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# **Site and chirality selective chemical modifications of boron nitride nanotubes (BNNTs) via Lewis acid/base interactions**

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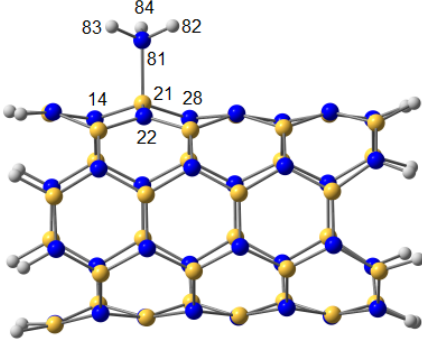
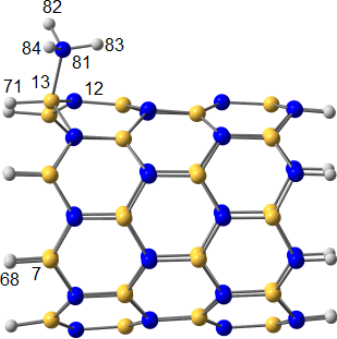
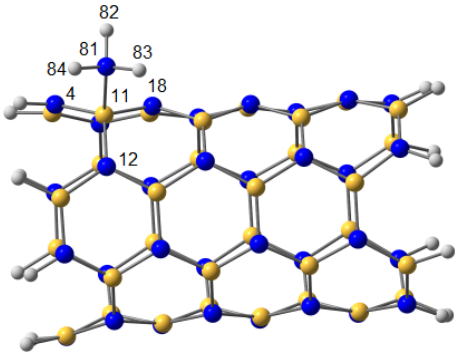
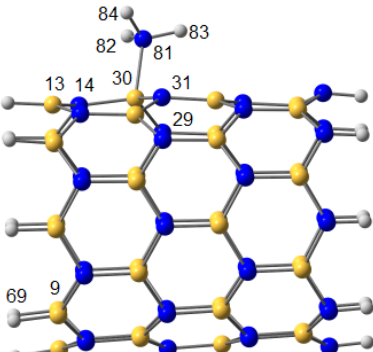
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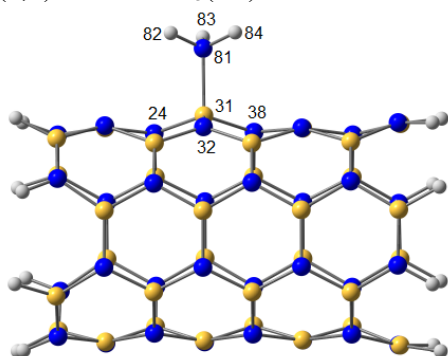
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Table S1. Comparison of B3LYP/6-31G\* and B3LYP/6-31+G\* adsorption energies of BNNT-NH<sub>3</sub> and BNNT-BH<sub>3</sub> (see Figure 1-4 for structures). All energies are in kcal/mol.

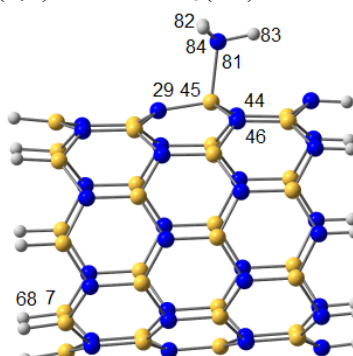
|                            | E <sub>ad</sub> |         |
|----------------------------|-----------------|---------|
| (4,4)-BNNT-NH <sub>3</sub> | 6-31G*          | 6-31+G* |
| 1A                         | 9.86            | 7.58    |
| 1B                         | 9.70            | 7.64    |
| 1C                         | 9.65            | 7.49    |
| 1D                         | 9.34            | 7.20    |
| (8,0)-BNNT-NH <sub>3</sub> |                 |         |
| 2A                         | 16.60           | 14.03   |
| 2B                         | 10.18           | 7.68    |
| 2C                         | 9.72            | 7.28    |
| 2D                         | 3.25            | 1.29    |
| (4,4)-BNNT-BH <sub>3</sub> |                 |         |
| 3A                         | 22.96           | 21.05   |
| 3B                         | 10.57           | 9.28    |
| 3C                         | 10.04           | 8.39    |
| 3D                         | 9.20            | 7.85    |
| (8,0)-BNNT-BH <sub>3</sub> |                 |         |
| 4A                         | 17.06           | 15.80   |
| 4B                         | 17.01           | 15.53   |
| 4C                         | 9.71            | 8.04    |
| 4D                         | 9.25            | 7.71    |

Table S2. Bond distances and changes in distances ( $\delta R$ ) of  $\text{BH}_3$  adsorbed BN-nanotubes at the adsorption sites as well as at their neighbors. All distances are in Å.

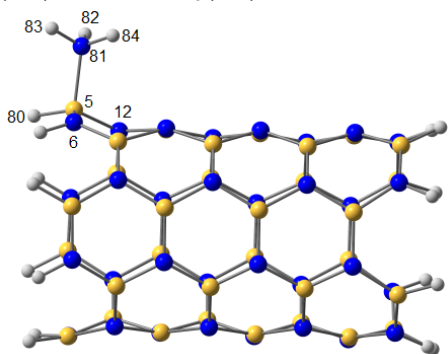
| Bond distance ( R )   |                |                        |                       |  |                |                        |                       |
|---|----------------|------------------------|-----------------------|--|----------------|------------------------|-----------------------|
| Name of the System  | 6-31+G* (BSSE) | 6-31+G* ( $\delta R$ ) | 6-31G* ( $\delta R$ ) | Name of the System   | 6-31+G* (BSSE) | 6-31+G* ( $\delta R$ ) | 6-31G* ( $\delta R$ ) |
| (4,4)-BNNT-NH <sub>3</sub> (1A)   |                |                        |                       | (8,0)-BNNT-NH <sub>3</sub> (2A)  |                |                        |                       |
|    |                |                        |                       |    |                |                        |                       |
| N81-H84   | 1.021          | -0.001                 | 0.000                 | N81-H84  | 1.020          | -0.001                 | -0.001                |
| N81-H83   | 1.021          | -0.001                 | -0.001                | N81-H83  | 1.023          | -0.001                 | -0.001                |
| N81-H82   | 1.021          | -0.001                 | -0.001                | N81-H82  | 1.020          | 0.001                  | 0.002                 |
| B21-N81   | 1.724          | 0.006                  | 0.000                 | N81-B13  | 1.691          | 0.007                  | 0.005                 |
| B21-N14   | 1.526          | 0.000                  | 0.003                 | B13-N12  | 1.534          | 0.000                  | 0.001                 |
| B21-N28   | 1.518          | 0.000                  | 0.002                 | B13-H71  | 1.204          | 0.001                  | 0.001                 |
| B21-N22   | 1.530          | 0.000                  | 0.002                 | H68-B7   | 1.192          | 0.000                  | 0.000                 |
| (4,4)-BNNT-NH <sub>3</sub> (1B)   |                |                        |                       | (8,0)-BNNT-NH <sub>3</sub> (2B)  |                |                        |                       |
|  |                |                        |                       |  |                |                        |                       |
| N81-H84   | 1.021          | -0.001                 | -0.001                | N81-H84  | 1.021          | 0.000                  | 0.000                 |
| N81-H83   | 1.022          | -0.001                 | 0.000                 | N81-H83  | 1.023          | 0.000                  | 0.000                 |
| N81-H82   | 1.021          | -0.001                 | 0.000                 | N81-H82  | 1.021          | -0.001                 | 0.000                 |
| B11-N81   | 1.727          | 0.010                  | 0.001                 | N81-B30  | 1.725          | 0.009                  | 0.005                 |
| B11-N4  | 1.510          | 0.000                  | 0.004                 | B30-N31  | 1.525          | 0.000                  | 0.001                 |
| B11-N18   | 1.517          | 0.000                  | 0.001                 | B30-N14  | 1.513          | 0.000                  | 0.002                 |
| B11-N12   | 1.535          | -0.001                 | 0.002                 | H69-B9   | 1.193          | 0.001                  | 0.000                 |

(4,4)-BNNT-NH<sub>3</sub>(1C)

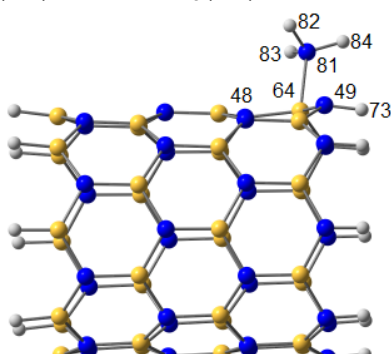
|         |       |        |        |
|---------|-------|--------|--------|
| N81-H84 | 1.021 | -0.001 | -0.001 |
| N81-H83 | 1.021 | -0.001 | 0.000  |
| N81-H82 | 1.021 | -0.001 | -0.001 |
| B31-N81 | 1.733 | 0.010  | 0.003  |
| B31-N24 | 1.516 | -0.001 | 0.001  |
| B31-N38 | 1.516 | 0.000  | 0.002  |
| B31-N32 | 1.526 | 0.000  | 0.002  |

(8,0)-BNNT-NH<sub>3</sub>(2C)

|         |       |        |        |
|---------|-------|--------|--------|
| N81-H84 | 1.020 | -0.001 | -0.001 |
| N81-H83 | 1.024 | -0.001 | -0.001 |
| N81-H82 | 1.020 | -0.001 | -0.001 |
| N81-B45 | 1.724 | 0.009  | 0.006  |
| B45-N46 | 1.527 | -0.001 | 0.001  |
| B45-N29 | 1.508 | 0.000  | 0.002  |
| H68-B7  | 1.193 | 0.001  | 0.000  |

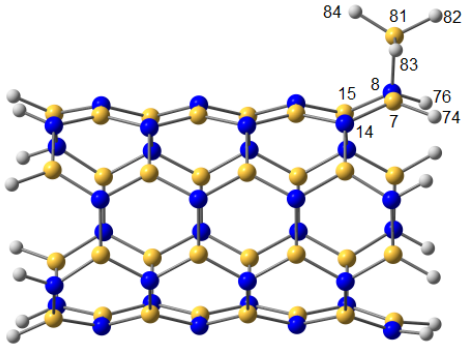
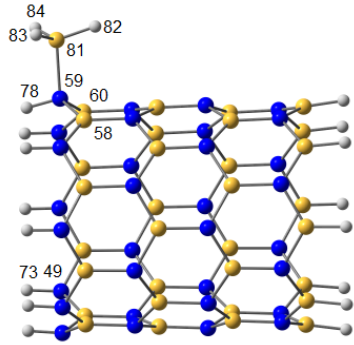
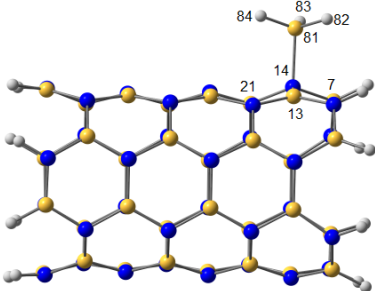
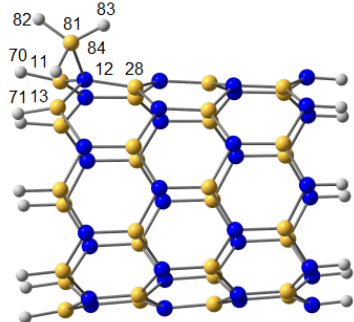
(4,4)-BNNT-NH<sub>3</sub>(1D)

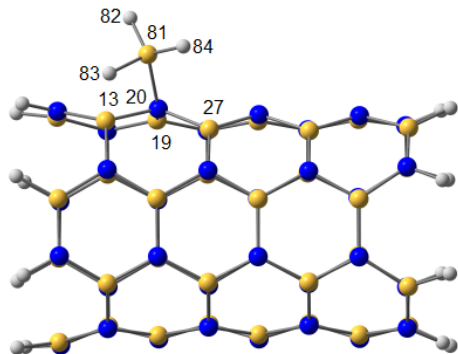
|         |       |        |        |
|---------|-------|--------|--------|
| N81-H84 | 1.022 | -0.001 | -0.001 |
| N81-H83 | 1.020 | -0.001 | 0.000  |
| N81-H82 | 1.021 | -0.001 | 0.000  |
| B5-N81  | 1.724 | 0.010  | 0.004  |
| B5-N80  | 1.213 | 0.000  | 0.000  |
| B5-N12  | 1.524 | -0.001 | 0.001  |
| B5-N6   | 1.510 | 0.000  | 0.003  |

(8,0)-BNNT-NH<sub>3</sub>(2D)

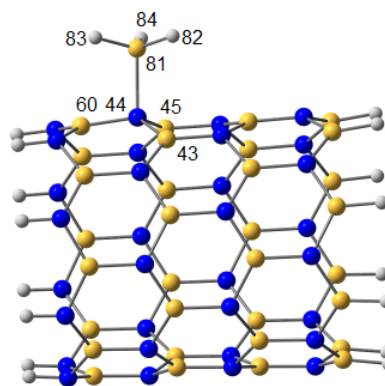
|         |       |        |        |
|---------|-------|--------|--------|
| N81-H84 | 1.021 | 0.000  | 0.000  |
| N81-H83 | 1.020 | 0.001  | 0.001  |
| N81-H82 | 1.020 | -0.001 | -0.001 |
| N81-B64 | 1.789 | 0.056  | 0.003  |
| B64-N48 | 1.506 | -0.001 | 0.001  |
| B64-N80 | 1.513 | -0.001 | 0.007  |
| N80-H73 | 1.016 | 0.000  | 0.000  |

Table S3. Bond distances and changes in distances ( $\delta R$ ) of  $BH_3$  adsorbed BN-nanotubes at the adsorption sites as well as at their neighbors. All distances are in Å.

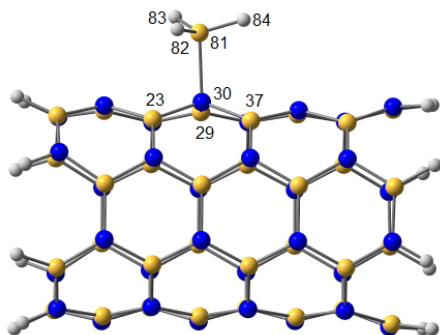
| Bond distance(R)  |                |                        |                       |  |                |                        |                       |
|---|----------------|------------------------|-----------------------|--|----------------|------------------------|-----------------------|
| Name of the System  | 6-31+G* (BSSE) | 6-31+G* ( $\delta R$ ) | 6-31G* ( $\delta R$ ) | Name of the System   | 6-31+G* (BSSE) | 6-31+G* ( $\delta R$ ) | 6-31G* ( $\delta R$ ) |
| (4,4)-BNNT-BH <sub>3</sub> (3A)    |                |                        |                       | (8,0)-BNNT-BH <sub>3</sub> (4A)    |                |                        |                       |
| B81-H84   | 1.194          | 0.001                  | 0.001                 | B81-H84  | 1.212          | 0.000                  | 0.001                 |
| B81-H83   | 1.197          | 0.001                  | 0.001                 | B81-H83  | 1.212          | 0.000                  | 0.001                 |
| B81-H82   | 1.326          | -0.001                 | -0.001                | B81-H82  | 1.207          | 0.000                  | 0.001                 |
| N8-B81  | 1.585          | 0.002                  | 0.003                 | N59-B81  | 1.740          | 0.004                  | 0.002                 |
| B7-H74  | 1.196          | 0.000                  | 0.000                 | N59-B60  | 1.503          | 0.000                  | 0.001                 |
| B7-B14  | 1.493          | 0.000                  | 0.001                 | N59-B58  | 1.503          | 0.000                  | 0.001                 |
| N8-H76  | 1.015          | 0.000                  | 0.000                 | N59-H78  | 1.019          | 0.000                  | 0.001                 |
| N8-B15  | 1.484          | 0.000                  | 0.000                 | N49-H73  | 1.015          | 0.000                  | 0.001                 |
| B7-N8   | 1.590          | 0.002                  | 0.002                 | -  | -              | -                      | -                     |
| (4,4)-BNNT-BH <sub>3</sub> (3B)  |                |                        |                       | (8,0)-BNNT-BH <sub>3</sub> (4B)  |                |                        |                       |
| B81-H84   | 1.204          | 0.003                  | 0.002                 | B81-H84  | 1.315          | -0.001                 | -0.001                |
| B81-H83   | 1.210          | 0.000                  | 0.000                 | B81-H83  | 1.196          | 0.000                  | 0.001                 |
| B81-H82   | 1.208          | 0.000                  | -0.001                | B81-H82  | 1.195          | 0.001                  | 0.001                 |
| N14-B81   | 1.804          | 0.008                  | 0.010                 | N12-B81  | 1.610          | 0.002                  | 0.004                 |
| N14-B7  | 1.500          | 0.000                  | 0.002                 | N12-B11  | 1.610          | 0.003                  | 0.004                 |
| N14-B21   | 1.501          | 0.000                  | 0.002                 | H71-B13  | 1.192          | 0.000                  | 0.000                 |
| N14-B13   | 1.501          | 0.000                  | 0.002                 | N12-B28  | 1.488          | 0.000                  | 0.000                 |
| -   | -              | -                      | -                     | H70-B11  | 1.187          | 0.000                  | 0.000                 |
| -   | -              | -                      | -                     | N12-B13  | 1.606          | 0.000                  | 0.000                 |

(4,4)-BNNT-BH<sub>3</sub>(3C)

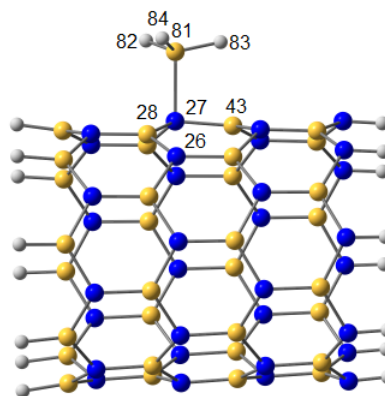
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|---------|-------|-------|-------|
| B81-H84 | 1.201 | 0.000 | 0.001 |
| B81-H83 | 1.299 | 0.000 | 0.001 |
| B81-H82 | 1.195 | 0.001 | 0.001 |
| N20-B81 | 1.629 | 0.002 | 0.004 |
| N20-B13 | 1.557 | 0.000 | 0.001 |
| N20-B27 | 1.501 | 0.000 | 0.0   |
| N20-B19 | 1.506 | 0.000 | 0.001 |

(8,0)-BNNT-BH<sub>3</sub>(4C)

|         |       |        |       |
|---------|-------|--------|-------|
| B81-H84 | 1.209 | 0.000  | 0.000 |
| B81-H83 | 1.209 | 0.000  | 0.000 |
| B81-H82 | 1.204 | 0.001  | 0.001 |
| N44-B81 | 1.808 | 0.008  | 0.014 |
| N44-B60 | 1.490 | 0.000  | 0.002 |
| N44-B43 | 1.499 | -0.001 | 0.002 |
| N44-B45 | 1.499 | 0.000  | 0.001 |

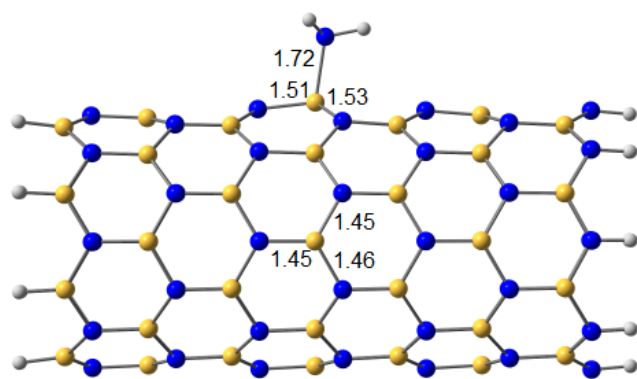
(4,4)-BNNT-BH<sub>3</sub>(3D)

|         |       |        |        |
|---------|-------|--------|--------|
| B81-H84 | 1.202 | 0.000  | 0.000  |
| B81-H83 | 1.206 | 0.000  | 0.000  |
| B81-H82 | 1.209 | 0.000  | 0.000  |
| N30-B81 | 1.828 | 0.008  | 0.015  |
| N30-B37 | 1.494 | -0.001 | -0.002 |
| N30-B23 | 1.497 | -0.001 | 0.004  |
| N30-B29 | 1.498 | 0.000  | -0.001 |

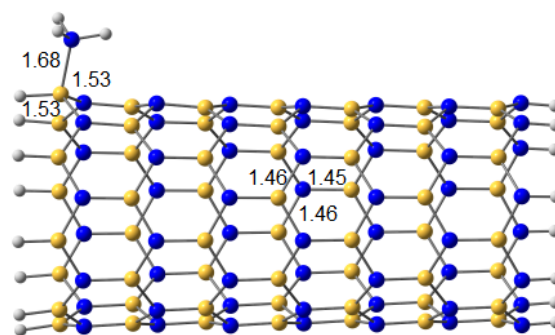
(8,0)-BNNT-BH<sub>3</sub>(4D)

|         |       |        |        |
|---------|-------|--------|--------|
| B81-H84 | 1.205 | 0.002  | 0.002  |
| B81-H83 | 1.216 | 0.003  | 0.002  |
| B81-H82 | 1.204 | -0.002 | -0.002 |
| N27-B81 | 1.783 | -0.004 | 0.002  |
| N27-B26 | 1.495 | 0.000  | 0.001  |
| N27-B43 | 1.489 | 0.001  | 0.001  |
| N27-B28 | 1.495 | 0.000  | 0.001  |

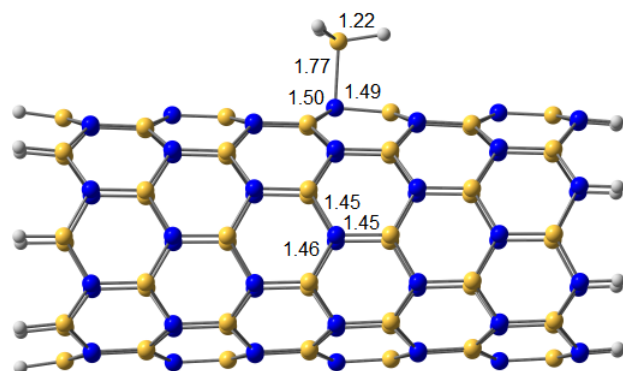
Fig. S4. B3LYP/6-31+G\* optimized structures of L-(8, 0)-BNNT-NH<sub>3</sub>/BH<sub>3</sub>. Blue, yellow and grey colors represent N, B and H atoms, respectively. m and e stand for middle and edge site respectively. Bond lengths are in Å. BSSE corrected adsorption energies (in kcal/mol) are given in parenthesis and a positive value indicates attractive interaction between two units,



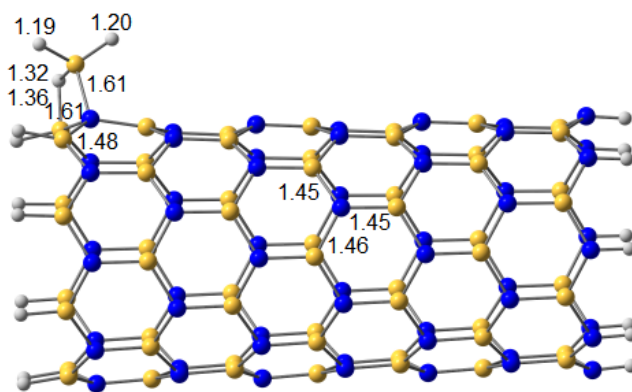
L-(8,0)-BNNT-NH3-m (**4.2**)



L-(8,0)-BNNT-NH3-e (**11.9**)



L-(8,0)-BNNT-BH3-m (**6.9**)



L-(8,0)-BNNT-BH3-e (**14.2**)

Fig. S5. B3LYP/6-31+G\* optimized structures of (12, 0)-BNNT-NH<sub>3</sub>/BH<sub>3</sub>. Blue, yellow and grey colors represent N, B and H atoms, respectively. m and e stand for middle and edge site respectively. Bond lengths are in Å. BSSE corrected adsorption energies (in kcal/mol) are given in parenthesis and a positive value indicates attractive interaction between two units,

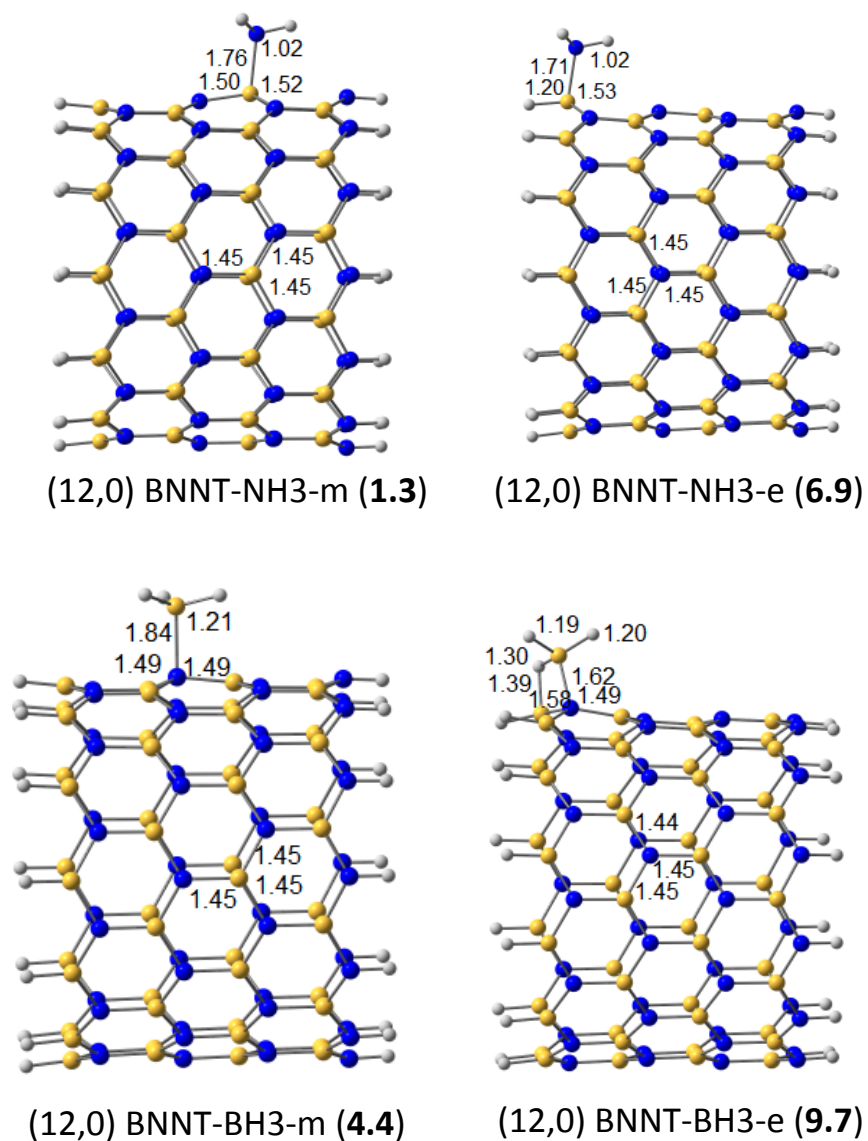




Fig. S6. Hybridization angle ( $\phi$  in degree) of  $\text{BH}_3$  ( $\text{sp}^2$ ),  $\text{NH}_3$  ( $\text{sp}^3$ ) and  $\text{H}_3\text{BNH}_3$  ( $\text{sp}^3$ ). The dummy atom (pink) is used to show angles.

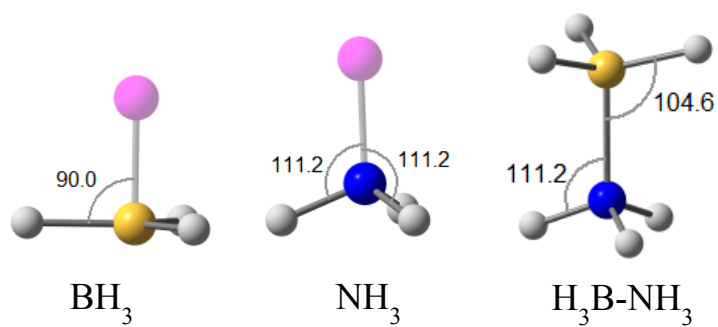


Fig. S7. Approximate hybridization angles ( $\phi$  in degree) of B (Lewis acid centers) and N (Lewis base centers) sites in (4,4) and (8,0) BNNTs, The dummy atom (pink) is used to show angles.

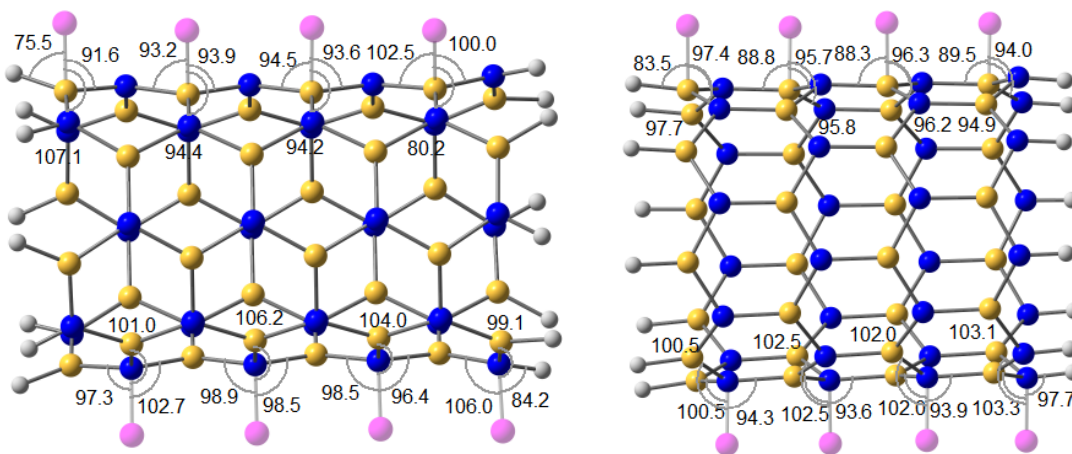
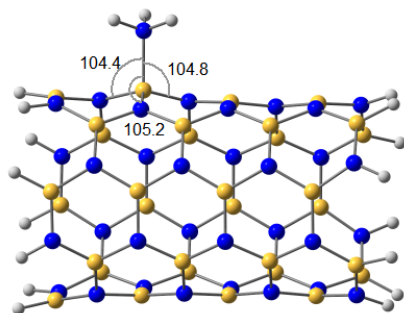
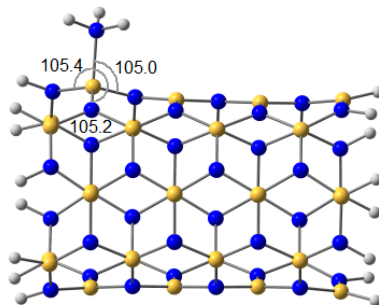


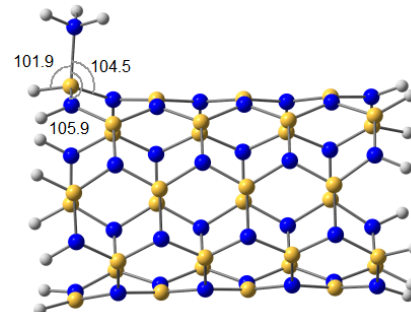
Fig. S8. Hybridization angles ( $\phi$  in degree) of BNNT-NH<sub>3</sub> complexes.



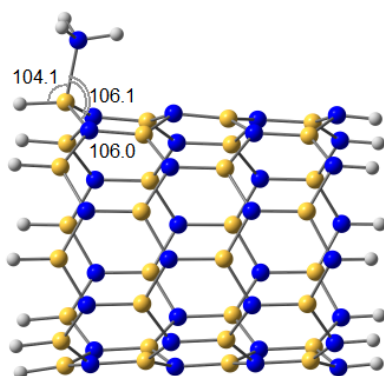
(4,4)-BNNT-NH<sub>3</sub> (1A)  
Middle site



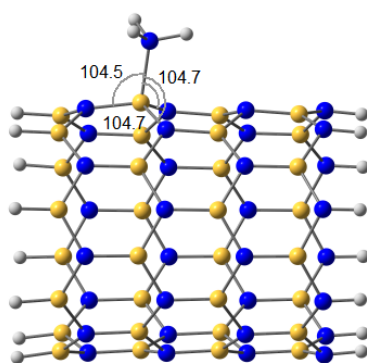
(4,4)-BNNT-NH<sub>3</sub> (1B)  
Near the edge site



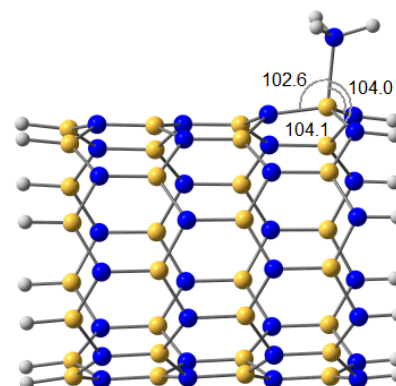
(4,4)-BNNT-NH<sub>3</sub> (1D)  
Edge site



(8,0)-BNNT-NH<sub>3</sub> (2A)  
Edge site



(8,0)-BNNT-NH<sub>3</sub> (2B)  
Middle site



(8,0)-BNNT-NH<sub>3</sub> (2D)  
Near the edge site

Fig. S9. Hybridization angles ( $\phi$  in degree) of BNNT-BH<sub>3</sub> complexes.

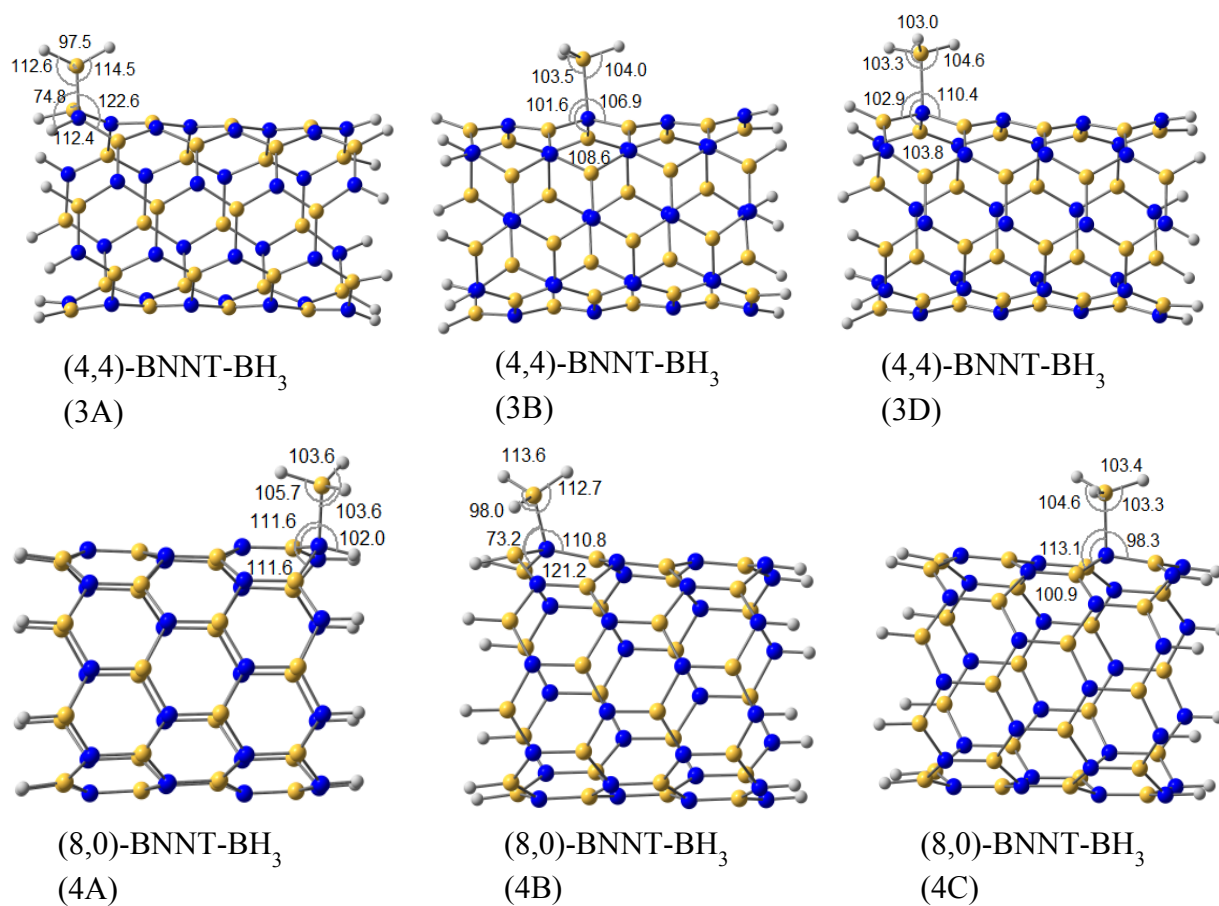


Fig. S10. Natural bond orbitals (NBO) of B-N and B-H-B 3c-2e bonds of most stable BNNT-NH<sub>3</sub>/BH<sub>3</sub> complexes. NBO representing multi-center B-H-B bond of B<sub>2</sub>H<sub>6</sub> are shown in the last diagram.

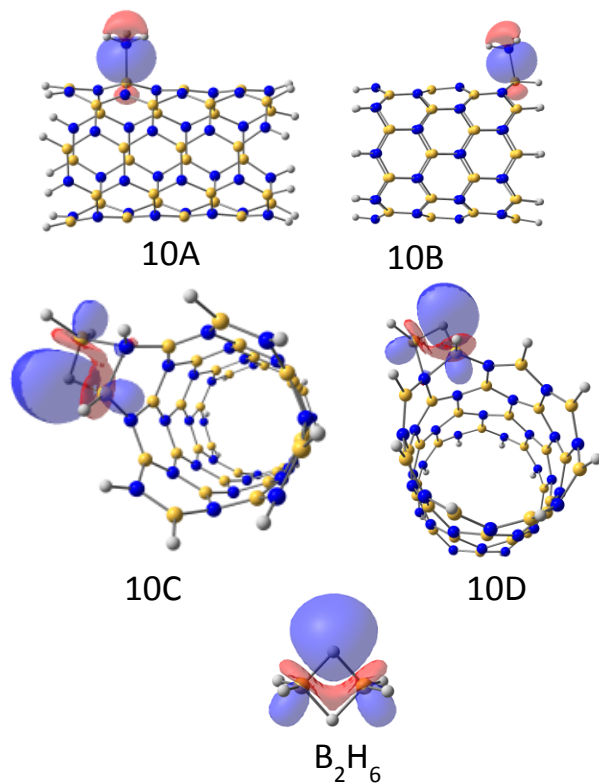
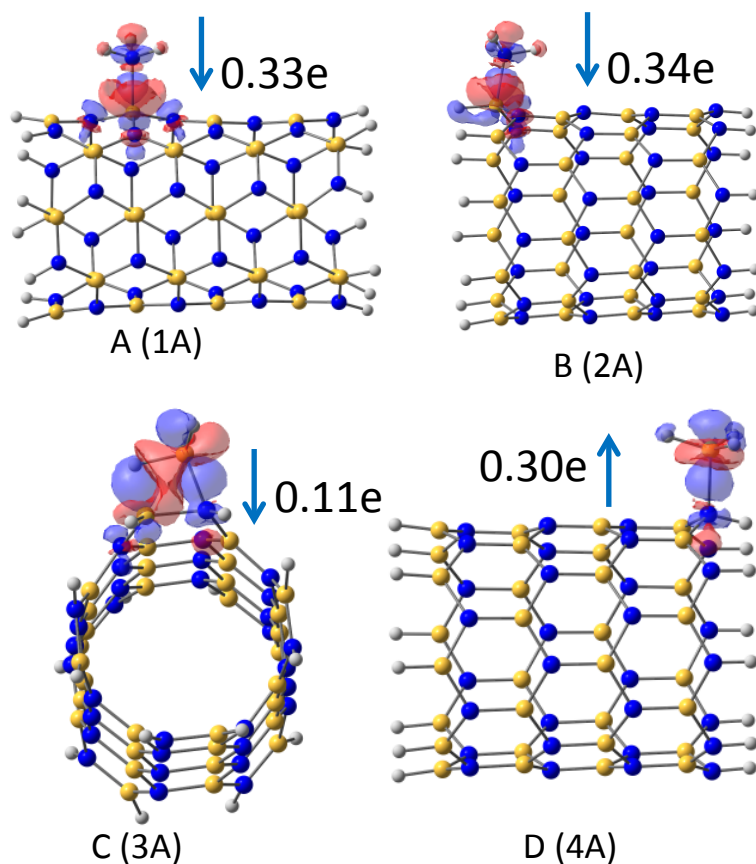


Fig. S11. Density difference plots of most stable BNNT-NH<sub>3</sub>/BH<sub>3</sub> structures. Blue and red regions indicate gain and loss of electron density, respectively, upon complex formation. Contour value is 0.005 au. Electron transfer from or to BNNTs are shown by arrow with number of electron.



Density difference plots clearly indicate charge gain (blue) or loss (red) is localized in the vicinity of the active site only. In all complexes, electron density builds between B and N atoms of host and guest, indicated by blue region and some gain is also noted on nitrogen of NH<sub>3</sub> in 1A and 2A. However, boron of BH<sub>3</sub> in 3A and 4A loses some density. Formation of 3c-2e bond in 3A (also in 4B, not shown) causes gain in density between H(BH<sub>3</sub>) and B(NT), while weakening of H<sub>2</sub>B-H bond is reflected by the loss of density.

Table S12. Electron exchange between BN-nanotubes and NH<sub>3</sub>/BH<sub>3</sub> in complexes. Natural (NPA) group charges are obtained from B3LYP/6-31G\* calculations.

| (4,4)-BNNT-NH <sub>3</sub> |  | (8,0)-BNNT-NH <sub>3</sub> |   |
|----------------------------|--|----------------------------|---|
| 1A                         | NH <sub>3</sub> $\xrightarrow{0.33\text{ e}}$ (4,4) BNNT | 2A                         | NH <sub>3</sub> $\xrightarrow{0.34\text{ e}}$ (8,0)BNNT |
| 1B                         | NH <sub>3</sub> $\xrightarrow{0.32\text{ e}}$ (4,4) BNNT | 2B                         | NH <sub>3</sub> $\xrightarrow{0.33\text{ e}}$ (8,0)BNNT |
| 1C                         | NH <sub>3</sub> $\xrightarrow{0.33\text{ e}}$ (4,4)BNNT  | 2C                         | NH <sub>3</sub> $\xrightarrow{0.33\text{ e}}$ (8,0)BNNT |
| 1D                         | NH <sub>3</sub> $\xrightarrow{0.33\text{ e}}$ (4,4)BNNT  | 2D                         | NH <sub>3</sub> $\xrightarrow{0.33\text{ e}}$ (8,0)BNNT |
|                            |  |                            |   |
| (4,4)-BNNT-BH <sub>3</sub> |  | (8,0)-BNNT-BH <sub>3</sub> |   |
| 3A                         | BH <sub>3</sub> $\xleftarrow{0.11\text{ e}}$ (4,4) BNNT  | 4A                         | BH <sub>3</sub> $\xrightarrow{0.30\text{ e}}$ (8,0)BNNT |
| 3B                         | BH <sub>3</sub> $\xrightarrow{0.26\text{ e}}$ (4,4)BNNT  | 4B                         | BH <sub>3</sub> $\xleftarrow{0.14\text{ e}}$ (8,0)BNNT  |
| 3C                         | BH <sub>3</sub> $\xleftarrow{0.10\text{ e}}$ (4,4)BNNT   | 4C                         | BH <sub>3</sub> $\xrightarrow{0.22\text{ e}}$ (8,0)BNNT |
| 3D                         | BH <sub>3</sub> $\xrightarrow{0.23\text{ e}}$ (4,4)BNNT  | 4D                         | BH <sub>3</sub> $\xrightarrow{0.19\text{ e}}$ (8,0)BNNT |

Fig. S13. B3LYP/6-31+G\* optimized structures of  $\text{H}_3\text{N}-\text{AlH}_3$  and corresponding BNNT- $\text{AlH}_3$ . Bond lengths are in Å and angles are in degree. e and m stand for edge and middle adsorption sites of the tube, respectively. BSSE corrected adsorption energies (in kcal/mol) are given in bold and a positive value indicates attractive interaction between two units, otherwise interaction is repulsive. Electron transfer from or to BNNTs are shown by arrow with number of electron.

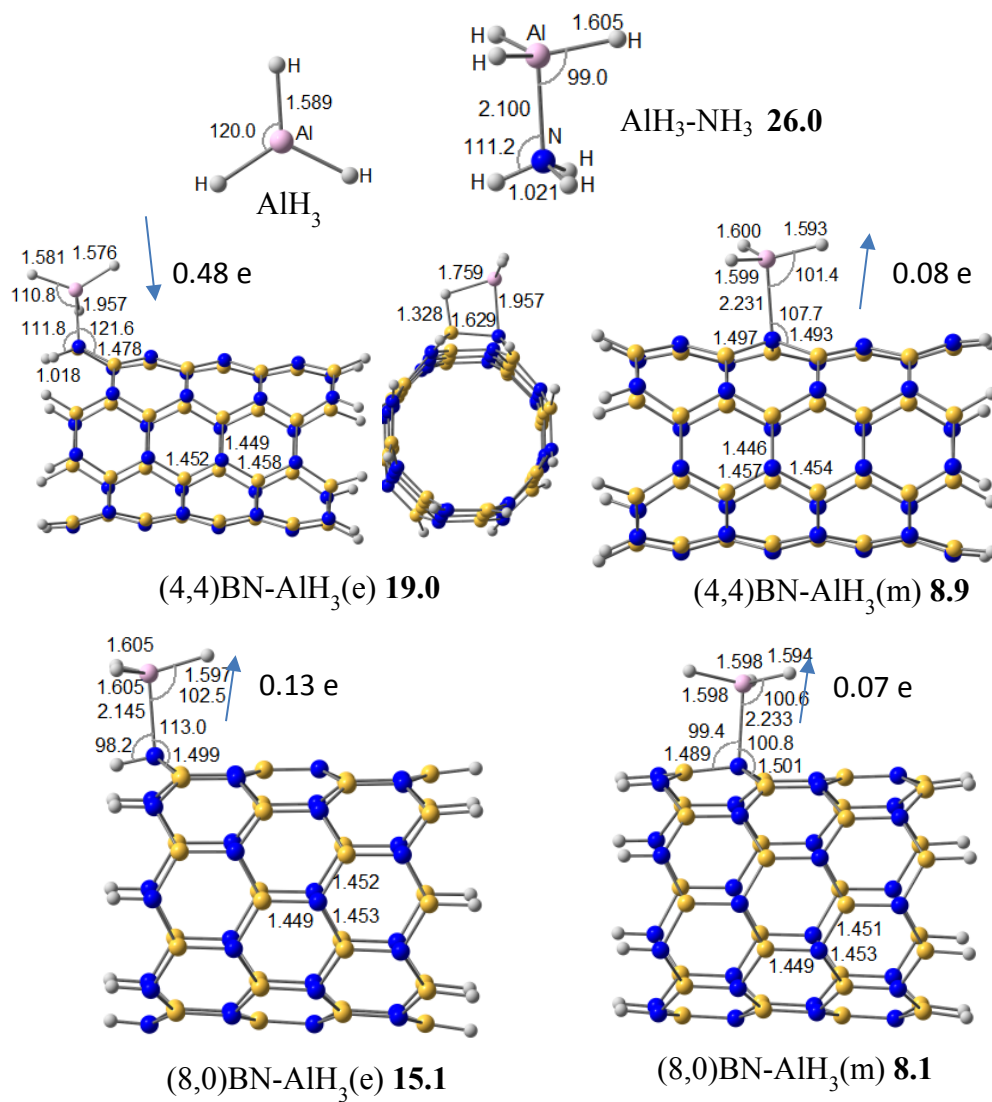


Fig. S14. B3LYP/6-31+G\* optimized structures of  $\text{H}_3\text{N}-\text{BF}_3$  and corresponding BNNT-  $\text{BF}_3$ . Bond lengths are in Å and angles are in degree. e and m stand for edge and middle adsorption sites of the tube, respectively. BSSE corrected adsorption energies (in kcal/mol) are given in bold and a positive value indicates attractive interaction between two units, otherwise interaction is repulsive. Electron transfer from or to BNNTs are shown by arrow with number of electron.

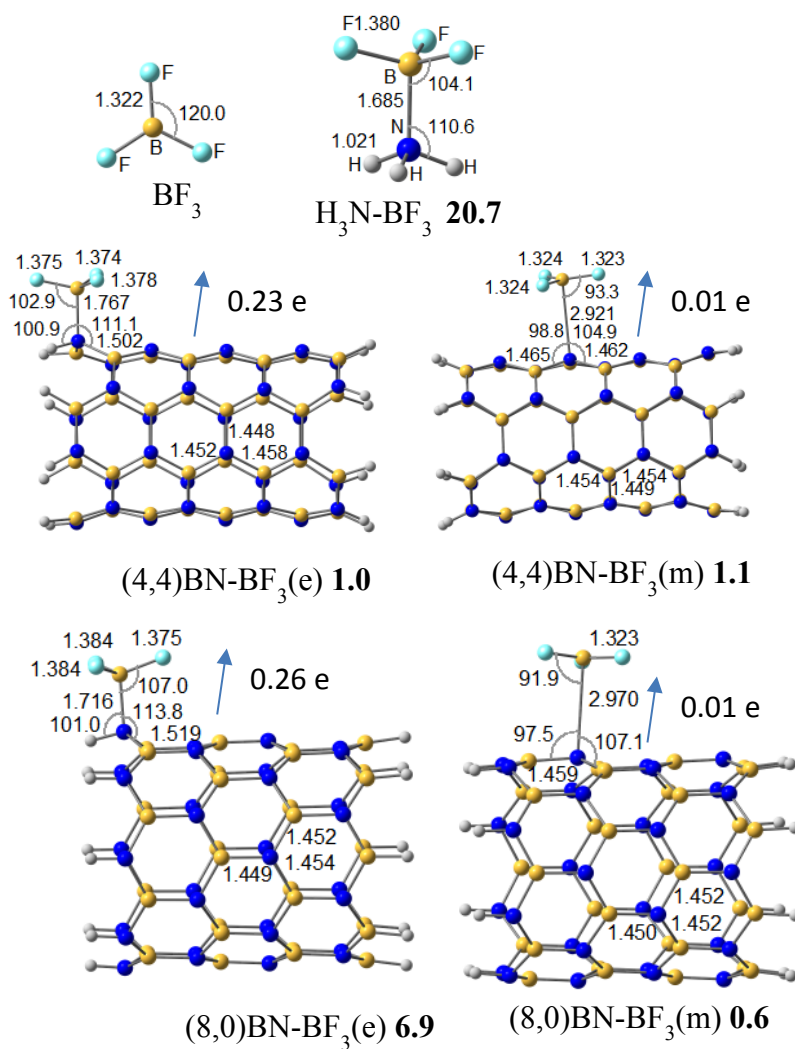




Fig. S15. B3LYP/6-31+G\* optimized structures of  $\text{H}_3\text{N}-\text{BCl}_3$  and corresponding BNNT-  $\text{BCl}_3$ . Bond lengths are in Å and angles are in degree. e and m stand for edge and middle adsorption sites of the tube, respectively. BSSE corrected adsorption energies (in kcal/mol) are given in bold and a positive value indicates attractive interaction between two units, otherwise interaction is repulsive. Electron transfer from or to BNNTs are shown by arrow with number of electron.

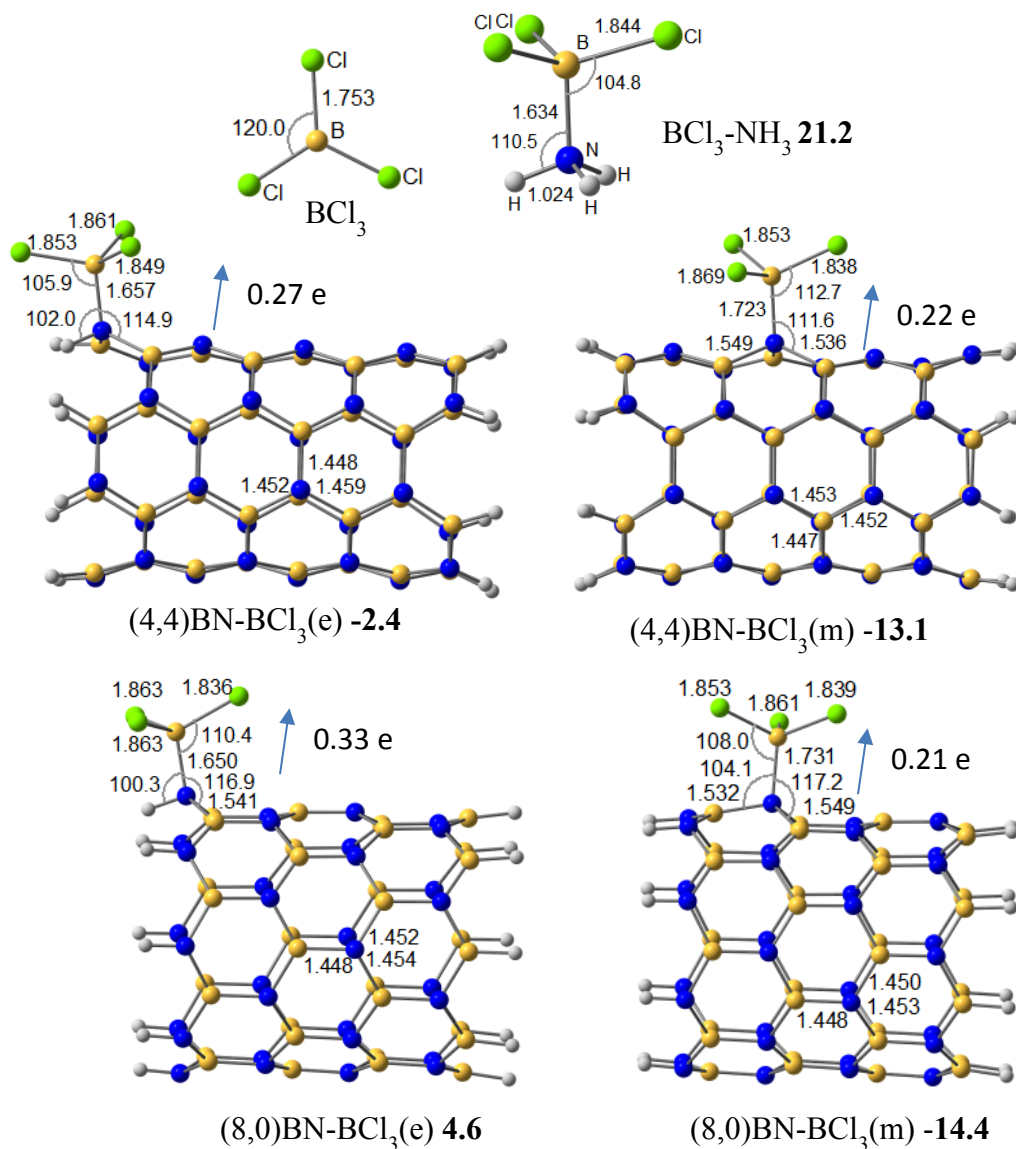


Fig. S16. B3LYP/6-31+G\* optimized structures of  $\text{H}_3\text{N}-\text{BH}_2\text{CH}_3$  and corresponding BNNT-  $\text{BH}_2\text{CH}_3$ . Bond lengths are in Å and angles are in degree. e and m stand for edge and middle adsorption sites of the tube, respectively. BSSE corrected adsorption energies (in kcal/mol) are given in bold and a positive value indicates attractive interaction between two units, otherwise interaction is repulsive. Electron transfer from or to BNNTs are shown by arrow with number of electron.

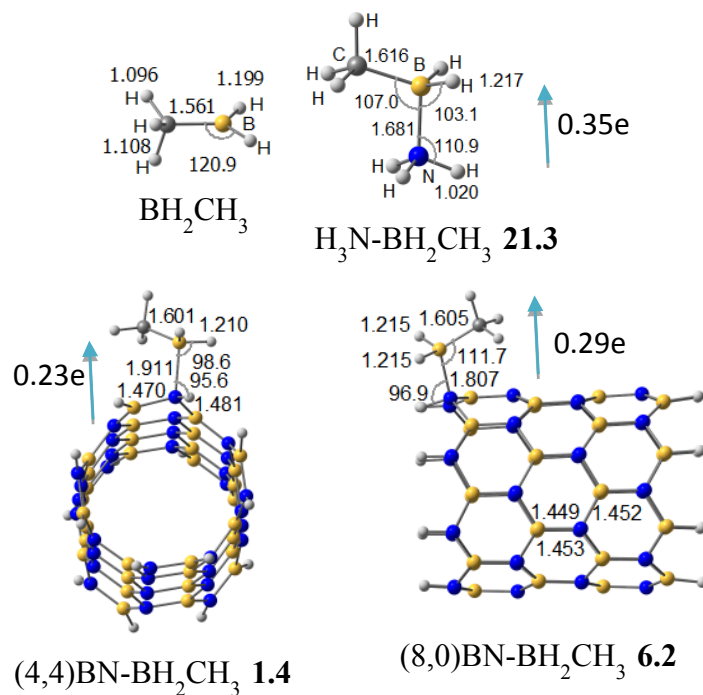


Table-S17. Effects of BSSE correction on adsorption energies; non-corrected energy ( $E_{ad}$ ) and CP-corrected adsorption energies ( $E_{ad(CP)}$ ). All energies are in kcal/mol.

|                                     | BNNT as Lewis Base (LB) |              |          |              | Models                |              |
|-------------------------------------|-------------------------|--------------|----------|--------------|-----------------------|--------------|
|                                     | (4,4)                   |              | (8,0)    |              | NH <sub>3</sub> as LB |              |
| Lewis acids                         | $E_{ad}$                | $E_{ad(CP)}$ | $E_{ad}$ | $E_{ad(CP)}$ | $E_{ad}$              | $E_{ad(CP)}$ |
| BH <sub>3</sub> (e)                 | 21.1                    | 20.0         | 15.8     | 14.7         | 30.1                  | 28.3         |
| BH <sub>3</sub> (m)                 | 9.3                     | 8.3          | 8.0      | 7.0          |                       |              |
| BF <sub>3</sub> (e)                 | 3.9                     | 1.0          | 10.1     | 6.9          | 23.6                  | 20.7         |
| BF <sub>3</sub> (m)                 | 3.3                     | 1.1          | 2.6      | 0.6          |                       |              |
| BCl <sub>3</sub> (e)                | 0.8                     | 2.4          | 8.2      | 4.6          | 24.3                  | 21.2         |
| BCl <sub>3</sub> (m)                | -8.9                    | -13.1        | -10.4    | -14.4        |                       |              |
| BH <sub>2</sub> CH <sub>3</sub> (e) | 2.7                     | 1.4          | 7.7      | 6.2          | 23.3                  | 21.3         |
|                                     |                         |              |          |              |                       |              |
| AlH <sub>3</sub> (e)                | 20.7                    | 19.0         | 16.8     | 15.1         | 27.7                  | 26.0         |
| AlH <sub>3</sub> (m)                | 10.8                    | 8.9          | 9.9      | 8.1          |                       |              |
| CH <sub>3</sub> <sup>+</sup> (e)    | 110.5                   | 109.2        | 115.4    | 113.8        | 112.2                 | 109.8        |
| CH <sub>3</sub> <sup>+</sup> (m)    | 112.0                   | 110.6        | 108.7    | 107.3        |                       |              |
|                                     |                         |              |          |              |                       |              |
|                                     |                         |              |          |              |                       |              |
| Lewis base                          | BNNT as Lewis Acid (LA) |              |          |              | BH <sub>3</sub> as LA |              |
| NH <sub>3</sub> (e)                 | 7.2                     | 4.6          | 14.0     | 11.4         | 30.1                  | 28.3         |
| NH <sub>3</sub> (m)                 | 7.6                     | 4.6          | 7.7      | 4.7          |                       |              |
| NH <sub>2</sub> CH <sub>3</sub> (e) | 8.9                     | 6.6          | 13.9     | 11.5         | 33.8                  | 32.4         |
| NH <sub>2</sub> CH <sub>3</sub> (m) | 9.6                     | 6.7          | 10.0     | 7.1          |                       |              |
| NH <sub>2</sub> COOH                |                         |              |          |              | 13.4                  | 12.2         |
|                                     |                         |              |          |              |                       |              |
| H-Bond                              |                         |              |          |              |                       |              |
| NH <sub>2</sub> COOH (e)            | 5.0                     | 2.8          | 8.9      | 7.8          | 12.8                  | 11.2         |
| NH <sub>2</sub> COOH (m)            | 5.0                     | 4.1          | 4.0      | 2.0          |                       |              |
| NH <sub>2</sub> COOH (m1)           |                         |              | 2.4      | 0.9          |                       |              |