

Molecular-dynamics study of detonation. II. The reaction mechanism

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In this work, we investigate mechanisms of chemical reactions that sustain an unsupported detonation. The chemical model of an energetic crystal used in this study consists of heteronuclear diatomic molecules that, at ambient pressure, dissociate endothermically. Subsequent association of the products to form homonuclear diatomic molecules provides the energy release that sustains the detonation. A many-body interaction is used to simulate changes in the electronic bonding as a function of local atomic environment. The consequence of the many-body interaction in this model is that the intramolecular bond is weakened with increasing density. The mechanism of the reaction for this model was extracted by investigating the details of the molecular properties in the reaction zone with two-dimensional molecular dynamics. The mechanism for the initiation of the reaction in this model is pressure-induced atomization. There was no evidence of excitation of vibrational modes to dissociative states. This particular result is directly attributable to the functional form and choice of parameters for this model, but might also have more general applicability.

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I. INTRODUCTION

Are the reactions that occur behind the shock front in a detonation due to rapid heating of molecules to dissociation after shock wave passage, or are the bonds mechanically broken after shock compression? Or do reactions result from a combination of both processes? What are the consequences for the properties of the detonation wave of each of these mechanisms? The mechanism of detonation remains unknown, despite numerous investigations over the past century [1]. The major obstacle in determining reaction mechanisms from experiment lies in the short time and spatial scales over which detonation occurs, and is complicated by the extreme pressure and energy releases accompanying the detonation.

Recent advances in spectroscopic methods are beginning to allow the direct measurements of molecular changes in a system that is shocked, toward the goal of determining the atomistic processes involved in the detonation [2]. Others are investigating energy transfer rates [3] and mechanisms [4] in energetic materials and possible relationships to the initiation of the detonation. These studies are providing critical pieces to a multifaceted chemical problem that is not completely understood. These measurements of molecular changes of explosives, however, have not been made under the extreme conditions associated with detonation. Direct measurement of chemical reactions occurring during this destructive and rapid event remains a formidable hurdle that has not been overcome.

As the development of diagnostic tools to measure these ultrafast events has progressed, computational resources and theoretical methods to model atomistic

processes involved in the detonation have advanced as well. Early reactive models used in molecular dynamics simulations described molecular crystals consisting of diatomics in metastable bound states that, when shocked, dissociated exothermically to form the more stable monatomic products [5,6]. A more recent study [7], in which molecular dynamics simulations of the detonation were compared to hydrodynamic predictions, used an isomerization of the reactant as the energy release reaction that drives the detonation. Although these models predicted reasonable pressure and temperature rises when compared to observed data from actual detonations, the chemistry that drives the detonation in these models is qualitatively unrealistic. A next step toward a more realistic description of the chemical processes occurring in a detonation would be one in which the energy release is due to endothermic dissociation of the reactants followed by exothermic association of the fragments to form more stable products. One must proceed in this process slowly, that is, introducing a minimum of additional complexity in the reaction mechanisms so as not to mask incremental information which might result.

Brenner and co-workers [8,9] in attempts to describe more realistically exothermic chemical reactions occurring in a shock wave, developed a series of models based on gas phase reactivity schemes. The models describe a heteronuclear diatomic molecular crystal that requires energy to break the molecular bonds in the low-pressure crystal, but releases substantial energy upon formation of homonuclear products. Two-dimensional molecular dynamics simulations using their first model [8] (denoted model 0 hereafter) appeared to sustain shock waves that were driven by exothermic reactions. However, the mac-

roscopic property profiles (pressure, density, and temperature) of the detonating system were flat topped and split [8]; this behavior has not been observed in experiment. The results of a second study [9] using a similar model (the only difference being in the molecular size parameter) showed macroscopic shock profiles that agreed well with the simplest model of detonation [1], in which reaction takes place immediately behind the shock front. We will denote this model hereafter as model I. Features of these models will be discussed below.

The studies described in this paper and our accompanying paper [10] were partially prompted after our attempts to reproduce simulation results (published in Ref. [9]) using model I failed. Additionally, model I (as well as model 0) had features in the potential energy surface that seemed undesirable to us, as described below. We made changes to model I by removing these undesired features (the revised model denoted as model II hereafter) and then performed two-dimensional molecular dynamics simulations of the detonation of this model energetic crystal initially at low temperature and low pressure. The equation of state and Hugoniot curve for this model were also obtained from two-dimensional molecular dynamics simulations at appropriate temperatures and pressures. These were used to make hydrodynamic predictions of detonation pressures, densities, and velocities. The hydrodynamic predictions were in good agreement with the results of our simulations of the detonation, indicating that our model and the method of molecular dynamics are consistent with the conservation laws of the process of detonation.

The focus of the study reported here is to examine the microscopic details of the detonation process and extract information concerning the mechanism of the detonation. We will first provide a detailed analysis of the interaction potentials for models 0, I, and II and show that the reaction mechanisms of these systems when subjected to shock are strongly dependent on the functional form of the models. Then, using the results from two-dimensional molecular dynamics simulations reported in our accompanying paper [10], we will dissect the reaction zone, and follow the changes in molecular properties throughout this region. This will reveal the mechanism of detonation for this model.

II. MODELS

A. Models 0 and I

Models 0 and I, used by Brenner and co-workers [8,9], are two-dimensional crystals of diatomic molecules arranged in a herringbone lattice. The interaction potential that is used to describe these crystals is

$$V = \sum_i^N \sum_{j>i}^N \{ f_c(r_{ij}) [V_R(r_{ij}) - \overline{B_{ij}} V_A(r_{ij})] + V_{\text{vdW}}^{(1)}(r_{ij}) \}, \quad (1)$$

the sums being over the N atoms comprising the model. The functional forms and parameters for the terms in Eq. (1) are given in Table I. The expression in Eq. (1) is based

on ideas proposed and developed by Abell [11] and Tersoff [12]. The leftmost terms within the summations in Eq. (1) make up the intramolecular interaction potential of diatomic molecules, and the second term within the sums, V_{vdW} , is an intermolecular interaction term. The function $f_c(r_{ij})$ smoothly attenuates the molecular bonding interaction to zero at 3.0 \AA . The bond-order function $\overline{B_{ij}}$, which ranges in value from zero to 1 depending on the local atomic environment, introduces many-body effects by modifying the attractive term of the molecular bonding portion of Eq. (1), $V_A(r_{ij})$. For an isolated diatomic, the value of $\overline{B_{ij}}$ is 1, and $V_A(r_{ij})$ will have a maximum contribution to Eq. (1) for the ij pair at the specified internuclear distance. If the diatomic is closely surrounded by many other atoms, as in the high-density region of a shock-compressed crystal, the value of $\overline{B_{ij}}$ decreases, depending on the number and location of the nearest neighbors surrounding the ij pair. This decrease in $\overline{B_{ij}}$ will correspondingly decrease the attractive interaction for the ij pair. This rubric is intended to mimic modifications of the bonding character upon increased density. As is made clear by the notation for the parameters given in Table I, this form of the potential is capable of describing diatomic molecules formed by atoms of types A - B , A - A , and B - B as well as intermolecular interactions between these various atom types. The curves in Fig. 1(a) illustrate the effect of the $\overline{B_{ij}}$ term on the intramolecular term for an A - B pair. The 11 curves in this figure correspond to the A - B intramolecular interaction

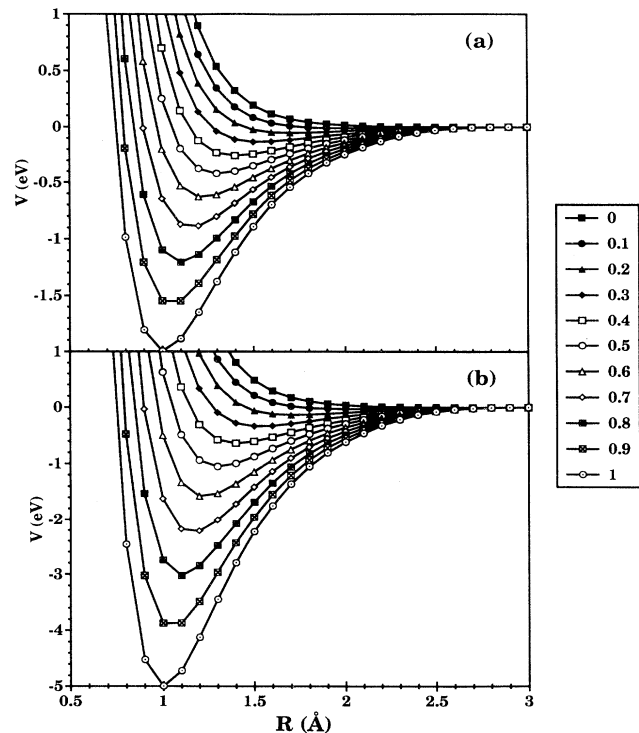


FIG. 1. Intramolecular interaction potentials of model 0 for different $\overline{B_{ij}}$ values [Eq. (1)] as functions of internuclear distance for (a) A - B interactions and (b) A - A interactions. Magnitude of the $\overline{B_{ij}}$ values are denoted in the legend.

TABLE I. Parameters and functional forms used for the potential energy expressions in Eqs. (1) and (2).

Parameter	Model 0 ^a Ref. [8]	Model I Ref. [9]	Model II	Functional Form
D_e^{AA} (eV)	5.0	5.0	5.0	$V_R(r) = \frac{D_e}{S-1} \exp[-\alpha\sqrt{2S}(r-r_e)]$
D_e^{BB} (eV)	5.0	5.0	2.0	
D_e^{AB} (eV)	2.0	2.0	1.0	$V_A(r) = \frac{SD_e}{S-1} \exp[-\alpha\sqrt{2/S}(r-r_e)]$
r_e^{AA} (Å)	1.0	1.2	1.2	
r_e^{BB} (Å)	1.0	1.2	1.5	$\bar{B}_{ij} = \frac{1}{2}(B_{ij} + B_{ji})$
r_e^{AB} (Å)	1.0	1.2	1.35	
α (Å ⁻¹)	1.8	1.8	1.8	$B_{ij} = \left\{ 1 + G \sum_{k \neq i,j} f_c(r_{ik}) \exp[m(r_{ij} - r_{ik})] \right\}^{-n}$
G	2.7	2.7	2.7	
m (Å ⁻¹)	5.0	5.0	7.5	
n	2.25	2.25	3.5	
ε (eV)	0.5	0.5	0.5	
σ (Å)	0.005	0.005	0.005	$f_c(r_{ij}) = \begin{cases} 1, & r < 2.0 \\ \frac{1}{2}\{1 + \cos(\pi[r-2])\}, & 2.0 \leq r < 3.0 \\ 0, & 3.0 \leq r \end{cases}$
mass_A (amu)	2.988	2.988	2.988	
mass_B (amu)	14.0	14.0	15.0	$V_{\text{vdW}}^1 = \begin{cases} 0, & r < 1.75 \\ P_0 + r[P_1 + r(P_2 + rP_3)], & 1.75 \leq r < 2.91 \\ 4\epsilon \left[\left(\frac{\sigma}{r}\right)^{12} - \left(\frac{\sigma}{r}\right)^6 \right], & 2.91 \leq r < 7.32 \\ 0, & 7.32 \leq r \end{cases}$
P_0 (eV)	14.0	14.0	46.0	
P_1 (eV Å ⁻¹)	0.4727	0.4727	0.4854	
P_2 (eV Å ⁻²)	-0.6996	-0.6996	-0.7184	
P_3 (eV Å ⁻³)	0.3364	0.3364	0.3455	
c_3 (eV Å ⁻³)	-0.0520	-0.0520	-0.05344	$V_{\text{vdW}}^2 = \begin{cases} 0, & r < 1.75 \\ P_0 + r[P_1 + r(P_2 + rP_3)], & 1.75 \leq r < 2.91 \\ 4\epsilon \left[\left(\frac{\sigma}{r}\right)^{12} - \left(\frac{\sigma}{r}\right)^6 \right], & 2.91 \leq r < 7.31 \\ \sum_{i=3}^5 c_i (r-7.32)^i, & 7.31 \leq r \leq 7.32 \\ 0, & 7.32 \leq r \end{cases}$
c_4 (eV Å ⁻⁴)			925.4631	
c_5 (eV Å ⁻⁵)			138 743.7872	
			5 548 241.6326	

^aThere is a typographical error in Table I of this reference for the B_{ij} definition [D. W. Brenner (private communication)]. It is corrected in Ref. [9].

with \bar{B}_{ij} values ranging from 0 to 1. This figure shows that the ij attraction is decreased by over one-half for $\bar{B}_{ij}=0.7$ and the molecular interaction appears to be completely repulsive for $\bar{B}_{ij}<0.1$. Also, the positions of the potential energy minima increase with decreasing \bar{B}_{ij} . The behavior of this function with decreasing \bar{B}_{ij} indicates that, as the local atomic environment becomes increasingly dense around an ij pair, the i and j atoms will be “forced apart” by the increased repulsion that can be attributed to this functional form.

Figure 1(b) shows the effect of the \bar{B}_{ij} term on the intramolecular term for an A - A (or B - B) pair. (The masses and homonuclear interactions of A and B atoms are the same in model 0.) As in Fig. 1(a), the intramolecular attraction is decreased by over one-half for $\bar{B}_{ij}=0.7$ and becomes completely repulsive for $\bar{B}_{ij}<0.1$. However, due to the differences in well depths for the intramolecular potential for A - A (or B - B) pairs and A - B pairs, the attractive interactions for A - A pairs are considerably greater than for A - B pairs for most values of \bar{B}_{ij} .

Brenner and co-workers found that model 0 produced

a “flat-topped, split shock wave structure” of the detonating crystal that is due to a “high-pressure, dissociative phase transition unintentionally introduced through our initial parametrizations” [9]. In our preliminary studies using model 0, we were able to reproduce each feature of the potential energy surface that was reported [8], including sound speed, equilibrium nearest-neighbor distance, barrier to collinear reaction of $A + A \rightarrow B$, and shock wave velocities. We also reproduced the flat-topped split shock wave structures alluded to in footnote 13 of Ref. [9].

Brenner and co-workers subsequently published results from simulations using model I in which the “high-pressure, dissociative phase transition” does not occur, and the detonation profiles show a single shock front that is followed by a reacting flow and rarefaction wave [9]. The results using model I were those that we could not reproduce.

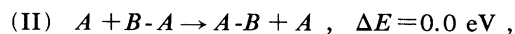
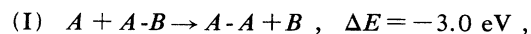
Model I as reported in Ref. [9] is the same as model 0 with one difference; that is, the value of r_e . In model 0, r_e had a value of 1.0 Å; r_e in model I is 1.2 Å. Thus the changes in intramolecular interaction potential with \bar{B}_{ij}

for model I are similar to those of model 0, shown in Fig. 1, with the equilibrium distances shifted by 0.2 Å for each curve.

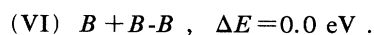
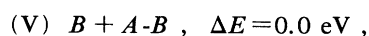
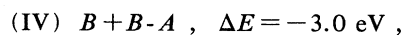
Using the parameters of model I, we have performed molecular dynamics simulations in an attempt to reproduce results of Ref. [9]. However, we once again saw the flat-topped, split shock wave structure that was seen in model 0. Figure 2 is a snapshot of the detonating system at 15 ps resulting from our simulations using model I. This snapshot shows that four distinct regions exist: the undisturbed crystal, a compressed crystal in which the molecules are reoriented, but do not react; the “high-pressure, dissociative phase transition” [9] in which molecular identity is lost, and the rarefaction region, consisting of vibrationally excited homonuclear products. The velocities of the two leading discontinuities are 8.1 and 6.8 km/s, respectively, whereas Ref. [9] reported the velocity of the single shock front as 9.3 km/s.

We do not know the source of the discrepancies between our calculations and those of Ref. [9]. Several details about the calculations, particularly regarding the initial conditions, were not given in Ref. [9], including the equilibrium molecular orientation, lattice constants, temperature of the undisturbed crystal, and number of atoms in the system. We therefore had to attempt to reproduce the simulations reported in Ref. [9] with an incomplete set of information. We determined the equilibrium low-pressure (1 atm) low-temperature (20 K) crystal structure for models 0 and I using *NPT* Metropolis Monte Carlo sampling as described in our accompanying paper [10]. The low-temperature, ambient pressure lattice parameters of the crystal defined by model I in the *x* and *y* directions are $a = 4.16$ Å and $b = 6.25$ Å, respectively. The center-of-bond (COB) fractionals for the two molecules (1 and 2) in the unit cell are at (0.25,0.25) and (0.75,0.75), respectively. The equilibrium *A-B* bond length is 1.199 45 Å, and the molecular orientation of molecules 1 and 2 is 30.51° and -30.51°, respectively, relative to the crystal *x* axis. The details of the molecular dynamics simulations are given in Sec. III D of our accompanying paper [10].

Upon finding these discrepancies in our calculations when compared to those published in Ref. [9], we attempted to understand the source of the “dissociative phase transition” which seemed to cause the problem of the flat-topped shock structure. We began our analyses by examining the three-body molecular interaction term that includes \bar{B}_{ij} for model I. We generated contour plots, shown in Fig. 3, of the collinear reactions



Note that, because the masses and bond energies of *A-A* and *B-B* are the same for model I, reactions (I)–(III) have the same potential energy features as the following collinear reactions:



As stated above, we were able to reproduce the contour plot and barrier to reaction (I) for model 0, given in Ref. [8]. Figure 3(a), the contour of reaction (I) for model I, is similar in feature to the contour of model 0 given in Ref. [8]; we calculate a barrier to reaction upon collinear approach as 0.10 eV, whereas the value reported in Ref. [9] is 0.08 eV (this may not be significant).

However, as is evident in Figs. 3(b) and 3(c), the trimers *ABA* and *AAA* exist upon linear approach. (Due to the features of the potential, this indicates that the trimers *BAB* and *BBB* also exist.) We characterized these species through normal mode analyses, after finding the equilibrium structures of these trimers using the Newton-Raphson energy minimization method [13]. The geometries and vibrational frequencies of these species are given in Table II. Also, we characterized the structures corresponding to the saddle points apparent in these figures. We were able to reproduce the features of the *A-A* and *A-B* diatomic molecules that were reported in Ref. [9], as well as the crystalline binding energy of 0.04 eV per molecule and nearest-neighbor distances of 3.3 Å. We were unable to locate any nonlinear trimers *AAA* or *ABA* or any of the trimers *AAB* or *BBA* using Eq. (1) for either of models 0 or I. Higher-order multimers were not found.

The relative energy features of the potential energy surface for model I are illustrated in Fig. 4. These schematics show relative energies of the trimers and the saddle points leading to their formation and/or decomposition. The shallow minima of the trimers relative to the saddle points leading to dissociation suggest that the trimers would be short-lived in a high-energy environment such as that which exists behind the shock front. Additionally, the trimer minima will be affected by the densi-

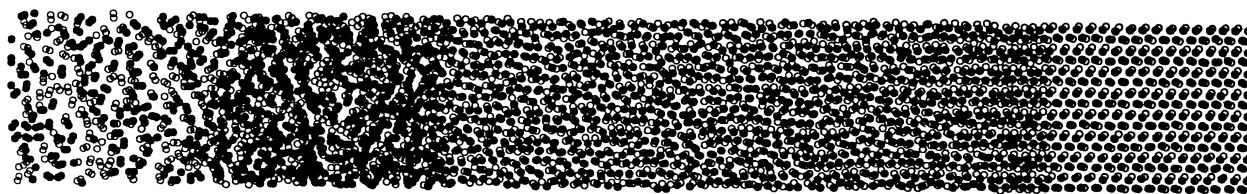


FIG. 2. Snapshot of chemically sustained shock waves at 15 ps initiated by a four-layer flier plate with an impact velocity of 6 km/s. The model used in this simulation is model I (Ref. [9]). The two types of atoms are depicted by filled and empty circles. Dimensions of the portion of the crystal shown in this snapshot are 387×50 Å².

ty, since trimer existence is due to the intramolecular interaction term in Eq. (1). The effect on the trimer minima due to the changes in density as the shock front travels through the region will affect the lifetime of the trimer

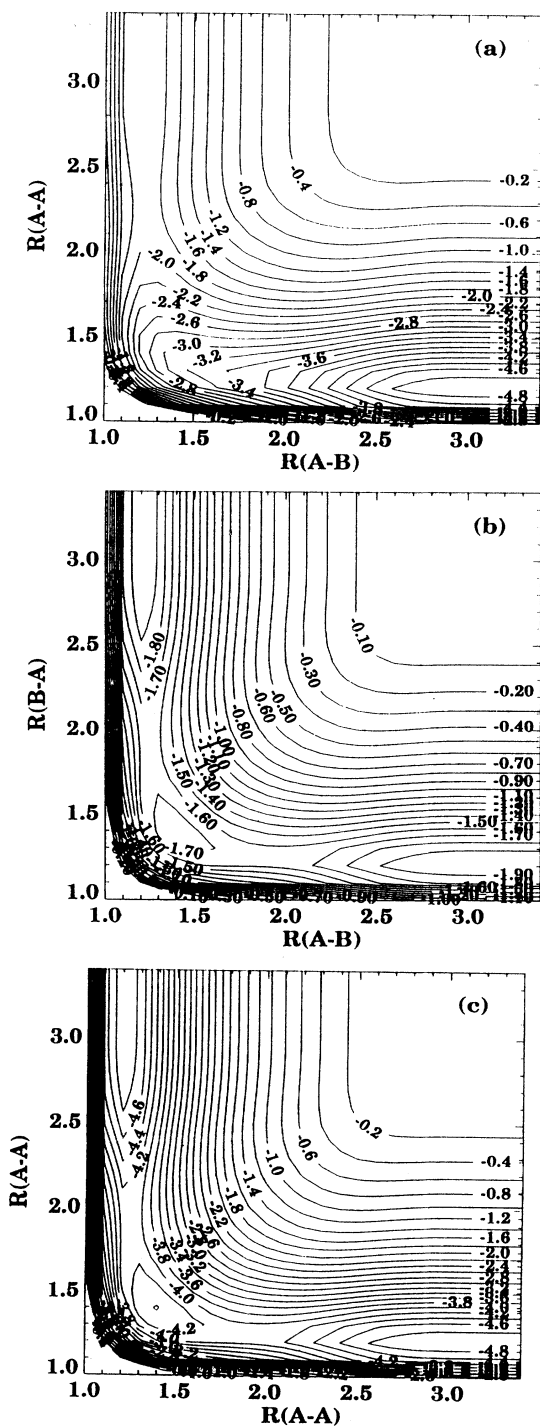


FIG. 3. Potential energy contours (in eV) for the following collinear three-body reactions using model I: (a) $A + A-B \rightarrow A-A + B$; (b) $A + BA \rightarrow A-B + A$; (c) $A + A-A \rightarrow A-A + A$. Units for the internuclear distances along both axes are Å.

intermediate. The ability of Eq. (1) to form trimers leads us to suspect this as the cause of the region in the detonating system that Brenner *et al.* call the “dissociative phase transition” [9]. We think it might be more appropriate to label that region an *associative* phase transition, where trimer formation occurs, leading to subsequent decomposition to the more stable homonuclear diatomic species. In any case, changing the value of the parameter r_e from 1.0 to 1.2 Å did not remove the source of the “flat-topped, split shock wave structure” [9] which we think is due to this ability to form trimers.

B. Model II

The ability of trimer formation using models 0 and I indicated to us that saturation of the molecular bond was not correctly accounted for in Eq. (1) with those sets of parameters. It seems appropriate to us that not only should the attractive portion of the potential be modified according to the local atomic environment, but that the repulsive wall should be affected as well, to take into account saturation of the bonds. We have therefore modified Eq. (1) to

$$V = \sum_i^N \sum_{j>i}^N \{ f_c(r_{ij}) [(2 - \bar{B}_{ij}) V_R(r_{ij}) - \bar{B}_{ij} V_A(r_{ij})] + V_{\text{vdW}}^{(2)} \} \quad (2)$$

where the term \bar{B}_{ij} has the same description as in Eq. (1).

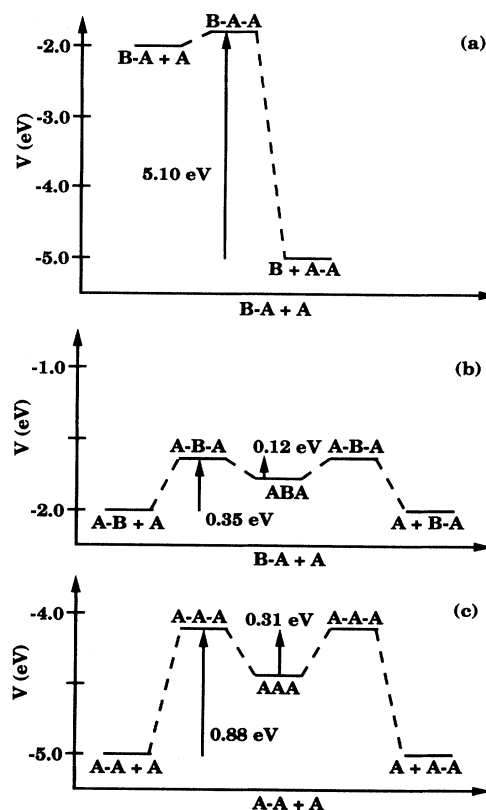


FIG. 4. Schematic of the potential energy for three-body collinear reactions (I)–(III) for model I.

TABLE II. Geometric parameters and harmonic vibrational frequencies of critical points on the potential energy surfaces for models 0, I, and II.

Species	Internuclear distances (Å)	Harmonic frequencies (cm ⁻¹)	Energy (eV)
Model 0			
<i>A-B</i>	$R(A-B)=1.000\ 00$	1064	-2.0
<i>A-A (B-B)</i>	$R(A-A)=1.000\ 00$	1682	-5.0
<i>A-A-A (B-B-B)</i>	$R(A-A)=1.240\ 82$	200,296,739	-3.940 69
<i>A-B-A (B-A-B)</i>	$R(A-B)=1.238\ 22$	150,184,466	-1.581 14
<i>A + A-B → A-A + B</i>	$R(A-A)=2.150\ 00$	217i,982	-1.894 57
	$R(A-B)=1.012\ 98$		
<i>A + B-A → A-B-A</i>	$R(A-B)=1.376\ 25$	158i,171,570	-1.578 84
	$R(B-A)=1.149\ 01$		
<i>A + A-A → A-A-A</i>	$R(A-A)=1.355\ 66$	225i,280,866	-3.937 64
	$R(A-A)=1.161\ 41$		
Model I			
<i>A-B</i>	$R(A-B)=1.200\ 00$	1064	-2.0
<i>A-A (B-B)</i>	$R(A-A)=1.200\ 00$	1683	-5.0
<i>A-A-A (B-B-B)</i>	$R(A-A)=1.386\ 09$	195,690,863	-4.423 61
<i>A-B-A (B-A-B)</i>	$R(A-B)=1.385\ 54$	122,439,545	-1.770 19
<i>A + A-B → A-A + B</i>	$R(A-A)=2.283\ 75$	257i,974	-1.902 25
	$R(A-B)=1.213\ 10$		
<i>A + B-A → A-B-A</i>	$R(A-B)=1.870\ 13$	230i,847	-1.646 94
	$R(B-A)=1.245\ 72$		
<i>A + A-A → A-A-A</i>	$R(A-A)=1.870\ 13$	365i,1339	-4.117 34
	$R(A-A)=1.245\ 72$		
Model II			
<i>A-B</i>	$R(A-B)=1.350\ 00$	727	-1.0
<i>A-A</i>	$R(A-A)=1.200\ 0$	1626	-5.0
<i>B-B</i>	$R(B-B)=1.500\ 0$	587	-2.0
<i>A + B-A → A-B + A</i>	$R(A-B)=1.649\ 67$	300i,275	-0.572 37
<i>B + A-B → B-A + B</i>	$R(A-B)=1.649\ 67$	171i,157	-0.572 37
<i>A-B + B → A + B-B</i>	$R(A-B)=1.684\ 60$	367i,365	-0.821 74
	$R(B-B)=1.798\ 43$		
<i>B + B-B → B-B + B</i>	$R(B-B)=1.799\ 67$	242i,222	-1.144 74
<i>A + A-A → A-A + A</i>	$R(A-A)=1.499\ 69$	670i,631	-2.861 86

The parameters and functional forms for the terms in Eq. (2) are given in Table I. We found two discontinuities in the weak long-range van der Waals interaction for models 0 and I; one at $r=2.91$ Å and the other at $r=7.32$ Å. We have corrected the discontinuity at $r=2.91$ Å by determining the cubic spline coefficients using the requirements that at $r=1.75$ Å the energy and energy first derivatives must equal 0.0, and at $r=2.91$ Å the energy and energy first derivatives must equal those produced by the Lennard-Jones potential function. We also added a quintic spline to the Lennard-Jones potential in the region of 7.31–7.32 Å, to allow a continuous cutoff of the intermolecular interaction. The quintic spline coefficients were determined in the same manner as the cubic spline coefficients described above; however, energy second derivatives were required to match at the functional interfaces at 7.31 and 7.32 Å. We have only provided four significant figures for the spline coefficients used in Eq. (2); we suggest that machine accuracy of these coefficients be determined by the interested modeler.

We have also changed the model parameters to allow different exothermic reactions, and set the masses of par-

ticles *A* and *B* to 15 and 46 amu, respectively, in order to introduce mass effects. Also, the molecular sizes of the *A-B*, *A-A*, and *B-B* diatomics were set to 1.35, 1.2, and 1.5 Å, respectively. The reactions that this model describes are

- (VII) $A-B + A \rightarrow A + B-A$, $\Delta E=0.0$ eV ,
 (VIII) $B-A + A \rightarrow B + A-A$, $\Delta E=-4.0$ eV ,
 (IX) $A-B + B \rightarrow A + B-B$, $\Delta E=-1.0$ eV ,
 (X) $B-A + B \rightarrow B + A-B$, $\Delta E=0.0$ eV ,
 (XI) $A-A + A \rightarrow A + A-A$, $\Delta E=0.0$ eV ,
 (XII) $B-B + B \rightarrow B + B-B$, $\Delta E=0.0$ eV .

Geometric parameters and vibrational frequencies of the stable species for these reactions are given in Table II. We also show contour plots of the collinear reactions using Eq. (2), assuming only three atoms, in Fig. 5. We did not find trimer or other multimer formation using Eq. (2). For the collinear approach of *A* to *A-B*, there is no bar-

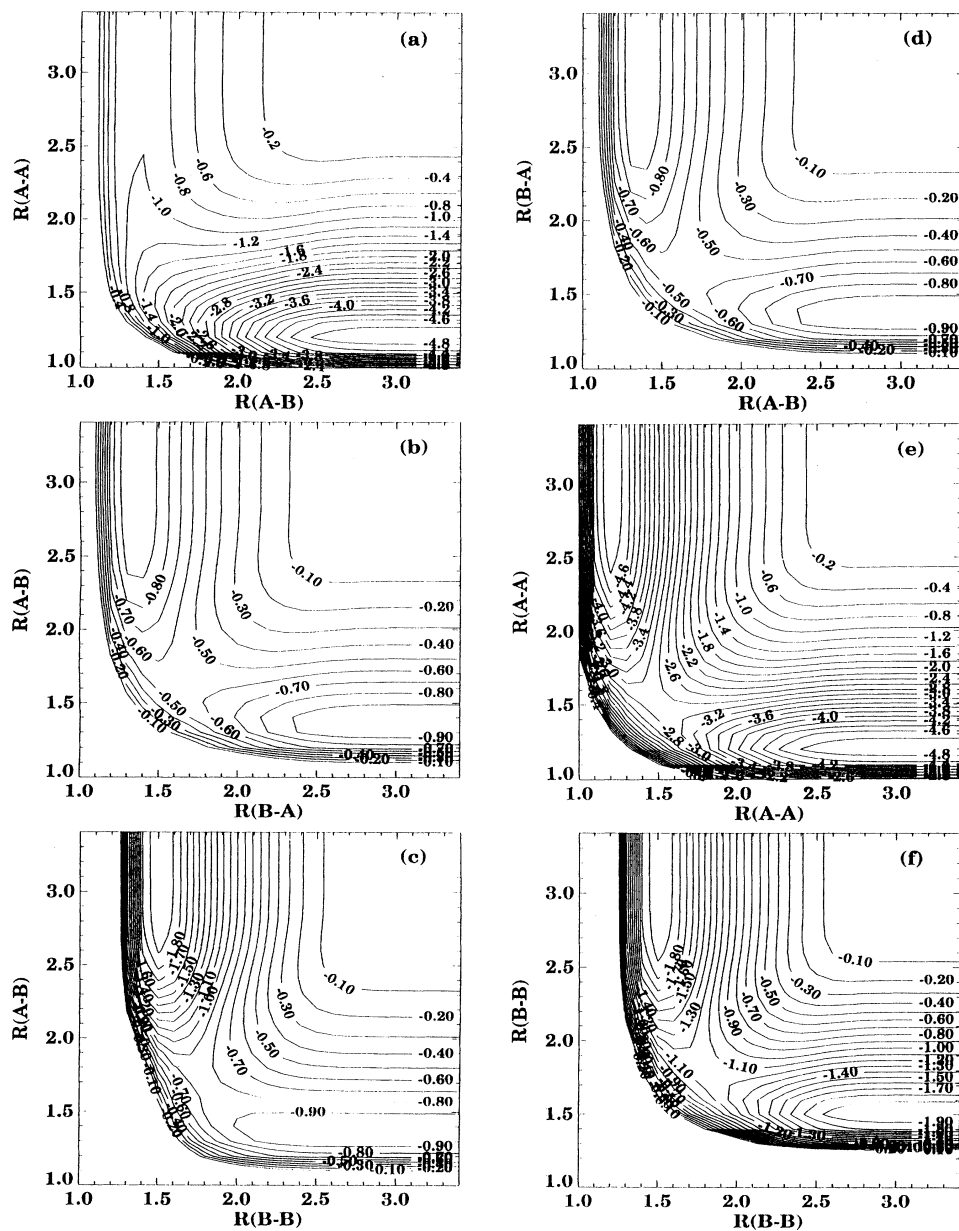


FIG. 5. Potential energy contours (in eV) for the following collinear three-body reactions using model II: (a) $A + A-B \rightarrow A-A + B$; (b) $A + B-A \rightarrow A-B + A$; (c) $A + B-B \rightarrow A-B + B$; (d) $B + A-B \rightarrow B-A + B$; (e) $A + A-A \rightarrow A-B + A$; (f) $B + B-B \rightarrow B-B + B$. Units for the internuclear distances along both axes are Å.

rier to approach. The transition state structures and vibrational frequencies for the collinear reactions are also given in Table II.

We have given considerable attention to describing the features of the potential energy surface for model II assuming only three atoms, primarily because such attention was given in the previous studies for models 0 and I [8,9] and we wished to provide direct comparison. However, in the high-density region behind the shock wave, the potential energy interactions (specifically the intramolecular interactions) will be affected by many more atoms than three. Therefore, as we did for model 0, we show the effect of the intramolecular interaction with different values of \bar{B}_{ij} in Fig. 6, which correspond to varying degrees of density of atoms surrounding an ij pair.

These curves differ from those of model 0 (Fig. 1) in

that the well depths are decreased by one-half for $\bar{B}_{ij}=0.8$ rather than $\bar{B}_{ij}=0.7$. Additionally, the repulsive interactions for these \bar{B}_{ij} values are much greater than those for model 0, and the positions of the minima are at larger distances with decreasing \bar{B}_{ij} than those for model 0. This model is analogous to models 0 and I in that the atoms are forced apart with decreasing \bar{B}_{ij} .

The two-dimensional crystal modeled by Eq. (2) also has an equilibrium configuration consistent with a herringbone arrangement. We determined the 1 atm, 20 K orientation of the crystal using the *NPT* Monte Carlo calculations (described in Ref. [10]). The low-temperature, ambient pressure lattice parameters of the crystal in the x and y directions, respectively, are $a=4.34$ Å and $b=6.27$ Å. The COB fractionals for the two molecules (1 and 2) in the unit cell are at (0.25,0.25) and (0.75,0.75), respectively. The $A-B$ equilibrium bond length is 1.349

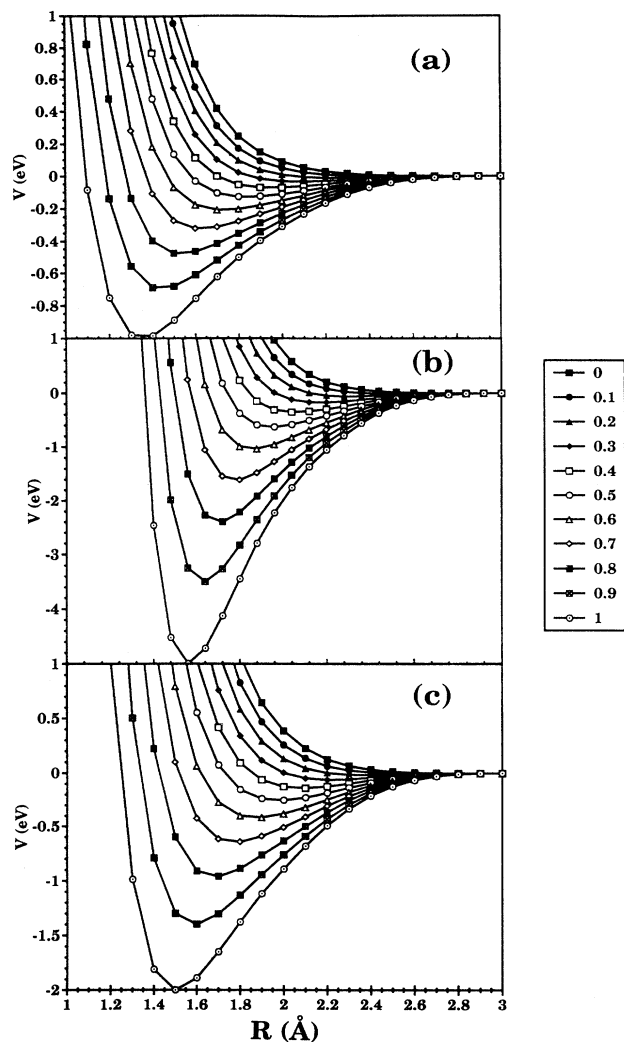


FIG. 6. Intramolecular interaction potentials of model II for different B_{ij} values [Eq. (2)] as functions of internuclear distance for (a) $A-B$ interactions, (b) $A-A$ interactions, and (c) $B-B$ interactions. Magnitude of the B_{ij} values is denoted in the legend.

Å, and the molecular orientation of molecules 1 and 2 is 29.0° and -29.0° , respectively, relative to the crystal x axis.

NPT Metropolis Monte Carlo calculations of equilibrium crystal structure as a function of pressure at 20 K were performed to determine the sound speeds of the different models. Results of the density as a function of pressure for models I and II are shown in Table III. These results for P ranging from 0.0 to $0.015 \text{ eV}/\text{\AA}^2$ were fitted to a cubic polynomial, and the slope of the P - V curve was extracted, which is directly proportional to the square of the sound speed of the crystal [1]. The sound speeds for models I and II are 2.0 and 1.2 km/s, respectively.

III. DETAILS OF THE CALCULATIONS

Details of the molecular dynamics simulations are given in the accompanying paper on the comparison of molecular dynamics simulations to hydrodynamic predictions [10]. The results that are analyzed in this paper are from the simulation in which shock is initiated by a plate of $A-A$ molecules striking the simulation cell with an impact velocity of 12 km/s.

Because our interest is to examine the mechanistic details involved in the reactions initiating the detonation, we will focus our analysis on a region we call the reaction zone. The reaction zone here is defined as the region between the shock front and the point at which the number of reacted $A-B$ molecules equals or exceeds the number of unreacted $A-B$ molecules. It is difficult to determine whether a molecule has reacted or not within this zone in an unambiguous manner, due to the effect on Eq. (2) of the higher density of this region. As shown earlier, there exists a density of nuclei [reflected by the B_{ij} term of Eq. (2)] that eliminates any attraction between the $A-B$ pair, even though the atoms are within the 3.0 \AA cutoff for the intramolecular term. However, we know unambiguously that some sort of molecular interaction exists if the $A-B$ internuclear distance is less than 3.0 \AA according to Eq. (2). Therefore, for simplicity, we have used the following simple geometric test to determine reactivity. We have

TABLE III. Lattice parameters, density, and molecular geometries versus pressure for models I and II.

Pressure ($\text{eV}/\text{\AA}^2$)	Model I					Model II				
	$\langle a \rangle$ (\AA)	$\langle b \rangle$ (\AA)	$\langle \rho \rangle$ ($\text{amu}/\text{\AA}^2$)	$\langle r_{ij} \rangle$ (\AA)	$\langle \theta_{ij} \rangle$ (deg)	$\langle a \rangle$ (\AA)	$\langle b \rangle$ (\AA)	$\langle \rho \rangle$ ($\text{amu}/\text{\AA}^2$)	$\langle r_{ij} \rangle$ (\AA)	$\langle \theta_{ij} \rangle$ (deg)
0.0	4.16	6.25	2.1538	1.200	30.3	4.34	6.27	4.4833	1.349	29.1
0.000 05	4.17	6.23	2.1556	1.199	29.7	4.33	6.28	4.4865	1.349	29.3
0.000 1	4.16	6.24	2.1573	1.199	30.0	4.33	6.27	4.4937	1.349	29.1
0.000 5						4.32	6.25	4.5185	1.349	29.1
0.001	4.16	6.18	2.1782	1.199	29.2	4.31	6.22	4.5508	1.349	29.0
0.002 5						4.24	6.19	4.6484	1.349	29.8
0.005	4.10	6.03	2.2651	1.199	28.7					
0.01	4.04	5.94	2.3336	1.198	28.9					
0.012 5						4.13	5.94	4.9731	1.347	29.0
0.015						4.11	5.90	5.0311	1.346	29.0

defined that the original A - B pair has undergone reaction (is dissociated, or is bound to an atom that is not its original molecular partner) if the distance between the atomic centers exceeds 3.0 Å. Figure 8(d) of our accompanying paper [10] shows the number of unreacted A - B molecules and dissociated A - B molecules at 7.8 ps in the simulation. The point at which the number of dissociated A - B molecules exceeds the number of unreacted A - B molecules falls slightly behind the shock fronts in the density, pressure, and temperature profiles. As shown in the accompanying paper [10], the width of the reaction zone is approximately constant throughout the simulations of unsupported detonations and is, on average, 14 Å wide.

We will examine the translational, rotational, and vibrational kinetic energy distributions of all unreacted A - B molecules within the reaction zone. Before the kinetic energy components for each molecule are calculated, the local mass flow velocity is removed as described in our accompanying paper [10]. Additionally, we will calculate the distributions of A - B bond lengths and orientation angles of the A - B molecules relative to the crystal x axis in this region. We will also calculate the distributions of the intramolecular interaction potential and the B_{ij} term for the unreacted A - B molecules. The calculation of the translational energy of the unreacted A - B molecule is straightforward:

$$[\text{KE}(\text{translation})] = \frac{P_{\text{c.m.}}^2}{2M}, \quad (3)$$

where M is the total mass of the A - B molecule and $P_{\text{c.m.}}$ is the center-of-mass momentum of the molecule. As an approximation, we assume that the rotational energy is the kinetic energy due to motion of the atoms in the molecule that are perpendicular to the molecular bond

$$[\text{KE}(\text{rotation})] = \frac{1}{2} \sum_{i=A}^B m_i v_{i(\text{perpendicular})}^2, \quad (4)$$

where $v_{i(\text{perpendicular})}$ denotes the velocity components perpendicular to the molecular bond (center-of-mass velocity removed). The vibrational kinetic energy is the kinetic energy due to motion of the atoms in the molecule along the molecular bond

$$[\text{KE}(\text{vibration})] = \frac{1}{2} \sum_{i=A}^B m_i v_{i(\text{parallel})}^2, \quad (5)$$

where $v_{i(\text{parallel})}$ denotes the velocity components parallel to the molecular bond (center-of-mass velocity removed).

Although these calculations of the internal energy distributions are based on a simple idea, especially in light of the fact that the molecules might not even be bound in this zone due to the high-density effects on the interaction potential, it represents adequately energy flow into the directional components of the system.

IV. RESULTS

Each of the calculations that simulated unsupported detonations discussed in our accompanying paper (flier plate impact velocities ≥ 4.7 km/s) provides the same temperature, density, and pressure profiles as well as the same microscopic behavior in the reaction zone once the

detonation wave reaches the steady-state velocity [10]. Therefore we are presenting the results of only one simulation, the simulation with plate impact velocity of 12 km/s.

We have examined in detail the various properties within the reaction zone in order to detect any variations occurring there. The reaction zone is partitioned into seven regions, labeled 0–6. Each region has a 2.17 Å width in the crystal x direction ($\frac{1}{2}$ the size of the unit cell in the x direction), and because the shock profile is uniform in the y direction the length in the y direction is the entire crystal length. Region 0 is the area directly behind the shock front to 2.17 Å behind the shock front; region 1 begins at the end of region 0, and ends at 4.34 Å behind the shock front; region 2 ends 6.51 Å behind the shock front; and so on, concluding with region 6, which ranges from 13.02 to 15.19 Å behind the shock front. Distributions of properties of the system were calculated at every 100th integration step throughout the 12 km/s trajectory; cumulative distributions of these properties were compiled, normalized, and analyzed. The normalization is such that the integral of the property over the independent variable is 1. Within the statistical uncertainty resulting from this limited sampling, none of the properties changed in time after the first 0.25 ps following flier plate impact. In order to obtain better statistical sampling, we therefore report cumulative normalized distributions and corresponding time averages for all times later than 0.5 ps after flier plate impact. We have calculated energy and orientational distributions of the unreacted A - B molecules that are in each region. Additionally, we have calculated the distributions of the intramolecular interactions and corresponding B_{ij} terms for the unreacted A - B pairs in these regions. Averages obtained from the normalized distributions of these properties for each of the seven regions within the reaction zone are shown in Table IV.

Figures 7–9 show the cumulative translational, rotational, and vibrational kinetic energy distributions, respectively, of the unreacted A - B pairs in the seven zones throughout the 12 km/s simulation. It is evident that the translational modes of the A - B molecules in the region directly behind the shock front (region 0) are substantially excited as the shock wave passes through it but the internal modes are excited to only one-quarter of the value of the translational excitation. The average translational energy of molecules within region 1 is less than half that of region 0, while the average energy of the internal modes has doubled, indicating that substantial energy transfer among molecular modes has occurred by the time the molecules have moved from region 0 to region 1. Since the average residence time of the molecules in each 2.17 Å region in the reaction zone is only 3.3×10^{-14} s (assuming local mass flow velocity of 6.6 km/s, the speed of the detonation wave in this simulation [10]) energy transfer among molecular modes is rapid. Beyond region 1, the regional averages reach plateau values, consistent with equipartitioning of energy among translational and internal modes. It is evident that substantial vibrational excitation does not occur due to

TABLE IV. Averages of properties of unreacted $A-B$ molecules in reaction zone throughout 12 km/s simulation of unsupported detonation.

Region	$\langle E(\text{trans}) \rangle$ (eV)	$\langle E(\text{rot}) \rangle$ (eV)	$\langle E(\text{vib}) \rangle$ (eV)	$\langle V_{ij} \rangle$ (eV)	$\langle \bar{B}_{ij} \rangle$	$\langle r_{ij} \rangle$ (Å)	$\langle \theta_{ij} \rangle$ (deg)
0	1.1745	0.1336	0.1512	-0.3302	0.6767	1.4462	40.3
1	0.5027	0.2492	0.2821	0.0245	0.3070	1.8209	51.4
2	0.4077	0.2601	0.2472	0.0564	0.1622	2.0568	49.4
3	0.4334	0.2368	0.2387	0.0572	0.1553	2.0920	46.9
4	0.4578	0.2347	0.2420	0.0470	0.1425	2.1322	45.1
5	0.4608	0.2298	0.2289	0.0430	0.1411	2.1540	45.1
6	0.4805	0.2235	0.2208	0.0387	0.1413	2.1625	46.0

shock impact. Therefore the mechanism of vibrational excitation to dissociation of $A-B$ molecules from shock impact (thermal decomposition) is not consistent with this model.

Figures 10 and 11 show the distributions of molecular bonding energies for atom pairs and the \bar{B}_{ij} terms within the seven regions. Figures 12 and 13 show the distributions of the internuclear $A-B$ distances and rotational orientations within the seven regions. The distribution of the \bar{B}_{ij} values in region 0 is spread across the range of values, reflecting high density behind the shock front, but has a broad maximum at 0.9. The average \bar{B}_{ij} value for region 0 is 0.7. Figure 6 shows that the maximum $A-B$ intramolecular attraction corresponding to this \bar{B}_{ij} value is -0.3 eV when $R_{AB} = 1.6$ Å. The corresponding distribution for internuclear distances of the $A-B$ molecules within region 0 (Fig. 12), however, is a narrow distribution centered at 1.35 Å. $A-B$ molecules with this internuclear distance and a \bar{B}_{ij} value of 0.7 (or less) experience no attraction and different degrees of repulsion, depending on the actual \bar{B}_{ij} value. This is reflected in the intramolecular potential energy distribution for the $A-B$ pairs shown in Fig. 10. The distribution has a peak at $V_{ij} = 0$ eV for region 0, but also has a shoulder extending from $V = 0$ to -1 . The shoulder is due to the significant number of $A-B$ molecules that have \bar{B}_{ij} values > 0.7 within region 0. The distribution of \bar{B}_{ij} values within re-

gion 1 is substantially different from that of region 0, with its peak at 0.1, but with a broad shoulder extending to 0.9. The average \bar{B}_{ij} value within this region is 0.3, and the corresponding average internuclear distance is 1.8 Å. The distribution of R_{AB} 's, however, shows a peak at 1.35 Å, with a broad shoulder extending to the dissociation limit of 3.0 Å. The intramolecular attraction for a \bar{B}_{ij} value of 0.3 is slight [< -0.05 eV at $R(\text{eq}) = 2.0$ Å], and the distribution of internuclear distances suggests that a substantial number of $A-B$ molecules experience no attraction and some degree of repulsion within this zone, also reflected in the distribution of the intramolecular interactions in Fig. 10 for region 1. Beyond region 1, the \bar{B}_{ij} distributions are similar, with peaks at 0.05. Likewise, the internuclear distance distributions are similar, and are broad bands extending from 1.2 Å to the dissociation limit of 3.0 Å.

Figure 13 shows the distribution of the orientation angle of the $A-B$ molecules throughout the reaction zone. Structure in the orientational distributions exists in regions 0-2. In the undisturbed crystal, the molecular bond orientation is $\pm 29^\circ$ relative to the crystal x axis. The distribution of the angular orientation in region 0 is peaked at 30° , indicating, for the most part, no rotational reorientation upon shock impact. This, coupled with the internuclear distance distribution for the $A-B$ molecules (Fig. 12) and the density of the system for this region

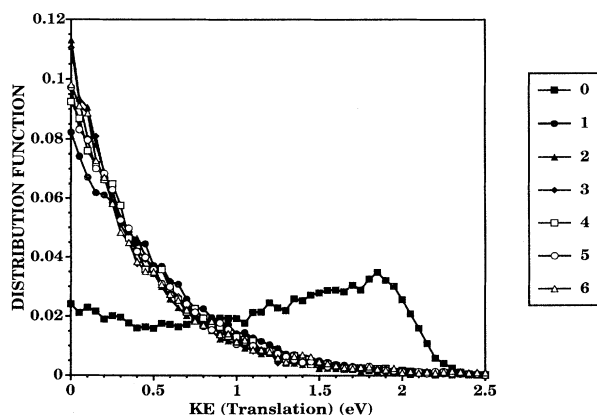


FIG. 7. Normalized translational energy distributions of original $A-B$ pairs for seven regions directly behind the shock front through the 12 km/s simulation (up to 7.8 ps). See text for definition of regions.

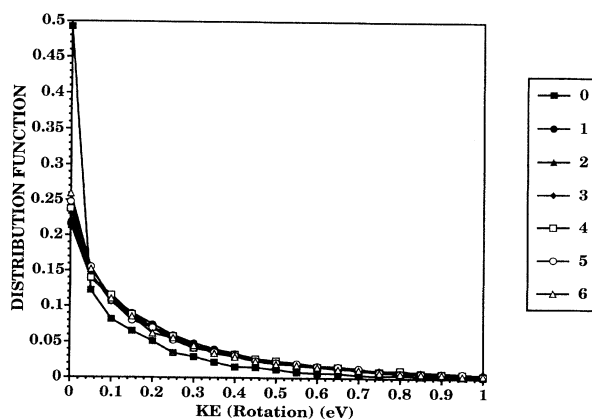


FIG. 8. Normalized rotational energy distributions of original $A-B$ pairs for seven regions directly behind the shock front through the 12 km/s simulation (up to 7.8 ps). See text for definition of regions.

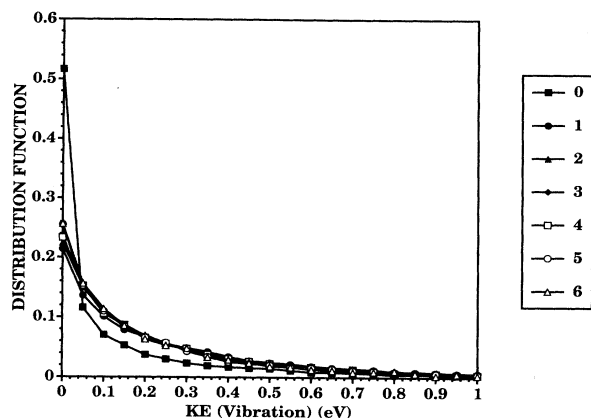


FIG. 9. Normalized vibrational kinetic energy distributions of original $A-B$ pairs for seven regions directly behind the shock front through the 12 km/s simulation (up to 7.8 ps). See text for definition of regions.

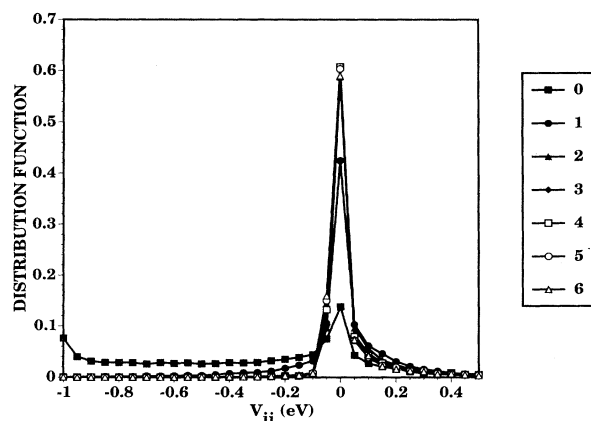


FIG. 10. Normalized distribution of intramolecular terms of Eq. (2) of original $A-B$ pairs for seven regions directly behind the shock front through the 12 km/s simulation (up to 7.8 ps). See text for definition of regions.

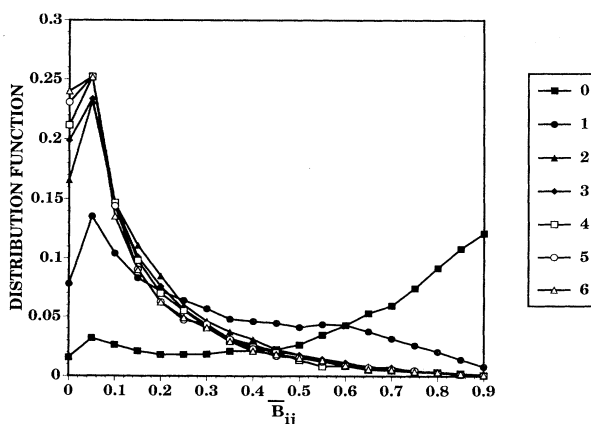


FIG. 11. Normalized distribution of $\overline{B_{ij}}$ terms of Eq. (2) of original $A-B$ pairs for seven regions directly behind the shock front through the 12 km/s simulation (up to 7.8 ps). See text for definition of regions.

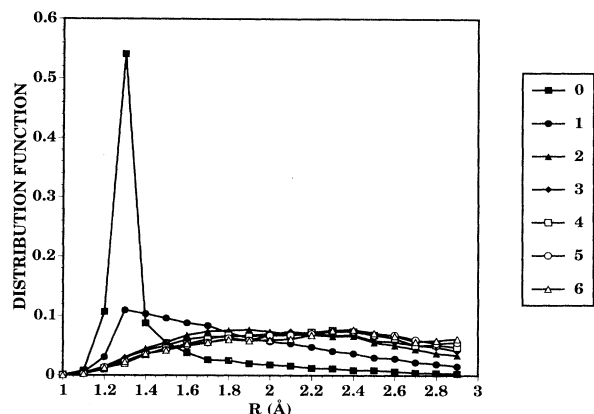


FIG. 12. Normalized distribution of internuclear distances of original $A-B$ pairs for seven regions directly behind the shock front through the 12 km/s simulation (up to 7.8 ps). See text for definition of regions.

($\sim 7 \text{ amu}/\text{\AA}^2$) [10], indicates that the first geometric parameters affected by the shock wave are the intermolecular distances (they are reduced). By region 1, however, the angular orientational distribution has a broad peak at 60° , showing that the $A-B$ molecules are lining up such that the molecular bond axes are nearly perpendicular to the direction of the shock wave propagation. By region 3, the orientational distribution is flat, indicating that the original $A-B$ pairs have experienced complete rotational disorder. By this region, atomization of the species has occurred and the atoms are energetically free to associate with potential molecular partners. We wish to point out that compression of the herringbone lattice in the direction of shock propagation, followed by realignment of the molecular bond axes approximately perpendicular to this direction, places the unreacted $A-B$ molecules in an optimum arrangement to form the homonuclear products. As shown above, the properties of the intramolecular interaction potentials for both heteronuclear and homonu-

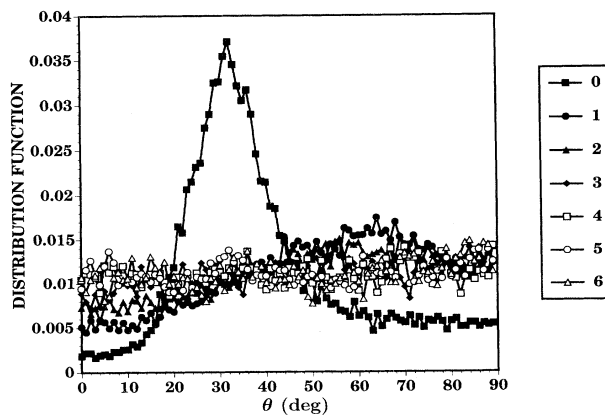


FIG. 13. Normalized distribution of orientation angles of original $A-B$ pairs relative to the crystal x axis for seven regions directly behind the shock front through the 12 km/s simulation (up to 7.8 ps). See text for definition of regions.

clear interactions are such that at any degree of density homonuclear attractions are greater than heteronuclear and homonuclear repulsions are less than heteronuclear. It is possible that the compression followed by realignment of the molecules initially behind the shock front might play some role in the rapidity with which energy redistribution among all modes and completion of reactions have occurred.

These results clearly indicate that A - B molecules within the reaction zone are in transition to an atomic phase due to the changes in the intramolecular interactions caused by the high density behind the shock front. From as close as 2.0 Å behind the shock front, the A - B molecules experience, on average, repulsive interactions. Therefore we have a picture of A atoms and B atoms that are in close proximity to one another, but are not bound. For the regions beyond 0 and 1 the average B_{ij} values and corresponding distributions show little internuclear attraction whatsoever among the atoms, indicating complete atomization of the products. As the shock front proceeds through the crystal, the rarefying flow becomes less dense, increasing the B_{ij} values. Correspondingly, the repulsive intramolecular interactions decrease while attractions increase. Because the well depths of the homonuclear species are larger than those of the heteronuclear molecules, homonuclear product formation is energetically favored; the realignment of the molecules in region 1 puts the system in optimum orientation for homonuclear product formation. All of these factors result in complete reaction of the shocked A - B molecular crystal to exothermically form the homonuclear diatomic products.

Atomization of a molecular solid subjected to hydrostatic high pressure is not unknown. Such phenomena have been observed in high-pressure studies of atomization of solid iodine [14–16], HI [17], and IBr [18]. Although pressure-induced atomization was not specifically investigated in a shock-loading experiment of iodine [19], x-ray spectra taken from these experiments are consistent with the x-ray spectra of the atomized phase of solid iodine at pressures above 21 GPa [14]. *Ab initio* pseudo-potential calculations have shown pressure-induced changes in the electronic structure of molecular crystals, including loss of covalency leading to molecular dissociation with increasing pressure [20]. The electron densities of these materials (Br_2 , I_2 , IBr) determined from x-ray diffraction studies as a function of pressure also show delocalization of the covalent electrons [21]. These rearrangements of the electrons are consistent with the present study and provide the mechanism for the transition from a molecular solid to the atomized solid in several halogen materials. Experimental studies of energetic materials subjected to high pressure have shown that pressure can drastically affect thermal decomposition rates and reaction mechanisms [22,23]. For example, the activation energy for the thermal decomposition of β -HMX (octahydro-1,3,5,7-tetranitro-1,3,5,7-tetrazocine), one of the most widely used explosives, approaches zero at very high pressures [22]. This was attributed to changes in the interaction potential (i.e., electronic structure) of the system under compression. Simi-

larly, the thermal rate of decomposition of nitromethane, a prototypical explosive, and its chemical reactivity were enhanced with increased pressure [23]. These studies, both experimental and theoretical, on real molecular crystals indicate changes in the electronic structure of molecular bonds due to pressure and suggest to us that the pressure-induced atomization mechanism of detonation of our model energetic crystal is not unreasonable. We stress, however, that the interaction potential for our system was not developed or parametrized to yield measured high-pressure behavior. Rather, the reaction mechanism (pressure-induced atomization) is a result of the functional form and choice of parameters for the potential. It may be that this model fortuitously describes correctly a significant reaction mechanism occurring in a detonation. The molecules used in the present study are very simple, containing but a single degree of vibrational motion. It will be interesting to investigate, with these techniques, the effect of large molecules (many vibrational modes) on the energy sharing process and their importance on the reaction mechanisms. It is imperative that either new experimental techniques be developed to probe microscopically the state of the system behind a detonation wave or first principles calculations such as those done by McMahan [20] must be carried out to determine quantitatively the changes in the electronic structure of explosives when subjected to pressures consistent with those measured in detonations. Only then will it be possible to state whether models such as that presented in this work are adequate to represent correctly the chemical reactions leading to detonation.

V. CONCLUSIONS

We have presented a comparative study of models that produce features of an unsupported detonation. The three interaction potentials, based on Abell-Tersoff [11,12] ideas and developed by Brenner and co-workers [8,9], incorporate many-body effects on the intramolecular terms. A many-body term attenuates the molecular attractions with increasing density while increasing the repulsive portion of the term. This changes the equilibrium molecular size and binding energy of a molecule as density increases. When the density becomes large enough, the atoms in a molecule are pushed apart and experience no attraction. In other words, this model predicts pressure-induced atomization.

Results of a two-dimensional molecular dynamics simulation of a detonation in an accompanying study show that, for flier plates moving with velocities greater than 4.7 km/s, unsupported detonations are sustained, and have similar macroscopic properties once the detonation waves reach the steady-state velocity [10]. The detonation wave propagates through the quiescent crystal at 6.6 km/s; immediately behind the detonation wave, there is, on average, a 14 Å wide region in which microscopic and macroscopic properties of the crystal are time independent. It is within this region that the heteronuclear molecules are undergoing reaction; we have labeled this the reaction zone.

We examined in detail seven regions within the 14 Å

reaction zone over time; the cumulative distributions of molecular properties including translational and internal kinetic energies, geometric parameters, and intramolecular potential energies were calculated. The average values of properties of the reaction zone converge to time-independent, constant values approximately 4 Å behind the shock front. Within the 4 Å region immediately behind the shock front, the shock wave first compresses the crystal and rotates the molecules such that the molecular bonds are almost perpendicular to the shock wave propagation. Translational modes of the molecules are excited, but the internal modes of the molecules are cold. Thereafter, energy rapidly transfers into internal modes until an equipartitioning among these kinetic degrees of freedom is established. By this point, the atoms experience an almost completely repulsive intramolecular interaction; in other words, they become atomized. As the shock wave passes, the density in the rarefying flow becomes lower; concurrently the molecular bonding attractions increase, resulting in the association of the atoms to diatomic products. The attractive portions of the homonuclear interaction potentials are considerably larger than the heteronuclear interaction potential; thus homonuclear product formation is energetically favored over heteronuclear recombination. The homonuclear association reactions release substantial chemical

energy which drives the detonation wave. Vibrational excitation of the reactant heteronuclear molecules to dissociation (thermal decomposition) does not contribute to the detonation mechanism. Rather, the mechanism of detonation for this model is pressure-induced atomization of the molecules behind the shock front, followed by association of the atoms to form homonuclear products. This mechanism is attributed directly to the functional form of the many-body term of the interaction potential; subsequent first principles calculations must be done on systems subjected to high pressures to determine if this interaction potential correctly describes a significant mechanism for initiation reactions in detonation.

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- [1] W. Fickett and W. D. Davis, *Detonation* (University of California Press, Berkeley, 1979); W. Fickett, *Introduction to Detonation* (University of California Press, Berkeley, 1985).
- [2] Y. M. Gupta, G. I. Pangilinan, J. M. Winey, and C. P. Constantinou, *Chem. Phys. Lett.* **232**, 341 (1995).
- [3] L. E. Fried and A. J. Ruggiero, *J. Phys. Chem.* **98**, 9786 (1994).
- [4] D. D. Dlott and M. D. Fayer, *J. Chem. Phys.* **92**, 3798 (1990); A. Tokmakoff, M. D. Fayer, and D. D. Dlott, *J. Phys. Chem.* **97**, 1901 (1993); S. Chen, W. A. Tolbert, and D. D. Dlott, *ibid.* **98**, 7759 (1994); S. Chen, X. Hong, J. R. Hill, and D. D. Dlott, *ibid.* **99**, 4525 (1995).
- [5] A. M. Karo, J. R. Hardy, and F. E. Walker, *Acta Astronaut.* **5**, 1041 (1978).
- [6] D. H. Tsai and S. F. Trevino, *Phys. Rev. A* **24**, 2743 (1981).
- [7] T. Kawakatsu and A. Ueda, *J. Phys. Soc. Jpn.* **57**, 2955 (1988); **58**, 831 (1989); T. Kawakatsu, T. Matsuda, and A. Ueda, *ibid.* **57**, 1191 (1988).
- [8] D. W. Brenner, in *Shock Compression of Condensed Matter*, edited by S. C. Schmidt, R. D. Dick, J. W. Forbes, and D. G. Tasker (Elsevier, Amsterdam, 1992), p. 115; D. H. Robertson, D. W. Brenner, M. L. Elert, and C. T. White, *ibid.*, p. 123; C. T. White, D. H. Robertson, M. L. Elert, and D. W. Brenner, in *Microscopic Simulations of Complex Hydrodynamic Phenomena*, edited by M. Mareschal and B. L. Holian (Plenum, New York, 1992), p. 111; C. T. White, S. B. Sinnott, J. W. Mintmire, D. W. Brenner, and D. H. Robertson, *Int. J. Quantum Chem. Symp.* **28**, 129 (1994).
- [9] D. W. Brenner, D. H. Robertson, M. L. Elert, and C. T. White, *Phys. Rev. Lett.* **70**, 2174 (1993).
- [10] B. M. Rice, W. Mattson, J. Grosh, and S. F. Trevino, preceding paper, *Phys. Rev. E* **53**, 611 (1996).
- [11] G. C. Abell, *Phys. Rev. B* **31**, 6184 (1985).
- [12] J. Tersoff, *Phys. Rev. Lett.* **61**, 2879 (1988).
- [13] H. Margenau and G. M. Murphy, *The Mathematics of Physics and Chemistry* (Van Nostrand, New York, 1943), pp. 476–477.
- [14] K. Takemura, S. Minomura, O. Shimomura, and Y. Fujii, *Phys. Rev. Lett.* **45**, 1881 (1980).
- [15] K. Takemura, S. Minomura, O. Shimomura, Y. Fujii, and J. D. Axe, *Phys. Rev. B* **26**, 998 (1982).
- [16] Y. Fujii, K. Hase, N. Hamaya, Y. Ohishi, A. Onodera, O. Shimomura, and K. Takemura, *Phys. Rev. Lett.* **58**, 796 (1987).
- [17] J. van Straaten and I. F. Silvera, *Phys. Rev. Lett.* **57**, 766 (1986).
- [18] Y. Fujii, Y. Ohishi, A. Onodera, K. Takemura, and R. L. Reichlin, *Jpn. J. Appl. Phys.* **24S**, 606 (1985).
- [19] B. J. Alder and R. H. Christian, *Phys. Rev. Lett.* **4**, 450 (1960); A. K. McMahan, B. L. Hord, and M. Ross, *Phys. Rev. B* **15**, 726 (1977).
- [20] A. K. McMahan, *Physica B* **139-140**, 31 (1986).
- [21] Y. Fujii (unpublished).
- [22] G. J. Piermarini, S. Block, and P. J. Miller, *J. Phys. Chem.* **91**, 3872 (1987).
- [23] G. J. Piermarini, S. Block, and P. J. Miller, *J. Phys. Chem.* **93**, 457 (1989).