

Christopher A. Brown · Michael L. McKee

Rearrangements in icosahedral boranes and carboranes revisited

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Abstract The structure, stability, and intermolecular rearrangements between *ortho*-, *meta*-, and *para*-C₂B₁₀H₁₂ and B₁₂H₁₂²⁻ were investigated using the hybrid density functional B3LYP/6-31G(d) for vibrational frequencies, as well as B3LYP/6-311+G(2d,p) for single-point electronic energies. The general trends in free energies of rearrangement between *ortho*-C₂B₁₀H₁₂ to *meta*-C₂B₁₀H₁₂ and *meta*-C₂B₁₀H₁₂ to *para*-C₂B₁₀H₁₂ presented here are consistent with experimental reaction temperatures. In addition, the majority of the rearrangements can be viewed in terms of concerted diamond–square–diamond steps and triangular face rotations.

Keywords Deltahedron · Carboranes · Icosahedral · Triangular face rotation · Diamond–square–diamond

Introduction

The roots of modern boron cluster chemistry originated in a few seminal works by Wade [1, 2], Hart and Lipscomb [3–6], and Williams [7–9]. These compounds, which are characterized by their electron deficiencies, form non-classical three-center two-electron bonds and have been the topic of numerous texts [10–14]. This leads to the very familiar deltahedron geometries for *closo*-C_xB_{n-x}H_{n-x+2} with $n+1$ skeletal electron pairs, *nido*-C_xB_{n-x}H_{n-x+4} with

$n+2$ skeletal electron pairs, and *arachno*-C_xB_{n-x}H_{n-x+6} with $n+3$ skeletal electron pairs, determined empirically by Wade's Rules [1, 2, 15, 16]. Although a great deal of time has been spent in comparing and contrasting carborane systems of varying size in terms of stability, rearrangement, and reactivity [7, 8, 17–23], a special effort has been expended in understanding *closo*-B₁₂H₁₂²⁻, *closo*-C₂B₁₀H₁₂, and their respective derivatives [21, 24–29], which is due primarily to their unusually high stability. An eclectic array of mechanisms have been quite naturally suggested for the intermolecular rearrangement of *closo*-C₂B₁₀H₁₂, from those involving highly symmetric cubeoctahedral and anticubeoctahedral transition states to those with relatively simple triangular face rotations (TFRs) and diamond–square–diamond (DSD) steps [4, 5, 20, 25, 30–36]. Rearrangements in substituted icosahedral C₂B₁₀H₁₂ have been studied experimentally by way of ¹⁰B-labeling, halogen substitution, methyl substitution, and even tethered linkages [3, 24, 28, 36, 37] in the hopes that the electronic structure of the cage will be preserved.

Despite previous theoretical work on the stability of large carboranes and their derivatives and the aforementioned experimental work on intermolecular rearrangements in *closo*-C₂B₁₀H₁₂, there has been relatively little rigorous computational work in verifying whether any of the proposed mechanisms play a role in these intermolecular rearrangement reaction pathways. The present work confirms the presence of a TFR pathway in both the carborane and the borane case. Another pathway, which consists of concerted DSD steps and the opening of the cage into a pseudo-*nido* intermediate, is also described for both cases.

Methodology

The following calculations were done using the Gaussian03 program [38]. Species were optimized and vibrational frequencies were calculated at the B3LYP/6-31G(d) level, leading to their characterization as stationary points. In

Dedicated to Professor Dr. Paul von Ragué Schleyer on the occasion of his 75th birthday.

This article has already been published online first (DOI: <http://dx.doi.org/10.1007/s00894-005-0039-1>). Due to an oversight, this version contained several mistakes. The article is herewith republished in its entirety as an "erratum".

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C. A. Brown · M. L. McKee (✉)
Department of Chemistry and Biochemistry,
Auburn University,
179 Chemistry Building,
Auburn, AL 36849, USA
e-mail: mckeeml@auburn.edu
e-mail: cabrown@fas.harvard.edu

¹ From this point on, carborane will be used synonymously with icosahedral C₂B₁₀H₁₂ and borane with icosahedral B₁₂H₁₂²⁻.

addition, zero-point energies, integrated specific heat corrections at constant pressure, and entropies were calculated at this level. Once a continuous reaction pathway was determined, single-point electronic energies were calculated for the pertinent species at the B3LYP/6-311+G(2d,p) level. The enthalpies for the species at 500 and 800 K were approximated to be the same as the enthalpy at 298 K. These values were then used to determine the absolute free energy at 298, 500, and 800 K. The change in free energies at 298, 500, and 800 K for the species in the carborane rearrangement are relative to *ortho*-C₂B₁₀H₁₂ and, similarly, the species in the borane rearrangement are relative to B₁₂H₁₂²⁻.

Results and discussion

The reaction pathways between *ortho*-, *meta*-, and *para*-C₂B₁₀H₁₂, and B₁₂H₁₂²⁻ are shown in Figs. 1 and 2, with representative geometries in Figs. 3 and 4, respectively. At first glance, some common features between Figs. 1 and 2 should be apparent. The arrangement of the vertices and

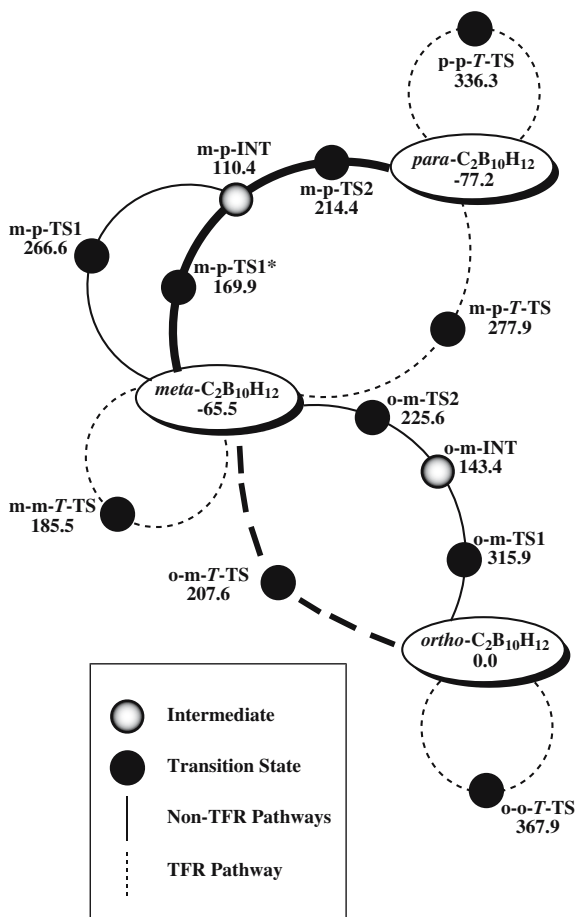


Fig. 1 Rearrangement pathways for C₂B₁₀H₁₂. Values indicate free energies in kilojoules per mole (kJ/mol) at 800 K. **Bold lines (dashed and solid)** indicate the lowest free energy pathway from *ortho*- to *meta*-C₂B₁₀H₁₂ (**bold and dashed**) and from *meta*- to *para*-C₂B₁₀H₁₂ (**bold and solid**)

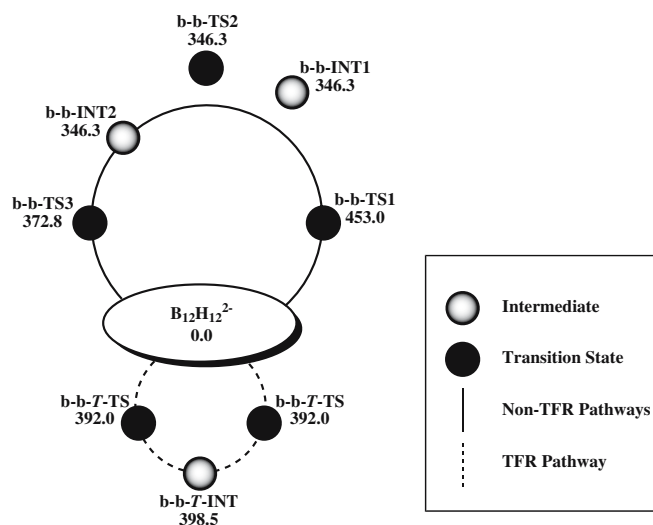


Fig. 2 Rearrangement pathways for B₁₂H₁₂²⁻. Values indicate free energies in kilojoules per mole (kJ/mol) at 800 K. The stationary points *b-b*-TS2 and *b-b*-INT1 are above the curve because the free energy barrier is essentially zero, which suggests that B₁₂H₁₂²⁻ is converted directly to *b-b*-INT2 (through *b-b*-TS1)

edges for both C₂B₁₀H₁₂ and B₁₂H₁₂²⁻ are very similar (Figs. 1 and 2). Both have two distinct pathways operating the TFR pathway and the non-TFR pathway (the non-TFR pathway is actually quite complicated and cannot be succinctly described). In addition to the graphical similarities in the schemes, the representative geometries in Figs. 3 and 4 are analogs. The *ortho*- to *meta*-C₂B₁₀H₁₂ case mimics the *meta*- to *para*-C₂B₁₀H₁₂ case, which mimics the B₁₂H₁₂²⁻ case. These similarities will prove to be useful when discussing the mechanisms of the pathways in detail.

General trends

Before discussing rearrangement mechanisms, it is important to establish some confidence by way of general trends between the stabilities and geometries of the three carborane isomers. Pertinent electronic and thermodynamic data can be found in Tables 1 and 2. Notice that the ΔG (298 K) for the three isomers decreases from *ortho*- to *meta*- to *para*-C₂B₁₀H₁₂, which is consistent with experimental observation [39]. Furthermore, the approximate transition temperatures of 450 °C for the *ortho*- to *meta*-C₂B₁₀H₁₂ rearrangement and 620 °C for the *meta*- to *para*-C₂B₁₀H₁₂ rearrangement are qualitatively consistent with the barrier heights in Fig. 1 [39–41]. Besides trends in stability, two recent gas-phase electron diffraction (GED) studies have made it possible to compare the calculated unique bond distances in the three isomers to the

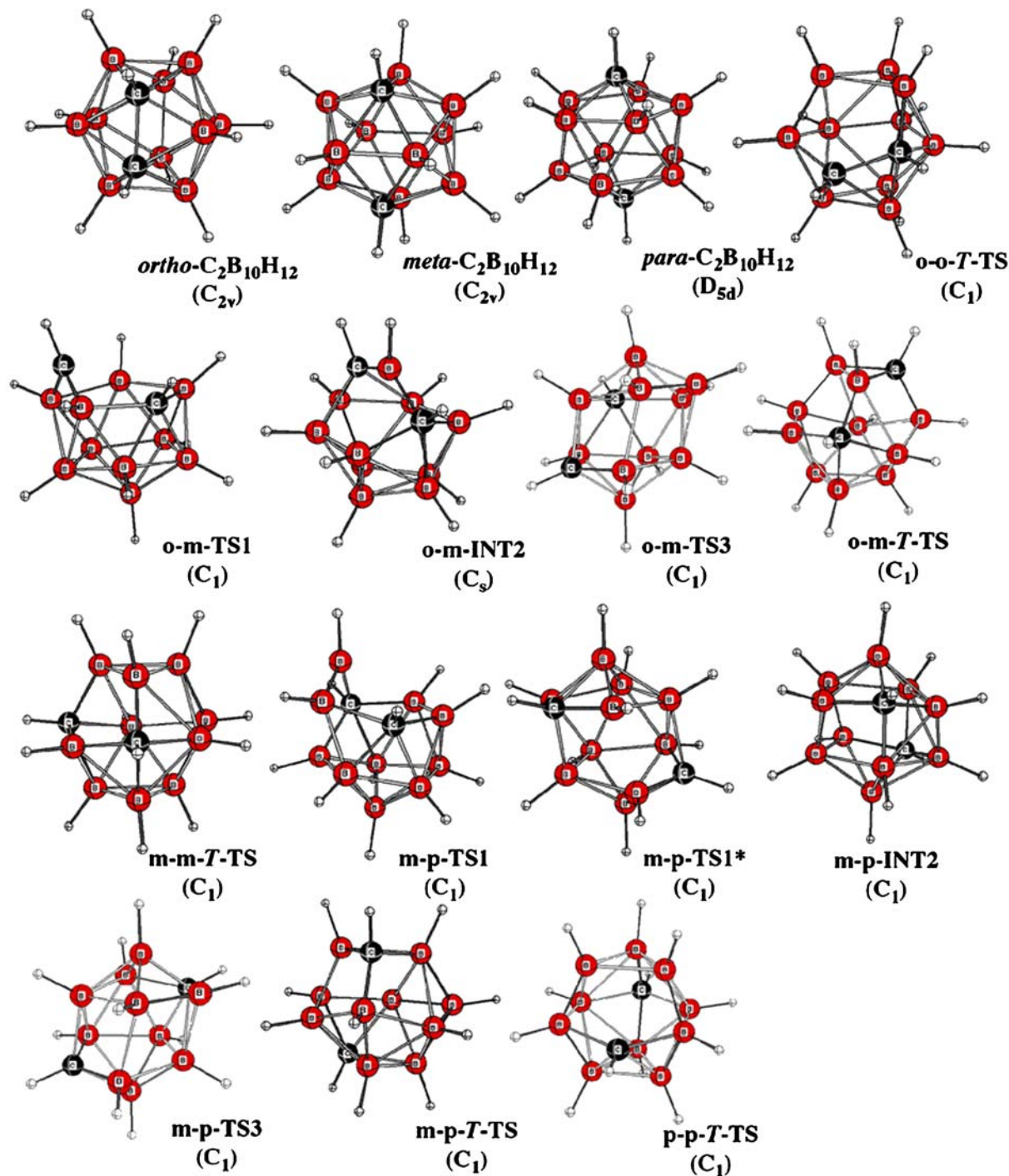


Fig. 3 Geometries for *ortho*-, *meta*-, and *para*-C₂B₁₀H₁₂. The nomenclature is as follows: the *first two letters* separated by a *dash* indicate which carboranes are being rearranged (e.g., m-p

corresponds to a *meta* to *para* rearrangement species). An *italicized T* indicates a TFR pathway species. The *last letters* (and possible *number*) indicates the type of stationary point of the species

experimentally determined unique bond distances (see Table 3 and Fig. 5) [42, 43]. In almost all of the bonds, the

difference between the calculated distance and the most recent experimentally determined distance (GED at 463 K) is around 0.01 Å.

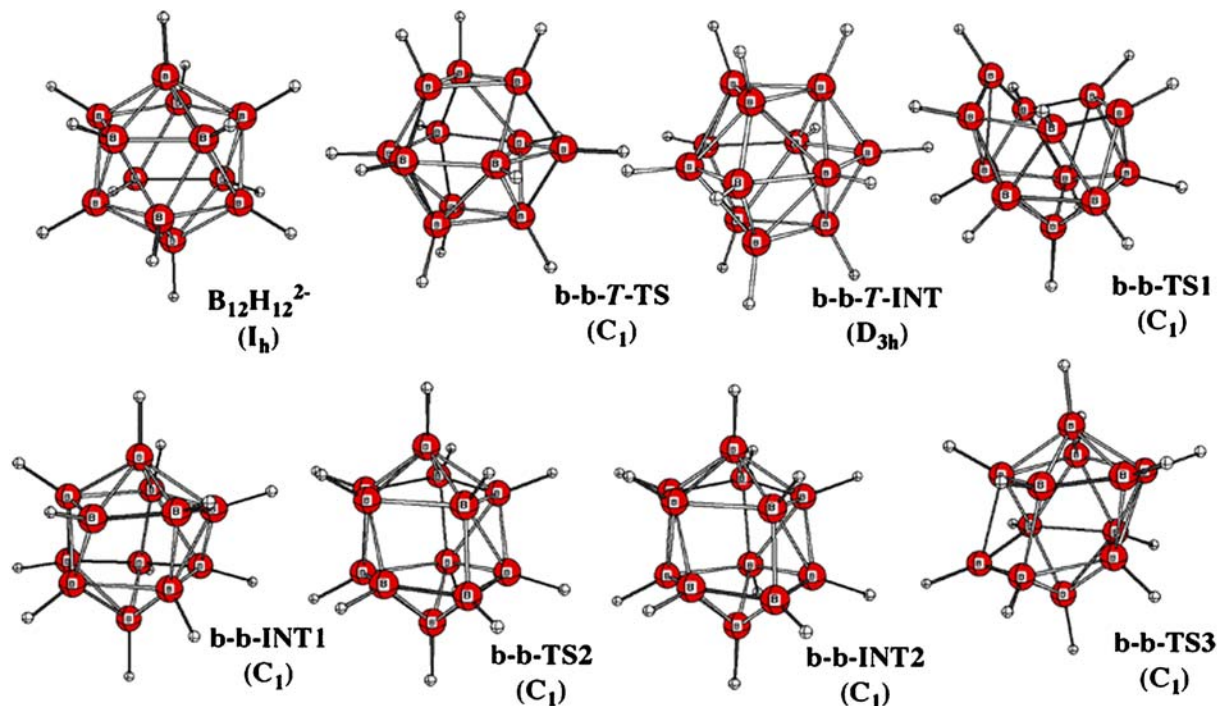


Fig. 4 Geometries of $B_{12}H_{12}^{2-}$ species. The nomenclature system is analogous to the carborane case. Consult Fig. 3 for a description

Table 1 Total electronic energies (in hartrees), zero-point energies (kJ/mol), heat capacity corrections to 298 K, and entropies (J/mol K) for species on the $C_2B_{10}H_{12}$ and $B_{12}H_{12}^{2-}$ potential energies surfaces

Structures	Total energy	Zero-point energy	$\int Cp$	Entropy
<i>ortho</i> - $C_2B_{10}H_{12}$	-332.17455	466.1	22.7	347.1
<i>meta</i> - $C_2B_{10}H_{12}$	-332.20059	468.0	22.4	345.5
<i>para</i> - $C_2B_{10}H_{12}$	-332.20500	468.0	22.4	345.7
<i>o-m</i> -TS1	-332.04031	445.7	26.6	372.2
<i>o-m</i> -INT	-332.10921	455.1	26.8	373.7
<i>o-m</i> -TS2	-332.07899	451.5	25.4	363.8
<i>o-m</i> -T-TS	-332.08224	448.5	26.7	373.5
<i>o-o</i> -T-TS	-332.01780	444.1	27.6	380.3
<i>m-m</i> -T-TS	-332.09144	449.5	26.7	372.2
<i>m-p</i> -TS1	-332.06051	448.2	26.3	370.2
<i>m-p</i> -TS1*	-332.10091	452.5	25.1	362.4
<i>m-p</i> -INT	-332.12258	455.5	26.6	371.5
<i>m-p</i> -TS2	-332.08274	451.5	25.6	365.8
<i>m-p</i> -T-TS	-332.05341	446.7	27.3	378.8
<i>p-p</i> -T-TS	-332.03032	444.5	27.7	379.4
$B_{12}H_{12}^{2-}$	-305.77394	439.1	23.1	348.7
<i>b-b</i> -T-TS	-305.60815	418.6	28.3	383.6
<i>b-b</i> -T-INT	-305.60974	419.7	29.6	373.2
<i>b-b</i> -TS1	-305.58691	419.4	27.4	377.1
<i>b-b</i> -INT1	-305.63168	423.9	26.0	367.4
<i>b-b</i> -TS2	-305.63167	423.9	26.0	367.4
<i>b-b</i> -INT2	-305.63167	423.9	26.0	367.5
<i>b-b</i> -TS3	-305.61911	422.4	26.6	374.5

Table 2 Relative enthalpies (kJ/mol), entropies (J/K·mol), and free energies (kJ/mol) for species on the C₂B₁₀H₁₂ and B₁₂H₁₂²⁻ potential energy surfaces

Structures	ΔH (0 K)	ΔH (298 K)	ΔS	ΔG (298 K)	ΔG (500 K)	ΔG (800 K)
<i>ortho</i> -C ₂ B ₁₀ H ₁₂	0.0	0.0	0.0	0.0	0.0	0.0
<i>meta</i> -C ₂ B ₁₀ H ₁₂	-66.5	-66.8	-1.5	-66.3	-66.0	-65.5
<i>para</i> -C ₂ B ₁₀ H ₁₂	-78.0	-78.3	-1.4	-77.9	-77.6	-77.2
<i>o-m</i> -TS1	332.1	336.0	25.1	328.5	323.4	315.9
<i>o-m</i> -INT	160.6	164.7	26.6	156.8	151.4	143.4
<i>o-m</i> -TS2	236.3	239.0	16.7	234.0	230.6	225.6
<i>o-m-T</i> -TS	224.7	228.8	26.4	220.9	215.5	207.6
<i>o-o-T</i> -TS	389.6	394.5	33.3	384.6	377.9	367.9
<i>m-m-T</i> -TS	201.6	205.6	25.1	198.1	193.1	185.5
<i>m-p</i> -TS1	281.6	285.1	23.1	278.2	273.6	266.6
<i>m-p</i> -TS1*	179.8	182.2	15.4	177.6	174.5	169.9
<i>m-p</i> -INT	125.9	129.9	24.4	122.6	117.7	110.4
<i>m-p</i> -TS2	226.5	229.4	18.7	223.8	220.0	214.4
<i>m-p-T</i> -TS	298.7	303.3	31.7	293.8	287.4	277.9
<i>p-p-T</i> -TS	357.1	362.1	32.3	352.5	346.0	336.3
B ₁₂ H ₁₂ ²⁻	0.0	0.0	0.0	0.0	0.0	0.0
<i>b-b-T</i> -TS	414.7	419.9	34.9	409.5	402.5	392.0
<i>b-b-T</i> -INT	411.6	418.2	24.5	410.8	405.9	398.5
<i>b-b</i> -TS1	471.4	475.7	28.4	467.2	461.5	453.0
<i>b-b</i> -INT1	358.3	361.3	18.7	355.7	351.9	346.3
<i>b-b</i> -TS2	358.3	361.3	18.7	355.7	351.9	346.3
<i>b-b</i> -INT2	358.3	361.3	18.8	355.7	351.9	346.3
<i>b-b</i> -TS3	389.8	393.4	25.8	385.7	380.5	372.8

*Ortho- to meta-carborane rearrangement*²*TFR pathway*

Figure 6 illustrates the *ortho- to meta*-C₂B₁₀H₁₂ TFR mechanism where the structures of the stationary points are similar to those in the *ortho*-C₂B₁₀H₁₂ to *ortho*-C₂B₁₀H₁₂ degenerate TFR, the *meta*-C₂B₁₀H₁₂ to *para*-C₂B₁₀H₁₂ TFR, and the borane TFR rearrangement.³ The left column in Fig. 6 displays the geometric progression through the reaction and the right column shows a two-dimensional projection of the pertinent atoms and bonds of the corresponding geometry on the left. As can be seen in the left column, a TFR acts by taking an equilateral triangle of three atoms, lifting them up slightly, rotating them by 120° (counter-clockwise in this case), and lowering them back down. The projective graphs display unequivocally that the mechanism is, in fact, a sequence of concerted diamond-square-diamonds. The transition state graph corresponding to *o-m-T*-TS shows the three simultaneous square faces.

²David Wales has two examples of carborane rearrangements that illustrate the necessary movement: (1) rearrangement from the icosahedral 1,2-carborane to a higher energy C_S isomer (<http://www.ch.cam.ac.uk/SGTL/teach/djw1.html>) and (2) rearrangement from the 1,7-icosahedral carborane to a higher energy isomer based on an anticuboctahedron (<http://www.ch.cam.ac.uk/SGTL/teach/djw2.html>).

³All of these pathways operate in essentially the same manner, except for slight deviations in geometry due to juxtaposition of boron with carbon and vice versa.

Non-TFR pathway

Figure 7 illustrates the non-TFR mechanism. The first step in this mechanism is for one of the *ortho*-C₂B₁₀H₁₂ carbons, situated at the apex of a pentagonal cap, to break three of its bonds, while performing a DSD transition on the lower diamond. The square face in the DSD transition can be seen in the lower part of the graph corresponding to *o-m*-TS1. As the carbon migrates and forms the bond in the DSD, the species approaches a pseudo-*nido* geometry, which is displayed graphically in brackets. This geometry, in which the carbon lies approximately in a hexagonal plane, is never realized because other stabilizing forces exert themselves and propel the species towards the geometry in *o-m*-INT. The geometry displays two pentagonal faces in lieu of the hexagonal face of the pseudo-*nido* geometry. The last transition state *o-m*-TS2 is due to two adjacent DSD transitions on the opposite side of the cluster. In one direction, the species minimizes to *o-m*-INT. In the other direction, the molecule minimizes to *meta*-C₂B₁₀H₁₂.

*Meta- to para-carborane rearrangement**TFR pathway*

See “*TFR pathway*” subsection under “*Ortho- to meta-carborane rearrangement*”.

Table 3 Comparison of calculated distances [B3LYP/6-31G(d)] with distances from two gas-phase electron diffraction studies for *ortho*-, *meta*-, and *para*-C₂B₁₀H₁₂

Bond	B3LYP/6-31G(d)	GED [42]	GED [43]
<i>ortho</i> -C ₂ B ₁₀ H ₁₂			
C(1)–C(2)	1.625	1.624(8)	1.653(49)
C(1)–B(3)	1.720	1.734(7)	1.711(14)
C(1)–B(4)	1.700	1.700(6)	1.711(14)
B(3)–B(4)	1.780	1.788(6)	1.802(13)
B(4)–B(5)	1.786	1.794(8)	1.802(13)
B(3)–B(8)	1.765	1.774(9)	1.789(9)
B(5)–B(9)	1.778	1.787(6)	1.789(9)
B(7)–B(8)	1.781	1.796(6)	1.789(9)
B(8)–B(9)	1.793	1.808(8)	1.789(9)
B(9)–B(12)	1.784	1.787(9)	1.789(9)
<i>meta</i> -C ₂ B ₁₀ H ₁₂			
C(1)–B(2)	1.694	1.678(5)	1.720(9)
C(1)–B(4)	1.715	1.730(4)	1.720(9)
C(1)–B(5)	1.712	1.704(8)	1.720(9)
B(2)–B(3)	1.789	1.786(9)	1.831(52)
B(2)–B(6)	1.768	1.771(6)	1.791(15)
B(4)–B(5)	1.783	1.801(5)	1.791(15)
B(4)–B(9)	1.781	1.780(6)	1.817(13)
B(5)–B(9)	1.778	1.783(6)	1.817(13)
B(4)–B(8)	1.771	1.778(9)	1.817(13)
B(9)–B(10)	1.793	1.795(9)	1.817(13)
C(1)⋯C(7)	2.608	2.575(9)	
<i>para</i> -C ₂ B ₁₀ H ₁₂			
C–B	1.708	1.698(3)	1.710(11)
B–B tropical	1.790	1.784(1)	1.792(7)
B–B meridional	1.767	1.774(4)	1.772(13)
C(1)⋯C(12)	3.057	3.029(5)	3.05(5)

Non-TFR pathway

The non-TFR mechanism for the *meta*- to *para*-rearrangement (Fig. 8) mimics the *ortho*- to *meta*-C₂B₁₀H₁₂ rearrangement except that there is a second possible transition state *m-p*-TS1*, which is lower in energy than the analogous *ortho*- to *meta*-C₂B₁₀H₁₂ transition state (*m-p*-TS1, Table 1). The structural change from *meta*-C₂B₁₀H₁₂ to *m-p*-TS1 is exactly the same as in the *ortho*- to *meta*-C₂B₁₀H₁₂ case, except that only a boron vertex moves from the apex of a pentagonal cap opening a face in the cluster. Just as with the *ortho*- to *meta*-C₂B₁₀H₁₂ rearrangement, a “*nido*” geometry is never reached, leading instead to *m-p*-INT. The more thermodynamically favorable pathway is through *m-p*-TS1* which can be described as two adjacent concerted DSD steps. The second half of the rearrangement is analogous to the *ortho*-C₂B₁₀H₁₂ to *meta*-C₂B₁₀H₁₂ rearrangement: two concerted adjacent DSD steps occur on the opposite side of the cluster (see *m-p*-TS2) and *para*-C₂B₁₀H₁₂ results.

Borane rearrangement

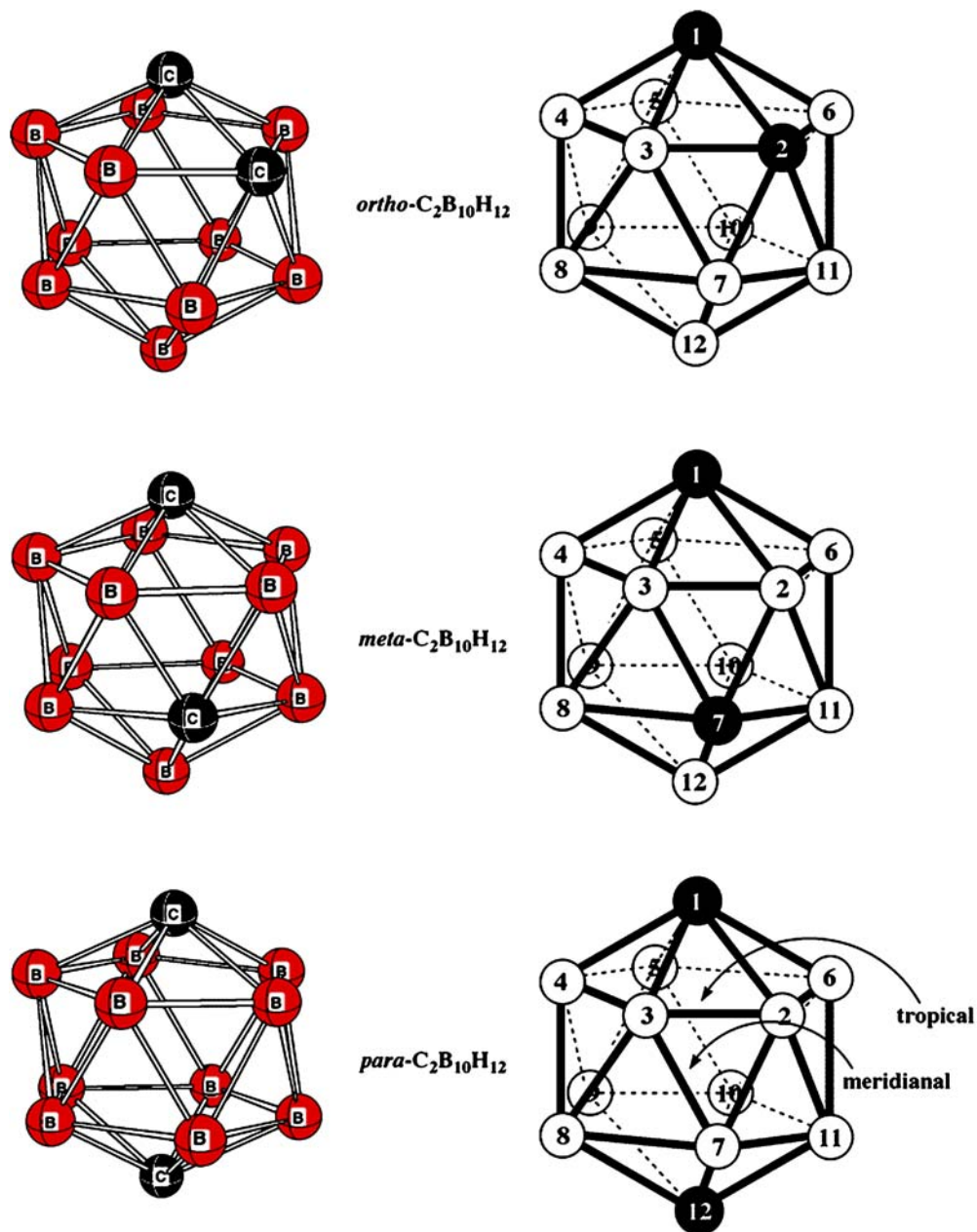
TFR pathway

See “TFR pathway” subsection under “Ortho- to meta-carborane rearrangement”. There is one exception in the B₁₂H₁₂²⁻ case. There is a D_{3h} intermediate *b-b-T*-INT that is lower than *b-b-T*-TS in enthalpy at 298 K by 1.7 kJ/mol. When entropic effects are taken into account, the intermediate *b-b-T*-INT is actually higher in free energy at 298, 500, and 800 K.

Non-TFR pathway

Figure 9 illustrates the non-TFR mechanism in the B₁₂H₁₂²⁻ rearrangement. The first transition state *b-b*-TS1 is essentially the same as *o-m*-TS1 and *m-p*-TS1 except that there are no carbon atoms present. A boron atom located at

Fig. 5 Numbering system with respective geometries for Table 3



the apex of a pentagonal cap migrates, breaking three bonds, and performs a DSD step on the lower diamond. The geometry again progresses towards a pseudo-*nido* geometry but does not quite achieve it. Another bond begins to form, which acts to prevent the cluster from opening, resulting in *b-b*-INT1. An interesting feature of the pathway is that there is an additional minimum (*b-b*-TS2) and transition state (*b-b*-INT1) along the reaction pathway that disappear when zero-point corrections are included. From *b-b*-TS2, two adjacent concerted DSD steps are needed to reach *b-b*-TS3 which then leads directly to icosahedral B₁₂H₁₂²⁻.

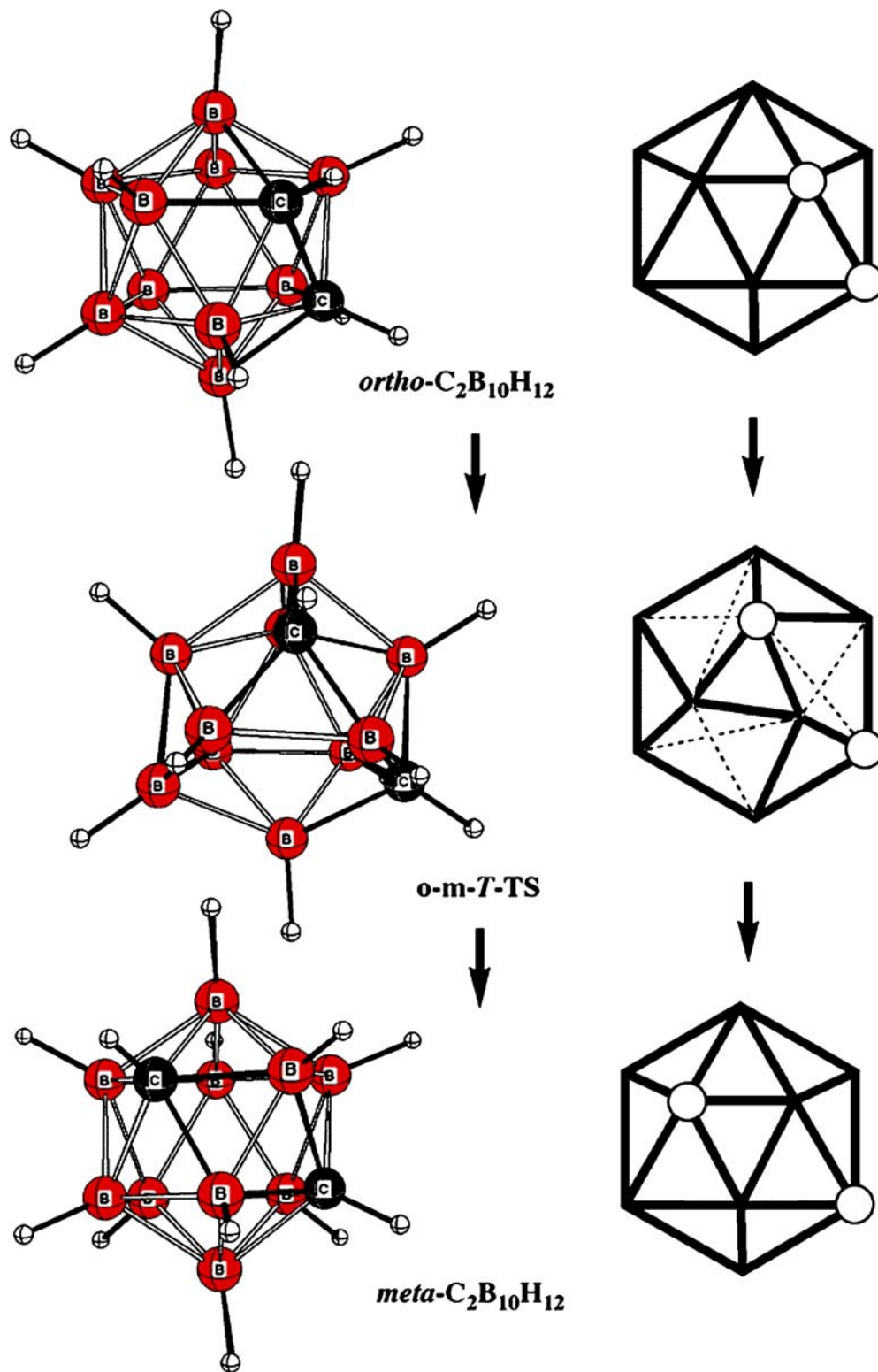
The free energy of activation in the gas phase for rearrangement of the boron vertices at 800 K is 398.5 kJ/mol via *b-b*-*T*-INT (Fig. 2). The free energy barrier is much higher than calculated for rearrangement in the isoelec-

tronic carboranes (Fig. 1). In fact, the barrier is higher than for fragmentation of a B–H bond, which implies that B₁₂H₁₂²⁻ should undergo decomposition rather than rearrangement. In a recent study of the boron hydride monoanions, B₁₂H₁₂²⁻ was determined to be the smallest boron hydride dianion with a positive adiabatic ionization potential [22]. It is still unlikely that both electrons would remain attached at elevated temperatures.

Experimental verification

There is one very interesting anomaly that emerges with respect to carborane rearrangements. The lowest free

Fig. 6 A localized graphical picture that describes the progression of the TFR pathway. The *ortho*- to *meta*- $C_2B_{10}H_{12}$ TFR pathway is shown



energy carborane barrier is a degenerate rearrangement *meta*- to *meta*- $C_2B_{10}H_{12}$, which is 185.5 kJ/mol at 800 K. This process may prove useful in rationalizing products obtained from rearrangement of the substituted carborane. In addition, because the barrier is lower than the barrier for rearrangement to *ortho*- and *para*- $C_2B_{10}H_{12}$, if one were

able to synthesize a substituted *meta*- $C_2B_{10}H_{12}$ with a methyl group on a boron atom adjacent to carbon, then, at a moderate temperature, one would expect to see migration of the methyl group to a boron position non-adjacent to carbon. We offer this prediction as a means of testing the rearrangement mechanism in the gas phase.

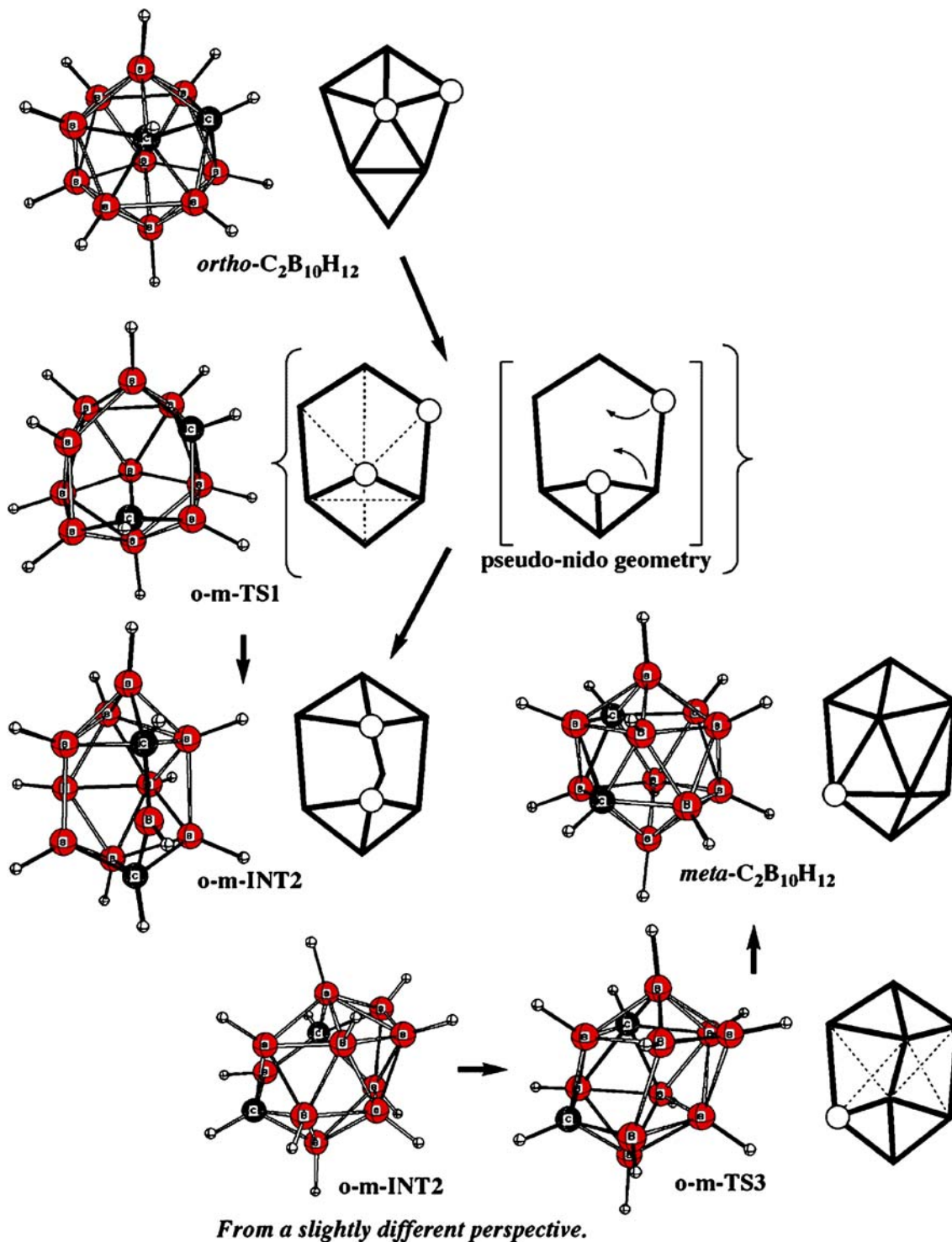


Fig. 7 The *ortho*- to *meta*-C₂B₁₀H₁₂ non-TFR rearrangement pathway

Conclusion

The rearrangement mechanism in C₂B₁₀H₁₂ and B₁₂H₁₂²⁻ has been explored using density functional theory methods. From *ortho*-C₂B₁₀H₁₂, the lowest free energy pathway to *meta*-C₂B₁₀H₁₂ is via a triangular-face-rotation mechanism

with a barrier of 207.6 kJ/mol. From *meta*- to *para*-C₂B₁₀H₁₂, the lowest-energy rearrangement mechanism is non-TFR which might loosely be described as a *closo* to *nido* to *closo* mechanism as the transition state is much more open than the reactant or product. The free energy barrier is 279.9 kJ/mol, 72.3 kJ/mol lower in free energy

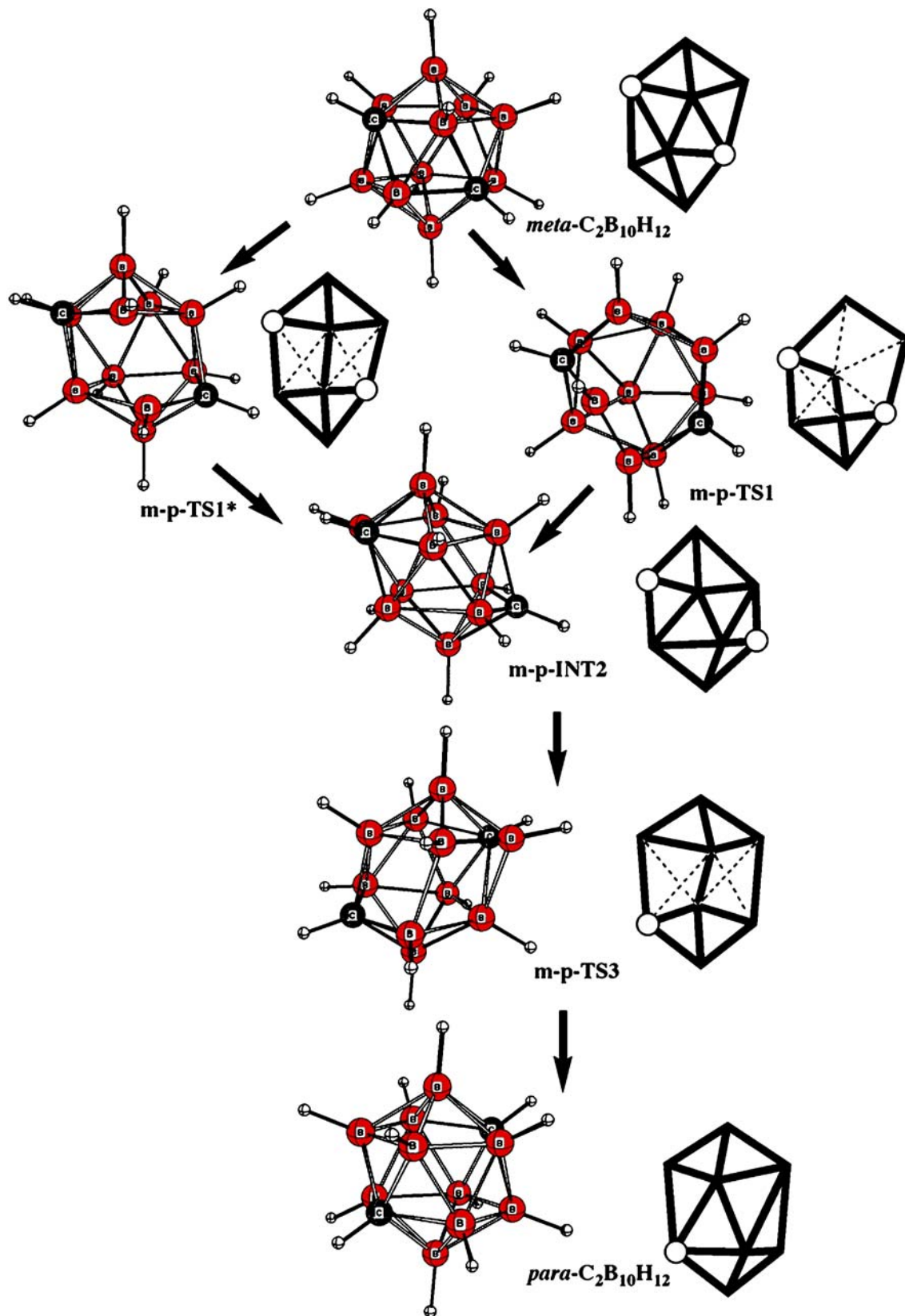


Fig. 8 The *meta*- to *para*- $C_2B_{10}H_{12}$ non-TFR rearrangement pathway

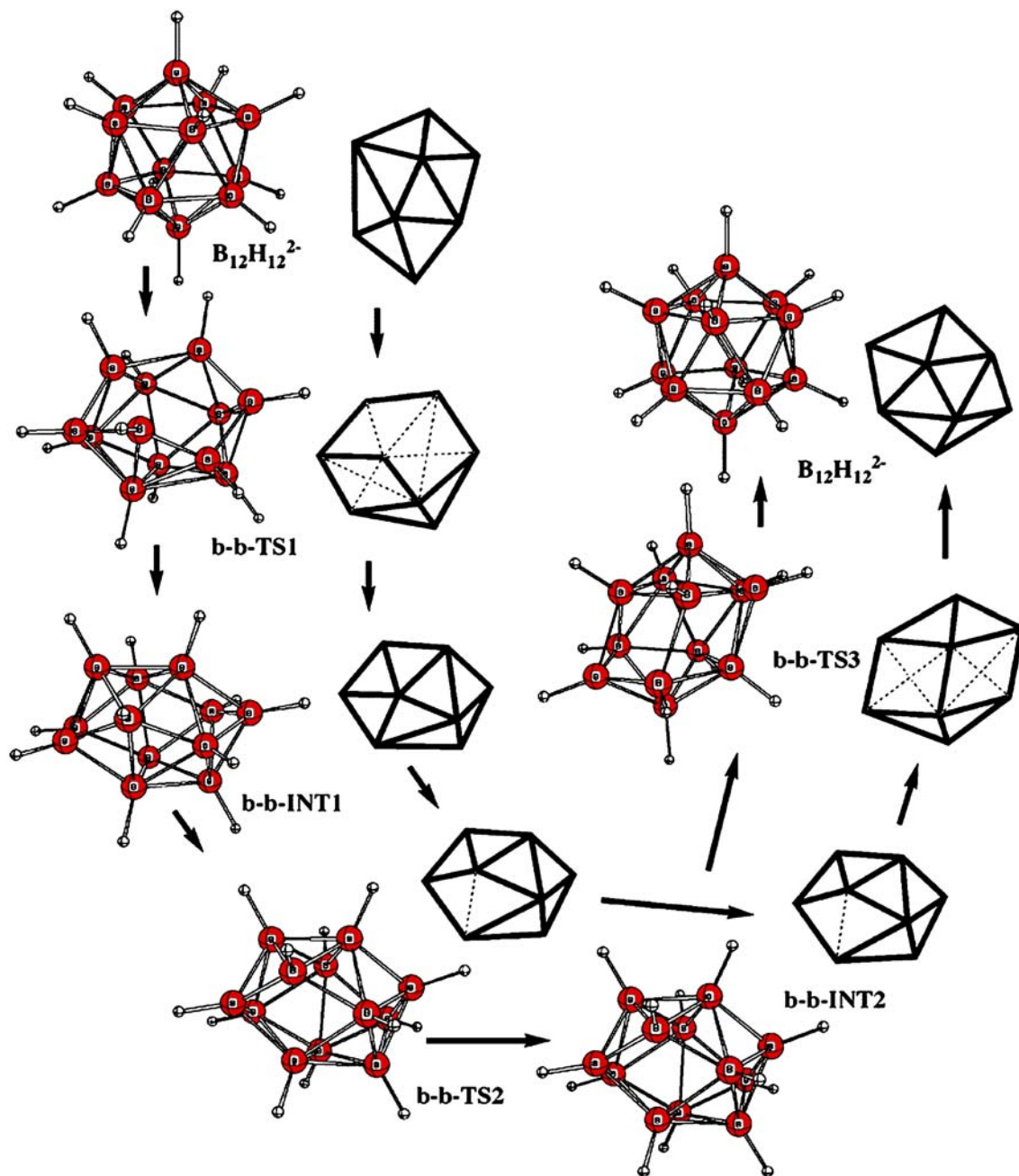


Fig. 9 The $B_{12}H_{12}^{2-}$ to $B_{12}H_{12}^{2-}$ non-TFR rearrangement pathway

than the *ortho* to *meta* process. The rearrangement of $B_{12}H_{12}^{2-}$, which would scramble labeled vertices, has a free energy barrier of 398.5 kJ/mol. For $B_{12}H_{12}^{2-}$, the preferred process in the gas phase would be to emit an electron or cleave a B–H bond.

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