

What Effect Does Fully Equilibrating After Fine-Graining Have on the Charge Transport Properties?

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December 14, 2017

1 Mobilities

ID	Simulation Name	Density (g cm ⁻³)	Anisotropy (Arb. U.)	Stacks (Arb. U.)	Stack Threshold (Å)	Mobility (cm ² V ⁻¹ s ⁻¹)
1	equilP3HT_1.5	1.676	0.2201	1	7.0845	1.18×10^1
2	equilP3HT_1.75	1.061	0.0105	1	4.67	8.05×10^{-1}
3	equilP3HT_2.0	0.892	0.0068	4	4.8388	5.17×10^{-1}
4	equilP3HT_2.25	0.787	0.0050	42	4.6085	3.77×10^{-1}
5	equilP3HT_2.5	0.685	0.0097	64	4.8188	2.43×10^{-1}
6	origP3HT_T1.5	1.676	0.1282	1	7.3947	1.17×10^1
7	origP3HT_1.75	1.061	0.0197	19	4.4702	3.48×10^{-1}
8	origP3HT_2.0	0.892	0.0085	1	5.0683	4.46×10^{-1}
9	origP3HT_2.25	0.787	0.0114	5	4.7747	5.52×10^{-1}
10	origP3HT_2.5	0.685	0.0188	25	4.8465	4.03×10^{-1}

Table 1: The results from MorphCT for the usual P3HT morphologies, with Voronoi neighbours. Runs **1-5** are the previous Voronoi systems (where the systems were run for 1E5 timesteps during the final fine-graining phase), whereas runs **6-10** have been equilibrated for 1E7 timesteps during the final fine-graining phase.

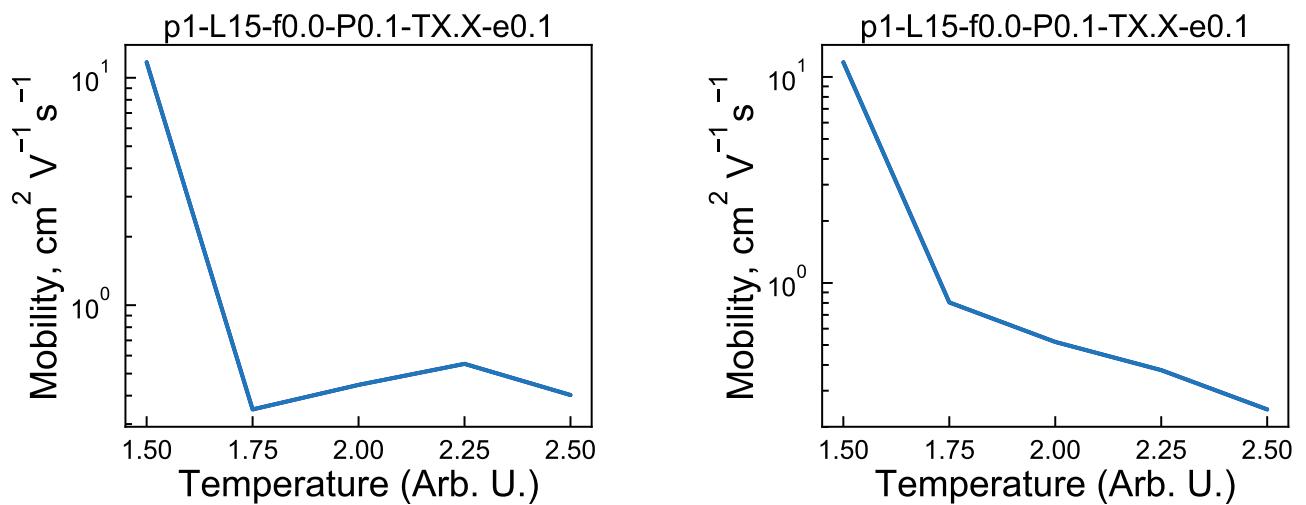


Figure 1: The evolution of the mobility of the $p1\text{-L15-f0.0-P0.1-TX.X-e0.5}$ systems.

2 Equilibration

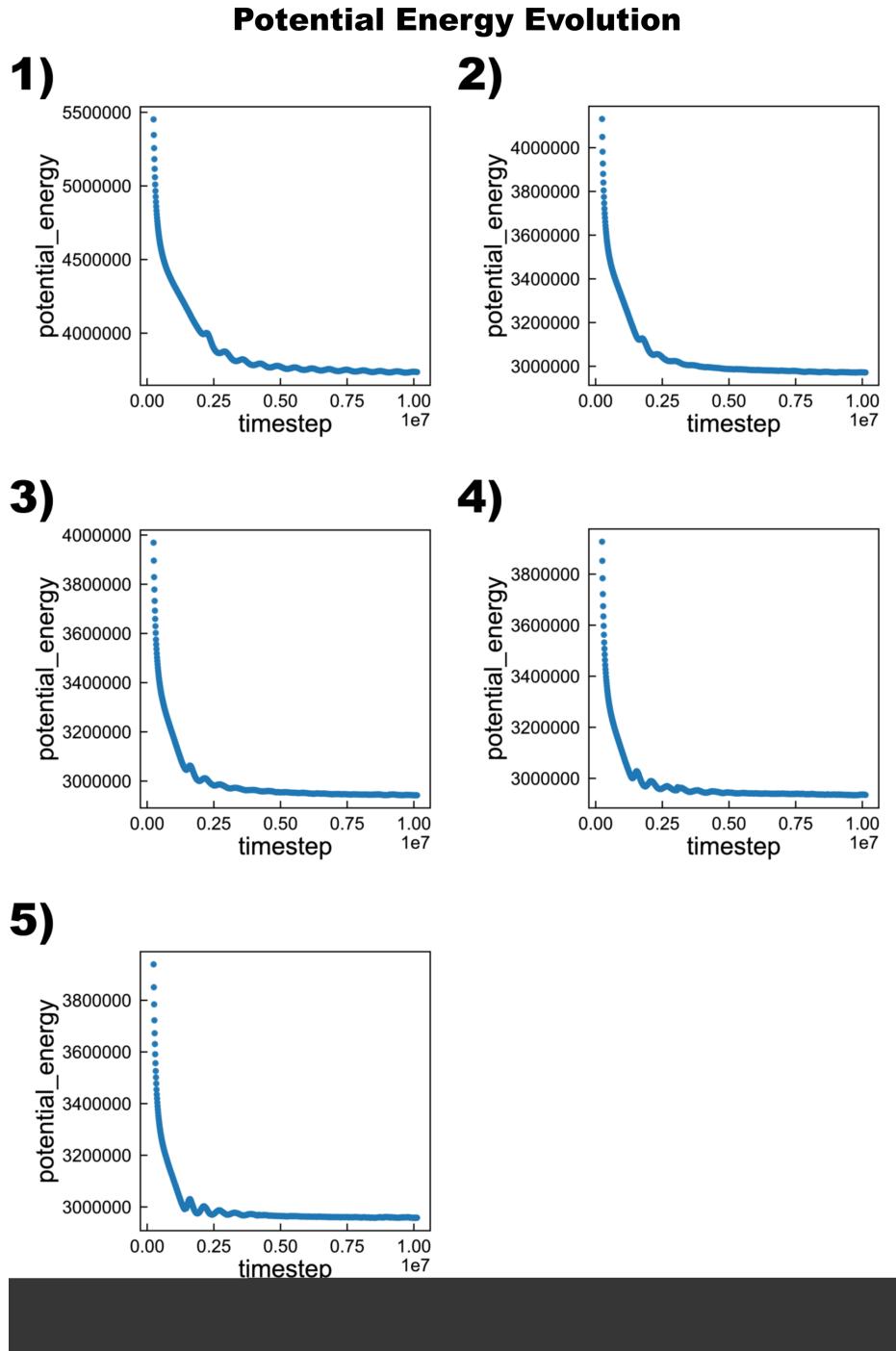


Figure 2: The evolution of the potential energy of the p1-L15-f0.0-P0.1-TX.X-e0.5 systems that have been run until equilibration. Numbers correspond to the IDs given in table 1: **1)** $T = 1.5$, **2)** $T = 1.75$, **3)** $T = 2.0$, **4)** $T = 2.25$, **5)** $T = 2.5$. The temperature was dumped 1000 times for each phase, and only the final 990 energy values recorded are shown here. The final phase ran $1E7$ timesteps of $1E-5s$ each.

3 Results

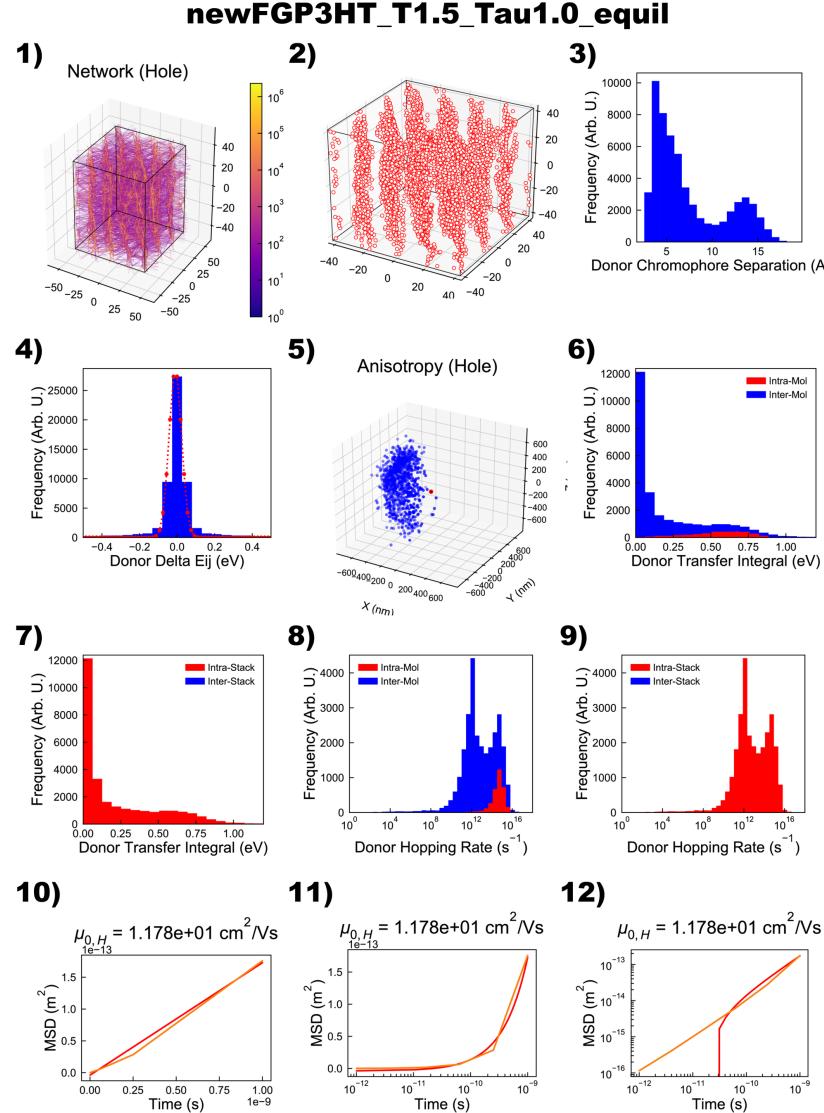


Figure 3: 1) Chromophore connectivity network, 2) Location of ‘stacks’, 3) Distribution of connected chromophore separations (defines stacks), 4) Density of states of Frontier molecular orbital (δE_{ij}), 5) KMC Carrier termination locations (defines anisotropy), 6) Histogram of molecular transfer integrals, 7) Histogram of stack transfer integrals, 8) Histogram of molecular hopping rates, 9) Histogram of stack hopping rates, 10) Linear MSD plot, 11) Semi-log-x MSD plot, 12) Logarithmic MSD plot.

newFGP3HT_T1.75_Tau1.0_equil

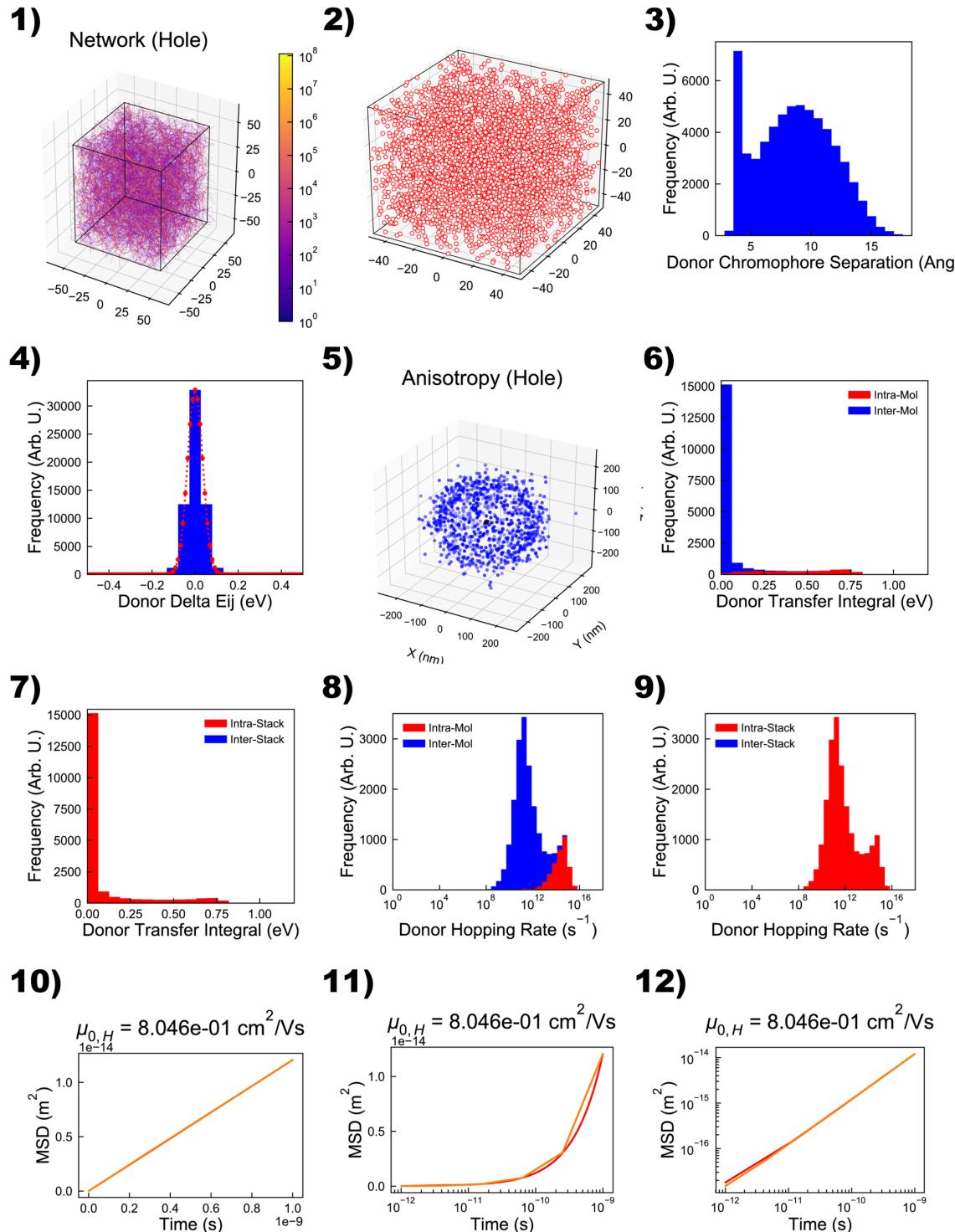


Figure 4: 1) Chromophore connectivity network, 2) Location of ‘stacks’, 3) Distribution of connected chromophore separations (defines stacks), 4) Density of states of Frontier molecular orbital (ΔE_{ij}), 5) KMC Carrier termination locations (defines anisotropy), 6) Histogram of molecular transfer integrals, 7) Histogram of stack transfer integrals, 8) Histogram of molecular hopping rates, 9) Histogram of stack hopping rates, 10) Linear MSD plot, 11) Semi-log-x MSD plot, 12) Logarithmic MSD plot.

newFGP3HT_T2.0_Tau1.0_equil

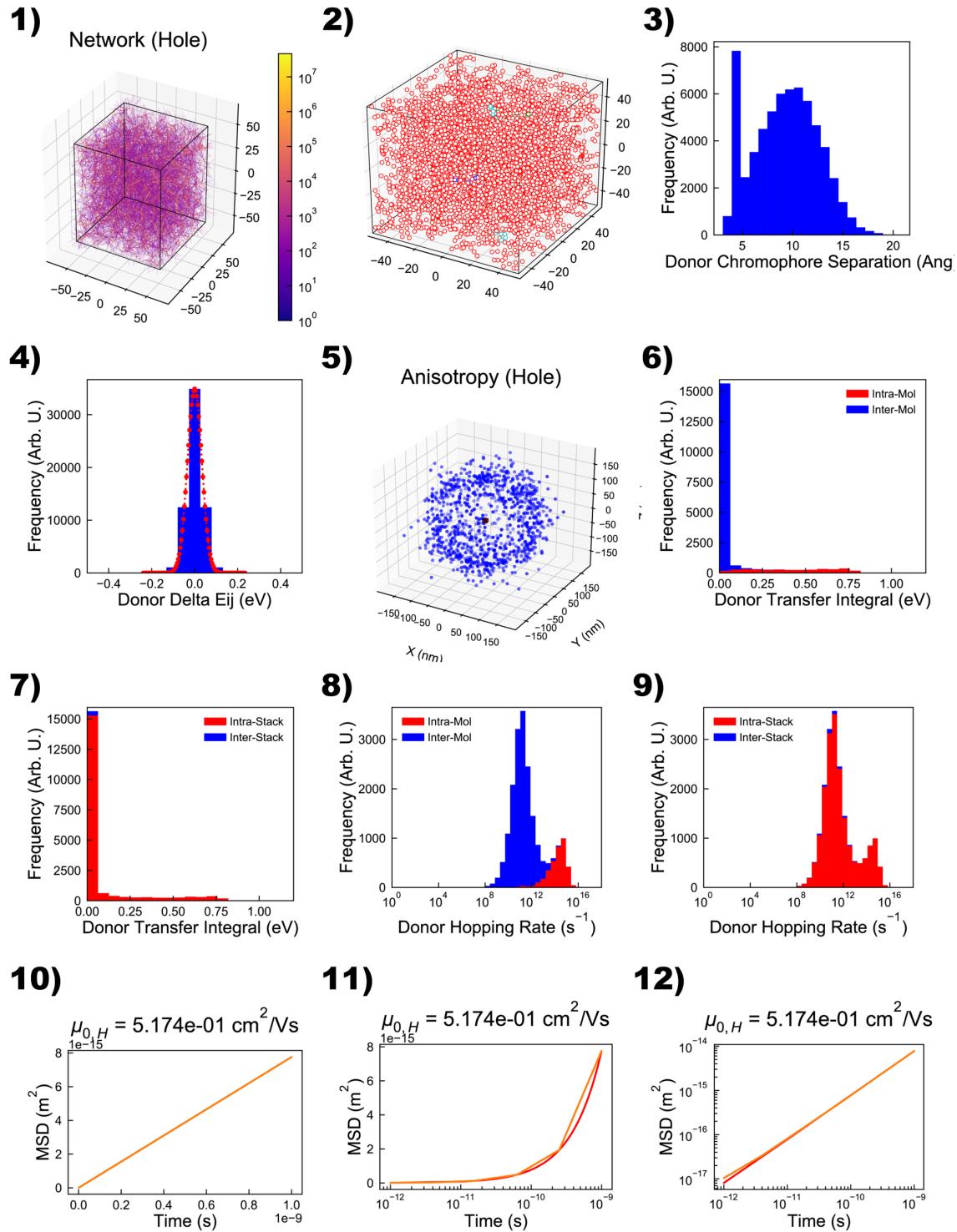


Figure 5: 1) Chromophore connectivity network, 2) Location of ‘stacks’, 3) Distribution of connected chromophore separations (defines stacks), 4) Density of states of Frontier molecular orbital (ΔE_{ij}), 5) KMC Carrier termination locations (defines anisotropy), 6) Histogram of molecular transfer integrals, 7) Histogram of stack transfer integrals, 8) Histogram of molecular hopping rates, 9) Histogram of stack hopping rates, 10) Linear MSD plot, 11) Semi-log-x MSD plot, 12) Logarithmic MSD plot.

newFGP3HT_T2.25_Tau1.0_equil

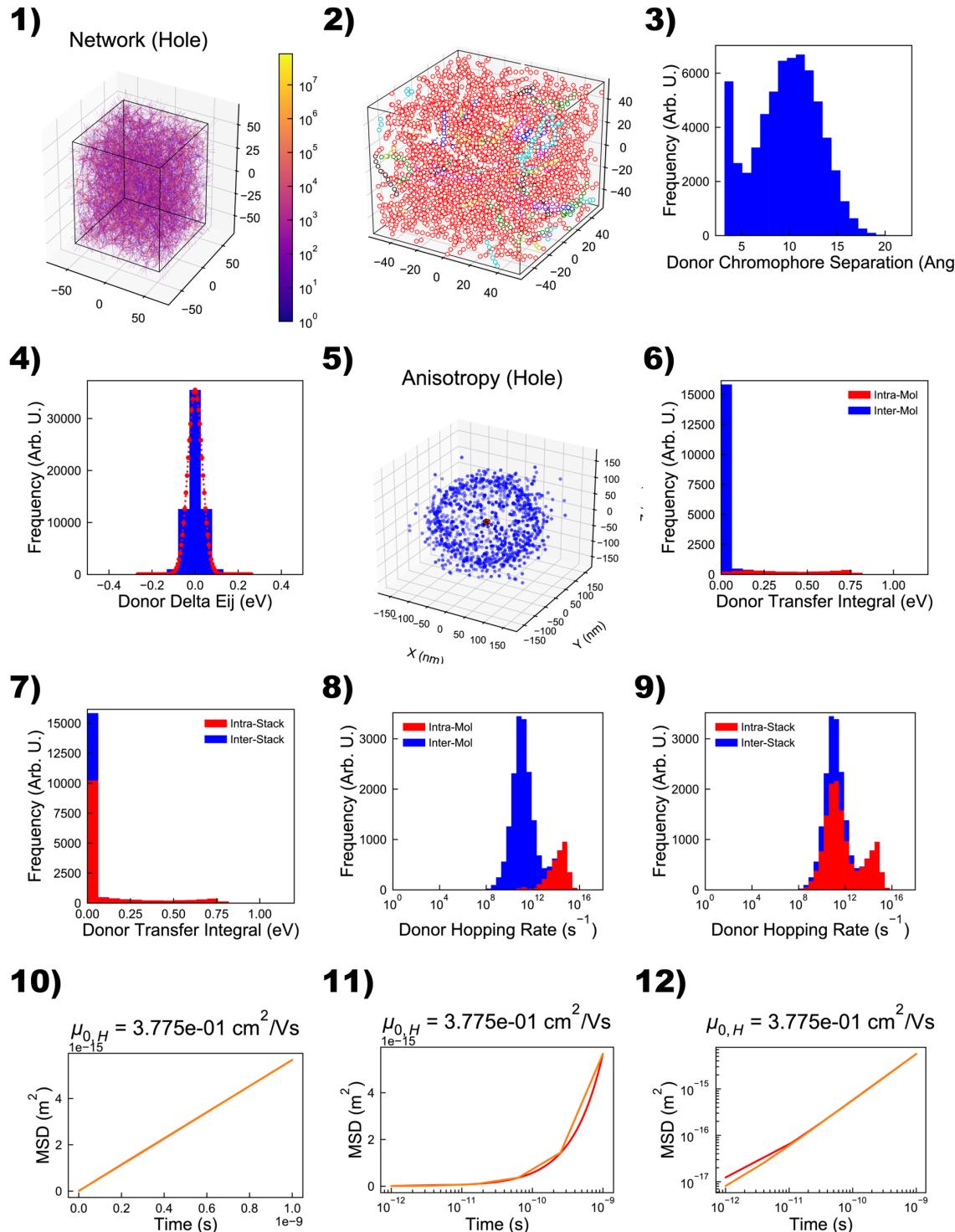


Figure 6: 1) Chromophore connectivity network, 2) Location of ‘stacks’, 3) Distribution of connected chromophore separations (defines stacks), 4) Density of states of Frontier molecular orbital (δE_{ij}), 5) KMC Carrier termination locations (defines anisotropy), 6) Histogram of molecular transfer integrals, 7) Histogram of stack transfer integrals, 8) Histogram of molecular hopping rates, 9) Histogram of stack hopping rates, 10) Linear MSD plot, 11) Semi-log-x MSD plot, 12) Logarithmic MSD plot.

newFGP3HT_T2.5_Tau1.0_equil

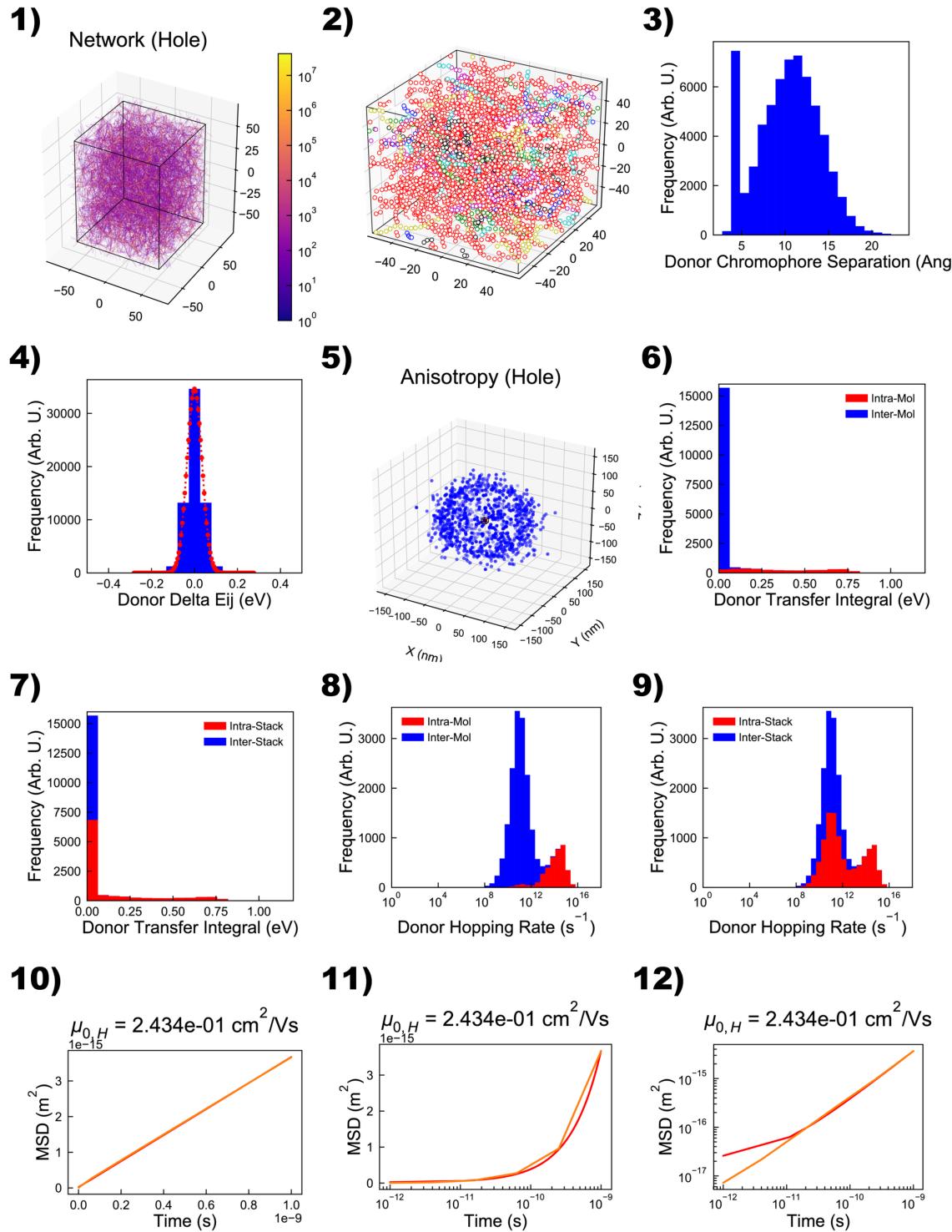


Figure 7: 1) Chromophore connectivity network, 2) Location of ‘stacks’, 3) Distribution of connected chromophore separations (defines stacks), 4) Density of states of Frontier molecular orbital (ΔE_{ij}), 5) KMC Carrier termination locations (defines anisotropy), 6) Histogram of molecular transfer integrals, 7) Histogram of stack transfer integrals, 8) Histogram of molecular hopping rates, 9) Histogram of stack hopping rates, 10) Linear MSD plot, 11) Semi-log-x MSD plot, 12) Logarithmic MSD plot.