

Silicon-Based Molecular Switch Junctions

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Received: 4 May 2009 / Revised: 3 June 2009 / Accepted: 3 June 2009

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ABSTRACT

In contrast to the static operations of conventional semiconductor devices, the dynamic conformational freedom in molecular devices opens up the possibility of using individual molecules as new types of devices such as a molecular conformational switch or for molecular data storage. Bistable molecules—such as those having two stable *cis* and *trans* isomeric configurations—could provide, once clamped between two electrodes, a switching phenomenon in the non-equilibrium current response. Here, we model molecular switch junctions formed at silicon contacts and demonstrate the potential of such tunable molecular switches in electrode/molecule/electrode configurations. Using the non-equilibrium Green function (NEGF) approach implemented with the density-functional-based tight-binding (DFTB) theory, a series of properties such as electron transmissions, current–voltage characteristics in the different isomer conformations, and potential energy surfaces (PESs) as a function of the reaction coordinates along the *trans* to *cis* transition were calculated for two azobenzene-based model compounds. Furthermore, in order to investigate the stability of molecular switches under ambient conditions, molecular dynamics (MD) simulations at room temperature were performed and time-dependent fluctuations of the conductance along the MD pathways were calculated. Our numerical results show that the transmission spectra of the *cis* isomers are more conductive than *trans* counterparts inside the bias window for both model compounds. The current–voltage characteristics consequently show the same trends. Additionally, calculations of the time-dependent transmission fluctuations along the MD pathways have shown that the transmission in the *cis* isomers is always significantly larger than that in their *trans* counterparts, showing that molecular switches can be expected to work as robust molecular switching components.

KEYWORDS

Molecular electronics, molecular switches, charge transport, silicon–molecule interface

Introduction

Since the successful measurement of current flow through individual molecules in the past few decades, various kinds of molecular devices have been demonstrated by many groups, showing interesting properties such as negative differential resistance [1]

and memory effects [2]. The realization of electronic function at the molecular scale, envisioned by Aviram and Ratner in 1974 [3], is gradually finding experimental support due to the development of scanning probe technologies [4, 5]. Nevertheless, many challenges still remain. The conductance of a molecule connected between contacts changes

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dramatically depending on many parameters such as the coupling strength between the molecule and the contacts, the electronic structure of the molecule and the contacts, and molecular conformational changes. Therefore, it is highly desirable to understand the relationship between these parameters and transport properties in the design of controllable molecular devices.

In contrast to the static operations of conventional semiconductor devices, the dynamic conformational freedom in molecular devices opens up the possibility of using individual molecules as a new type of device such as a conformational molecular switch or for molecular data storage. The ability to control the conductance at the single molecule scale will have a great impact in nanoscale electronics as well as in other fields such as chemical sensors [6] or medical diagnostics [7]. Photochromic organic molecules having bistable characteristics are attractive materials for the design of molecular switches since a great variety of properties, such as their molecular conformations and their electronic structures, can be reversibly altered from one state to another by external stimulations such as light irradiation and current pulses. Azobenzene, diarylethene, and their derivatives are representative photochromic molecules. Recently, there have been many theoretical and experimental studies of molecular switches [8]. Theoretical studies have explored the use of azobenzene as a molecular switch when connected by Au electrodes [9, 10] or carbon nanotubes (CNTs) [11]. Promising results were obtained, as a significant difference in conductance was seen between the two conformations. Additionally, the study of diarylethene derivatives connected to Au contacts also has shown large changes in conductance between open or closed forms [12].

Several groups have addressed the experimental fabrication of molecular switches consisting of molecules with bistable characteristics sandwiched between contacts. Dulic et al. have created diarylethene derivative-based molecular switch junctions connected between gold contacts by means of the mechanically controlled break junction (MCBJ) technique and measured the current flow through the junction before and after UV irradiation [13]. The

current flow through the molecule was suddenly increased when subjected to UV irradiation. Additionally, the current flow did not decay after stopping the irradiation. Although reversible switching has not been observed, conformational switching and the control of current flow through the single molecular device have been realized.

Lörtscher et al. have fabricated molecular junctions where single bipyridyl-dinitro oligophenylene dithiole molecules have been connected between gold contacts with the MCBJ technique [14]. They have demonstrated reversible molecular switching by applying current pulses across the gold contacts. Additionally, Henzl et al. [15, 16] and Choi et al. [17] have reported reversible *cis–trans* isomerization of a single azobenzene derivative on Au(111) surfaces induced by the inelastic excitation of vibrational states by an STM tip. Furthermore, Alemani et al. have reported mechanical reversible isomerization of a single azobenzene derivative on the Au(111) surface induced by an electric field applied from an STM tip not in contact with the molecule [18–20].

In this article, we explore the potential of azobenzene-based molecules as reversibly tunable molecular switches in electrode/molecule/electrode configurations. We have modeled two kinds of molecular switch junctions formed at silicon contacts. In order to improve the molecule-contact coupling, the molecules are covalently attached to Si contacts. We present a detailed analysis of the energetics, potential energy surface (PES) as a function of the reaction coordinate along the *trans* to *cis* transition, the stability along molecular dynamics (MD) pathways, and transport properties in the two conformations using the non-equilibrium Green function (NEGF) approach implemented with the density-functional-based tight-binding (DFTB) method.

This paper is organized as follows. In Section 1, we present details of the computations used in our study. In Sections 2.1 and 2.2, we show the molecular structures, electronic structures, and PESs of molecular switches in the gas phase and connected between Si contacts. In Section 2.3, we show the transport properties and the stabilities of the molecules between Si contacts. We conclude in Section 3.



1. Method

The relaxation of the molecular structures, total energy calculations, and electronic structure calculations are based on a second-order expansion of the Kohn–Sham total energy in density functional theory with respect to charge density fluctuations [21]. The zeroth order approach is equivalent to a common standard non-self-consistent tight-binding scheme, while at second order a transparent, parameter-free, and readily calculable expression for generalized Hamiltonian matrix elements can be derived. The resulting DFTB method provides accurate results comparable to the results obtained with *ab initio* DFT methods or first principles calculations, while requiring fewer computational resources. The method involves modification by self-consistent redistribution of Mulliken charges. The computations in the DFTB+ implementation employed use a minimal basis set, a two-center approximation, neglect of three centers and a crystal field. The atomic pair potentials are computed at a pre-computation stage using a DFT code. The Slater-Koster integrals are stored (as a function of the interatomic distance) in look-up tables which are used during the actual simulation. DFTB+ has been extensively validated in the past years by many applications related to, e.g., semiconductor surface reconstructions [22–25], reaction of organic molecules [22, 26, 27], and biomolecules [22, 28–30].

Conductance calculations were done via *gDFTB* code [31, 32]. The *gDFTB* code is an extension of the DFTB method to include the NEGF formalism combined with the Landauer formalism. The *gDFTB* code is based on a tight-binding representation of the wavefunction and a density functional formulation of the Coulomb and exchange correlation potentials. The conductance at low or zero bias (equilibrium) is obtained from electronic transmission probability T as $G = G_0 T(E, V=0)$, where $G_0 = 2e^2/h$ is the quantum conductance, e is the electron charge, and h is the Plank constant. The electronic transmission probability is related to the retarded/advanced Green function of the molecular region $G^{R/A}(E)$ and the self-energies of the electrodes $\Sigma^{R/A}(E)$ by the Fisher and Lee relation [33] $T(E, V) = 4\text{Tr}[G^R(E)\text{Im}\cdot\Sigma_L(E)G^A(E)\text{Im}\Sigma_R(E)]$. We emphasize that when calculating the current $I(V) = (2e/h)$

$\int dE [f_L(E) - f_R(E)] T(E, V)$, the full voltage-dependent transmission, $T(E, V)$ is calculated through a self-consistent Poisson-NEGF cycle in *gDFTB*. Here, $f_{L/R}(E)$ are the Fermi functions of the left/right electrodes. The minimal basis set used in *gDFTB* could be responsible for inaccuracies in the calculation of the current. This may result in problems describing transport mediated via the tails of the wavefunctions. This problem could be solved by using an *ab initio* code with a richer basis set, but would dramatically increase the computational requirements without adding any qualitatively new results to our treatment.

2. Results and discussion

Figure 1(a) shows the two model systems considered in this work. Two different pairs of linkers are attached to the azobenzene molecules. We refer to these two model systems as system A and system B. In each system, the two linkers covalently connecting the molecule to silicon contacts are substituted at the *meta*- and *ortho*-positions of the phenyl rings. Large movements of one phenyl ring involved with the *cis*–*trans* isomerization are unfavorable for the stable operation of molecular switching since such large movements prevent stable coupling between the molecule and Si contacts. However, via the

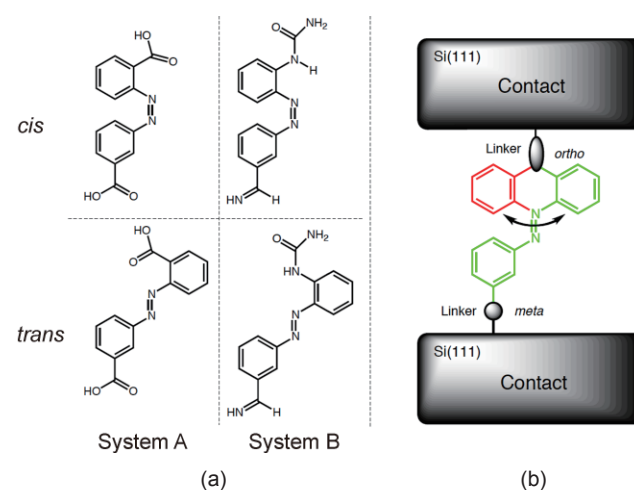


Figure 1 (a) The two azobenzene derivatives modeled in this work and (b) a schematic illustration of an azobenzene-based molecular switch connected to Si contacts via linkers. The two linkers are substituted at the *meta*- and *ortho*- positions of the phenyl rings. These substitution positions reduce the movement of contact atoms during *cis*–*trans* isomerization

linkers at the *meta*- and *ortho*-positions, unfavorable displacement at the molecule-contact interface can be reduced as shown in Fig. 1(b). Additionally, the change in molecular length on going from one isomeric state to the other is small (see Table 1) so that the molecules do not tilt on undergoing the *cis*–*trans* isomerization. The atoms at the contacts do not move throughout the *cis*–*trans* isomerization. Thus, the choice of these positions for the substituents is preferable for the stable operation of molecular switching.

Table 1 Molecular length of two molecular switches in gas phase: The molecular lengths are defined by the atomic pairs in parentheses which make bonds with silicon surfaces at top and bottom. These values are obtained after structure optimization by DFTB+ calculations. For the detail see the text.

	System A (O–O)	System B (N–N)
<i>cis</i>	10.208 Å	11.383 Å
<i>trans</i>	11.000 Å	10.685 Å

2.1 Molecules in the gas phase

For the determination of the stable configurations of the molecules between the Si contacts, we first performed a geometry optimization for the isolated molecules by means of the conjugate-gradient technique. The geometry optimization was carried out until the absolute value of the interatomic force reduced to less than 10 atomic units. We used the Slater–Koster (SK) parameters developed by Elstner et al. for the C, H, N, O atoms [22] and those of Szuets et al. for the Si atom [24, 25] as basis sets. Figures 2(a) and 2(c) show the optimized structures, the frontier orbitals, and the molecular energy levels of systems A and B, respectively. The *trans* isomers are characterized by planar structures while in each *cis* isomer two phenyl rings are twisted relative to each other by 12.9° in both systems A and B.

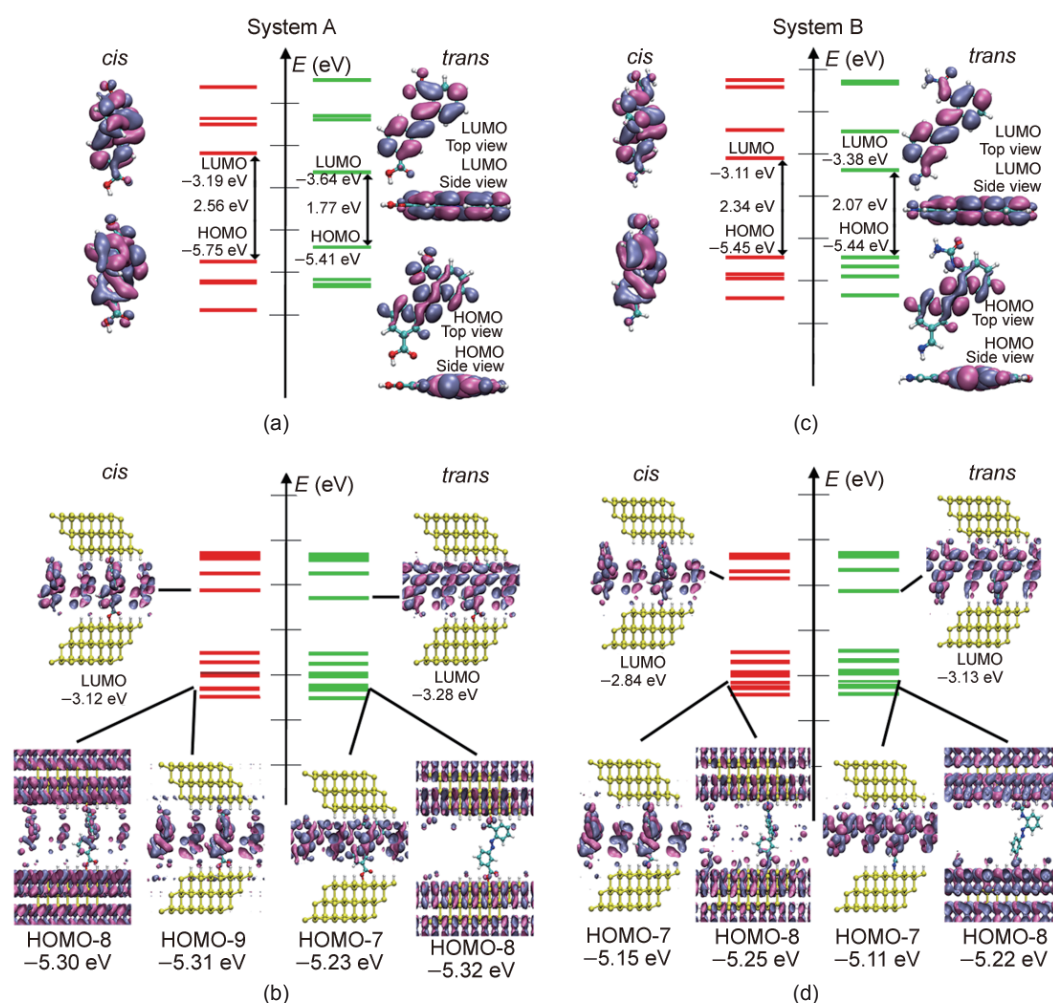


Figure 2 Energy level diagram and frontier orbitals of system A (a) in the gas phase and (b) between Si contacts. Energy level diagram and frontier orbitals of system B (c) in the gas phase and (d) between Si contacts

Figure 3(a) shows the PES of total energy as a function of the reaction coordinates along the *trans* to *cis* transition for the two model systems in the gas phase. The total energy calculations showed that the *trans* isomers in systems A and B are energetically more stable than their *cis* counterparts by 0.193 eV and 0.277 eV, respectively. The *trans* to *cis* potential energy barriers for systems A and B are 1.060 eV and 1.433 eV, respectively. These features are similar to the azobenzene molecule in the gas phase where its *trans* isomer is experimentally reported to be more stable than the *cis* counterpart by 0.6 eV and the potential energy barrier from the *trans* to the *cis* counterpart is 1.0 eV [34].

2.2 Molecules between Si contacts

Let us consider the model systems connected between Si contacts. Hereafter we refer to these coupled systems as “extended molecules”. The

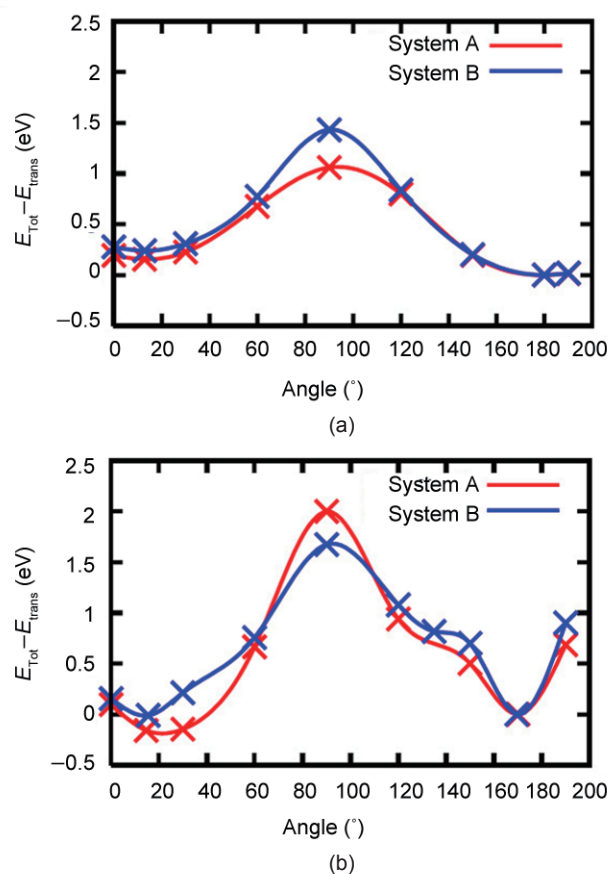


Figure 3 Potential energy surfaces of systems A and B (a) in the gas phase and (b) between Si contacts. The total energies of both systems are plotted as a function of the dihedral angle between the two bonds connecting nitrogen atoms and phenyl rings. The curves interpolate points, which are indicated by the crosses

complexity of the systems leads to a very complicated PES with many metastable states; for simplicity we only consider the vertical arrangements of the molecules with respect to the hydrogen-passivated Si(111) surface. The structure-optimized molecules in the gas phase are covalently connected to hydrogen-passivated Si(111) surfaces and relaxed again using DFTB+ with periodic boundary conditions. The lateral spacing between azobenzene molecules is set large enough to exclude lateral interactions between them. In each unit cell, the top/bottom silicon contacts are comprised of 54 Si atoms forming three silicon layers and 8 hydrogen atoms covering the Si(111) surface. In order to include the reconstruction of the silicon atoms on the surfaces, the first two silicon layers of each electrode from the surface are relaxed, while the third layer remains fixed, which is required for the subsequent conductance calculations. The Si–Si bond lengths in the fixed layers are set as 2.35 Å, which is the experimental value for bulk Si crystal. Figure S-1 in the Electronic Supplementary Material (ESM) shows the total energy of the extended molecules as a function of the separation of two Si surfaces. Figure 4 shows the optimized unit cells of two model systems at the separation of Si surfaces where the total energies of *trans* isomers between Si contacts are their local minima, which are shown by vertical dotted lines in Fig. S-1 (in the

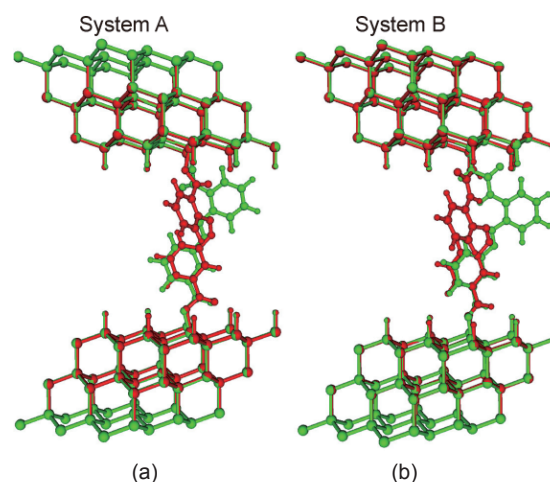


Figure 4 Optimized structure (unit cell) of the two molecular switch junctions modeled in this work. *cis* conformations are shown in red and *trans* conformations in green. For comparison, the two isomer conformations are superimposed. Optimizations are carried out using the conjugated-gradient technique with DFTB+ under periodic boundary conditions. The silicon layers at the bottom and top are fixed

ESM). The PES calculations for extended molecules and subsequent conductance calculations are carried out at these separations.

The PESs of total energies in the extended molecules are shown in Fig. 3(b). The potential energy barriers of extended molecules along the reaction coordinate from *trans* to *cis* isomers in systems A and B are 2.001 eV and 1.675 eV, respectively. The potential energy barriers are increased compared with those in the gas phase (Fig. 3(a)). This is due to the reduction of the conformational degree of freedom of molecules confined between Si contacts.

2.3 Transport properties

Once the relaxed geometries have been determined, we first focus on the electronic transmission probabilities of molecular switch junctions for *cis* and *trans* isomer conformations at equilibrium. The structure-optimized unit cell of the extended molecules is put between semi-infinite Si bulk layers and then transmission calculations are carried out. Figure 5 shows the electronic transmission probabilities at equilibrium. The Fermi energies are near the edge of the valence band (for the Fermi energies, see Table 2). Thus, the conductance in these model systems is driven by hole transport under small bias. Our results show that the *cis* isomers are more conductive than the *trans* counterparts around the Fermi energies in both of the model systems. The low conductance between the resonant peaks indicates that the transport can only take place through tunneling. The conductance around the Fermi energy in *cis* isomers is higher than that for the *trans* counterparts by a few orders of magnitude. This suggests that switching behavior may be expected at low bias. The low transmissions around the Fermi energies are due to the weak coupling between the molecules and the Si contacts giving rise to narrow resonant peaks. As shown in the density of states (DOS) profiles of both system A and B in Fig. 6, the narrow resonant peaks due to the HOMO and LUMO of isolated molecules are located at about -5.0 eV and -3.0 eV, respectively. The Fermi energies lie in the center of gaps between these narrow resonant peaks due to the HOMO and LUMO of isolated molecules. Hence, the DOS on the molecules around the Fermi energies are very low.

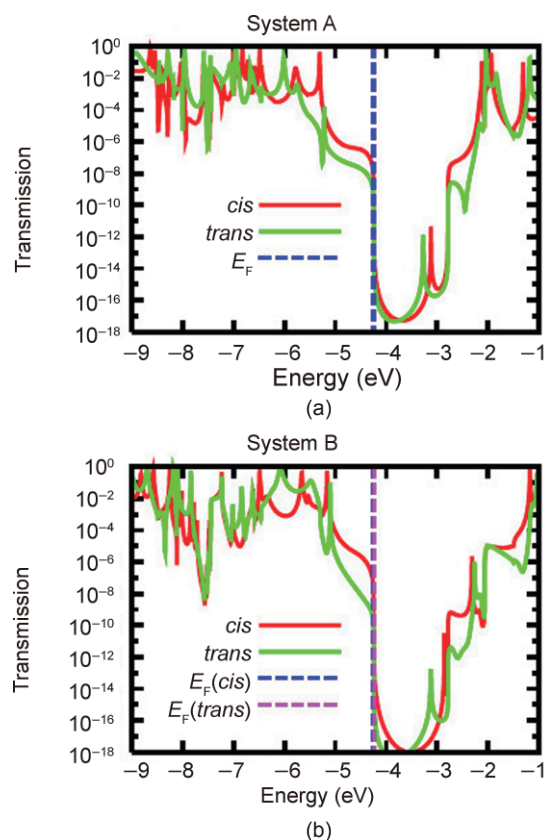


Figure 5 Electronic transmission probabilities for systems A and B at equilibrium. Transmission spectra for *cis* conformations are shown in red and for *trans* conformations in green. Broken lines show the Fermi energies (E_F). In (a), the Fermi energies for the two isomers are same. Thus, only one broken line is shown. For details of the Fermi energies, see Table 2

Table 2. Fermi energies of two molecular switching junctions.

	System A	System B
<i>cis</i>	-4.259 eV	-4.255 eV
<i>trans</i>	-4.259 eV	-4.253 eV

Previous theoretical studies of azobenzene derivatives connected between Au [9, 10] or CNT electrodes [11] have reported that *cis* isomers are less conductive than their *trans* counterparts, since the π -conjugation in the *trans* isomers along the molecular framework is broken by the torsion involved with *cis*–*trans* isomerizations. However, the same discussion does not hold true in our case since the HOMO of the *trans* isomers does not consist of p_z orbitals perpendicular to the molecular framework, but non-binding orbitals mainly localized around nitrogen atoms [35, 36], as shown in Figs. 2(a) and 2(c). The eigenstates consisting of p_z orbitals of the



trans isomers lie in the HOMO-3 levels. Thus, these states do not contribute to the transport at low bias. This counter-intuitive result ($I(\text{cis}) > I(\text{trans})$) can also be understood by analysis of the resulting mixture states of the extended molecular systems as follows.

Figure 5 shows that the transmission curves for *cis* isomers near the Fermi energies are higher than those of *trans* counterparts, implying there will be a higher current through the *cis* isomers than the *trans* counterparts at low bias. The difference between the transmission curves of the *cis* and *trans* isomers near the Fermi energies comes from the difference between the height of the resonant peaks of the *cis* and *trans* isomers around -5.0 eV. The height of the transmission resonant peaks in the two different isomer conformations can be interpreted from the molecular orbitals of the extended molecules. The HOMOs of the *cis* isomers interact with surface states of the Si contacts and their orbitals hybridize. The resulting mixed states, corresponding to the HOMO-8 of system A in Fig. 2(b) and of system B in Fig. 2(d), have a non-negligible orbital distribution bridging across the Si states. These states can contribute to transport as shown in the resonance peaks (in red) of system A at -5.31 eV in Fig. 5(a), and of system B at -5.17 eV in Fig. 5(b).

On the other hand, the HOMOs of the *trans* isomers have neither orbital amplitude around the linkers nor interact strongly with the surface states of the Si contacts. This leads to less orbital hybridization as can be seen for HOMO-7 of Figs. 2(b) and 2(d). These states do not contribute to transport as shown by the sharper and smaller resonant peaks (in green) for system A at -5.22 eV in Fig. 5(a) and for system B at -5.10 eV in Fig. 5(b). Despite the delocalized molecular orbitals along the molecular framework, the LUMOs of both the *trans* and *cis* isomers in both model systems do not interact with the surface state of the Si surface. This is because the energetic difference between the LUMOs of both *cis* and *trans* isomers and the conduction state of the Si surface (the conduction band is above -2.70 eV and the valence band is below -4.51 eV) is large. Additionally, in both model systems, there are no state available in the silicon contacts at the energy around the LUMOs of either the *trans* and *cis* isomers. Thus, these

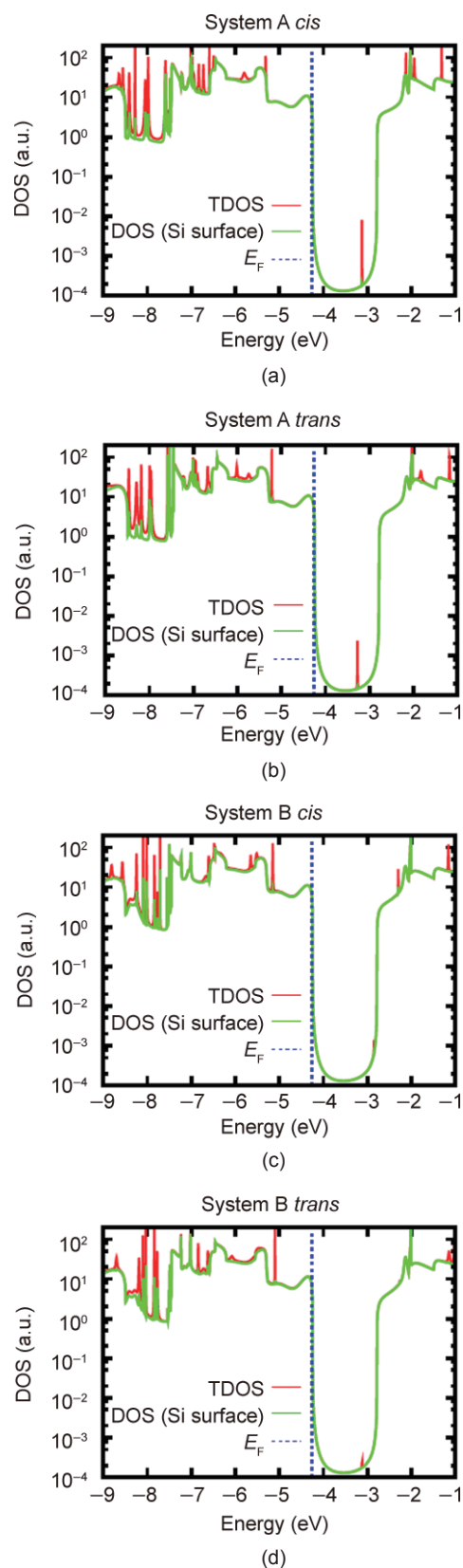


Figure 6 Total densities of states (TDOS) for systems A and B at equilibrium. TDOS are shown in red curves. The contributions from Si surfaces to TDOS are shown in green curves. Broken lines show the Fermi energies (E_F)

states lead to much lower peaks around -3.0 eV in transmission spectra as shown in Fig. 5.

We have to pay attention not only to the π -conjugation but also to the interaction between the molecules and the contacts. These results support the conclusion that the electronic structure of both molecule and contact are crucial in determining the transport properties at low bias.

In order to confirm the possible switching behavior already suggested by the analysis of transmission probabilities at equilibrium, we have calculated the current–voltage (I – V) characteristics for the two different isomer conformations of the two model systems. The I – V characteristics for *cis* and *trans* conformations for the two model systems are shown in Fig. 7. The charge transport in *trans* isomers is strongly suppressed compared to that in the *cis* counterparts for both of the two model systems. If the bias voltage is less than 1.0 V, the resonant peaks do not enter the Fermi bias window so that the I – V curves are smooth within the bias window. Therefore, we can expect constant on–off ratios as a response to the change or fluctuation of bias voltage, which are preferable for the robust control of the molecular switching. The bottom panels in Figs. 7(a) and 7(b) show the on–off ratios (the ratio of current in the *cis* isomer to that in the *trans* counterpart). The on–off ratio reaches as high as 647 for system B at 0.2 V

(bottom panel of Fig. 7(b)). Although the on–off ratios decrease with higher bias voltage, all the molecular switches recorded on–off ratios of over 10 within 1.0 V. Higher on–off ratios for lower bias voltage are advantageous both for power saving and for the suppression of unexpected breakdown triggered at higher bias voltage.

2.4 Time dependent fluctuation of conductance

For the reliable operation of molecular switching one needs to investigate the stability of the model systems between Si contacts and the fluctuation of the current through them under ambient conditions. For this purpose, we have performed MD simulations of the model systems between Si contacts and calculated the conductance of the two different isomer conformations along the MD pathways. The calculation setting was as follows. We used the Verlet algorithm for the MD simulation in DFTB+ with periodic boundary conditions. We used the same SK parameters used for the structure optimization. The MD duration is set for 10 ps, i.e., 10,000 steps at intervals of 1.0 fs. The nuclear and electronic temperatures are set as 300 K. After performing MD simulations (animations of the results of the MD simulations are shown in the ESM), the coordinates in every 100 steps are extracted from MD trajectories. Finally, we repeated conductance

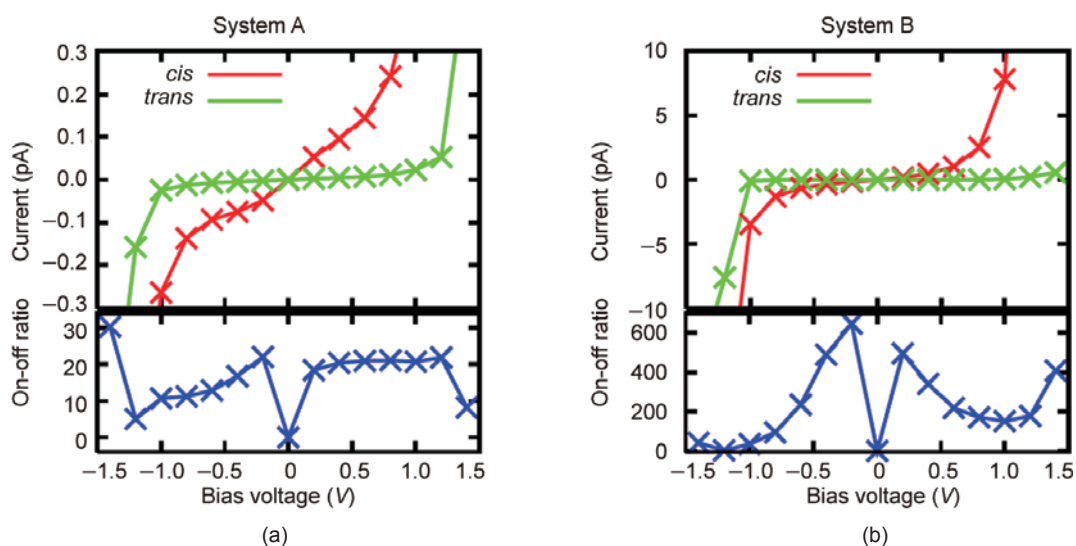


Figure 7 I – V characteristics of molecular switch junctions in the two different conformations; the bottom panels show the on–off ratios (the ratio of current between *cis* and *trans* isomers). In all cases, the *cis* conformations (red) are more conductive than their *trans* counterparts (green)



calculations for the collected 100 coordinates using the *gDFTB* code, both with and without applied bias voltage. We then analyzed the fluctuation of the transmission probabilities and *I-V* curves along the MD pathways.

The transmission probabilities along the MD pathways are plotted in Fig. S-2 (in the ESM). For each isomer, 100 transmission spectra are plotted. We can see that the main features of the transmission spectra in Fig. 5 remain unchanged. Hence, we can expect that the fluctuation of current flow under ambient conditions is small. The dispersion of the current along the MD pathways as a function of bias voltage is plotted in Fig. 8. The dispersion of the current in the two isomers is distinguishable, with the *cis* isomers always being more conductive than the *trans* counterparts. The currents in the two isomers as a function of time under bias (0.2 V)

shown in Fig. 9 also support this conclusion. The on-off ratio is over 15 (the minimum value is 15.5 at 4.7 ps in Fig. 9(b)) for all model systems. Therefore, the molecular switch junctions modeled on this study may be expected to work as switching components under ambient conditions.

Finally, we would like to comment on the possible experimental realization of molecular conformational switching devices consisting of azobenzene derivatives and Si contacts. The attachment of aromatic compounds to the Si surface should be possible using the methodology developed by Murata et al. [37–39]. The realization of two terminal Si/molecule/Si devices by covering the functionalized Si surfaces from the top with a Si contact might present a formidable challenge due to the intrinsic asymmetry in molecular junctions. However, first attempts at such a difficult task have already proved

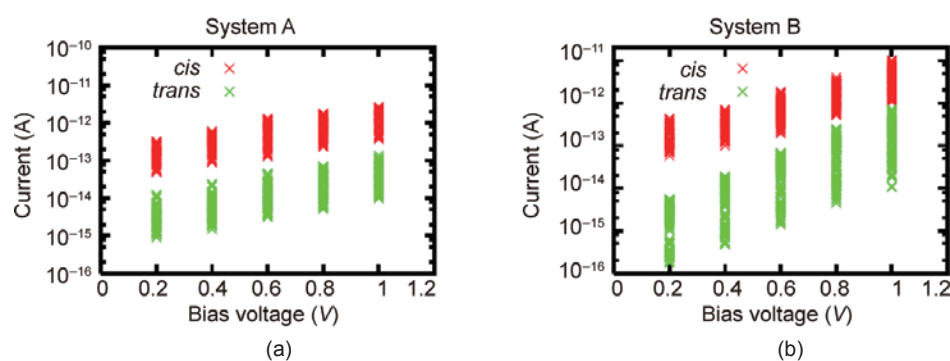


Figure 8 Dispersion of currents along MD pathways as a function of bias voltage. For each bias voltage 100 values of the current in the *cis*-isomers along the MD pathways are plotted in red crosses and those for the *trans*-isomers are plotted in green crosses

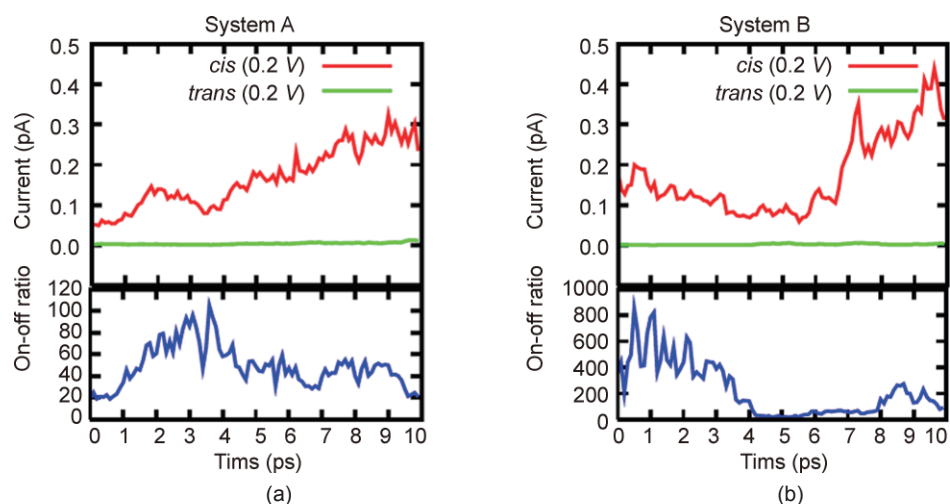


Figure 9 Time-dependent fluctuation of current (top) and on-off ratio (bottom) along MD pathways at fixed bias voltage in the molecular switch junctions

successful [40, 41]. For further understanding and control of the molecular switching, systematic studies on the effect of contacts [42, 43], internal scattering effects [44], surface roughness, and impurities [45] will be required. The spacing between the two Si surfaces will also be an important factor since it will influence the PES, molecular and electronic structure, and the resulting conductivity of the molecule embedded between Si contacts. Concerning the design of azobenzene derivatives, further investigation is required to determine which linkers and functional groups lower the PES barrier for *cis*–*trans* isomerization. One possibility to achieve this is to increase the flexibility of the molecule by using long linkers which should not, however, increase the fluctuation of current due to the flexibility of the molecules.

3. Conclusions

We have modeled two molecular switch junctions consisting of azobenzene derivatives and silicon contacts and explored their potential as molecular switches by analysis of the energetics, PES, and transport properties in the two different isomer conformations: *cis* and *trans*. Modeling of the molecular switches demonstrated that the attachment of linkers at the *meta*- and *ortho*- positions could reduce the steric repulsion between the molecules and silicon contacts during *cis*–*trans* isomerization to a value which is reasonable for the robust operation of molecular switching. Our numerical results show that *cis* isomers are more conductive than the *trans* counterparts inside the bias window for both model systems. *I*–*V* characteristics also show the same trend. Although the π -conjugation through the molecular framework is reduced due to the torsion of the two phenyl rings, the *cis* isomers interact much more strongly with the surface states of the silicon contacts than their *trans* counterparts, resulting in larger conductivities. Additionally, the calculation of currents through the two model systems along the MD pathways under ambient conditions revealed that the currents in the *cis* isomers are always significantly larger than in the *trans* counterparts leading to high on–off ratios. Importantly, the

dispersion of the current as a function of time is small enough to distinguish the two different isomer conformations. Thus, azobenzene derivatives between Si contacts are promising for applications as robust molecular switching components or as single molecular data storage devices.

Acknowledgements

This work has been partially funded by the Volkswagen Foundation by the Deutsche Forschungsgemeinschaft (DFG) under Contracts No. CU 44/5-2, CU 44/8-1, and CU 44/3-3, and by the WCU (World Class University) program through the Korea Science and Engineering Foundation funded by the Ministry of Education, Science and Technology (Project No. R31-2008-000-10100-0). We acknowledge the Center for Information Services and High Performance Computing (ZIH) at the Dresden University of Technology for computational resources. We thank Rafael Gutiérrez, Haldun Sevinçli, Florian Pump and Cormac Toher for fruitful discussions.

Electronic Supplementary Material: Supplementary material is available in the online version of this article at <http://dx.doi.org/10.1007/s12274-009-9067-y> and is accessible free of charge.

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