

## Supplementary Material

### DFT ANALYSIS OF FENETHYLLINE (CAPTAGON): INVESTIGATING ITS INTERACTION WITH GRAPHENE AS A POTENTIAL ADSORBENT

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Table S1. Computational details: A comprehensive summary of all simulation settings and parameters (including those from DFT, Monte Carlo, MD, and GULP optimizations)

| DFT calculations (DMol3)     |  |
|------------------------------|--|
| Parameter                    | Value  |
| Functional                   | M11L   |
| Basis set                    | DNP (Double Numerical with Polarization basis set) |
| Spin treatment               | Unrestricted                                       |
| Total charge                 | 0  |
| SCF density convergence      | $1 \times 10^{-6}$ [e/Å <sup>3</sup> ]             |
| Geometry optimization cutoff | $1 \times 10^{-5}$ [Ha]                            |
| Global orbital cutoff        | 3.7 [Å]  |
| Max displacement             | 0.300 [Å]  |
| Integration grid             | Fine (~9000 pts/atom)                              |

| <b>Monte Carlo</b>          |   |
|-----------------------------|---|
| <i>Parameter</i>            | <i>Value</i>  |
| Force field                 | Dreiding  |
| Electrostatic cutoff        | 18.5 [Å]  |
| van der Waals cutoff        | 18.5 [Å]  |
| Simulation type             | Simulated annealing   |
| Steps per cycle             | 100000 (Number of MC steps per heating/cooling cycle)                             |
| Heating cycles              | 30 (Number of temperature cycles in annealing)                                    |
| <b>Molecular Dynamic</b>    |   |
| <i>Parameter</i>            | <i>Value</i>  |
| Ensemble                    | NVT (Constant number of particles, volume, and temperature)                       |
| Temperature                 | 298 K (System temperature maintained by the Nose thermostat)                      |
| Thermostat                  | Nose (Temperature control method)   |
| Timestep                    | 1 fs (Time increment per MD step)   |
| Simulation duration         | 2000 ps [Total run time (2,000,000 steps × 1 fs)]                                 |
| Force field                 | Dreiding (Empirical potential for intramolecular and intermolecular interactions) |
| Electrostatic & vdW cutoff  | 18.5 [Å] (Cutoff distance for non-bonded interactions)                            |
| Initial velocities          | Random (Velocities assigned based on random distribution at start of simulation)  |
| <b>GULP</b>                 |   |
| <i>Parameter</i>            | <i>Value</i>  |
| Forcefield                  | ReaxFF 6.0 (Reactive force field for bond formation/breaking)                     |
| Charges                     | Forcefield assigned (Charges taken from ReaxFF potential)                         |
| Energy (function) tolerance | $1.0 \times 10^{-5}$ [eV] (Threshold for energy change per step)                  |
| Max. gradient tolerance     | $1.0 \times 10^{-4}$ [eV] (Maximum allowed force on atoms)                        |