

Excimontec v1.0 User Manual

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I. INTRODUCTION

Kinetic Monte Carlo simulations are a powerful computational tool that have been used in concert with experiments and more detailed theoretical methods to understand and optimize organic semiconductor materials and devices.[1] However, despite over 30 years of applying KMC tools to organic semiconductors, no widespread or standardized software tools have taken hold in the community. Instead, many research groups around the world have maintained private codebases of varying complexity, efficiency, and reliability. As a result, there have been large barriers to entry for new researchers and a lot of repeated effort throughout the community that would be much better off applied to pushing the capabilities of the technique and further refining the physical models.

Excimontec represents an honest effort to bring the community together around a well-tested, optimized, reliable, and accessible open-source tool for performing KMC simulations of organic electronic devices. The current version (v1.0) implements a cubic lattice-based model for the organic semiconductor layer and is focused on simulation tests for neat single carrier diodes and binary donor-acceptor organic photovoltaic devices. Future extension of the software to simulate light emitting diodes and field-effect transistors is planned. The software is written in modern C++ and is optimized for efficient pleasingly parallel execution on high performance computing clusters using MPI. This software package uses object-oriented design and extends the KMC_Lattice framework.[2] It is also designed to work with the Ising_OPV software tool for creating and importing bulk heterojunction morphologies.[3]

II. SIMULATION TESTS AND THEIR PARAMETERS

Excimontec is designed to allow users to select and run a particular simulation test with a specific set of input parameters. Currently there are five different simulations tests available:

1. Exciton Diffusion Test
2. Time-of-Flight Charge Transport Test
3. Internal Quantum Efficiency Test
4. Dynamics Test
5. Steady State Charge Transport Test

Choosing a particular test and setting the input parameters is made within the parameter file that is then passed as an input argument to the program. An example parameter file (parameters_default.txt) is provided in the root repository directory to be used as a starting point for customization.

A single parameter file is used to run the KMC simulation test for a single set of parameters. To do parameter sweeps, one must create separate parameter files for each condition and then run separate simulations for each. While this can be tedious when manually creating and modifying the parameter files, shell scripts can be used to quickly create a series of parameter files for a parameter sweep, and simulation job arrays can be used to submit all of the simulations in a parameter sweep and group the individual simulations together.

The parameter file has a fixed format and the program parses the parameter values in a set order to define the internal variables specified in the comments that follow each value. The comments starting with `"/"` that trail each parameter definition are ignored by the parser and are simply provided to inform the user of what each line represents and give information about the expected data type or other restrictions. Moving variable definitions to different positions in the file will cause an error or incorrect assignment of parameter values. Parameter sections headings that begin with `"##"` and hyphen line breaks are ignored by the parser. The program should generate an error message if you accidentally mess up the formatting or enter an invalid value for any of the parameters.

The Test Parameters section of the parameters file is used to choose a particular simulation test and set any options specific for each test. In the subsections below, the basics of each simulation test are explained and several examples of the types of physical phenomena that can be simulated are provided. If you have further questions about the capabilities and options for any of the simulation tests, please post a question in Issues section on the GitHub website.

Test Parameters
N_tests
Enable_exciton_diffusion_test
Enable_ToF_test
ToF_polaron_type
ToF_initial_polarons
Enable_ToF_random_placement
Enable_ToF_energy_placement
ToF_placement_energy
ToF_transient_start (s)
ToF_transient_end (s)
ToF_pnts_per_decade
Enable_IQE_test
IQE_time_cutoff (s)
Enable_extraction_map_output
Enable_dynamics_test
Enable_dynamics_extraction
Dynamics_initial_exciton_conc (cm^{-3})
Dynamics_transient_start (s)
Dynamics_transient_end (s)
Dynamics_pnts_per_decade
Enable_steady_transport_test
Steady_carrier_density (cm^{-3})
N_equilibration_events

A. Exciton Diffusion Test

This simulation mode is selected by turning on the `Enable_exciton_diffusion_test` option and is designed to allow users to test exciton diffusion behavior under a variety of different conditions. The simulation is performed by randomly placing excitons one at a time on the lattice, letting them diffuse via their normal hopping mechanism, and finally recording the displacement distance upon exciton recombination back to the ground state. The simulation is performed independently on each processor core until the number of tested excitons reaches the specified `N_tests` value from the parameter file. The independent results from each processor are then aggregated and the mean and standard deviation of the exciton diffusion length are calculated and output to the `analysis_summary` file.

When performed on a neat material, one can determine how properties like energetic disorder, temperature, exciton hopping rate, and exciton lifetime affect the diffusion length. For example, Fig. 1(a) shows how the exciton diffusion length increases as the singlet exciton hopping rate prefactor increases in a neat material with an uncorrelated Gaussian density of states. Then, Fig. 1(b) shows how the exciton diffusion length decreases as the energetic disorder increases and slows down the diffusion process. These types of simulations are a good way to determine the materials parameters required to reach a desired exciton diffusion length or to probe more fundamental diffusion phenomena.

B. Time-of-Flight Charge Transport Test

This simulation mode is designed to produce an ideal time-of-flight (ToF) charge transport experiment. In the ToF experiment, a very thin sheet of charge carriers is generated at one surface of film by a laser pulse, and then the transient current response is measured as the charge carriers move through the film and are eventually extracted at the opposite electrode. ToF experiments have been widely used to probe the

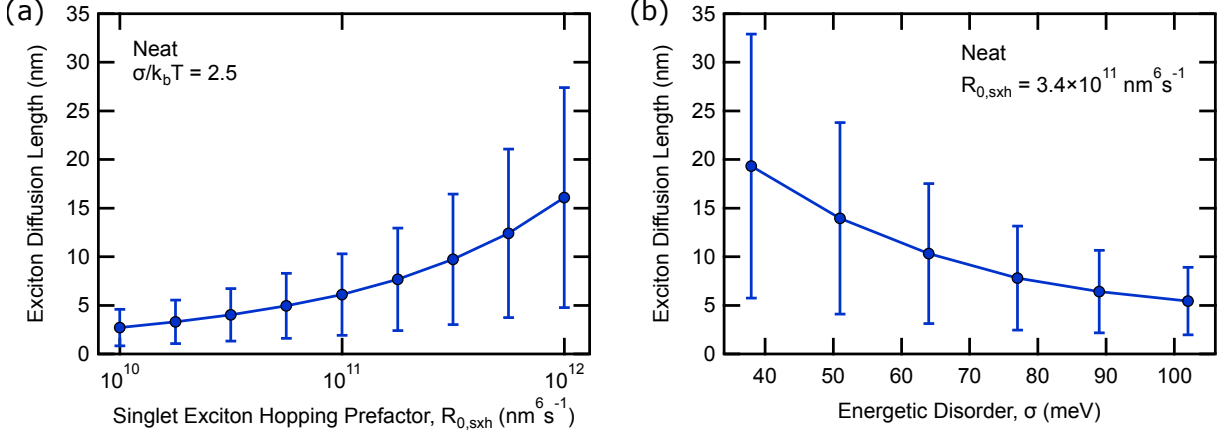


FIG. 1 Exciton diffusion test data showing how the exciton diffusion length depends on (a) the exciton hopping rate prefactor and (b) the energetic disorder of the material. Error bars show the first standard deviation of the diffusion length data.

fundamental charge transport behavior of organic semiconductors as a function of electric field, temperature, charge carrier density, etc.[4] Whereas the experiment is limited in that information can only be extracted from analysis of the current transient, KMC simulations of the same test can provide much more detailed information.

To perform this test, one must turn on the `Enable_ToF_test` option. The ToF test simulates the transport of an ultrathin sheet of charge carriers (electrons or holes) through the lattice, and the carrier type is set using the `ToF_polaron_type` parameter. The number of charge carriers in the sheet is set by the `ToF_initial_polarons` parameter, and they hop through the lattice under the influence of the defined `Internal_potential` until they are extracted at the opposite side. Transient data is recorded starting at the time defined by `ToF_transient_start` until `ToF_transient_end` at an interval specific by `ToF_pnts_per_decade`. The transit time of each extracted charge carrier is recorded, and any charge carriers stuck in the lattice longer than the `ToF_transient_end` cutoff time are removed from lattice and not included in the collection of transit times. The process is repeated until the total number of carriers simulated equals the `N_tests` parameter. The simulation is performed independently on each processor, and the results are aggregated and averaged at the end. In addition to an average transit time and mobility, the simulation provides the current density transient, charge carrier density transient, average occupation energy transient, average mobility transient, and transit time probability histogram. Examples of these results are provided in Fig. 2.

The ToF charge transport test provides users with a detailed tool to probe the transient mobility relaxation phenomena and how it is affected by the density of states, temperature, electric field, charge carrier density, material morphology, etc. For a neat material, Fig. 2(a) shows relaxation of the current density and average mobility due to charge carrier thermalization into the tail of the DOS until reaching nondispersive, steady state transport after approximately 10 ns. Figure 2(b) shows how the transition from dispersive to nondispersive transport shifts with electric field in a neat material and how the steady state mobility increases as the electric field increases. Figure 2(c) demonstrates how the electric-field dependence of mobility can also be quantified in bulk heterojunction blends to determine the complex interplay between energetic disorder (σ) and domain network tortuosity (τ) as shown previously.[5]

As more advanced features, users can also create the initial charge carriers at a specific energy position within the density of states instead of randomly populating it to probe how the starting energy of the charge carriers affects the transient behavior. To do this, one must turn off the `Enable_ToF_random_placement` option, turn on the `Enable_ToF_energy_placement` option, and set the `ToF_placement_energy`. In addition, one can produce charge extraction maps to probe for fluctuations in current density across the area of the film that may arise due to energetic fluctuations from the correlated disorder model or morphological features in a bulk heterojunction film by turning on the `Enable_extraction_map_output` option.

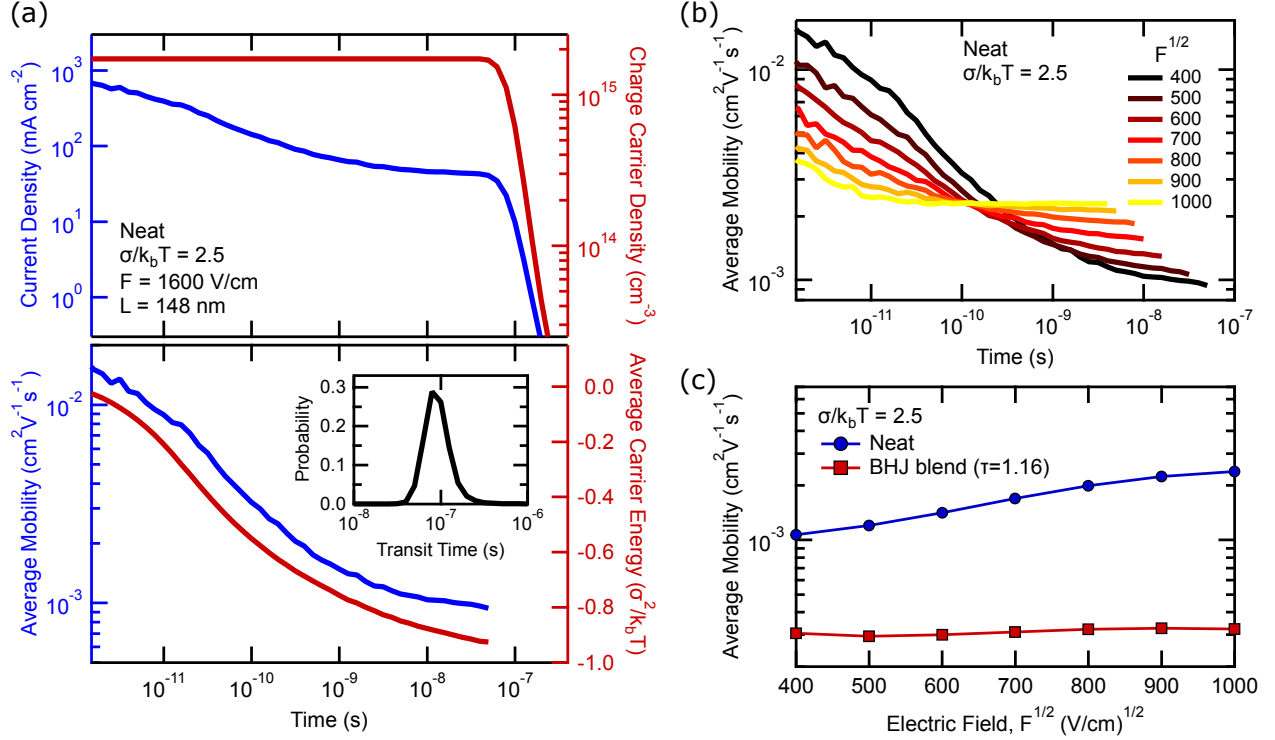


FIG. 2 Time-of-flight charge transport test results showing (a) neat material transients with transit time distribution inset, (b) neat material electric-field-dependent mobility relaxation, and (c) electric-field-dependent mobility in a neat material and a bulk heterojunction blend.

C. Internal Quantum Efficiency Test

The internal quantum efficiency (IQE) test allows users to simulate a common photovoltaic device characterization technique. It is a steady state measurement that quantifies the fraction of absorbed photons that are successfully converted into extracted charge carriers. The measurement is commonly done under short-circuit conditions, but one can also apply a bias. The Excimontec IQE test additionally provides a detailed breakdown of the various loss mechanisms that produce the final IQE value, including the exciton dissociation yield, the charge separation yield, and the bimolecular recombination loss fraction.

The IQE test is selected by turning on the `Enable_IQE_test` option and proceeds by randomly creating excitons on donor or acceptor sites at a rate specified by the `Exciton_generation_rate` parameter of each material. Exciton generation stops once the number of excitons created equals the `N_tests` parameter. The simulation finishes once all charge carriers are extracted from the lattice or once the `IQE_time_cutoff` is reached. The simulation is performed independently on each processor, and the results are aggregated and averaged at the end.

Figure 3 provides some examples for how the IQE test can be used. Figure 3(a) shows the exciton dissociation yield in a donor-acceptor bulk heterojunction blend with a domain size of 16 nm and demonstrates how the yield increases as the exciton hopping rate increases. Since these parameters were also tested using the exciton diffusion test, the increase in dissociation yield can also be related to the corresponding increase in the exciton diffusion length. The IQE test can also be used to probe charge separation phenomena and reproduce behavior that has been widely studied in the literature.[1] For example, Figure 3(b) shows the electric-field dependence of the charge separation yield for a donor-acceptor bilayer architecture with different degrees of polaron delocalization, following previously observed trends.[6] One can also explore many other dependencies such as temperature, energetic disorder, charge hopping rates, or various morphological features using a bulk heterojunction architecture.

For more detailed analysis, one can produce charge extraction maps to probe for fluctuations charge extraction across the area of the film that may arise due to energetic fluctuations from the correlated disorder model

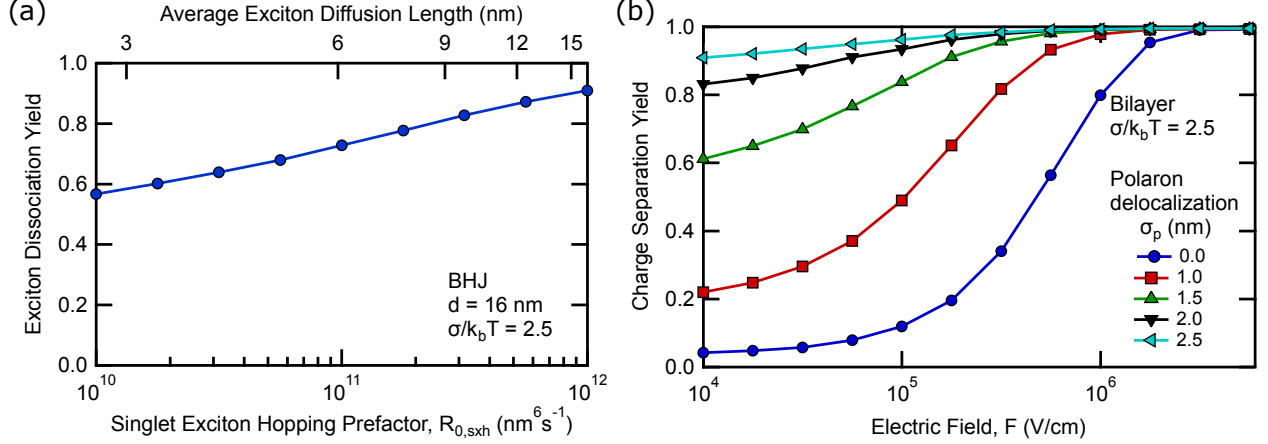


FIG. 3 Internal quantum efficiency test results showing (a) exciton dissociation yield in a donor-acceptor bulk heterojunction morphology and (b) charge separation yield in a donor-acceptor bilayer device architecture.

or morphological features in a bulk heterojunction film by turning on the `Enable_extraction_map_output` option.

D. Dynamics Test

Another popular type of experiment is a pump-probe experiment, where a laser pulse (pump) creates an initial population of excitons in the film that then evolves over time. Experimentally, the probe can be electrical or optical, but ideally one would like to measure how the populations of the different excited states evolve over time to determine the kinetics of the various mechanisms. The Excimontec dynamics test provides a wide variety of information about how the singlet exciton, triplet exciton, electron, and hole populations change over time.

The dynamics test is selected by turning on the `Enable_dynamics_test` option and begins by creating the initial excitons randomly throughout the lattice until reaching the concentration specified by the `Dynamics_initial_exciton_conc` parameter. The excitons are then allowed to diffuse, dissociate, or recombine as they will, and transient data is recorded starting at the time defined by `Dynamics_transient_start` until `Dynamics_transient_end` at an interval specific by `Dynamics_pnts_per_decade`. Polaron extraction at the top and bottom electrodes can be enabled using the `Enable_dynamics_extraction` option. The simulation is terminated once all of the excitons and charge carriers have disappeared from the lattice or once the `Dynamics_transient_end` cutoff time has been reached. The cycle is then repeated until the total number of excitons tested equals the `N_tests` parameter. The simulation is performed independently on each processor, and the results are aggregated and averaged at the end.

Figure 4 shows several examples of results from dynamics tests performed on a donor-acceptor bulk heterojunction blend. Figure 4(a) shows how an initial population of singlet excitons diffuses and relaxes into the tail of the DOS and mostly dissociates to form polarons (electrons and holes). This example also shows a small population of triplet excitons forming due to intersystem crossing from the singlet to triplet excitonic states. The polaron population then also relaxes into the tail of the DOS and decays at several timescales due to competing charge recombination mechanisms, with geminate recombination starting at the 1 ns timescale and bimolecular recombination starting at the 1 μ s timescale. Figure 4(b) focuses on singlet exciton dissociation dynamics, showing how exciton dissociation occurs faster when the exciton hop rate increases. Figure 4(c) focuses on charge carrier recombination dynamics showing how increasing polaron delocalization reduces the amount of geminate recombination and slows down the bimolecular recombination process. The dynamics test is particularly powerful for investigating how domain size and mobility together cause deviation from the Langevin model for encounter-limited bimolecular recombination.[7; 8] It can also be used to probe the impact of energetic disorder, density of states shape, and the charge-transfer state lifetime on the bimolecular recombination rate and recombination order.[9]

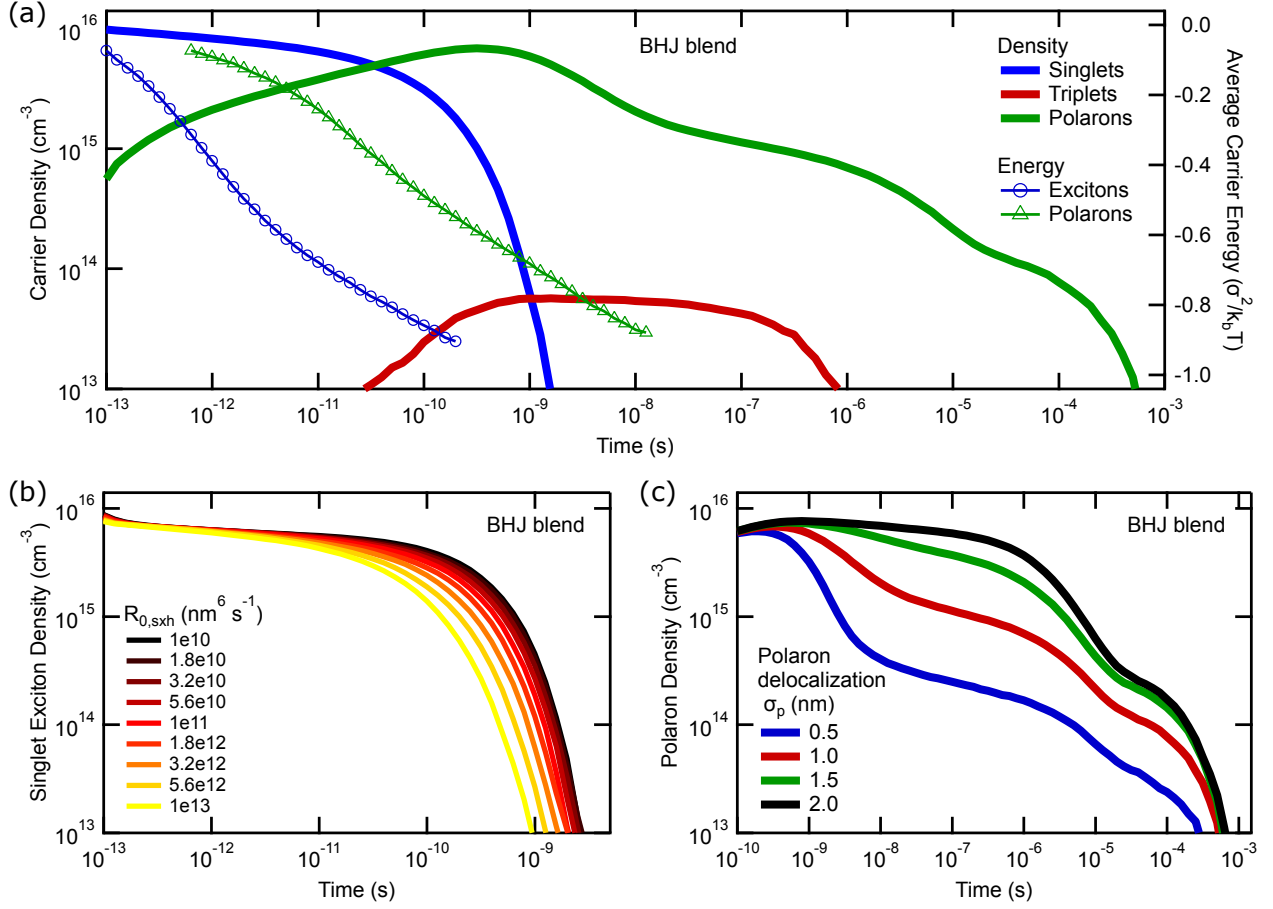


FIG. 4 Dynamics test results showing (a) full carrier density and energy transients, (b) exciton dissociation dynamics, and (c) charge carrier recombination dynamics in a BHJ blend.

E. Steady State Charge Transport Test

While the time of flight charge transport test is a good way to study transient effects such as mobility relaxation, one may also want to focus on steady state quasi-equilibrium transport behavior that can be relevant for different applications. The steady state transport test is designed to probe hole transport in a neat donor material or in donor-acceptor blends. Transport in doped semiconductors can also be simulated using the custom density of states import feature.

This test chosen using the `Enable_steady_transport_test` parameter and begins by creating holes randomly on donor sites until reaching the `Steady_carrier_density` value defined in the parameter file. The simulation then enters the equilibration phase where holes are allowed to undergo hopping events under the applied bias specified by the `Internal_potential` parameter until the number of executed events equals the `N_equilibration_events` parameter value. The simulation then enters the measurement phase and holes continue hopping until the number of events executed in the measurement phase equals the `N_tests` parameter. Periodic z-direction boundaries must be used and allow charge carriers to continue hopping indefinitely under the applied bias.

In addition to the state state mobility, this test calculates the transport energy and equilibration energy. It also provides detailed information about the density of states (DOS) and the density of occupied states (DOOS) under the given simulation conditions. The DOS and DOOS are both calculated with and without including electrostatic interactions to show how the state energies are modified by the presence of the charge carriers. The ratio of the DOOS to the DOS can then be fit to the Fermi-Dirac function to determine the Fermi energy and the effective temperature, and the Fermi energy can be combined with the transport energy

to determine the Seebeck coefficient.[10]

Fig. 5 shows example data for steady state charge transport of holes traveling through a neat material with a Gaussian density of states ($\sigma/k_bT = 2.5$) and $p = 1 \times 10^{16} \text{ cm}^{-3}$. As expected, this example shows electric-field activated mobility that roughly follows the Poole-Frenkel model ($\ln \mu \propto F^{1/2}$). In addition, at low electric-field, the calculated equilibration energy is very close to the true equilibration energy that resides at $-\sigma^2/k_bT$ relative to the average site energy. The electric field perturbs the carriers so that they sit at higher energies than they would under true zero-field equilibrium conditions. At higher electric field, the DOOS shifts to higher energy and the equilibration energy also increases in an analogous fashion to the effect of an increase in temperature, as expected from the effective temperature model.[11] Another interesting feature captured here is the appearance of a high energy shoulder in DOS due to the electrostatic interactions between charge carriers. When the system is occupied with carriers at $p = 1 \times 10^{16} \text{ cm}^{-3}$, the DOS does not have simple Gaussian shape. Sites near existing charge carriers have energies that are significantly shifted upward due to electrostatic repulsion. The steady state charge transport test allows users to probe the complex impacts of electric field, carrier density, temperature, density of states, and even morphology on the steady state bulk charge carrier mobility that can then be compared with experimental results.

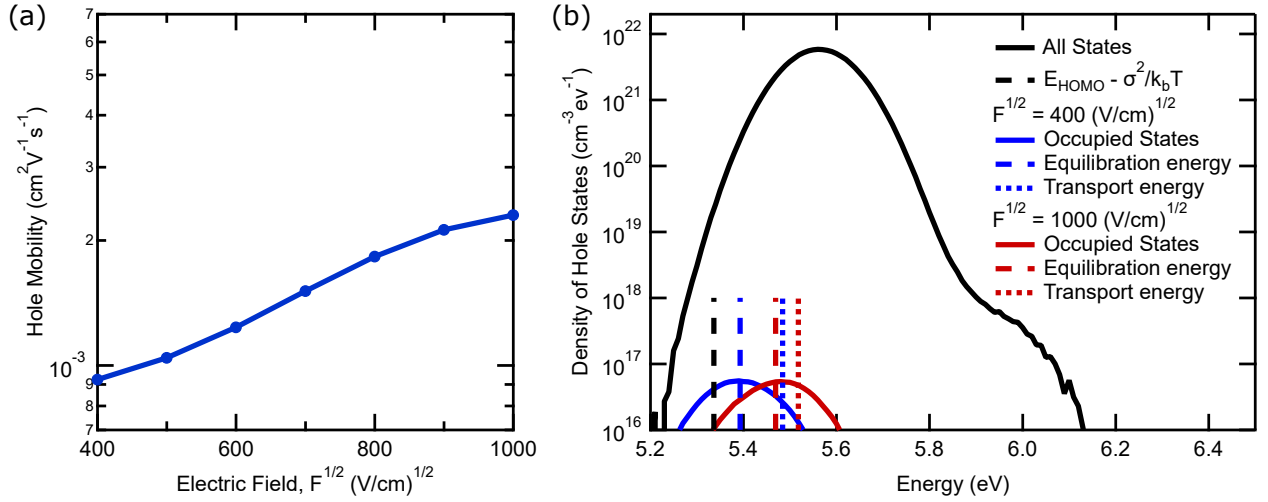


FIG. 5 Steady state charge transport test results showing (a) electric-field dependent mobility and (b) density of states data for hole transport using a Gaussian density of states ($\sigma/k_bT = 2.5$) and $p = 1 \times 10^{16} \text{ cm}^{-3}$.

III. OTHER SIMULATION PARAMETERS AND MODEL DETAILS

Specifying which device architecture to simulate on, which mechanism models to use, and all of the electronic materials properties and parameters is done using other sections of the parameter file.

This part of the user manual will go through each section of the parameter file and explain what each input parameter is used for.

A. KMC Algorithm Parameters

At any given time, there are a number of different mechanisms that can occur in an organic semiconductor device under operational conditions. Excitons and polarons can move, dissociate, recombine, etc. In a KMC simulation, each time the state of the system changes in any way, we say that an event has occurred. The goal of a KMC simulation is to simulate the evolution of the system over time, one event at a time, to determine the transient behavior or to gather statistics about the steady state behavior.

1. First-reaction Method:

Here, the formalism of the Gillespie first reaction method (FRM) is implemented to determine which event will occur next.[12] In a system containing numerous excitons and polarons, there are a number of different possible events that can occur at any given time. To determine when each will occur, the wait time for each possible event is calculated,

$$t = \frac{-\ln X}{R},$$

where X is a random number between 0 and 1 chosen from a uniform distribution and R is the rate of the particular event. Excimontec uses the Mersenne twister pseudorandom number generator implemented in the C++ 11 standard library for high quality random number generation. With the wait time calculated for all possible events, the event with the shortest wait time is executed. An extension of this method is the assumption that the majority of excitons and polarons are independent and that the calculated events can be put into a sorted schedule queue to be executed one by one as their scheduled time comes. After each iteration, only a small number of newly disabled events must be removed from the queue and only a small number of newly enabled events must be calculated and added to the queue. As a result, all possible events do not need to be recalculated after each iteration.

2. BKL Algorithm:

One also does not need to store all possible events for each exciton or polaron since only one of them will be executed before calculating the next event for that exciton or polaron. To efficiently determine the chosen event among many possibilities for a single exciton or polaron, the Bortz-Kalos-Lebowitz algorithm is used.[13] Instead of drawing a random number to calculate the wait time for each possible mechanism, the event can be chosen with only a single random number. One first calculates the rates for all possible (N) events, places them in an array, and adds them up,

$$R_{\text{sum}} = \sum_{i=0}^N R_i$$

One then generates a single random number and then finds the smallest event index ($i = n$) such that,

$$\sum_{i=0}^n R_i > R_{\text{sum}}X.$$

The n^{th} event becomes the selected event, and the wait time for the event is calculated using R_{sum} . With the BKL algorithm, the event queue remains relatively small since only one event is queued per exciton or polaron in the simulation.

3. Event Recalculation

The KMC algorithm section of the parameter file allows the user to choose from several different algorithms for event recalculation.

KMC Algorithm Parameters	
Enable_FRM	
Enable_selective_recalc	
Recalc_cutoff (nm)	r_{recalc}
Enable_full_recalc	

When the standard first-reaction method is enabled, as discussed previously, only events for directly affected objects in the simulation are recalculated. However, polarons specifically have long-range electrostatic interactions with each other, and their events are not truly independent from each other. The motion of one or several polarons modifies the electrostatic potential landscape, which should have an impact on the next event of other nearby polarons. When using the first-reaction method, this effect is ignored to save computing time. If one wants to check the significance of this simplifying assumption, the full recalculation algorithm can be enabled. With the full recalculation method, all events for all excitons and polarons are recalculated after each iteration of the simulation, thereby ensuring that the next event takes into full consideration all changes to the system.

As an alternative, a selective recalculation method is also available, which allows users to selectively update the events of all objects near the previous event within a user defined recalculation cutoff radius (r_{recalc}).^[14] By choosing the optimal recalculation cutoff radius, users can ensure that there is minimal impact from the simplifying assumptions of the first-reaction method, while also saving considerable computational time relative to the full recalculation method.

B. Lattice Parameters

The Lattice section of the parameter file is where the user defines the general simulation environment.

Lattice Parameters	
Enable_periodic.x	
Enable_periodic.y	
Enable_periodic.z	
Length (a)	L
Width (a)	W
Height (a)	H
Unit_size (nm)	a
Temperature (K)	T
Internal_potential (V)	V_{int}

The simulation volume is a cuboid shape and users must identify the dimension in the x, y, and z dimensions, which are the length (L), width (W), and height (H), respectively. In addition to the lattice size, users can set boundary conditions as either hard boundaries or periodic boundaries. In all of the simulation tests, the z-direction is treated as the charge transport direction and hard z-direction boundaries can represent the electrodes or material surface/interface. Periodic boundaries will often be desired in the x- and y-directions to simulate a thin film, but users can apply hard boundaries to investigate transport confinement effects. Users must also set the lattice unit size (a), which determines the real space distance between sites on the lattice and the total density of states. Another important general property of the lattice determine here is the temperature (T). Finally, the internal potential parameter (V_{int}) determines the voltage drop across the height (z-direction) of the lattice and is used to control the electric field in the simulation.

C. Device Architecture Parameters

The Device Architecture section allows users to choose from a number of different device architectures for the simulation box.

Device Architecture Parameter
Enable_neat
Enable_bilayer
Thickness_donor (a)
Thickness_acceptor (a)
Enable_random_blend
Acceptor_conc (vol frac)
Enable_import_morphology_single
Morphology_filename
Enable_import_morphology_set
Morphology_set_format
N_test_morphologies
N_morphology_set_size

One can implement a single neat material simulation that takes on the specified donor materials properties. One can create a donor-acceptor bilayer structure and specify the thickness of each sub-layer. One can use a random donor-acceptor blend in which sites in the lattice are randomly assigned as acceptor sites until reaching the specified acceptor concentration. Instead of using one of these simple built-in device architecture models, users can also import donor-acceptor morphologies as a single morphology that will be used for all simulations or a set of morphologies that will randomly assigned to each simulation. In particular, the morphology import feature is designed to allow users to import bulk heterojunction morphologies created using the Ising-OPV simulation tool.[3] However, any morphology conforming to the Ising-OPV morphology file format can be imported. When importing a set of morphologies, one must specify how many of the morphologies to test from the set and the total number of morphologies in the set.

D. Exciton Parameters and Mechanisms

The Exciton Parameters section contains all parameters used by the exciton physics models. Below, each of the exciton mechanisms currently implemented and their associated parameters are detailed. For most parameters, the parameter file allows users to specify different values for the donor and acceptor sites even though explicitly defined separate variables for each are not shown below.

Exciton Parameters	
Exciton_generation_rate ($\text{cm}^{-3} \text{s}^{-1}$)	G
Singlet_lifetime (s)	τ_{sx}
Triplet_lifetime (s)	τ_{tx}
R_singlet_hopping ($\text{nm}^6 \text{s}^{-1}$)	$R_{0,\text{sxh}}$
Singlet_localization (nm^{-1})	γ_{s}
R_triplet_hopping (s^{-1})	$R_{0,\text{txh}}$
Triplet_localization (nm^{-1})	γ_{t}
Enable_FRET_triplet_annihilation	
R_exciton_exciton_annihilation ($\text{nm}^6 \text{s}^{-1}$ or s^{-1})	$R_{0,\text{xxa}}$
R_exciton_polaron_annihilation ($\text{nm}^6 \text{s}^{-1}$ or s^{-1})	$R_{0,\text{xpa}}$
FRET_cutoff (nm)	$r_{\text{FRET,cut}}$
E_exciton_binding (eV)	E_{B}
R_exciton_dissociation (s^{-1})	$R_{0,\text{xd}}$
Exciton_dissociation_cutoff (nm)	$r_{\text{xd,cut}}$
R_exciton_isc (s^{-1})	R_{isc}
R_exciton_risc (s^{-1})	$R_{0,\text{risc}}$
E_exciton_ST (eV)	ΔE_{ST}

1. Exciton Generation

To simulate the creation of optically generated excitons, singlet excitons are created randomly on donor or acceptor sites based on the generation rates specified in the parameter file. Currently, the exciton generation model does not include any interference effects or extinction effects that can cause generation rate gradients in real devices. As a result, generation rates are uniform in the thickness direction, which is a reasonable first-order approximation for thin films (≈ 100 nm) on a non-reflecting substrate. For calculating the singlet exciton generation rate,

$$R_{\text{sxg}} = GV.$$

where, G is the exciton generation rate of the material and V is the total volume of the material in the lattice. It is assumed that no triplet excitons are generated by illumination due to the fact that it is a spin forbidden transition,

$$R_{\text{txg}} = 0.$$

2. Exciton Hopping

Transport of excitons from site to site can occur by two different mechanisms, Förster resonance energy transfer (FRET)[15] or Dexter electron exchange.[16] Under different conditions, one of the two mechanisms usually dominates, so both mechanisms do not need to be calculated for every iteration of the simulation. Singlet excitons are generated optically and can also readily emit a photon. Due to this property, singlet excitons can move from site to site by emitting and then absorbing a virtual photon, and this longer-range process typically dominates over the Dexter electron exchange mechanism. Conversely, triplet excitons have spin forbidden optical transitions, thereby making photon emission and absorption very slow processes. As a result, the energy transfer mechanism is negligible, and the shorter-range Dexter electron exchange mechanism dominates. With Dexter electron exchange, electrons on neighboring sites are exchanged via a charge transfer mechanism. The excited electron moves to a neighboring site, and the exciton effectively moves.

Regardless of the mechanism, hopping events are calculated for all destination sites within the FRET cutoff radius, $r_{\text{FRET,cut}}$, and the driving force of the transition is based on the initial and final state energies. For singlet excitons, the state energy is defined

$$E_s = E_{\text{HOMO}} - E_{\text{LUMO}} - E_B$$

where E_{HOMO} is the highest occupied orbital energy, E_{LUMO} is the lowest unoccupied orbital energy, and E_B is the exciton binding energy. For triplet excitons,

$$E_t = E_s - \Delta E_{\text{ST}}$$

where ΔE_{ST} is the singlet-triplet splitting energy.

Förster resonance energy transfer (FRET) model:

The rate for FRET-based singlet exciton hopping is defined

$$R_{\text{sxh,FRET},ij} = R_{0,\text{sxh}} \left(\frac{1}{d_{ij}} \right)^6 f_B(\Delta E_{\text{FRET},ij}),$$

where $R_{0,\text{sxh}}$ is the singlet exciton hopping prefactor, d_{ij} is the distance between sites, and

$$\Delta E_{\text{FRET},ij} = E_{s,j} - E_{s,i}.$$

f_B is the Boltzmann function,

$$f_B(\Delta E_{ij}) = \begin{cases} \exp\left(\frac{-\Delta E_{ij}}{k_B T}\right) & \Delta E_{ij} > 0 \\ 1 & \Delta E_{ij} \leq 0 \end{cases}$$

where k_B is the Boltzmann constant and T is the temperature.

Dexter electron exchange model:

The rate for Dexter-based exciton hopping is defined

$$R_{\text{txh,Dex},ij} = R_{0,\text{txh}} \exp(-2\gamma_t d_{ij}) f_B(\Delta E_{\text{Dex},ij}),$$

where $R_{0,\text{txh}}$ is the triplet exciton hopping prefactor, γ_t is the triplet localization parameter, and

$$\Delta E_{\text{Dex},ij} = E_{t,j} - E_{t,i}.$$

3. Exciton Dissociation

Exciton dissociation is essentially a transition from an exciton state to a charge-transfer (CT) state that spans a donor-acceptor interface. The event proceeds by a charge transfer mechanism. For an exciton located on a donor site, an electron transfers from the donor site to an acceptor site. For an exciton located on an acceptor site, a hole transfers from the acceptor site to a donor site. As a result, the mechanism is simulated using a charge transfer model. Excimontec implements two options for charge transfer models, the Miller-Abrahams phonon-assisted tunneling model[17] and a simplified Marcus model [18] that implements a polaronic mechanism.

Regardless of the model used, dissociation events are calculated for all nearby destination sites within the exciton dissociation cutoff radius, $r_{\text{xd,cut}}$, and the driving force for this transition depends on the energies of the initial final states. For singlet excitons,

$$\Delta E_{\text{sxd},ij} = E_{\text{CT},ij} - E_{s,i},$$

and for triplet excitons,

$$\Delta E_{\text{txd},ij} = E_{\text{CT},ij} - E_{t,i},$$

where $E_{\text{CT},ij}$ is the CT state energy. For the dissociation of an exciton starting from a donor site,

$$E_{\text{CT},ij} = E_{\text{HOMO}_D,i} - E_{\text{LUMO}_A,j} + E_{\text{B,CT}}(d_{ij}) + E_C(q_i) + E_C(q_j) - e\mathbf{d}_{ij} \cdot \mathbf{F},$$

where $E_{\text{B,CT}}$ is the CT state binding energy, $E_C(q)$ is the electrostatic interaction energy term for the newly formed polarons interacting with all other polarons in the system, e is the elementary charge constant, \mathbf{d}_{ij} is the CT state dipole displacement vector, and \mathbf{F} is the internal electric field vector. An analogous expression is used for dissociation of an an acceptor exciton. It is worth noting here that the different spin states of a CT state are not modeled here, and CT states with singlet or triplet spin are treated as the same and therefore assumed to have the same energies and the same mechanism rates. Moving on, the CT state binding energy is defined

$$E_{\text{B,CT}}(d_{ij}) = \frac{-e^2}{4\pi\epsilon\epsilon_0 d_{ij}},$$

where ϵ is the dielectric constant and ϵ_0 is the vacuum permittivity constant. The electrostatic interaction energy between polarons is calculated via the summation

$$E_C(q_k) = \sum_{l,l \neq k} \frac{q_k q_l}{4\pi\epsilon\epsilon_0 d_{kl}}, \quad d_{kl} \leq r_{\text{C,cut}},$$

which adds together the interaction energy between the new polaron on site k having charge q_k with all other polarons in the lattice, where the distance between the two polarons on site k and site l (d_{kl}) is less than the Coulomb cutoff radius ($r_{C,\text{cut}}$).

However, users can also enable the Gaussian polaron delocalization model[19] to reduce the CT state binding energy. With this model enabled,

$$E_{B,CT,d}(d_{ij}) = \frac{-e^2}{4\pi\epsilon\epsilon_0 d_{ij}} \text{erf}\left(\frac{d_{ij}}{\sigma_p\sqrt{2}}\right)$$

where σ_p is the polaron delocalization length. The Gaussian polaron delocalization model also modifies the polaron electrostatic interaction energy function, so that

$$E_{C,d}(q_k) = \sum_{l,l \neq k} \frac{q_k q_l}{4\pi\epsilon\epsilon_0 d_{kl}} \text{erf}\left(\frac{d_{ij}}{\sigma_p\sqrt{2}}\right), \quad d_{kl} \leq r_{C,\text{cut}}.$$

This model effectively reduces the interaction strength of the polarons when they are close to each other and converges to the standard Coulomb model for larger separation distances.

Miller-Abrahams Model:

The rate for the Miller-Abrahams dissociation model for singlet excitons is defined,

$$R_{\text{sxd},\text{MA},ij} = R_{0,\text{xd}} \exp(-2\gamma_s d_{ij}) f_B(\Delta E_{\text{sxd},ij}),$$

where $R_{0,\text{xd}}$ is the exciton dissociation prefactor, γ_s is the singlet delocalization parameter, and $\Delta E_{\text{sxd},ij}$ is the energetic driving force. For triplet excitons the rate is calculated,

$$R_{\text{txd},\text{MA},ij} = R_{0,\text{xd}} \exp(-2\gamma_t d_{ij}) f_B(\Delta E_{\text{txd},ij}).$$

Marcus Model:

The rate for the Marcus-based dissociation model of singlet excitons is defined,

$$R_{\text{sxd},\text{Marcus},ij} = R_{0,\text{xd}} \frac{1}{\sqrt{4\pi\lambda k_B T}} \exp(-2\gamma_s d_{ij}) \exp\left[-\frac{(\Delta E_{\text{sxd},ij} + \lambda)^2}{4\lambda k_B T}\right],$$

where λ is the reorganization energy. Similarly, for triplet excitons it is defined,

$$R_{\text{txd},\text{Marcus},ij} = R_{0,\text{xd}} \frac{1}{\sqrt{4\pi\lambda k_B T}} \exp(-2\gamma_t d_{ij}) \exp\left[-\frac{(\Delta E_{\text{txd},ij} + \lambda)^2}{4\lambda k_B T}\right].$$

4. Exciton Intersystem Crossing and Reverse Intersystem Crossing

Excitons can also undergo intersystem crossing events where they change between singlet and triplet spin states. Here, we assume that intersystem crossing, which is the transition from singlet to triplet, is always energetically favorable, and as a result, can simply be defined by the single parameter, R_{isc} . However, reverse intersystem crossing is typically characterized by an energy barrier. As a result, the rate is defined,

$$R_{\text{risc}} = R_{0,\text{risc}} \exp\left(\frac{-\Delta E_{\text{ST}}}{k_B T}\right),$$

where $R_{0,\text{risc}}$ is the reverse intersystem crossing prefactor and the energy barrier is given by the previously defined singlet-triplet splitting energy (ΔE_{ST}).

5. Exciton-Exciton Annihilation

Excitons can also "react" with other excitons through a mechanism called exciton-exciton annihilation. In this mechanism, one exciton transfers its excited state energy to another nearby exciton. Here, we assume this mechanism leads to the loss of the exciton initiating the event. The receiving exciton obtains additional energy, but here we assume that this energy is rapidly lost through vibrational relaxation transitions that dissipate the energy into the lattice and that this dissipation does not significantly alter the temperature of the system. We assume that these excited vibrational states relax much faster than any of the other exciton mechanisms and therefore have no impact on promoting any other events. As a result, we completely ignore these vibrational states and assume that the excess energy from the initiating exciton is immediately lost.

Similar to the exciton hopping models, we implement both a FRET-based and Dexter-based model for annihilation. If the initiating exciton has a singlet spin state, the FRET model is used, and if it has a triplet spin state, the Dexter model is used. Regardless of the model used, annihilation events are calculated for all destination sites within the FRET cutoff radius, $r_{\text{FRET,cut}}$.

FRET model:

The rate for the FRET-based model is defined,

$$R_{\text{xxa,FRET},ij} = R_{0,\text{xxa}} \left(\frac{1}{d_{ij}} \right)^6,$$

where $R_{0,\text{xxa}}$ is the exciton-exciton annihilation prefactor.

Dexter model:

The rate for the Dexter-based model is defined,

$$R_{\text{xxa,Dex},ij} = R_{0,\text{xxa}} \exp(-2\gamma_{\text{t}} d_{ij}).$$

6. Exciton-Polaron Annihilation

Excitons can also transfer their excited state energy to nearby polarons. Similar to the assumptions of the exciton-exciton annihilation model, we assume that this excess energy is rapidly dissipated into the lattice and has no impact on the receiving polaron's subsequent events. Again, if the exciton has a singlet spin state, the FRET-based model is used, and if it has a triplet spin state, the Dexter-based model is used. Regardless of the model used, annihilation events are calculated for all destination sites within the FRET cutoff radius, $r_{\text{FRET,cut}}$.

FRET model:

The rate for the FRET-based model is defined,

$$R_{\text{xpa,FRET},ij} = R_{0,\text{xpa}} \left(\frac{1}{d_{ij}} \right)^6,$$

where $R_{0,\text{xpa}}$ is the exciton-polaron annihilation prefactor.

Dexter model:

The rate for the Dexter-based model is defined,

$$R_{\text{xpa,Dex},ij} = R_{0,\text{xpa}} \exp(-2\gamma_{\text{t}} d_{ij}).$$

7. Exciton Recombination

Excitons can also lose their energy due to natural relaxation of the excited electron back to the ground state. For singlet excitons, this relaxation can often cause emission of a photon. For example, this is how organic light emitting diodes generate light. However, Excimontec does not currently account for emitted photons, and exciton recombination events simply cause the exciton to be removed from the system.

For singlet excitons the rate is defined,

$$R_{\text{sxr}} = \frac{1}{\tau_{\text{sx}}},$$

where τ_{sx} is the singlet exciton lifetime. For triplet excitons, the rate is defined,

$$R_{\text{txr}} = \frac{1}{\tau_{\text{tx}}},$$

where τ_{tx} is the triplet exciton lifetime. Since the ground state has a singlet spin state, triplet recombination to the ground state is a spin forbidden transition and will often be much slower than for singlet excitons.

E. Polaron Parameters and Mechanisms

The Polaron Parameters section contains all parameters used by the polaron physics models. Below, each of the polaron (electron or hole) mechanisms currently implemented and their associated parameters are detailed. For most parameters, the parameter file allows users to specify different values for the donor and acceptor sites even though explicitly defined separate variables for each are not shown below.

Polaron Parameters	
Enable_phase_restriction	
R_polaron_hopping (s^{-1})	$R_{0,\text{ph}}$
Polaron_localization (nm^{-1})	γ_{p}
Enable_miller_abrahams	
Enable_marcus	
Reorganization (eV)	λ
R_polaron_recombination (s^{-1})	$R_{0,\text{pr}}$
Polaron_hopping_cutoff (nm)	$r_{\text{ph,cut}}$
Enable_gaussian_polaron_delocalization Polaron_delocalization_length (nm)	σ_{p}

1. Polaron Hopping

Polarons can hop between sites on the lattice, and Excimontec implements two options for polaron hopping models, the Miller-Abrahams phonon-assisted tunneling model[17] and a simplified Marcus model [18] that implements a polaronic mechanism. In the parameter file, users must choose which model to use in their simulation. Regardless of the model used, hopping events are calculated for all available destination sites within the polaron hopping cutoff radius, $r_{\text{ph,cut}}$. Users can restrict holes to donor sites and electrons to acceptor sites using the Enable_phase_restriction parameter.

Miller-Abrahams Model:

The rate for the Miller-Abrahams model is defined,

$$R_{\text{ph,MA},ij} = R_{0,\text{ph}} \exp(-2\gamma_{\text{p}}d_{ij})f_{\text{B}}(\Delta E_{\text{ph},ij}),$$

where $R_{0,\text{ph}}$ is the polaron hopping prefactor and $\Delta E_{\text{ph},ij}$ is the change in the electrostatic potential energy for a polaron moving from site i to site j . For electrons, this is calculated,

$$\Delta E_{\text{ph},ij} = E_{\text{LUMO},j} - E_{\text{LUMO},i} + E_{\text{C}}(q_j) - E_{\text{C}}(q_i) + e\mathbf{d}_{ij} \cdot \mathbf{F},$$

where \mathbf{d}_{ij} is the polaron displacement vector.

Marcus Model:

The rate for the Marcus-based model is defined,

$$R_{\text{ph},\text{Marcus},ij} = R_{0,\text{ph}} \frac{1}{\sqrt{4\pi\lambda k_{\text{B}}T}} \exp(-2\gamma_{\text{p}}d_{ij}) \exp\left[-\frac{(\Delta E_{\text{ph},ij} + \lambda)^2}{4\lambda k_{\text{B}}T}\right].$$

2. Polaron Recombination

When an electron and a hole are near each other, a recombination event can occur whereby the two carriers "react" and both are removed from the system. For simplicity, only a Miller-Abrahams model is implemented for this mechanism. A recombination event is calculated for all electron-hole pairs that have a separation distance that is less than the polaron hopping cutoff radius, $r_{\text{ph},\text{cut}}$.

The rate for the polaron recombination event is defined,

$$R_{\text{pr},ij} = R_{0,\text{pr}} \exp(-2\gamma_{\text{p}}d_{ij}),$$

where $R_{0,\text{pr}}$ is the polaron recombination prefactor.

3. Polaron Extraction

For simulation tests where polarons are extracted at assumed electrode contacts at the top and bottom of the lattice (steady state transport test and dynamics test), when polarons arrive near the contacts, a polaron extraction event is enabled. The electrode contacts are assumed to be blocking for the opposite carrier type but allow barrier-less extraction of the proper carrier types. The cathode is assumed to be placed on the bottom interface near $z = 0$ where electrons are extracted, and the anode is at the top interface near $z = H$ where holes are extracted.

For simplicity, only a Miller-Abrahams model is implemented for this mechanism. An extraction event is calculated only when the distance between the polaron and the electrode interface is less than the polaron hopping cutoff radius, $r_{\text{ph},\text{cut}}$. The rate for the polaron extraction event is defined,

$$R_{\text{pe},i} = R_{0,\text{ph}} \exp(-2\gamma_{\text{p}}d_i),$$

where d_i is the shortest distance between the polaron and the electrode interface.

F. Site Energy Parameters

Organic semiconductors are typically energetically disordered materials in which fluctuations in the molecular environment cause the molecular electronic state energies to vary from molecule to molecule or from segment to segment in a polymer. The Site Energy Parameters section allows users to control the energetic disorder model used in the simulation. For most parameters, the parameter file allows users to specify different values for the donor and acceptor sites even though explicitly defined separate variables for each are not shown below.

Site Energy Parameters	
Homo (eV)	E_{HOMO}
Lumo (eV)	E_{LUMO}
Enable_gaussian_dos	
Energy_stdev (eV)	σ
Enable_exponential_dos	
Energy_urbach (eV)	E_{U}
Enable_correlated_disorder	
Disorder_correlation_length (nm)	ξ_{d}
Enable_gaussian_kernel	
Enable_power_kernel	
Power_kernel_exponent	p
Enable_interfacial_energy_shift	
Energy_shift (eV)	
Enable_import_energies	
Energies_import_filename	

The simplest and most common model to capture disorder is the well-known Gaussian disorder model, in which there are random fluctuations in site energies.[4] Later work demonstrated that in some systems, energetic disorder is not completely random but spatially correlated.[20; 21] In some materials, it has also been demonstrated that device behavior is better described using low energy tail states with an exponential distribution.[22; 23; 24] With Excimontec, users can choose from several different energetic disorder models described below. However, even more complex density of states distributions with or without correlations could also be possible, so users can also import custom site energies from a text file if desired. In each of these cases, the median of the density of states is given by the HOMO and LUMO energies defined for the donor and acceptor sites. And in all cases, a single energy is assigned to each site that is relative to the HOMO and LUMO. This keep memory usage low and allows larger lattice sizes to be simulated, but the simplifying assumption here is that energies for HOMO and LUMO states on the same site are correlated.

1. Uncorrelated Gaussian Disorder Model

When the the uncorrelated Gaussian disorder model is enabled, it is implemented by simply drawing each site energy from a Gaussian (normal) distribution with a mean value of 0 and a specified standard deviation (σ) also often called the energetic disorder parameter.

2. Correlated Gaussian Disorder Models

The original correlated disorder model explained the origin of the correlations to be due to interactions of the polaron with randomly oriented dipoles.[20; 21] However, there can be other causes of correlations, such as correlations in molecular spacing or orientation that impart differences in the correlation function.[25] To modify both the length scale of the correlation and the functional form of the correlation decay, a kernel algorithm is implemented, similar to work done previously by Gartstein and Conwell.[20] There are currently three kernel models to choose from, a Gaussian kernel and power kernel with an exponent of -1 or -2. The different kernels can be used to modify the functional form of how the energy correlation decays with distance. Users can inspect the final disorder correlation from the data generated in the "DOS_correlation_data" files.

3. Exponential Disorder Model

The standard exponential distribution, unlike a Gaussian, is discontinuous at zero and does not contain a high energy tail as would be expected in real materials. To create a more appropriate overall density

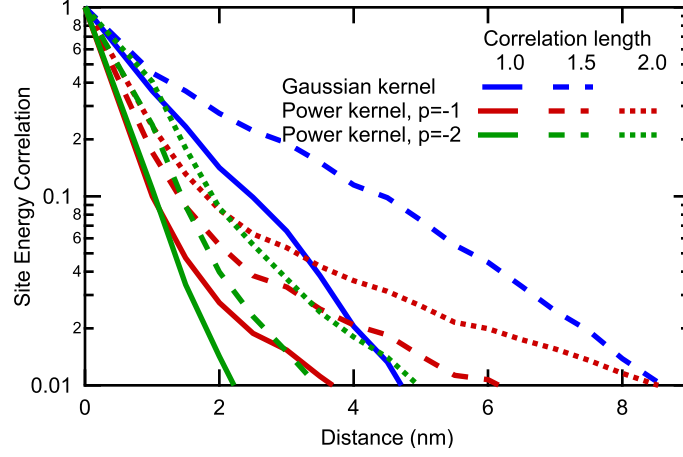


FIG. 6 Correlated Gaussian disorder model site energy correlation examples showing how the different kernel functions cause the correlation to decay as a function of distance.

of states distribution, while retaining the exponential tail states, a hybrid exponential-Gaussian density of states model is implemented, such that the median state energy remains at zero. Half of the sites above zero (high energy sites) are assigned using a Gaussian distribution, and the other half of low energy sites below zero are assigned using an exponential distribution. The Gaussian half is constrained such that its peak is continuous with the maximum of the exponential distribution. As a result, the overall distribution is defined only by the exponential distribution's Urbach energy (E_u) and the standard deviation of the Gaussian is automatically set to meet the continuity constraint. Figure 7 shows an example of the hybrid exponential-Gaussian density of states distribution compared to the standard Gaussian shape.

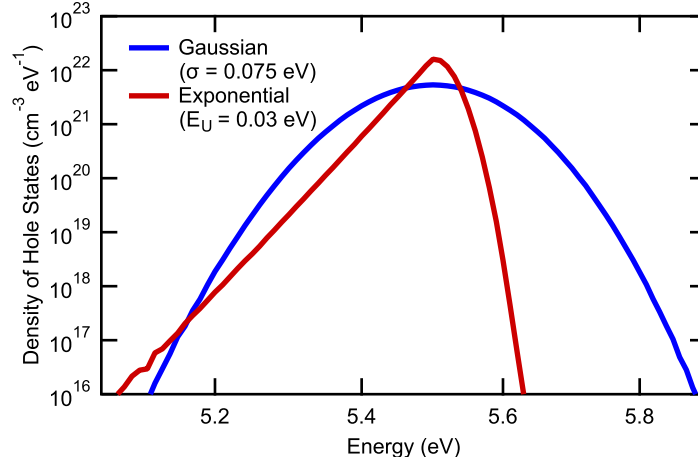


FIG. 7 Density of states distributions for the exponential disorder model compared to the Gaussian disorder model.

4. Interfacial Energy Model

In addition to the effects captured by the other models, it has also been found that sites near the donor-acceptor interface can have shifted energy levels that form an interfacial energy cascade.[26] To model this, a simple phenomenological model is implemented here where the energy shift depends on the purity of a given region of material. The assumption here is that the presence of the dissimilar material (molecules) acts as a defect that disrupts the favorable interactions between molecules in the neat material and reduces the energetic stabilization of the local electronic states. As a result, for every lattice site, a site energy is

assigned from the density of states distribution of the pure material, and this energy is then shifted by an amount ($\Delta E_{s,i}$), where

$$\Delta E_{s,i} = N_{1,i}\epsilon_s + N_{2,i}\frac{\epsilon_s}{\sqrt{2}} + N_{3,i}\frac{\epsilon_s}{\sqrt{3}}. \quad (1)$$

Here, $N_{1,i}$ is the number of dissimilar first-nearest neighbors around site i , $N_{2,i}$ is the number of dissimilar second-nearest neighbors around site i , and $N_{3,i}$ is the number of dissimilar third-nearest neighbors around site i , and ϵ_s is the energy shift parameter. As a result, the site energy is shifted depending on the number of dissimilar neighbors in a distance normalized fashion, where the dissimilar sites farther away have less of an impact.

G. Electrostatic Interaction Parameters

Electrostatic Interaction Parameters	
Dielectric_constant	ϵ
Enable_Coulomb_maximum	
Enable_Coulomb_cutoff	
Coulomb_cutoff (nm)	$r_{C,cut}$

Electrostatic interactions between polarons are an important factor in determining polaron recombination rates[8] and transport behavior[27]. Users can set the dielectric constant (ϵ), which acts to screen these interactions. Coulomb interactions are included between each polaron pair as long as their separation distance is less than the Coulomb cutoff radius ($r_{C,cut}$). The Coulomb cutoff radius can be explicitly defined in the parameter file or it can be set to dynamically scale with the size of the lattice by setting it to half the smallest dimension via turning on the Enable_Coulomb_maximum parameter. It is not recommended to set the cutoff radius to be larger than half of the smallest dimension in order to prevent double counting interactions with the same polaron across the periodic boundaries.

IV. PROGRAM INSTALLATION

To install Excimontec, you must compile the executable from the source code on your machine. Detailed instructions are provided below for Windows and Ubuntu Linux operating systems. Once compilation is finished the executable is created in the root repository directory by default and can then be copied to another simulation job directory if desired.

A. Linux (Ubuntu)

1. Update the packages list:

```
sudo apt update
```

2. Install the GCC compiler and GNU Make:

```
sudo apt-get install build-essential
```

3. Install Open MPI:

```
sudo apt-get -y install -qq openmpi-bin libopenmpi-dev
```

4. Download Excimontec source files:

To download the entire repository, use

```
git clone --recurse-submodules https://github.com/MikeHeiber/Excimontec
```

By default, this will checkout the latest version on the master branch. To then checkout a specific tagged release (like v1.0.0), use

```
git checkout tags/v1.0.0
```

To only download a specific release (like v1.0.0), use

```
git clone --recurse-submodules https://github.com/MikeHeiber/Excimontec --branch=v1.0.0
```

5. Build and Test Excimontec:

When first using Excimontec on your computer, you should build and run the test suite to make sure everything is working correctly. Set the Excimontec directory as the current working directory,

```
cd Excimontec
```

and then use GNU Make to build the test executable.

```
make test
```

Once the build is complete, run the test suite

```
./test/Excimontec_tests.exe
```

If there are no test errors, build the main executable.

```
make
```

If everything builds correctly, you can run a simple test using the default parameters.

```
mpiexec -n 4 Excimontec.exe parameters_default.txt
```

This will execute a simple exciton diffusion simulation test, and when it finishes several output files will be created in the project directory. You can quickly check the results in the analysis_summary file with the command.

```
cat analysis_summary.txt
```

The default parameters should produce an exciton diffusion length of approximately 24 nm.

B. Windows 10

1. Install Ubuntu via Windows Subsystem for Linux:

Follow the instructions from

<https://docs.microsoft.com/en-us/windows/wsl/install-win10>

2. Follow the Linux installation instructions

C. Windows (older versions)

1. Install Microsoft Visual Studio 2019 IDE:

Download Microsoft Visual Studio IDE from

<https://visualstudio.microsoft.com/>

and make sure to install the C++ development tools.

2. Install Microsoft MPI:

Download the Microsoft MPI files from

<https://docs.microsoft.com/en-us/message-passing-interface/microsoft-mpi>

and then install both MS-MPI and the MS-MPI Software Development Kit.

3. Download (Clone) Excimontec Files:

Open Microsoft Visual Studio 2019 and select "Clone or checkout code".

Enter the repository address, <https://github.com/MikeHeiber/Excimontec> and then click Clone.

Once cloning is finished, the Excimontec repository will be loaded into the Solution Explorer pane on the left.

4. Load Microsoft Visual Studio Solution:

In the Solution Explorer, open the msvc\2019\ folder and then double click the Excimontec.sln solution file. Once loaded, the Solution Explorer should show three projects: Excimontec, Excimontec_tests, gtest.

The solution file is set to build Excimontec for 64 bit systems, but this can be changed to 32 bit by changing the build platform using the drop down box in the toolbar at the top.

5. Build and Test Excimontec:

When first using Excimontec on your computer, you should build and run the test suite to make sure everything is working correctly. To do this, build the gtest project first by right-clicking gtest in the Solution Explorer and selecting Build.

Once the gtest build is complete, right click the Excimontec_test project in the Solution Explorer and select Build.

Once the Excimontec_test build is complete, make it the active project by right clicking the Excimontec_test project and selecting Set as Startup Project, and then press Cntrl+F5 to run the test suite without the debugger.

If there are no test errors, build the main executable by right-clicking on Excimontec in the Solution Explorer and selecting Build.

If it builds correctly, you can run a simple test using the default parameters. First, open the command prompt and change the working directory to the Excimontec build directory.

```
cd %HOMEDRIVE%%HOMEPATH%\Source\Repos\Excimontec\msvc\2019\x64-Release
```

Then run Excimontec with the default parameters.

```
mpiexec -n 4 Excimontec.exe ../../parameters_default.txt
```

This will execute a simple exciton diffusion simulation test, and when it finishes several output files will be created in the build directory. You can quickly check the results in the analysis_summary file with the command.

```
type analysis_summary.txt
```

The default parameters should produce an exciton diffusion length of approximately 24 nm.

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