

Table 11. Geometrical parameters for the optimized [(CO)₂Re(C₅H₄)C≡C(C₅H₄)Re[(CO)₂](alkane) complexes, calculated at the RIMP2/ 6-31G*, LANL2DZ, RIMP2-VDZ (auxiliary basis set) level of theory

RIMP2	<i>Interacting</i>			<i>M-H-</i>	<i>C-H</i>	<i>Average</i>
	<i>Pair of</i>	<i>M-H</i>	<i>M-C</i>	<i>C</i>	<i>bound</i>	<i>C-H free</i>
	<i>Atoms</i>	<i>Distance</i>	<i>distance</i>	<i>angle</i>	<i>distance</i>	<i>Distance</i>
Methane	Re ₁ -C	1.98	2.62	111.28	1.15	1.09
	Re ₂ -C	1.99	2.64	111.93	1.15	1.09
Ethane	Re ₁ -C1	1.92	2.58	111.98	1.15	1.09
	Re ₂ -C2	1.92	2.58	111.57	1.15	1.09
Propane	Re ₁ -C1	1.91	2.70	121.88	1.15	1.09
	Re ₂ -C3	1.91	2.57	111.49	1.15	1.10
Butane	Re ₁ -C1	1.95	2.61	112.84	1.15	1.09
	Re ₂ -C4	1.95	2.61	112.42	1.15	1.09
Pentane	Re ₁ -C1	1.91	2.74	126.30	1.14	1.09
	Re ₂ -C5	1.90	2.66	118.82	1.15	1.09
Cyclopropane	Re ₁ -C1	1.94	2.50	105.18	1.14	1.08
	Re ₂ -C2	1.93	2.52	106.99	1.14	1.08
Cyclopentane	Re ₁ -C1	1.89	2.66	119.96	1.16	1.09
	Re ₂ -C2	1.89	2.62	116.19	1.16	1.09
Propane	Re ₁ -C1	1.91	2.70	121.88	1.15	1.09
	Re ₂ -C3	1.91	2.57	111.49	1.15	1.10
Butane	Re ₁ -C1	1.95	2.61	112.84	1.15	1.09
	Re ₂ -C4	1.95	2.61	112.42	1.15	1.09
Pentane	Re ₁ -C1	1.91	2.74	126.30	1.14	1.09
	Re ₂ -C5	1.90	2.66	118.82	1.15	1.09

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	Re ₂ -C2	1.93	2.52	106.99	1.14	1.08
Cyclopentane	Re ₁ -C1	1.89	2.66	119.96	1.16	1.09
	Re ₂ -C2	1.89	2.62	116.19	1.16	1.09

“M-H distance” and “M-C distance” are the distances between the designated Re atom (“M” = Re₁ or Re₂) and the H and C atoms, respectively, of the C-H σ -bond that interacts most strongly with that particular Re atom (i.e. the C-H σ -bond that lies in closest proximity to that Re atom). The designations “C1”, “C2”, etc. indicate the position along the alkane chain / ring of the CH₃ / CH₂ group that is interacting with the designated Re atom. “M-H-C angle” is the angle between the Re atom and the aforementioned C-H bond. “C-H bond distance” is the length of the aforementioned C-H bond. “Average C-H free distance” is the average length of the remaining C-H bonds that are located on the same CH₃ (linear alkanes)/ CH₂ (cyclic alkanes) group as the interacting bond (in the case of cyclic alkanes, “Average C-H free distance” is simply one bond length, not an average of two lengths).