-15.1	-172.6	-108.1	-55.6	179.0
taOoG14	ttb2p-te6	66.6	1.7	52.2	-14.4	-171.9	-152.3	-56.8	-176.3
taOoG15	ttb2p-ta5	67.1	-1.2	53.5	-14.1	-171.9	-106.1	-55.9	-74.2
taOoG16	te4dm-ta6	67.2	-19.6	58.1	-14.8	171.2	-107.0	-55.9	178.6
taOoG17	te3um-ta1	68.8	64.3	-69.0	-7.8	85.6	-75.0	57.0	82.4
taOoG18	te3dm-ta3	68.9	-56.8	68.9	-5.0	154.1	-80.8	54.4	177.7
taOoG19	te4dm-te3	69.7	-53.6	68.2	-6.8	155.0	-144.5	54.6	173.8
taOoG20	ttb2p-ta3-A	69.7	3.6	52.4	-12.8	-172.2	-83.7	55.1	174.7
taOoG21	ttb2p-te3	70.4	1.2	52.5	-14.3	-172.1	-146.5	55.1	172.4
taOoG22	ttb2p-a2	72.6	-7.1	56.8	-11.6	-162.3	-74.7	54.0	-107.1
taOoG23	te3um-ta6	73.5	66.6	-67.8	-7.7	83.0	-109.1	-55.3	178.8
taOoG24	te3um-te6	73.7	67.6	-67.9	-6.2	80.8	-149.8	-56.7	-176.4
taOoG25	te3um-ta5	74.0	66.9	-67.6	-7.8	82.4	-105.5	-56.3	-73.5

taOoG26	te3up-ta1	74.4	60.2	-66.0	8.0	-163.6	-79.6	60.2	84.0
taOoG27	tch1p-ta3	76.0	46.0	-55.4	44.9	-166.0	-72.7	59.7	-122.6
taOoG28	te3um-ta3	76.4	66.1	-68.0	-8.7	83.9	-81.9	51.5	164.3
taOoG29	te3um-te5	76.7	66.9	-67.8	-7.3	82.3	-144.4	-57.2	-79.1
taOoG30	te3um-ta2	77.0	65.1	-73.1	9.2	75.1	-73.7	52.7	-98.2
taOoG31	te3um-te3	77.7	67.7	-67.6	-6.8	80.8	-146.7	54.7	172.2
taOoG32	te4dm-ta1	77.8	-38.9	72.6	-13.4	142.6	-75.4	57.3	86.3
taOoG33	ttb2p-ta1-B	78.7	-0.6	56.9	-13.0	-177.8	-78.1	59.2	85.6
taOoG34	ttb2p-ta3-B	80.7	0.7	57.0	-11.8	-170.6	-82.2	56.0	-176.8
taOoG35	te4dm-ta3	82.7	-41.4	71.5	-15.1	148.3	-80.8	53.8	177.1
taOoG36	te4up-ta1	87.9	49.2	-68.7	17.6	-159.2	-79.9	60.4	84.3
taOoG37	ttb1m-ta1	89.1	20.2	-64.2	11.0	148.8	-75.9	56.5	88.0
caOoG1	ce3um-ca1	89.6	74.4	-64.6	-9.8	70.4	-71.5	51.8	89.6
taOoG38	ttw1m-ta3	89.9	-64.0	33.4	48.1	158.6	-71.3	59.5	-124.4
taOoG39	ttw1p-ta3	90.3	-54.6	30.2	48.3	-121.8	-71.2	60.0	-122.6
caOoG2	ce3um-ce8	92.2	73.8	-67.4	-5.1	69.0	-156.8	-176.2	-50.5
caOoG3	ce3um-ce1	92.5	72.6	-68.4	-3.9	70.1	-160.4	67.3	69.8
caOoG4	ctb2m-ca1-A	95.5	22.0	45.5	-8.3	55.0	-71.6	51.1	92.5
taOoG40	te3dm-ta9	96.6	-57.4	68.1	-6.2	156.0	-83.2	179.0	-170.8
taOoG41	ttb2m-ta3	97.7	19.3	48.8	-8.3	57.4	-83.4	54.3	177.5
taOoG42	ttb2p-ta9	98.5	0.7	52.4	-14.7	-170.6	-82.9	-179.5	-171.7
caOoG5	ce3um-ca6	99.8	75.1	-64.6	-9.7	69.4	-98.3	-55.2	177.7
caOoG6	ctb2m-ce8	100.5	25.1	43.5	-8.2	50.9	-156.5	-175.3	-51.2
caOoG7	ctb2m-ce1	100.7	24.3	44.1	-8.4	51.9	-159.9	67.4	71.0
caOoG8	ctb1p-ca2	101.0	-17.7	-54.4	5.9	-49.2	-94.3	68.1	-35.5
caOoG9	ce3um-ce6	101.0	74.7	-65.8	-7.3	68.9	-156.8	-56.3	-176.0
caOoG10	ce3um-ca5	101.4	74.8	-65.9	-7.6	68.8	-98.6	-55.8	-76.9
caOoG11	ce3um-ce5	105.1	74.2	-66.5	-6.5	69.2	-155.0	-58.0	-87.5
taOoG43	te3um-ta9	105.3	66.6	-68.9	-4.4	81.2	-82.7	-176.3	-163.4
caOoG12	ctb2m-ca6	106.3	24.9	43.6	-8.8	51.4	-98.0	-54.6	177.8
caOoG13	ctb2m-ce6-A	107.6	25.4	43.4	-8.0	50.5	-156.3	-55.7	-175.5
caOoG14	ce4dp-ca1-A	107.6	-45.2	70.1	-5.4	-118.7	-71.6	50.1	91.7

caOoG15	ctb2m-ca5	108.0	24.3	44.2	-8.4	51.7	-98.0	-55.3	-77.7
caOoG16	ce4dp-ca1-B	108.4	-29.9	68.3	-9.2	-117.7	-65.9	48.4	104.2
caOoG17	ce4dp-ca1-C	108.6	-22.3	63.3	-19.2	-116.6	-72.2	51.4	95.6
taOoG44	ttw1p-ta1	109.1	-48.0	22.5	50.3	-113.5	-87.9	80.1	75.0
caOoG18	ctb2m-ca1-B	109.6	17.5	50.2	-7.7	55.9	-71.6	52.0	91.8
caOoG19	ctb2p-ce8	111.1	-26.4	67.3	-6.9	-140.3	-155.8	-174.4	-51.1
caOoG20	ctb2m-ce5	112.0	24.9	43.7	-8.4	51.3	-152.5	-57.6	-88.8
caOoG21	ce4dp-ce1	112.7	-37.8	69.6	-4.2	-128.7	-158.8	67.6	70.7
caOoG22	ctb2m-ce6-B	113.7	6.2	53.7	-11.8	156.8	-159.4	-56.1	-176.7
caOoG23	ce4um-ca1	117.8	22.1	-66.6	15.1	129.9	-89.6	69.4	76.8
caOoG24	ctw1p-ca1	135.2	-65.3	16.3	46.1	-74.8	-73.2	55.8	85.0
caOoG25	ce4dm-ca2	141.3	4.9	51.4	-19.0	165.1	-80.1	89.6	-68.4

^a The conformers are depicted in Figure S1. The reference energy value V_0 is given in Table S1. The φ_4 , φ_5 , and φ_6 torsions are defined in Table S1. The torsion angles φ_7 , φ_8 , φ_9 , and φ_{10} are: Cu–O13–C13–C12, O13–C13–C12–C11, Cu–C12–C11–O11, HO13–O13–Cu–O11. The atom labels are defined in Figure 5-

Table S10. Characteristic torsion angles ($^{\circ}$) and B3LYP/BS0 relative electronic energy ΔV (kJ mol $^{-1}$) of 38 Cu(L-aThr) $_2$ conformers in the Oo-Oo coordination mode^a

name	conformer	ΔV	φ_7	φ_8	φ_9	φ_{10}	φ_{14}	φ_{15}	φ_{16}	φ_{17}
taOoOo1	te3dm-te3dm	99.7	-56.2	68.3	-6.8	153.7	-56.1	68.2	-6.8	153.7
taOoOo2	te3dm-ttb2p-A	102.6	2.2	51.9	-13.3	-173.1	-59.3	68.3	-5.2	155.7
taOoOo3	te3um-te4dm-A	109.0	63.1	-69.3	-7.2	87.9	-52.8	68.2	-8.4	150.5
taOoOo4	te3dm-te4dm-A	114.1	-56.7	68.8	-6.5	152.2	-46.3	71.5	-16.8	149.9
taOoOo5	te3dm-ttb2p-B	114.8	-59.8	68.3	-5.3	156.4	2.3	54.6	-13.9	-173.2
taOoOo6	te3dm-te4dm-B	116.2	-58.0	68.4	-5.9	154.1	-18.5	61.3	-15.8	167.6
taOoOo7	te4dm-tbt2p-A	117.9	3.7	51.2	-13.5	-176.4	-46.8	72.0	-15.5	150.5
taOoOo8	ttb2p-ttb2p	118.2	-3.7	54.8	-11.2	-168.7	-1.7	57.8	-11.9	-170.0
taOoOo9	te4dm-ttb2m	118.5	-13.1	56.4	-13.9	171.1	-47.0	72.2	-15.6	149.3
taOoOo10	te4dm-ttb2p-B	119.8	0.9	53.2	-11.8	-172.1	-20.1	61.7	-15.5	171.0
taOoOo11	tbt2p-te4up	122.1	-8.7	58.1	-7.8	-158.1	48.9	-68.5	19.0	-156.6
taOoOo12	te3dm-te4up	122.5	-60.2	66.8	-5.9	161.0	52.1	-68.7	16.4	-159.9
taOoOo13	te3um-te4dm-B	122.7	62.2	-69.7	-7.3	89.1	-42.8	70.7	-17.0	145.8
taOoOo14	te3um-ttb2p	122.8	63.6	-72.0	-0.2	82.9	3.8	54.8	-12.7	-172.6

taOoOo15	tbt1m-te4dm	124.5	-54.9	67.8	-8.1	154.8	19.1	-63.4	9.6	151.7
taOoOo16	ttb1m-ttb2p	125.5	0.9	53.0	-12.7	-168.0	10.7	-58.9	14.1	163.1
taOoOo17	te3um-te4up	132.0	71.6	-68.7	1.3	70.7	47.6	-68.1	18.4	-153.0
taOoOo18	tbt1m-te3um	134.1	66.8	-67.3	-9.0	83.8	16.8	-62.7	9.5	150.7
taOoOo19	ttb2p-ttw1m	137.8	0.1	53.7	-12.0	-165.2	-71.8	39.3	34.1	160.3
taOoOo20	te3dm-ttw1m	138.0	-55.9	67.7	-7.6	156.7	-72.2	43.9	30.8	158.0
caOoOo1	ce3um-ctb2m-A	141.1	-13.8	62.6	-11.2	129.1	75.8	-65.3	-8.2	67.4
caOoOo2	ce3um-ctb1p	152.8	61.8	-77.9	13.0	74.8	27.6	-66.2	11.8	-139.5
caOoOo3	ctb2p-ctb2p-A	154.4	1.5	55.7	-12.6	-170.2	-0.3	56.7	-12.4	-167.5
caOoOo4	ce3um-ctb2m-B	156.1	75.5	-66.0	-6.8	67.3	-20.0	68.7	-11.2	125.1
caOoOo5	ctb1p-ctw1p-A	156.8	-14.0	-58.7	1.2	-49.7	-59.7	29.2	49.8	-110.2
caOoOo6	ce3um-ctb2m-C	158.6	74.0	-64.8	-10.8	72.7	16.0	50.6	-11.7	64.0
caOoOo7	ctb2m-ctb2m-A	161.9	24.6	44.6	-8.1	50.6	-20.3	68.3	-12.8	127.9
caOoOo8	ctb2m-ctb2m-B	163.5	-10.4	60.6	-12.4	136.3	22.5	46.9	-9.5	51.7
caOoOo9	ctb1p-ctw1p-B	164.0	-13.8	-58.6	5.9	-173.4	-63.3	20.7	50.3	-84.2
caOoOo10	ce4up-ctb2m-A	164.5	25.3	43.8	-7.4	49.9	45.8	-67.4	9.8	-155.7
caOoOo11	ce4um-ce4um	165.8	59.0	-77.3	17.4	77.0	35.0	-69.8	19.1	104.3
caOoOo12	ctb2p-ctb2p-B	170.2	6.9	53.3	-12.4	-179.7	-25.9	72.3	-9.2	-138.8
caOoOo13	ce4up-ctb2p	170.7	3.4	54.1	-12.7	-176.5	31.6	-69.8	6.8	-125.3
caOoOo14	ctb2p-ctb2p-C	170.9	3.6	55.0	-12.5	-171.9	-2.8	59.8	-14.0	-167.6
caOoOo15	ce4up-ctb2m-B	177.7	23.6	44.9	-7.7	52.1	29.2	-66.4	17.2	-141.7
caOoOo16	ce4up-ctb2m-C	178.0	21.9	46.2	-7.5	53.8	21.3	-65.6	16.7	-133.3
caOoOo17	ce4um-ctb2m	179.7	21.0	46.5	-9.5	56.6	44.3	-70.6	14.3	112.8
caOoOo18	cbt2m-ce4um	182.9	-4.6	61.3	-9.8	105.6	60.0	-71.1	21.9	80.4

^a The conformers are depicted in Figure S1. The reference energy value V_0 is given in Table S1. The φ_7 , φ_8 , φ_9 , and φ_{10} torsions are defined in Table S2. The torsion angles φ_{14} , φ_{15} , φ_{16} , and φ_{17} are: Cu–O23–C23–C22, O23–C23–C22–C21, Cu–C22–C21–O21, HO23–O23–Cu–O21. The atom labels are defined in Figure 5.

Table S11. Characteristic torsion angles ($^{\circ}$) and B3LYP/BS0 relative electronic energy ΔV (kJ mol^{-1}) of 13 Cu(L-*a*Thr)₂ conformers in the Oo-No coordination mode^a

name	conformer	ΔV	φ_7	φ_8	φ_9	φ_{10}	φ_{18}	φ_{19}	φ_{20}
taOoNo1	te4dp-tap	108.5	-31.2	68.3	-8.1	-117.4	55.1	-31.1	-140.6
caOoNo1	ctb1p-cem	157.9	-9.8	-61.5	7.0	-176.3	145.3	-13.1	89.8

taOoNo2	te3um-tep	158.9	69.5	-72.9	5.9	68.9	144.7	-12.4	-152.5
caOoNo2	ce3um-cep	165.4	66.2	-76.0	11.1	70.2	150.9	-15.1	-151.5
taOoNo3	te4dp-tep	170.3	-43.2	68.9	-0.6	-129.2	140.1	-9.7	-156.0
taOoNo4	te4dm-tep	172.4	-48.7	68.4	-3.6	159.7	140.6	-9.8	-156.7
caOoNo3	ce3um-cem	173.7	62.8	-75.3	4.7	79.3	156.3	-15.5	112.0
taOoNo5	ttb2m-tep	177.6	21.7	44.6	-9.3	57.6	144.1	-12.5	-154.6
caOoNo4	ctb2m-cep	180.3	23.0	44.4	-10.7	55.9	152.1	-15.2	-150.5
caOoNo5	ce4dm-cem	181.3	-43.2	69.8	-5.0	145.5	150.7	-14.4	96.6
caOoNo6	ce4dp-cep	181.4	-44.8	70.0	-3.1	-121.9	156.7	-15.9	-142.3
taOoNo6	ttb2m-tem-A	182.4	-8.3	56.6	-14.2	178.3	153.6	-13.8	111.8
taOoNo7	ttb2m-tem-B	187.8	22.8	43.9	-10.4	57.3	157.5	-15.3	116.9

^a The conformers are depicted in Figure S1. The reference energy value V_0 is given in Table S1. The φ_7 , φ_8 , φ_9 , and φ_{10} torsions are defined in Table S2. The torsion angles φ_{18} , φ_{19} , and φ_{20} are: Cu–N21–C22–C21, N21–C22–C21–O21, HO23–O23–Cu–N21. The atom labels are defined in Figure 5.

Table S12. Characteristic torsion angles ($^{\circ}$) and B3LYP/BS0 relative electronic energy ΔV (kJ mol $^{-1}$) of 4 Cu(L- α Thr) $_2$ conformers in the No-No coordination mode^a

name	conformer	ΔV	φ_{11}	φ_{12}	φ_{13}	φ_{18}	φ_{19}	φ_{20}
caNoNo1	cep-cep	223.9	153.4	-15.3	-146.8	153.7	-15.4	-146.8
taNoNo1	tep-tep	227.1	150.5	-14.4	-152.0	150.5	-14.4	-151.7
caNoNo2	cem-cem	229.8	150.9	-14.5	100.5	152.0	-14.7	101.7
taNoNo2	tam-tam	257.0	108.0	1.4	171.2	104.3	3.7	170.7

^a The conformers are depicted in Figure S1. The reference energy value V_0 is given in Table S1. The φ_{11} , φ_{12} , and φ_{13} torsions are defined in Table S3. The φ_{18} , φ_{19} , and φ_{20} torsions are defined in Table S5.

Table S13. Means and standard deviations (in parentheses) of B3LYP/BS0 Cu–donor in-plane bond lengths (in Å) and six valence angles (in degree) around the copper atom in a number of conformers (*n*) of *trans*- and *cis*-Cu(L-Thr)₂ conformers in six possible coordination modes.

Internal coordinate	G-G		No-G		Oo-G		Oo-Oo		Oo-No		No-No	
	<i>trans</i>	<i>cis</i>										
<i>n</i>	35	22	16	16	39	23	13	15	7	6	1	3
Cu–N _{am}	2.04(1)	2.068(9)	2.00(3)	2.02(4)							2.000	2.04(3)
Cu–N _{am'}	2.033(5)	2.064(7)	2.05(1)	2.06(2)	2.03(1)	2.034(8)			1.992(3)	2.01(2)	1.998	2.04(4)
Cu–O	1.94(1)	1.942(1)			1.91(1)	1.90(2)	1.90(1)	1.92(4)	1.90(1)	1.92(1)		
Cu–O'	1.932(7)	1.936(7)			1.924(9)	1.95(2)	1.889(6)	1.90(3)				
Cu–O _h			1.94(1)	1.92(2)	2.03(1)	2.11(3)	2.02(2)	2.09(6)	2.04(1)	2.08(5)	2.087	2.11(4)
Cu–O _{h'}			2.05(6)	2.06(3)			2.022(7)	2.08(5)	2.03(1)	2.04(6)	2.087	2.03(8)
N _{am} –Cu–O	83.6(9)	82.3(8)										
N _{am'} –Cu–O'	84.2(6)	82.7(4)	83.7(9)	83(1)	83.9(8)	83.2(9)						
N _{am} –Cu–N _{am'}	177(3)	100(1)	175(3)	104(2)							177.1	104.3(8)
N _{am} –Cu–O'	96.1(6)	176(2)	95.3(9)	170(3)								
N _{am'} –Cu–O	96(1)	176(2)			94(1)	167(5)			166(3)	94.2(9)		
N _{am'} –Cu–O _{h'}									82.5(7)	83(1)	81.7	83(1)
N _{am} –Cu–O _h			99(2)	169(11)	171(2)	93(1)			96(1)	170(1)	98.7	172.2(9)
N _{am} –Cu–O _h			83(1)	82(1)							81.5	82(1)
N _{am} –Cu–O _{h'}											98.1	170(3)
O–Cu–O'	177(2)	95.5(4)			174(3)	98(3)	172(5)	100(4)				
O _h –Cu–O					93(1)	91(3)	93(1)	91(3)	93.9(8)	94(2)		
O _h –Cu–O'			173(3)	92(2)	90(2)	160(10)	89(3)	154(13)				
O'–Cu–O _{h'}							94(1)	94(2)				
O–Cu–O _{h'}							85(1)	150(17)	89.5(8)	173(5)		
O _h –Cu–O _{h'}							172(5)	87(5)	169(6)	89(2)	179.4	91(2)

Table S14. Means and standard deviations (in parentheses) of B3LYP/BS0 Cu–donor in-plane bond lengths (in Å) and six valence angles (in degree) around the copper atom in a number of conformers (*n*) of *trans*- and *cis*-Cu(L-*a*Thr)₂ conformers in six possible coordination modes

<i>n</i>	G-G		No-G		Oo-G		Oo-Oo		Oo-No		No-No	
	<i>Trans</i>	<i>cis</i>										
	68	45	15	15	44	25	20	18	7	6	2	2
Cu–N _{am}	2.04(1)	2.07(1)	2.00(3)	2.01(2)							2.002(1)	2.024(2)
Cu–N _{am'}	2.037(9)	2.068(9)	2.05(1)	2.06(2)	2.04(1)	2.04(1)			2.00(2)	2.007(2)	2.003(1)	2.025(3)
Cu–O	1.93(1)	1.94(1)			1.91(2)	1.90(2)	1.89(1)	1.91(2)	1.89(2)	1.92(2)		
Cu–O'	1.930(8)	1.935(8)	1.94(1)	1.92(2)	1.92(1)	1.96(5)	1.888(4)	1.91(2)	2.200			
Cu–O _h				2.06(3)	2.03(4)	2.12(6)	2.01(2)	2.09(5)	2.06(4)	2.08(5)	2.08(4)	2.126(7)
Cu–O _{h'}						2.02(1)	2.09(8)	2.07(5)	2.08(3)	2.08(4)	2.08(4)	2.126(6)
N _{am} –Cu–O	84.0(8)	82.4(6)	83.8(8)	83(1)								
N _{am'} –Cu–O'	84.2(8)	82.7(5)			83.8(8)	83(1)						
N _{am} –Cu–N _{am'}	177(2)	99.7(8)	177(2)	103(1)							175(1)	106.5(1)
N _{am} –Cu–O'	96.0(8)	176(2)	95.3(8)	171(2)	94(2)	165(9)						
N _{am'} –Cu–O	95.9(6)	176(2)							167(4)	95(2)		
N _{am'} –Cu–O _{h'}					171(8)	94(3)			82(2)	83.1(6)	81.6(1)	83(1)
N _{am'} –Cu–O _h			98(1)	167(13)					97.3(7)	170(2)	98.8(4)	166.6(7)
N _{am} –Cu–O _h			83(1)	82.4(6)							81.4(3)	83(1)
N _{am} –Cu–O _{h'}											98.8(6)	167.0(7)
O–Cu–O'	177(2)	95.5(4)			174(3)	100(6)	172(4)	101(4)				
O _h –Cu–O					92(2)	89(4)	92.9(8)	90(4)	92(3)	92(3)		
O _h –Cu–O'			172(3)	93(3)	90(3)	156(25)	89(2)	155(9)				
O'–Cu–O _{h'}							93(1)	92(3)				
O–Cu–O _{h'}							86(2)	152(18)	91(2)	173(6)		
O _h –Cu–O _{h'}							173(4)	89(7)	170(5)	89(1)	174(6)	89(1)

Table S15. The most stable vacuum *trans* and *cis* Cu(L-Thr)₂ and Cu(L-*a*Thr)₂ conformers in the six coordination modes, which were the starting structures for B3LYP/BS0 geometry optimization using PCM, their corresponding aqueous minima, and their relative vacuum electronic energies, and relative aqueous electronic and Gibbs free energies (ΔV and ΔG , in kJ mol⁻¹)^{a,b}

Cu(L-Thr) ₂					Cu(L- <i>a</i> Thr) ₂				
vacuum conformer	ΔV_{vacuum}	aqueous conformer / mode ^c	ΔV_{water}	ΔG_{water}	vacuum conformer	ΔV_{vacuum}	aqueous conformer / mode ^c	ΔV_{water}	ΔG_{water}
G-G									
te1-te1	0.0		0.0	0.0	te8-te8	7.4		-6.4	-0.3
te1-te8	7.0		1.5	3.1	te1-te8	8.6		2.2	6.0
te1-ta6	10.5		6.1	3.9	te1-te1	10.5		11.0	12.2
ta5-te1	10.9		4.3	4.6	te8-ta1	16.5		2.7	7.6
te3-ta5 (X-ray) ^d			6.4	2.9	te8-ta6	16.9		4.8	6.5
te1-te6	13.8		12.5	11.0	ca1-ca1	51.1		16.6	14.8
ce1-ce1	43.8		5.5	3.7	ce1-ca1	51.7		15.2	19.1
ce1-ca1	49.5		11.4	13.3	ce8-ca1	52.0		6.5	10.7
ce1-ca5	51.0		8.8	10.8	ce1-ce1	53.2		15.5	16.9
ce1-ca6	51.9		10.8	10.8	ce1-ce8	55.2		7.1	12.0
ce1-ce8	54.6		6.8	9.6	ce8-ce8	56.9		-1.1	4.4
ce8-ce8	65.0		8.7	13.9					
No-G									
tap-te1	41.3		44.8	41.1	tap-ta1	54.1		50.0	52.9
tap-ta1	56.1		53.5	55.3	tap-ta3	59.2		50.2	51.5
tap-ta2	61.4	tap-ta3	55.0	52.2	cap-ca3-A	81.4		50.3	54.8
cap-ca4	62.8		36.9	44.5	cap-ca3-B	85.6		52.4	60.2
cap-ca2-A	75.0	cap-ca2	65.8	76.0	cam-ca3	94.8		46.6	53.8
cap-ca3A	82.8	cap-ca3	44.8	55.6	cap-ca1	104.8		80.7	87.0
cap-ce6	84.1	ca1-ce6 / G-G	23.7	21.2	cep-ce1	119.0		57.2	60.3
cap-ca2-B	85.2	cap-ca3	44.8	55.6	cep-ce8	119.4		49.6	53.4
cap-ca3-B	89.7		54.5	64.1	cep-ca1	122.6		55.7	58.4
cap-ca3-C	91.7		52.6	53.1	tem-ta1	127.5		58.6	61.9
tem-te1	116.7		51.6	49.8	tep-te1	130.0		59.5	60.1
tep-te1	117.1		50.2	48.5	tem-te1	130.4		61.1	63.9
Oo-G									

te4dm-te1-A	49.4	ttw2m-te1	54.8	48.8	te4dm-te8	55.6		52.9	54.0
tbt2p-te1-A	53.5	tbt2p-te1	56.8	47.9	ttb2p-te8	56.8	ttb2m-te8	55.5	53.0
ttw1m-te1	56.6		57.6	50.9	te4dm-te1	57.1	te3dm-te1	61.3	59.4
te4dm-te8	57.2	ttw2m-te8	56.7	57.4	ttb2p-te1	57.3	ttb2m-te1	63.7	59.3
te4dm-ta5	60.1	ttw2m-ta5	59.1	56.2	ttb2m-te8	57.5	te4dm-te8	52.9	54.0
ctw1m-ce1	86.5		60.3	59.1	ce3um-ca1	89.6		64.7	64.6
cbt1m-ca1	86.7		69.3	75.9	ce3um-ce8	92.2		58.7	58.7
ctw1m-ca1	90.9		65.9	61.5	ce3um-ce1	92.5		66.9	64.3
ctw1m-ca6	94.2		66.0	62.9	ctb2m-ca1-A	95.5	ctb2m-ca1	71.9	67.9
ctw1m-ca5	94.6		64.1	62.2	ce3um-ca6	99.8		69.1	68.9

Oo-Oo

te4dm-te4dm-A	95.5	te3dm-te3dm	109.4	104.1	te3dm-te3dm	99.7		110.8	104.5
te4dm-te4dm-B	107.7	te3dm-te4dm	113.6	106.0	te3dm-ttb2p-A	102.6		114.0	105.4
tbt2p-te3dm	112.4	tbt2p-ttw1m	115.4	109.4	te3um-te4dm-A	109.0	te3dm-te3um	114.3	111.9
tbt2p-tbt2p	117.3	tbt2p-tbt2p	117.1	108.7	te3dm-te4dm-A	114.1		116.8	110.8
tbt4m-te4dm	117.3		116.6	113.9	te3dm-ttb2p-B	114.8		117.8	111.0
tbt2p-te3um	122.1	tbt2p-te3um	118.3	110.5	te3dm-te4dm-B	116.2		116.8	110.8
ce3um-ctb1m	126.8	ce3um-ca3 / Oo-G	65.7	63.7	ce3um-ctb2m-A	141.1	ce3um-ce4dm	123.3	112.7
ctb2m-ctw1m-A	137.7	ce3um-ce4dm	118.6	111.5	ce3um-ctb1p	152.8	ce3um-ce4up	123.1	120.1
cbt2m-ctb1m	150.7	ce4dm-ca3 / Oo-G	72.7	69.1	ctb2p-ctb2p-A	154.4	cbt2m-ctb2m	129.8	119.5
ctb2m-ctw1m-B	151.4	ce3um-ce3up	123.9	121.1	ce3um-ctb2m-B	156.1	ce3um-ce4dm	125.9	120.8
ce3um-ctb1p	154.2	ce3um-ce3up	123.9	121.1	ctb1p-ctw1p-A	156.8	ce4dp-ca1 / Oo-G	75.9	74.0
					ce3um-ctb2m-C	158.6	ce3um-ce4dm	125.9	120.7
					ctb2m-ce4um	182.9	ce4um-ctb2m	135.3	125.0

Oo-No

cbt2p-cap	96.2		98.9	98.2	te4dp-tap	108.5	ce3dp-ca1 / Oo-G	68.9	69.3
cbt1m-cem	151.2	tem-ta3 / No-G	56.0	56.5	ctb1p-cem	157.9	tem-ta3 / No-G	58.4	59.5
te3um-tep	151.5		104.9	102.4	te3um-tep	158.9		110.0	109.1
ttw1m-tem	159.9		109.4	109.4	ce3um-cep	165.4		111.2	110.3
te4dp-tep	162.3	ttw2p-tep	109.5	107.9	te4dp-tep	170.3	te3dp-tep	114.1	111.7
ctw1m-cem	162.7		108.9	107.1	te4dm-tep	172.4		114.3	113.3
te4dm-tep	163.4	ttw2m-tep	109.4	107.8	ce3um-cem	173.7		114.5	114.7
te4dm-tam	169.5		109.8	106.7	ttb2m-tep	177.6		116.8	112.8
ce4dm-cem	169.8		110.4	107.5	ctb2m-cep	180.3	cbt2m-cep	117.7	112.5
ctb2m-cep	173.0	ctb2m-cep	111.3	105.8	ce4dm-cem	181.3	ctw2m-cem	115.8	113.2

No-No					No-No			
<i>cap-cap</i>	99.7	<i>cap-ca1 / No-G</i>	57.6	58.3	<i>cep-cep</i>	223.9		109.7 112.9
<i>cep-cep</i>	211.7		101.9	101.2	<i>tep-tep</i>	227.1		112.6 113.4
<i>cem-cem</i>	212.8		105.5	105.1	<i>cem-cem</i>	229.8		113.9 112.4
<i>tep-tep</i>	212.8		104.0	103.7	<i>tam-tam</i>	257.0	<i>ta3-ta3 / G-G</i>	19.0 25.4

^a The conformers are defined in Figure S1 and Tables S1-S12. The reference energy values are those of the Cu(L-Thr)₂ te1-te1 conformer:

$V_{\text{vacuum}} = -1071.76257129$ a.u., $V_{\text{water}} = -1071.80072926$ a.u., $G_{\text{water}} = -1071.586956$ a.u.

^b The conformers with H_h in the carboxylate group are denoted in italics.

^c The names of the PCM optimized aqueous conformer and coordination mode are written only if the starting conformation and/or coordination mode changed during geometry optimization.

^d The starting conformer from the X-ray crystal and molecular structure (Ref. 17).

Table S16. The most stable B3LYP/BS0 vacuum *trans* and *cis* Cu(L-Thr)₂ and Cu(L-*a*Thr)₂ conformers in the six coordination modes, which were the starting structures for the B3LYP/BS1 geometry optimization in the gas phase and in aqueous solution using PCM, and their corresponding B3LYP/BS1 relative electronic and Gibbs free energies (ΔV and ΔG , in kJ mol⁻¹)^{a,b,c}

Cu(L-Thr) ₂				Cu(L- <i>a</i> Thr) ₂			
starting conformer	ΔV_{vacuum}	ΔV_{water}	ΔG_{water}	starting conformer	ΔV_{vacuum}	ΔV_{water}	ΔG_{water}
G-G							
te1-te1	0.0	0.0	0.0	<i>te8-te8</i>	6.0	-5.7	0.8
<i>te3-ta5</i> (X-ray) ^d	-	6.4	5.2	<i>ca1-ca1</i>	55.1	20.8	22.9
ce1-ce1	41.0	5.0	4.3	No-G			
No-G				<i>tap-ta1</i>	56.3	53.3	57.1
<i>tap-te1</i>	41.8	45.5	47.1	<i>tem-ta1</i>	134.7	67.6	73.0
cap-ca4	70.5	44.4	54.5	<i>cap-ca3-A</i>	84.4	57.5	62.6
Oo-G							
te4dm-te1-A	46.9	52.3	47.4	<i>te4dm-te8</i>	52.4	51.0	50.4
		ttw2m-te1					
ctw1m-ce1	84.5	60.4	59.9	<i>ce3um-ca1</i>	93.1	68.9	70.2
Oo-Oo							
te4dm-te4dm-A	90.7	104.3	99.6	<i>te3dm-te3dm</i>	95.3	106.7	100.8
		te3dm-te3dm		<i>ce3um-ctb2m-A</i>	139.2	120.0	114.2
ce3um-ctb1m	131.8	68.6	65.7			ce3um-ce4dm	

		ce3um-ca3/Oo-G					
Oo-No					Oo-No		
<i>ctb2p-cap</i>	93.9	98.0	98.1		<i>te4dp-tap</i>	114.3	70.7 72.4
<i>te3um-tep</i>	155.1	110.2	110.4		<i>ctb1p-cem</i>	134.7	ce3dp-ca1/Oo-G 67.1 70.2
No-No					<i>te3um-tep</i>	164.2	tem-ta1/No-G 116.2 115.7
<i>cap-cap</i>	89.9	59.8	60.7		<i>ce3um-cep</i>	172.9	118.0 117.6
		<i>cap-ca1/No-G</i>					
<i>cep-cep</i>	222.2	113.7	113.1		No-No		
<i>tep-tep</i>	220.8	114.3	116.5		<i>cep-cep</i>	237.1	122.4 125.1
					<i>tep-tep</i>	237.8	123.8 126.2

^a The conformers are defined in Figure S1 and Tables S1-S12. The reference B3LYP/BS1 energy values are of the Cu(L-Thr)₂ te1-te1 conformer: $V_{\text{vacuum}} = -2516.37645582$ a.u., $V_{\text{water}} = -2516.41165205$ a.u., $G_{\text{water}} = -2516.199097$ a.u.

^b The conformers with H_h in the carboxylate group are denoted in italics.

^c The names of the B3LYP/BS1 vacuum and/or PCM optimized aqueous conformers and coordination mode are written under the relative energy value only if the starting conformation and/or coordination mode changed during geometry optimization.

^d The starting conformer from the X-ray crystal and molecular structure (Ref. 17).

Table S17. Imaginary frequencies, ν (cm⁻¹), of the TS structures calculated by DFT/B3LYP and MP2 with the BS0, BS1, and BS2 basis sets, and the Q_{tunnel} factor calculated using Eq. 2.^a

TS	B3LYP/BS0		B3LYP/BS1		B3LYP/BS2		MP2/BS2	
	ν	Q_{tunnel}	ν	Q_{tunnel}	ν	Q_{tunnel}	ν	Q_{tunnel}
$\text{Cu}(\text{L}-\alpha\text{Thr})_2$								
TS1	78.9i	1.006	71.9i	1.005	78.2i	1.006		
TS2	78.1i	1.006	75.7i	1.006	76.8i	1.006		
TS3	31.9i	1.001	25.4i	1.001	23.7i	1.001		
TS4	35.0i	1.001	30.5i	1.001	35.7i	1.001		
TS' _{trans↔cis}	46.6i	1.002	67.8i	1.005	52.1i	1.003	43.7i	1.002
TS' _{trans↔cis,a}	68.9i	1.005	88.5i	1.008	71.9i	1.005		
TS6	39.6i	1.002	40.7i	1.002	34.0i	1.001		
TS5	43.9i	1.002	47.0i	1.002	46.0i	1.002		
TS" _{trans↔cis}	192.7i	1.037	177.3i	1.031	217.7i	1.048	222.5i	1.050
TS" _{trans↔cis,a}	213.7i	1.046	207.5i	1.043	238.8i	1.058		
TS _{Oo↔G,2}	76.2i	1.006	63.6i	1.004	65.2i	1.004	97.3i	1.009
TS _{Oo↔G,3}	80.8i	1.006	60.7i	1.004	67.5i	1.004		
$\text{Cu}(\text{L}-\text{Thr})_2$								
TS _{trans↔cis,1}	229.0i	1.053	210.2i	1.044	210.1i	1.044	230.1i	1.053
TS _{trans↔cis,2}	181.1i	1.033	179.8i	1.032	252.7i	1.065	225.7i	1.051
TS7	87.0i	1.007	83.4i	1.007	82.8i	1.007	87.0i	1.007
TS _{Oo↔G,1}	43.9i	1.002	33.3i	1.001	39.1i	1.002	61.4i	1.004

^a The TS structures are illustrated in Figures S2 and S3, and their corresponding minimum structures are given in Table 6.