# Many-body Reduced Vector Solution and Water Vibrations

A.S. ABDEL-RAHMAN\*

Physics Department, Faculty of Science, Cairo University, Egypt

Received: July 30, 2022. Accepted: October 12, 2022.

Reduced mass value and vector are well known for the two-body problem, but the many-body reduced vector problem is not solved yet. The study of many-body problems and their applications (such as vibrational spectroscopy) is one of the more important physical problems. Vibrational spectroscopy provides a powerful tool to perceive the molecular structures and atom motions of molecules. The water molecule is a threebody system stretching vibration that has been previously quantized; their frequencies were defined and showed the infrared (IR) absorption spectrum based on Morse potential. In this work, the reduced mass of the many-body problem is being solved and then used to study the intensity of the stretching vibration modes and show the ratio is in agreement with experiments. The molecule was studied in classical and quantum mechanics to determine its absorption intensity as an example of a reduced mass problem. The results show molecular atomic motions and changes in dipole and reduced mass vector. A Morse-like model for bending was predicted based on the spectroscopic vibration frequency and intensity, defining the bending potential depth of 93.5 kJ/mol.

Keywords: Many-body problem, reduced mass, morse potential, vibration, bending model, water

#### 1 INTRODUCTION

#### 1.1 Two-body system

In classical mechanics, the two-body problem is to predict the motion of two massive objects that are abstractly viewed as point particles. The problem assumes that the two objects interact only with one another; the only force

Corresponding author's e-mail: asabry@sci.cu,edu.eg

affecting each object arises from the other one, and all other objects are ignored.

Let  $r_1$  and  $r_2$  be the vector positions of the two bodies, and  $m_1$  and  $m_2$  be their masses. The goal is to determine the trajectories  $r_1(t)$  and  $r_2(t)$  for all time t, given the initial positions  $r_1(t=0)$  and  $r_2(t=0)$  and the initial velocities  $v_1(t=0)$  and  $v_2(t=0)$ .

When applied to the two masses, Newton's Second Law states that

$$F_{12}(r_1, r_2) = m_1 \ddot{r_1} \tag{1}$$

$$F_{21}(r_1, r_2) = m_2 \ddot{r_2} \tag{2}$$

where  $F_{12}$  is the force on mass 1 due to its interactions with mass 2, and  $F_{21}$  is the force on mass 2 due to its interactions with mass 1. The two dots on top of the r position vectors denote their second derivative with respect to time, or their acceleration vectors.

The vector  $r_{12}$  (the vector between the two bodies) may be used to rewrite Equations 1 and 2 together as:

$$\ddot{r}_{12} = \ddot{r}_2 - \ddot{r}_1 = \frac{F_{21}}{m_2} - \frac{F_{12}}{m_1} \tag{3}$$

The centre of mass, which is affected by all body masses, is written globally as:

$$\sum_{i=1}^{n} m_i \left( r_i - r_{cm} \right) = 0 \tag{4}$$

and is equal to:

$$r_{cm} = \frac{\sum_{i=1}^{n} m_i r_i}{\sum_{i=1}^{n} m_i}$$
 (5)

According to Newton's first law, the forces  $F_{12}$  and  $F_{21}$  are equal in magnitude and opposite in directions, so Equ. 3 may be written:

$$\ddot{r}_{12} = F_{21} \left( \frac{1}{m_2} + \frac{1}{m_1} \right) = \frac{F_{21}}{\mu} \tag{6}$$

where  $\mu$  is the reduced mass and can be written as follows for n-bodies globally:

$$\frac{1}{\mu} = \sum_{i=1}^{n} \frac{1}{m_i} \tag{7}$$

The problem is that for an n-body system, the division of total forces in the system by the system's reduced mass equals the second derivative of the reduced mass vector with respect to time. In a two-body system, the reduced mass vector is that between these bodies, but in a higher-body system, this vector is not defined.

#### 1.2 Molecular vibrations of water

Water molecule structure and vibration modes are discussed in many works. According to the group theory character table, the structure of the water molecule is a  $C_{2v}$  molecule (Table 1) with three vibration modes in the liquid case: (a)  $A_1$  symmetric stretching mode at 3656.65cm<sup>-1</sup> with 0.07 relative intensity, (b)  $B_1$  asymmetric stretching mode at 3755.79cm<sup>-1</sup> with 1.00 relative intensity, and (c)  $A_1$  symmetric bending mode at 1594.59cm<sup>-1</sup> with 1.47 relative intensity [1-7].

The infrared transmittance spectroscopy for vapour [10-15], liquid water [10, 16, 17] and ice [18-21] are reported and show the dependence of absorption lines on the temperature.

The rest structure of the water molecule according to ref [1-3], origin set at centre of mass (cm) below the oxygen atom at  $0.065\text{\AA}$  and above the hydrogen atoms at  $0.521\text{\AA}$  on the z-axiz. Both hydrogen atoms are located  $0.757\text{\AA}$  to the right and left of the z-axis. From here on out, the hydrogen atom located on the negative side of the x-axis will be denoted by  $H_1$  and the atom on the positive side will be denoted by  $H_2$ .

# 1.3 Morse potential

The separation between the hydrogen and oxygen atoms  $(r_e)$  is 0.957Å, with a Morse potential [22-27] (Equ. 8) of dissociation energy  $(D_d)$  and an O-H bond of 424.67kJ/mol [28, 29] is controlled by two coefficients; barrier depth

TABLE 1 C<sub>2v</sub> character table [8, 9].

C <sub>2v</sub>	Е	$C_{2}(z)$	$\sigma_v(xz)$	$\sigma_v(yz)$	linear functions and rotations	Quadratic functions
$A_1$	+1	+1	+1	+1	Z	$x^2, y^2, z^2$
$A_2$	+1	+1	-1	-1	$R_z$	xy
$\mathbf{B}_1$	+1	-1	+1	-1	$x, R_y$	XZ
$B_2$	+1	-1	-1	+1	y, R <sub>x</sub>	yz

(D) and potential width (a), which are related to the force constant ( $k = \mu\omega^2$ :  $\omega$  the angular frequency and equal to  $2\pi v_0$ ) as written in Equ. 9.

$$V = D\left(e^{-2a(r-r_e)} - 2e^{-a(r-r_e)}\right)$$
 (8)

$$a = \sqrt{\frac{k}{2D}} \tag{9}$$

The quantum solution of Morse potential eigenvalues  $(E_n)$  is discovered as described in Equ. 10, while n = 0,1,2,..., which determines the eigenstates, begins with the ground state n = 0 and proceeds to successive states n > 0 [30-32]. The quantum ground state  $(E_0)$  is slightly larger than the classical ground state (D).

$$E_n = h v_0 \left( n + \frac{1}{2} \right) - \frac{1}{4D} \left( h v_0 \left( n + \frac{1}{2} \right) \right)^2 - D \tag{10}$$

$$v_0 = \frac{a}{2\pi} \sqrt{\frac{2D}{\mu}} \tag{11}$$

$$\Delta E_n = E_{n+1} - E_n = h v_0 - (n+1) \frac{(h v_0)^2}{2D}$$
 (12)

## 1.4 Quantum harmonic oscillator

Although the simple harmonic oscillator is not applicable to spectroscopic potentials due to the anharmonicity found in real molecules, this potential was studied as an approximation in this work.

Equation 13 can be used to express the potential of a quantum harmonic oscillator. The solution of these potential eigenvalues  $(E_n)$  is found as described in Equation 14, as that of Morse One, but the harmonicity appears clearly where the difference between any two successive eigenstates is constant. The quantum ground state  $(E_0)$  is also larger than the classical ground state (D).

$$V = \left(\frac{1}{2}k(r - r_e)^2\right) - D = \left(2\pi^2 \mu v_0^2 (r - r_e)^2\right) - D \tag{13}$$

$$E_n = h v_0 \left( n + \frac{1}{2} \right) - D \tag{14}$$

The barrier depth of the harmonic oscillator is deeper than dissociation energy by  $(\frac{1}{2}hv_0)$  and the intercept of classical and quantum potentials is located at  $r_n$ ; and can be written as:

$$\delta_n = r_n - r_e = \pm \sqrt{\frac{2h\nu_0}{k} \left(n + \frac{1}{2}\right)} = \pm \sqrt{\frac{h}{2\pi^2 \mu \nu_0} \left(n + \frac{1}{2}\right)}$$
 (15)

where  $\delta_n$  is the change in the O-H bond length.

## 1.5 Classical and quantum mechanics

The forces between macroscopic systems are restricted by the classical equations of motion. This application is ideal because quantum mechanics equations, as well as the application of this mechanics to this range, are extremely difficult to solve. Going from the microscopic to the atomic level, the motion of electrons around the nucleus cannot be explained by classical mechanics, while quantum mechanics and its main point of quantized energy levels are the appropriate consideration for this level.

Many scientists believe that these two mechanics systems are actually the same, but this system is closer to the classical one for macroscopic ranges and the quantum one for microscopic ranges. One can imagine combining these two mechanics systems between these scales, such as the interatomic scale.

Harmonic oscillators are one of the important problems in mechanics. These oscillators can be found at the molecular level as vibrational interatomic atoms. The interpretation of these oscillator energies can be considered from either a classical or quantum perspective. In contrast to classical mechanics, which describes continuous energies, quantum mechanics depicts discrete energy levels. The disparity between these solutions to the same problem can be interpreted as a measurement agreement.

The quantum mechanics solution illustrates an energy level starting with ground level  $E_0$  and rising to successive higher levels. The transitions between these levels are governed by selection rules. According to classical fact, all atoms vibrate (at the ground energy level) continuously until they reach absolute zero temperature. At this temperature, one cannot expect the stopping of the motion of electrons around the nucleus where they obey quantum mechanics, but the stopping of vibrations is possible where it is classically interpretative.

From the quantum solution, the ground's vibrational energy level is located over the barrier potential. This difference can be linked to classical thermal energy as follows:

$$(E_0 - D) = (D_d - D) \propto k_B T \tag{16}$$

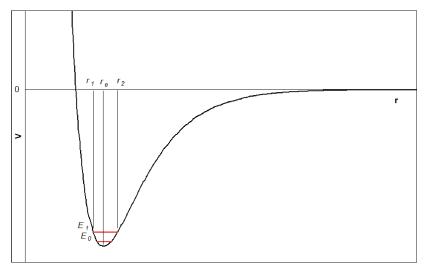


FIGURE 1 Morse potential and amplitudes  $r_1$  and  $r_2$ .

where  $k_B$  is the Boltzmann constant, where the amplitude of oscillation is a classical view, and the total energy of the oscillator system  $E_t$  (the sum of potential and kinetic energies) should meet the potential at the amplitude of oscillation (typically  $r_1$  and  $r_2$  in Fig. 1). From the point of view of quantum mechanics, there is no definite oscillation amplitude, but there is a probability of finding the particle of a certain length,  $\Delta r$  equal to the square of the eigenfunction.

The convergence between classical and quantum mechanics in the interatomic vibrations is to consider the oscillator total energy  $E_t$  equal to the quantized energy level  $E_1$  and hence intercepting at the oscillation amplitudes  $r_1$  and  $r_2$  in Fig. 1. So, the vibration oscillator energy will be treated as a quantum problem, while the vibration amplitude will be solved classically.

The zero temperature case makes the barrier depth equal to the ground energy level with zero vibrational amplitude ( $r_1 = r_2 = r_e$ ). This scenario is acceptable to discuss the vibrational modes of molecules.

Upon this assumption, the barrier width a and transition energy (and thus frequency  $v_0$ ) should be temperature-dependant. This also agrees with the experimental measurements of the IR transmittance spectra of water.

The IR vibration is caused by a change in the separation of the negative charge (the oxygen ion) from the net positive charge (midway between the two hydrogen ions) in the molecule. From this point, the importance of discussing the atomic locations through vibration is clear.

In this work, the n-body system's reduced mass vector is discussed, and the three vibration modes of water (as an example of a three-body problem) are also discussed by classical and quantum solutions, their intensities, and atom motion vectors. The total potential of the water molecule in this work is taken as the summation of the Morse potentials between  $H_1$ -O and  $H_2$ -O, while the weak Van der Waals potential between hydrogen atoms will be neglected (its values are very small compared to the Morse potential).

## 2 METHODOLOGY

## 2.1 Many-body system

To expand the two-body system to many body system, first of all, the force  $F_{21}$  in Equ. 6 will be replaced by the sum of all forces between each pair of bodies. After that, how to define the reduced mass vector?

To solve this question, the two-body unequally reduced mass vector can be written as:

$$r_{12} = r_2 - r_1 = (r_2 - r_{cm}) - (r_1 - r_{cm})$$
(17)

The reduced mass vector direction always takes one of the  $r_1$  and  $r_2$  directions (may be the larger one). According to this point, the unit vector of the reduced mass vector  $\hat{r}_{12}$  can be written as:

$$\widehat{r}_{12} = \frac{(r_2 - r_{cm}) + (r_1 - r_{cm})}{\left| (r_2 - r_{cm}) + (r_1 - r_{cm}) \right|}$$
(18)

and can be written globally as:

$$\hat{r} = \frac{\sum_{i=1}^{n} (r_i - r_{cm})}{\left| \sum_{i=1}^{n} (r_i - r_{cm}) \right|}$$
(19)

Now, one should differentiate between the reduced mass vector's magnitude and direction  $\hat{\eta}_1$ . Its magnitude is the summation of the magnitudes of the vectors from the two bodies to the centre of mass. Then Equ. 17 can be written in terms of  $\hat{\eta}_2$  as:

$$r_{12} = (r_2 - r_{cm}) - (r_1 - r_{cm}) = (|r_2 - r_{cm}| + |r_1 - r_{cm}|)\hat{r}_{12}$$
 (20)

and can be written globally as:

$$r = \left(\sum_{i=1}^{n} |r_i - r_{cm}|\right) \hat{r} = \frac{\sum_{i=1}^{n} |r_i - r_{cm}|}{\left|\sum_{i=1}^{n} (r_i - r_{cm})\right|} \sum_{i=1}^{n} (r_i - r_{cm})$$
(21)

This is the n-body system's reduced mass vector; its second time derivative equals the summation of all forces in the system divided by the system's reduced mass.

#### 3 RESULTS AND DISCUSSION

#### 3.1 Vibration and center of mass

According to the definition of vibration in molecules, the centre of mass is reserved. Where the movement of the centre of mass is achieved through vibration and rotation operations, this is solved by introducing the translation of molecules alongside the operations.

Herein, the centre of mass conservation sets two main equations for all vibration modes:

$$m_O Z_O' + m_H \left( -Z_{H1}' - Z_{H2}' \right) = 0$$
 (22)

$$m_O X_O' + m_H \left( -X_{H1}' + X_{H2}' \right) = 0$$
 (23)

The Z and X are the subscripted atoms' locations, the prime over the location parameters refer to the location via vibration, and the unprimed parameters refer to the rest of the molecule, and they are written in the introduction section.

The vibration operation is divided into some modes: stretching, bending (in-plane and out of plane), twisting, scissoring, and rocking. The fact that each of these modes is active alone means that the stretching vibration requires the changing of bond lengths without changing the bond angles [9, 33-35], and so on.

## 3.2 Symmetric stretching vibration of water

According to the  $C_{2\nu}$  character table, the  $A_1$  symmetric stretching vibration of water can be characterized. This vibration mode conserves all symmetry elements. This mode of vibration allows both hydrogen atoms to go away from

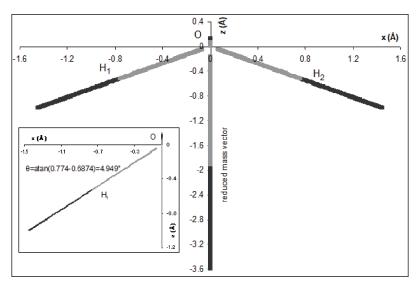


FIGURE 2 The water  $A_1$  symmetric stretching vibration.

the oxygen atom by the same displacement ( $\delta$ ) ( $Z'_{H1} = Z'_{H2}$  and  $X'_{H2} = -X'_{H1}$ ) and, of course, the oxygen atom goes up on the z-axis to conserve the centre of mass ( $X'_O = 0$ ). Also, the angle between O-H<sub>1</sub> and O-H<sub>2</sub> vectors is conserved, so the atomic motions of hydrogen atoms incline by about 5° (as shown in Figure 2, inset figure, where this motion path passes the origin or centre of mass) on the vector between oxygen and hydrogen atoms.

Figure 2 illustrates the water molecule's symmetric stretching vibration  $A_1$  along with the atomic motions and reduced mass vector. The darker points represent the stretching, while the lighter points represent the compression motion. If the  $H_1$  and  $H_2$  atoms move away from the oxygen atom in the mentioned direction by displacement ( $\delta$ ), all the atoms' equations of motion in terms of  $\delta$  are listed in Table 2 beside the reduced mass vector r and the dipole length d.

While both O-H bonds have synchronised motions, the total system potential has double the O-H dissociation energy as the ground state of the Morse potential, so:

$$\mu \ddot{r} = F = -\frac{\partial V}{\partial r} \hat{r} \tag{24}$$

or in scalar form

$$\ddot{\delta} = \frac{4aD}{1.9877\mu} \left[ e^{-2a\delta} - e^{-a\delta} \right]$$
 (25)

Atom	Description
$H_1$	$(-0.7569 - 0.7908\delta)x + (-0.5203 - 0.5436\delta)z$
$H_2$	$(0.7569 + 0.7908\delta)x + (-0.5203 - 0.5436\delta)z$
0	$(0.065554 + 0.068486\delta)z$
r	$(-1.9026 - 1.9877\delta)$
d	$(0.58587 + 0.6121\delta)$

TABLE 2 A<sub>1</sub> symmetric stretching vibration atomic motions.

This deferential equation may be solved numerically to get the function  $\delta(t)$ . But the importance of this problem is to get the values of D and a of the total system potential, which can be calculated by means of the eigenstates of the potential as follows:

$$-2D_d = E_0 = \frac{1}{2}hv_0 - \frac{1}{16D}(hv_0)^2 - D$$
 (26)

$$hv_{sy.str.} = \Delta E_0 = hv_0 - \frac{(hv_0)^2}{2D}$$
 (27)

$$v_0 = \frac{a}{2\pi} \sqrt{\frac{2D}{\mu}} \tag{28}$$

where  $v_{sy.str}$  the frequency of the symmetric stretching vibration of water molecules (=1.096 × 10<sup>14</sup>Hz). The solution of the Morse potential parameters (Figure 3) was achieved numerically and is listed in Table 3.

The IR vibration active mode is a result of the change of molecule dipole moment (or dipole length) through the vibration process.

As shown in Table III, the atomic vibration frequency and barrier potential are slightly larger than the symmetric stretching vibration frequency and double the dissociation energy, respectively. The reduced mass vector vibration amplitude has anharmonicity due to the oscillation's unequal displacements.

## 3.3 Asymmetric stretching vibration of water

The  $B_1$  asymmetric stretching vibration of water conserves the xz mirror and totally loses the  $C_2$  fold axis and xy mirror plane. This mode of vibration allows one hydrogen atom to go away from the oxygen atom while the other one moves towards the oxygen atom by the same displacement ( $\delta$ ) and, of course, the oxygen atom moves parallel to the x-axis to conserve the centre of

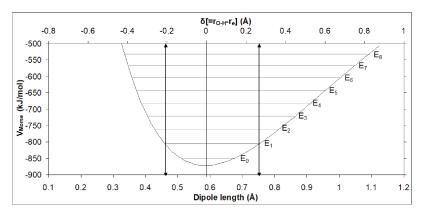


FIGURE 3
The water A<sub>1</sub> symmetric stretching vibration potential.

TABLE 3 A<sub>1</sub> symmetric stretching vibration parameters of water.

Parameter	Description	Value	Unit
$\nu_0$	Atomic vibration frequency	$1.1252 \times 10^{14}$	Hz
D	Barrier potential	871.66	kJ/mol
k	Vibration force constant	$4.055 \times 10^{2}$	N/m
a	Barrier width	$1.1835 \times 10^{10}$	$m^{-1}$
$h_{min}$	O-H minimum vibration bond length	0.75	Å
$h_{max}$	O-H maximum vibration bond length	1.228	Å
$\Delta r$	Reduced mass vector vibration amplitude	0.5387 - 0.4115	Å
$\Delta d$	Dipole length vibration amplitude	0.1659 - 0.1267	Å

mass ( $Z'_{O} = 0$ ). The angle between O-H<sub>1</sub> and O-H<sub>2</sub> vectors is conserved in any stretching vibration.

Figure 4 illustrates the water molecule's asymmetric stretching vibration  $B_1$  along with the atomic motions and reduced mass vector. The hydrogen atom moves in a straight line at an angle of nearly 5.57° (as shown in the inset figure of Figure 8), on the vector between oxygen and hydrogen atoms.

Table 4 shows the equation of motion of each atom beside the reduced mass vector under the influence of the hydrogen atom vibration amplitude ( $\delta$ ). One can notice that both the reduced mass vector and the dipole length are directly proportional to the square of the hydrogen vibration amplitude.

Since the motion of hydrogen atoms is out of phase (one atom moves towards the oxygen atom while the second one moves away), the sum of their O-H Morse potentials is close to the harmonic oscillator. This harmonic oscil-

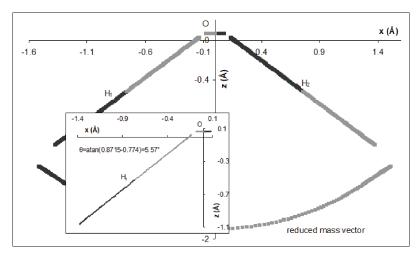


FIGURE 4 The water  $B_1$  asymmetric stretching vibration.

TABLE 4  $B_1$  asymmetric stretching vibration atomic motion.

Atom	Description
$H_1$	$(-0.7569 + 0.7023\delta)x + (-0.5203 + 0.6121\delta)z$
$H_2$	$(0.7569 + 0.7023\delta)x + (-0.5203 - 0.6121\delta)z$
0	$(-0.08848\delta)x + (0.065554)z$
r	$(1.90259 + 0.08623\delta^2)$
d	$(0.58587 + 0.04734\delta^2)$

lator depth should be double the O-H depth. The ground state energy (which equals the double of dissociation energy) should be larger than this depth by  $(\frac{1}{2}hv_0)$  and the separation between two successive energy states is quantized  $(hv_0)$ . The total potential of this system should be written as:

$$V_{tot} = D\left(e^{-2a\delta} + e^{2a\delta} - 2e^{-a\delta} - 2e^{a\delta}\right) \approx \frac{k'}{2}\delta^2 - 2D \tag{29}$$

where k' denotes the system's vibration force constant, which differs from the Morse potential force constant (mentioned in Equ. 9). After applying the spectroscopic coefficient ( $v_0 = 1.126 \times 10^{14}$  Hz), the potential depth D and the force constant k' are calculated to be 871.816 kJ/mol and  $4.06 \times 10^2$  N/m, respectively.

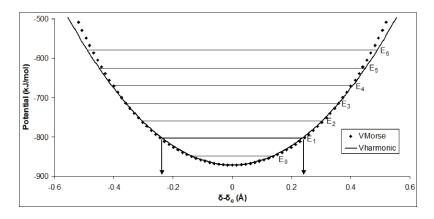


FIGURE 5 The water  $B_1$  asymmetric stretching potential.

The harmonic potential of this vibration (continuous curve in Figure 5) can be fitted to subtract two identical Morse potentials (dots curve in Figure 5), and then the Morse potential width a fitted to be  $1.13 \times 10^{10}$  m<sup>-1</sup>. The first six energy levels are set on the same figure, and  $E_1$  intercepts the potential at  $\pm 0.2348$ Å. Thus, the changes in reduced mass vector and dipole length are 0.00475Å and 0.0238Å, respectively.

The ratio of the dipole length change to the change in the reduced mass vector between symmetric and asymmetric vibrations is 0.0615:1. This ratio should represent the intensity of the experimental IR absorption lines and is found in good agreement with them [3, 36].

## 3.4 Symmetric bending vibration of water

The  $A_1$  symmetric bending vibration of water conserves all symmetry elements by synchronising the scissor motion of the hydrogen atoms while the oxygen atom moves on the z-axis to keep the centre of mass at the origin. Let's define  $\theta_e$  as the equilibrium bond angle; the vibration is mainly associated with the difference between the bond angle and its equilibrium value while the bond length is constant. Figure 6 illustrates the water molecule's symmetric bending vibration  $A_1$  along with the atomic motions and reduced mass vector. Table 5 shows these motions with respect to  $\varphi = (\theta - \theta_e)$ .

There is a definite potential model for bending [37-40], In this work; a simple model for bending is presented that resembles Morse potential in its shape. In bending vibration, while the bond lengths being constant, the effective term in the potential is directly proportional to  $\varphi$  (instead of  $\delta$  in stretching modes). The potential is really affected by the arc length in front of the bond angle, so the term should be written as  $r_e \varphi$ , and then the potential equation may be written as:

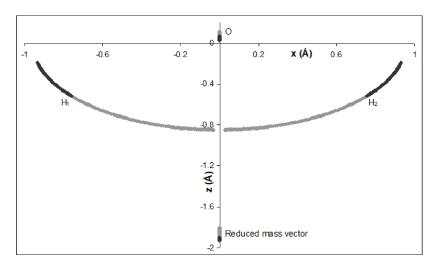


FIGURE 6 The water  $A_1$  symmetric bending vibration.

TABLE 5 A<sub>1</sub> symmetric bending vibration atomic motions.

Atom	Description
$H_1$	$(-0.75693 - 0.2808\phi + 0.0814\phi^2)x - (-0.5203 - 0.324\phi - 0.0735\phi^2)z$
$H_2$	$(0.75693 - 0.2808\phi - 0.0814\phi^2)x - (-0.5203 - 0.324\phi - 0.0735\phi^2)z$
0	$(0.065553 - 0.0408\phi - 0.0093\phi^2)z$
r	$(1.902594 - 0.0462\phi - 0.0107\phi^2)$
d	$(0.58587 - 0.3648\phi - 0.0827\phi^2)$

$$V = D\left(e^{-2ar_e\phi} - 2e^{-ar_e\phi}\right) \tag{30}$$

where both coefficients a and D still have their earlier definitions. Once applying the spectroscopic vibration frequency  $(4.78 \times 10^{13} \text{ Hz})$  and taking into account that the IR absorption line intensity should be 1.47 times the  $B_1$  stretching line, the dipole change to reduced mass vector change should be 7.365, and then the angle difference should be 0.2276 rad. This value is the intercept of the potential equation with the  $E_1$  energy state. Hence, one can estimate the potential barrier width and depth as shown in Table 6.

The water-bending potential model is being predicted and shown in Figure 7. The reason why the difference between ground state energy (29 kJ/mol) and potential depth is larger than that between ground and first energy states is the nearness between the vibrating energy (165.5 kJ/mol) and the barrier depth.

Parameter	Description	Value	Unit
$\nu_0$	Atomic vibration frequency	$4.147 \times 10^{14}$	Hz
D	Barrier potential	93.51	kJ/mol
k	Vibration force constant	$5.507 \times 10^{3}$	N/m
a	Barrier width	$1.332 \times 10^{11}$	$m^{-1}$
$\Theta_{min}$	H-O-H minimum vibration bond angle	101.55	۰
$\theta_{max}$	O-H maximum vibration bond angle	117.56	۰
$\Delta r$	Reduced mass vector vibration amplitude	1.8994 - 1.9147	Å
$\Delta d$	Dipole length vibration amplitude	0.6053 - 0.4961	Å

TABLE 6
A<sub>1</sub> symmetric bending vibration parameters of water.

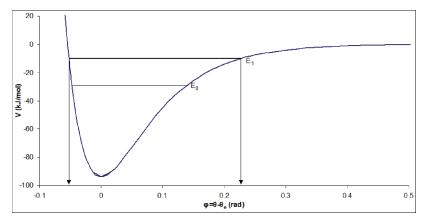


FIGURE 7 The water bending potential model.

#### CONCLUSIONS

Calculating the reduced mass vector is essential for describing the atomic motions of molecules and estimating the dipole moment, which describes the IR absorption of the molecule through vibration modes. The work shows the reduced mass vector of the many-body problem. Quantum and classical mechanics are used side by side to describe the vibration mechanisms in water molecules, which show the relative intensity of the IR absorption lines of water too close to the measured lines. The reason for the very high absorption of the  $B_1$  asymmetric stretching of water was clear, due to its higher stretching hormoticity than others. A novel prediction of the bending potential was presented and illustrates that the bending potential depth of water is 93.5 kJ/mol.

# **Conflict of Interest**

The authors declare no conflict of interest.

## **Data Availability Statement**

The datasets generated during and/or analyzed during the current study are available from the corresponding author on reasonable request.

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