

**Supporting information**

**for**

**Computational studies on the interactions of glycine amino acid  
with graphene, h-BN and h-SiC monolayers**

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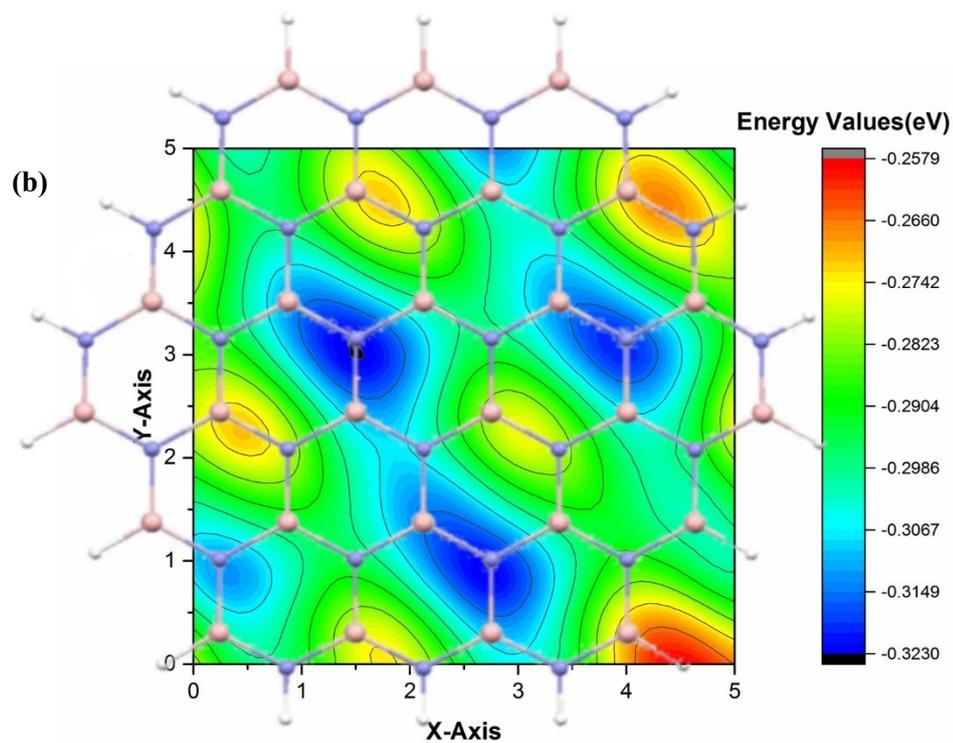
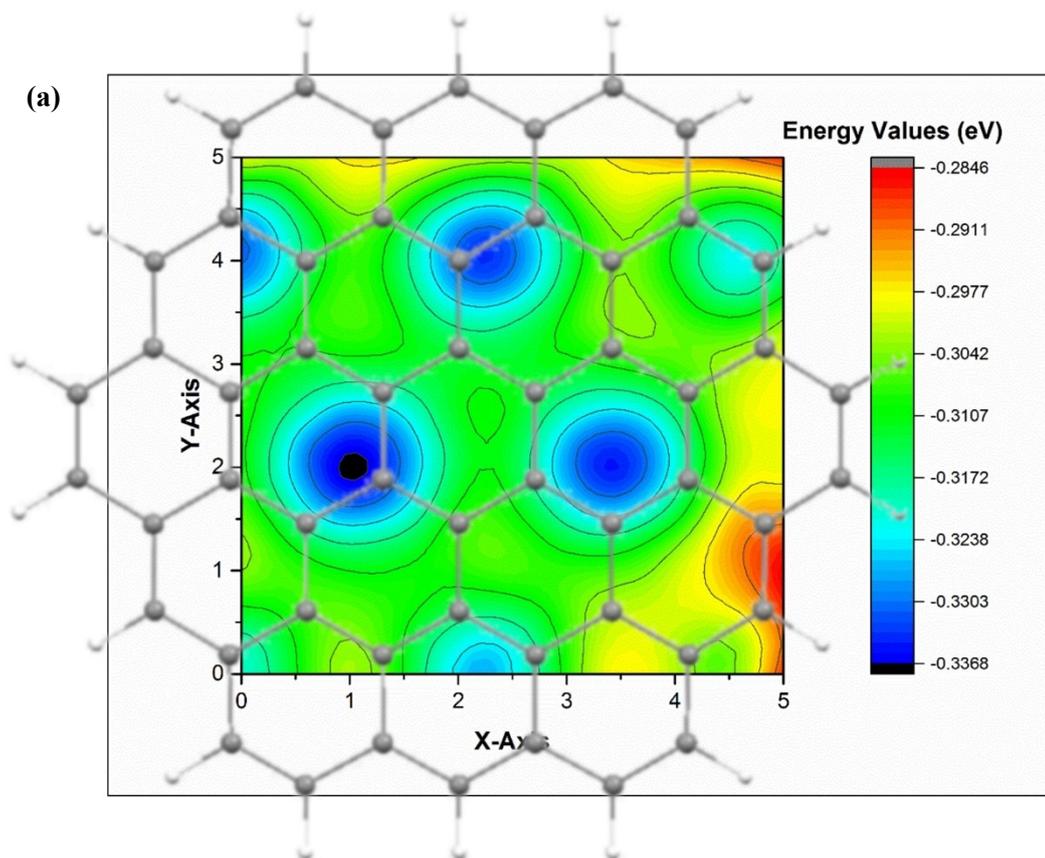
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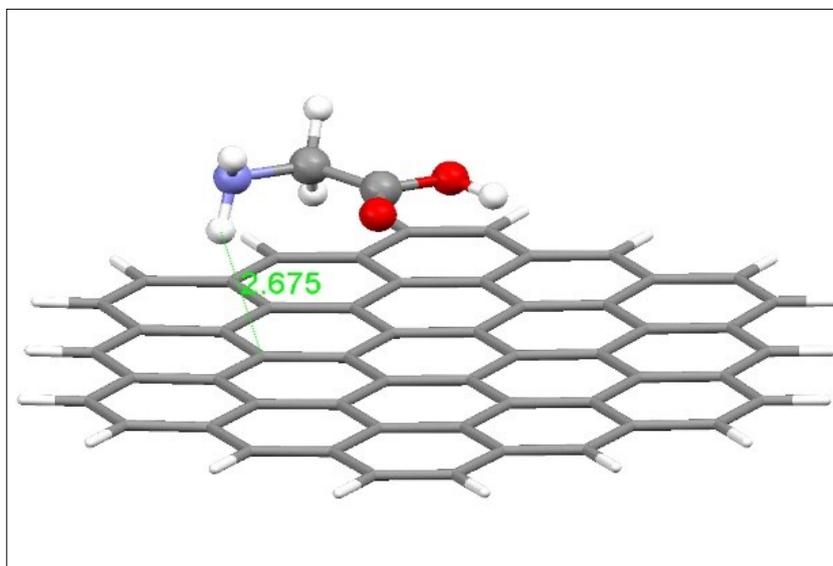
**Table S1. Binding energy value obtained from initial relaxation step for the glycine/h-BN system.**

<b>Configuration</b>	<b>Degree</b>	<b>Binding Energy (eV)</b>
1	0	-0.15933
2	30	-0.29867
3	60	-0.29815
4	90	-0.29683
5	120	-0.29559
6	150	-0.16123
7	180	-0.16463
8	210	-0.29698
9	240	-0.29702
10	270	-0.29924
11	300	-0.29807
12	330	-0.29248
NH <sub>2</sub>	---	-0.07969
OH	---	-0.08572

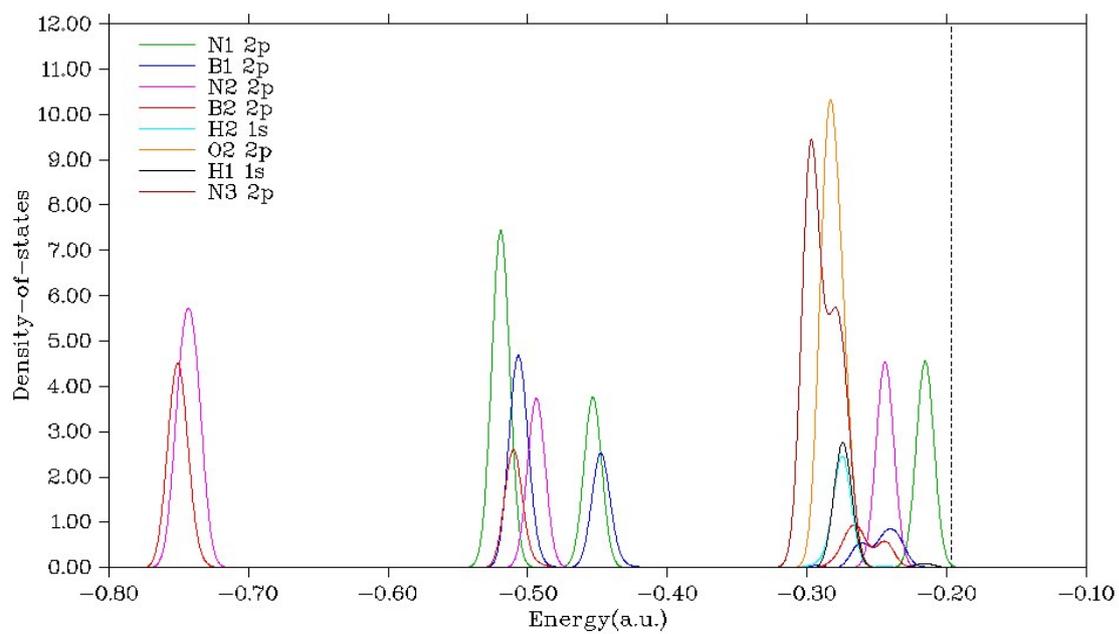
Figure S1. (a) and (b) PES plots with overlapped graphene and h-BN lattices, respectively.



**Figure S2. Optimized geometry of glycine/graphene complex at the B3LYP-D3 Level**



**Figure S3. PDOS plot for the glycine/h-BN system in wider energy range. The dashed vertical line corresponds to the HOMO level.**



**Figure S4. Visualization of (a) HOMO and (b) LUMO of the hexagonal SiC**

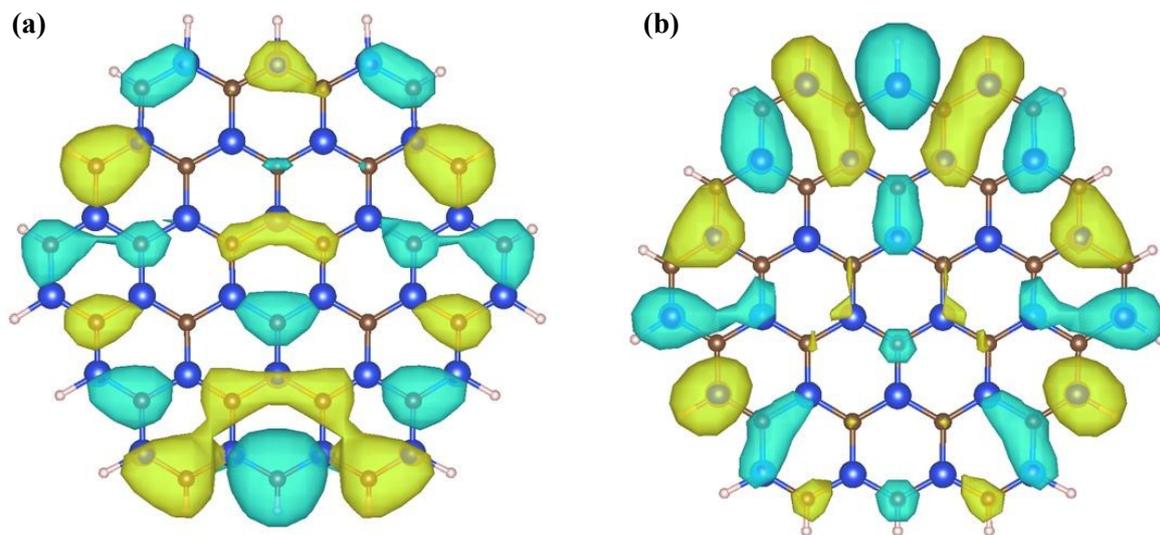
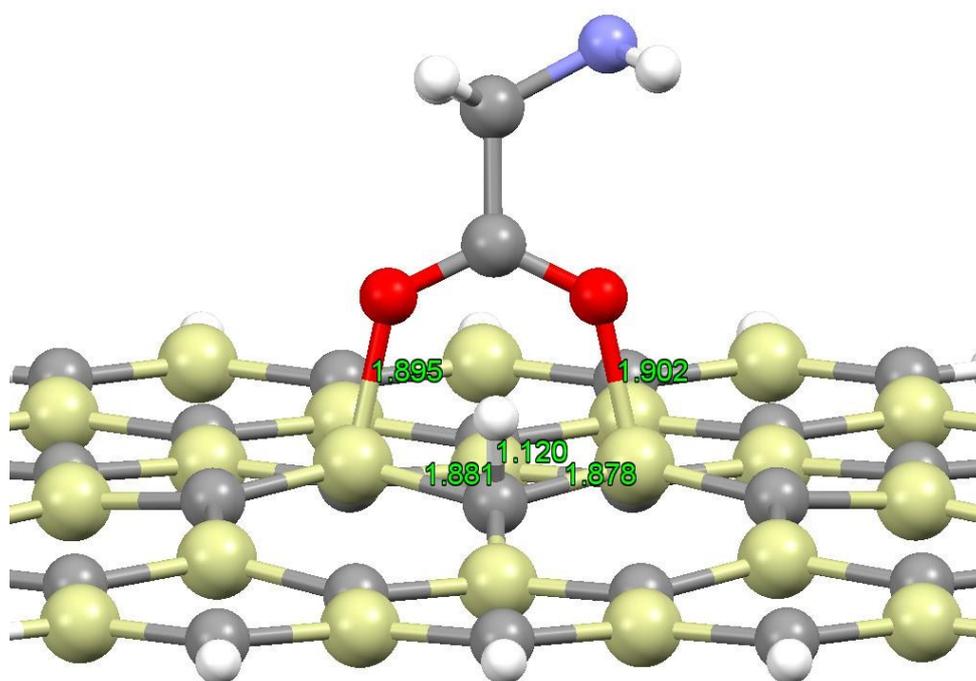


Figure S5. Optimized structure of the glycine/h-SiC system with constrained edge atoms.



**Figure S6. PDOS plot for the glycine/h-SiC system in wider energy range. The dashed vertical line corresponds to the HOMO level.**

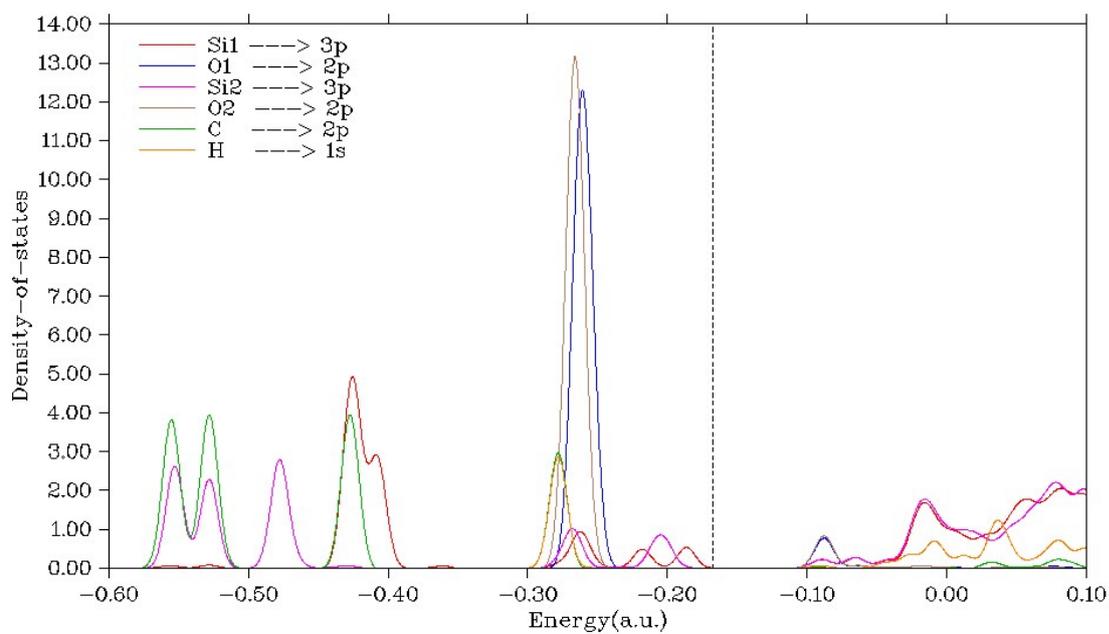


Figure S7. (a)-(c) Optimized structures of glycine/graphene, glycine/h-BN and glycine/h-SiC systems in solvent phase, respectively. (d)-(f) Optimized structures of zwitter/graphene, zwitter/h-BN and zwitter/h-SiC systems, respectively.

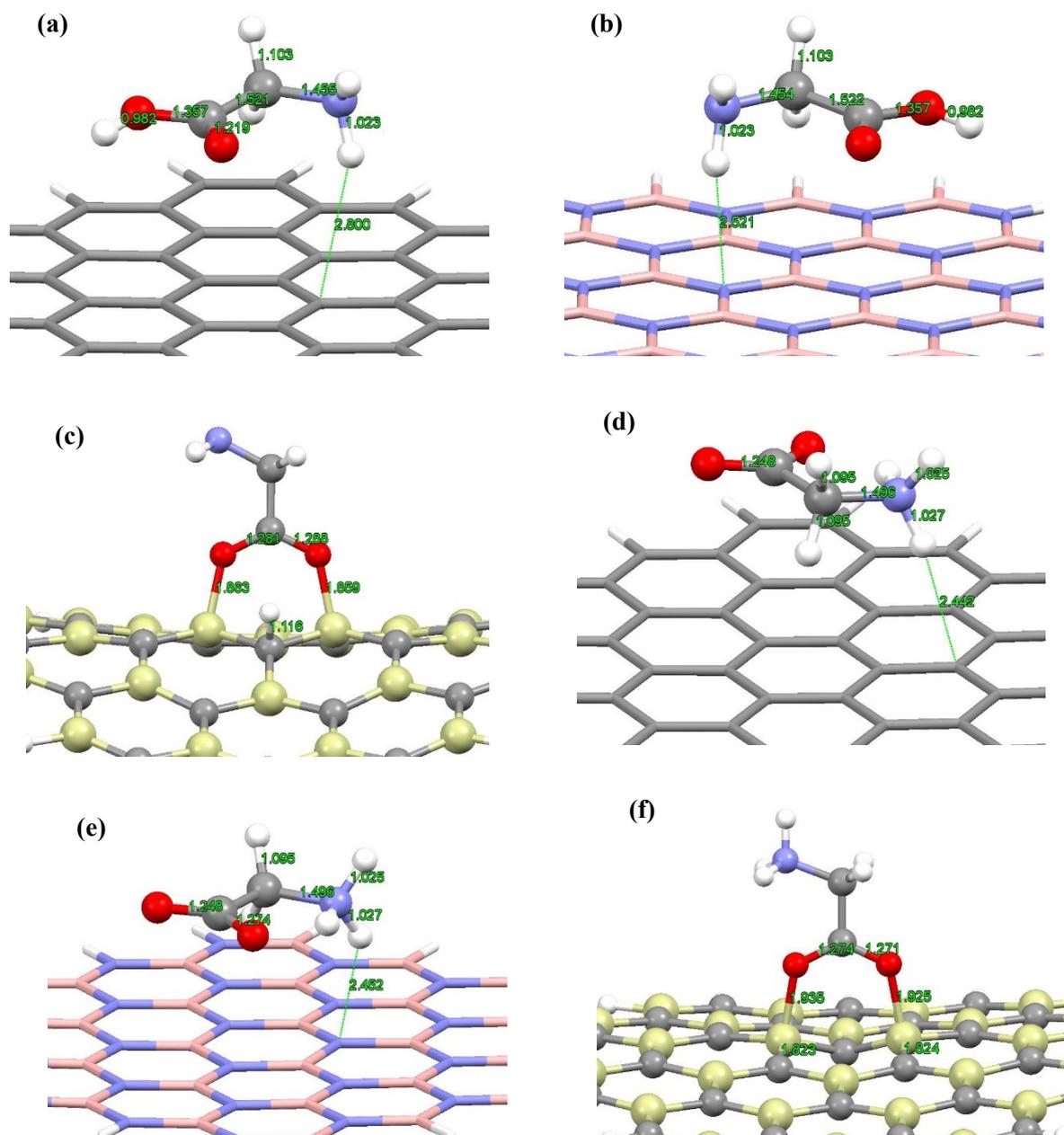


Figure S8. Optimized structures of the miniaturized system of (a) glycine/graphene, (b) zwitter/graphene, (c) glycine/h-SiC and (d) zwitter/h-SiC at the PBE-D3 Level.

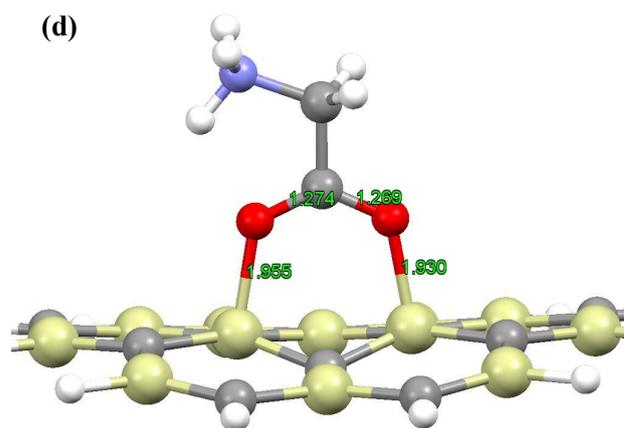
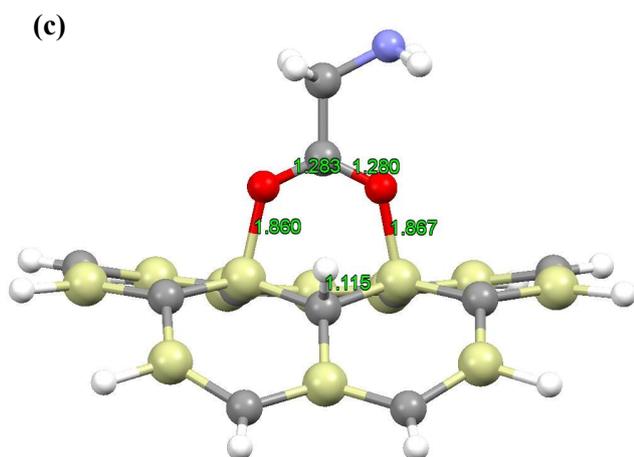
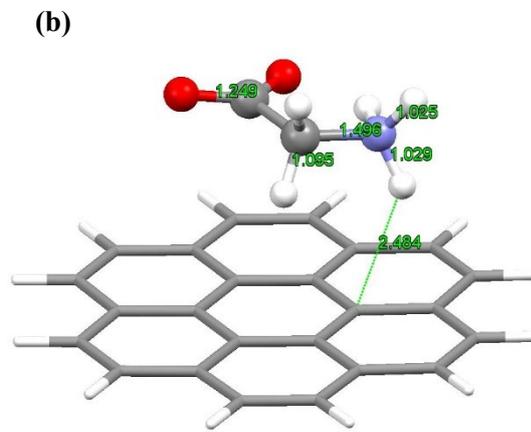
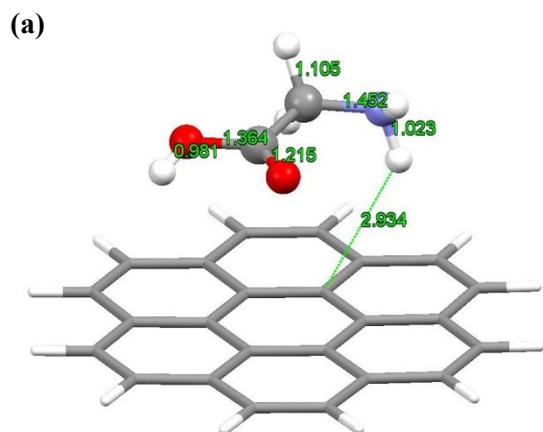


Figure S9. Optimized structures of the miniaturized system of (a) glycine/graphene, (b) zwitter/graphene, (c) glycine/h-SiC and (d) zwitter/h-SiC at the PW6B95-D3 Level.

