

## Supporting Information

### Computational Insights into Phthalimide-Core Non-Fullerene Acceptors for Next-Generation Organic Solar Cells

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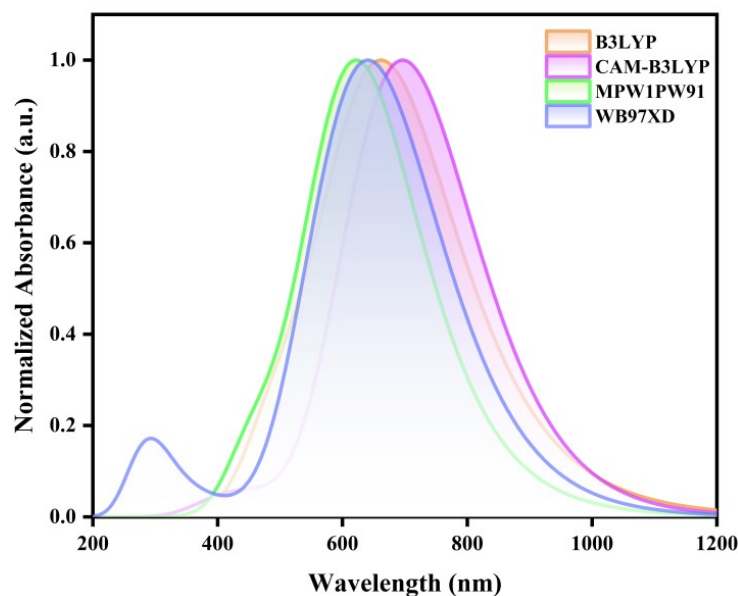


Figure S1 Comparison of four distinct functionals (B3LYP, CAMB3LYP, WB97XD, and MPW1PW91)

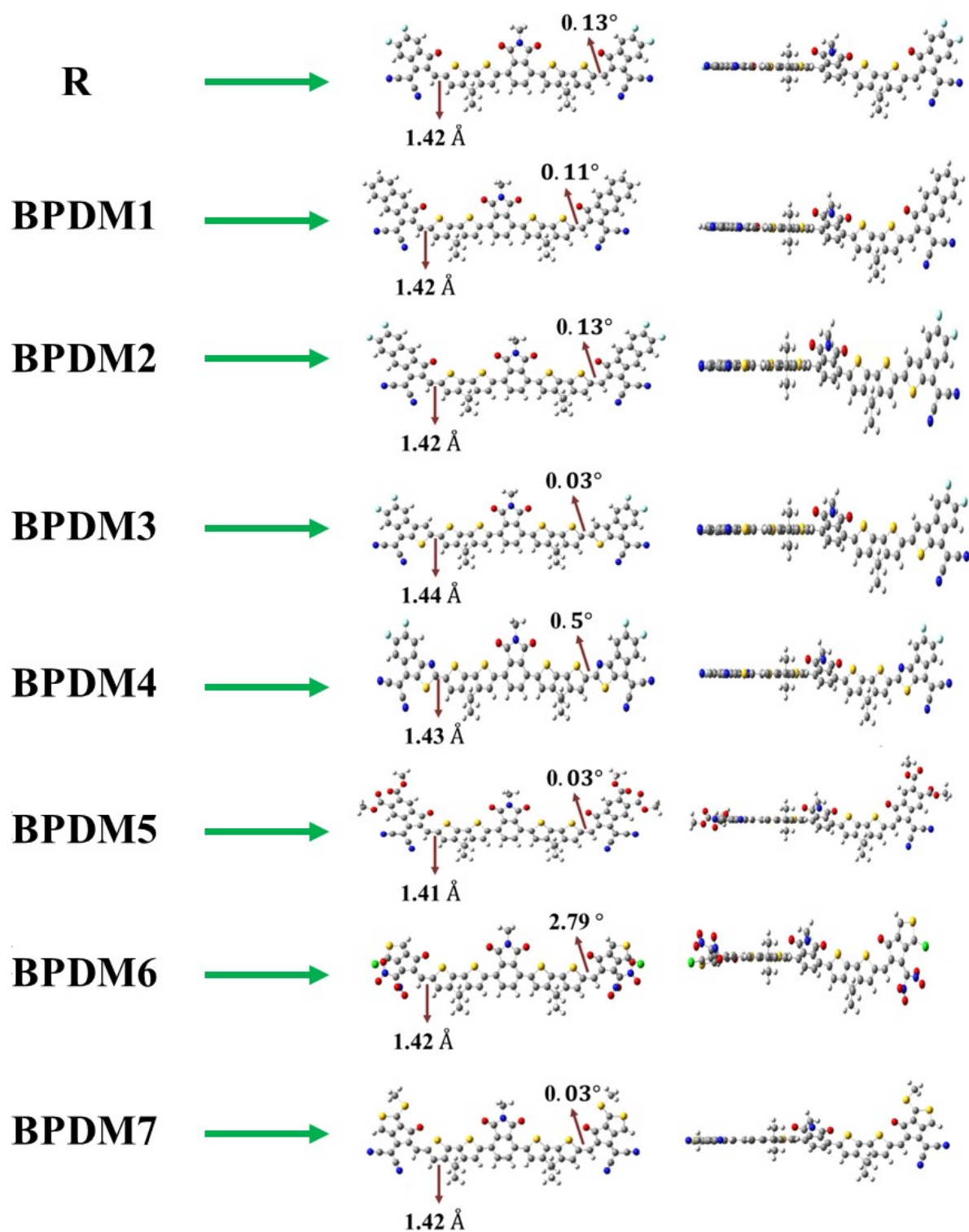


Figure S2 Optimized geometries of R and BPDM1-BPDM7 illustrating interatomic distances and torsional angles

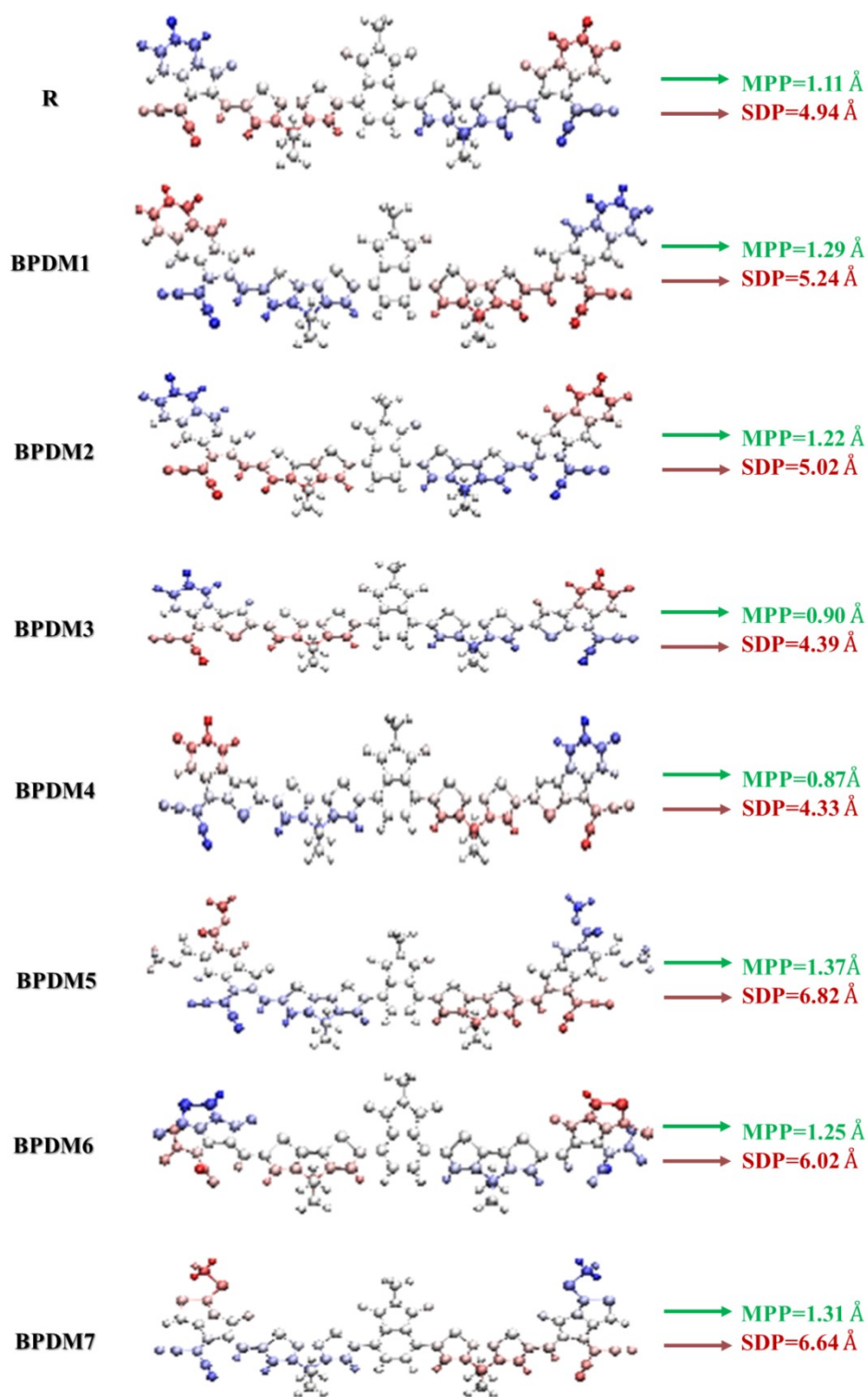


Figure S3 Visualization of molecular planarity in R and BPDM1-BPDM7 based on MPP (Molecular Planarity Parameter) and SDP (Span of Deviation from the Plane) values

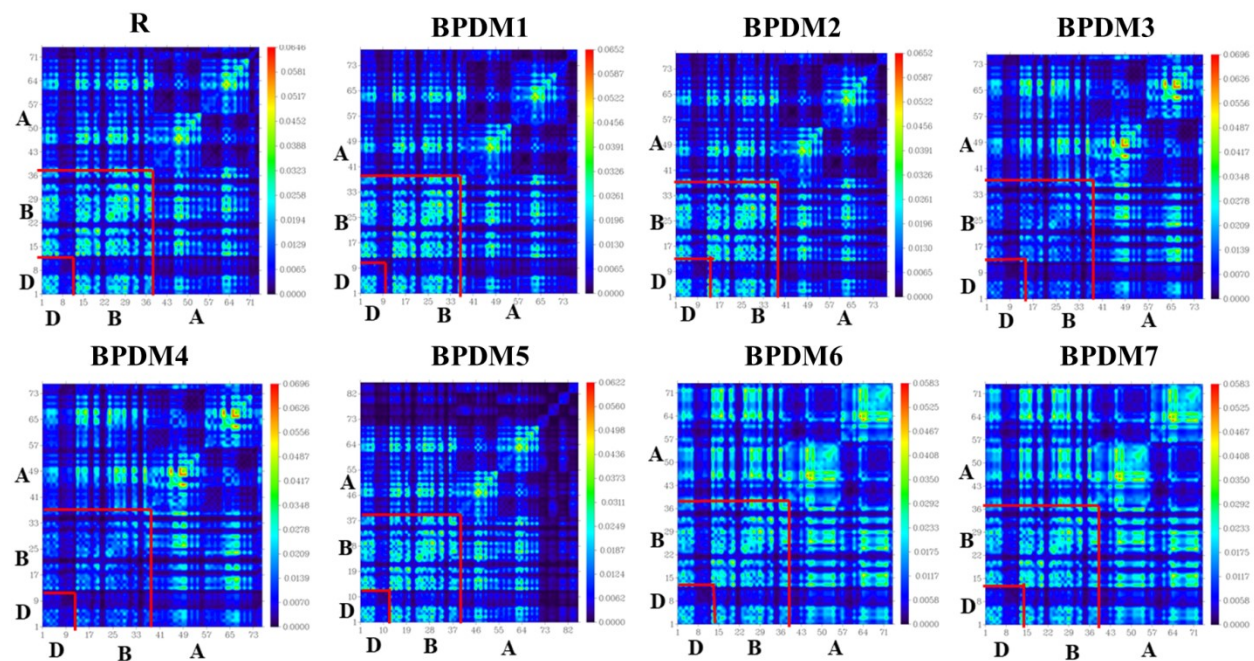


Figure S4 TDM plots of R and designed molecules

Table S1 Calculated absorption profile ( $\lambda_{\max}$ ), light harvesting efficiency (LHE), Excitation energy ( $E_X$ ), Oscillating strength ( $f$ ), and Dipole moments ( $\mu$ ) for R and deigned molecules in gas phase

| Molecules | $\lambda_{\max}$<br>Calculated<br>(nm) | $\lambda_{\max}$<br>Experimental<br>(nm) | $E_X$<br>(eV) | Dipole<br>moment<br>(D) | $f$  | LHE    | $E_b$<br>(eV) |
|-----------|----------------------------------------|------------------------------------------|---------------|-------------------------|------|--------|---------------|
| R         | 629                                    | 652                                      | 1.96          | 1.82                    | 2.78 | 0.9992 | 0.36          |
| BPDM1     | 642                                    | -                                        | 1.93          | 3.13                    | 3.03 | 0.9996 | 0.33          |
| BPDM2     | 649                                    | -                                        | 1.90          | 0.39                    | 3.04 | 0.9996 | 0.34          |
| BPDM3     | 758                                    | -                                        | 1.63          | 1.62                    | 1.82 | 0.9924 | 0.36          |
| BPDM4     | 752                                    | -                                        | 1.64          | 1.89                    | 1.70 | 0.9886 | 0.39          |
| BPDM5     | 639                                    | -                                        | 1.93          | 1.44                    | 2.89 | 0.9991 | 0.38          |
| BPDM6     | 671                                    | -                                        | 1.84          | 1.52                    | 2.55 | 0.9982 | 0.39          |
| BPDM7     | 640                                    | -                                        | 1.93`         | 3.88                    | 2.81 | 0.9993 | 0.34          |

Table S2 Calculated absorption profile ( $\lambda_{\max}$ ), light harvesting efficiency (LHE), Excitation energy ( $E_X$ ), Oscillating strength ( $f$ ), and Dipole moments ( $\mu$ ) for R and deigned molecules in solvent

| Molecules | $\lambda_{\max}$<br>Calculated<br>(nm) | $\lambda_{\max}$<br>Experimental<br>(nm) | $E_X$<br>(eV) | Dipole<br>moment<br>(D) | $f$  | LHE    | $E_b$<br>(eV) |
|-----------|----------------------------------------|------------------------------------------|---------------|-------------------------|------|--------|---------------|
| R         | 670                                    | 652                                      | 1.84          | 3.01                    | 3.07 | 0.9984 | 0.24          |
| BPDM1     | 681                                    | -                                        | 1.82          | 2.85                    | 3.37 | 0.9990 | 0.22          |
| BPDM2     | 689                                    | -                                        | 1.79          | 1.21                    | 3.37 | 0.9991 | 0.22          |
| BPDM3     | 829                                    | -                                        | 1.49          | 2.10                    | 2.11 | 0.9849 | 0.22          |
| BPDM4     | 825                                    | -                                        | 1.50          | 2.83                    | 1.94 | 0.9804 | 0.24          |
| BPDM5     | 686                                    | -                                        | 1.80          | 3.45                    | 3.04 | 0.9987 | 0.24          |
| BPDM6     | 737                                    | -                                        | 1.68          | 2.66                    | 2.73 | 0.9972 | 0.22          |
| BPDM7     | 678                                    | -                                        | 1.82`         | 3.62                    | 3.18 | 0.9985 | 0.24          |

Table S3 Theoretically calculated HOMO,LUMO and energy gap ( $E_g$ ) values of R and designed molecules using the B3LYP/6-31G(d,p) level of theory

| <b>Molecules</b> | <b>LUMO<br/>(eV)</b> | <b>HOMO<br/>(eV)</b> | <b><math>E_g</math><br/>(eV)</b> |
|------------------|----------------------|----------------------|----------------------------------|
| R                | -3.46                | -5.66                | 2.20                             |
| BPDM1            | -3.33                | -5.48                | 2.15                             |
| BPDM2            | -3.45                | -5.58                | 2.13                             |
| BPDM3            | -3.51                | -5.36                | 1.85                             |
| BPDM4            | -3.61                | -5.49                | 1.88                             |
| BPDM5            | -3.48                | -5.66                | 2.18                             |
| BPDM6            | -3.61                | -5.68                | 2.07                             |
| BPDM7            | -3.30                | -5.47                | 2.17                             |

Table S4 Percentage contribution of donors, acceptor and  $\pi$ -bridge in R and designed molecules

| <b>Molecules</b> | <b>Excitation energy<br/>states</b> | <b>Percentage<br/>contribution of<br/>donor core<br/>(%)</b> | <b>Percentage<br/>contribution of<br/>acceptor<br/>(%)</b> | <b>Percentage<br/>contribution of<br/>pi-bridge<br/>(%)</b> |
|------------------|-------------------------------------|--------------------------------------------------------------|------------------------------------------------------------|-------------------------------------------------------------|
| R                | HOMO                                | 10.3                                                         | 26.5                                                       | 63.5                                                        |
|                  | LUMO                                | 9.1                                                          | 52.8                                                       | 38.5                                                        |
| BPDM1            | HOMO                                | 9.9                                                          | 28.1                                                       | 62.5                                                        |
|                  | LUMO                                | 9.2                                                          | 53.1                                                       | 37.7                                                        |
| BPDM2            | HOMO                                | 10.2                                                         | 28.1                                                       | 61.7                                                        |
|                  | LUMO                                | 9.0                                                          | 53.9                                                       | 37.2                                                        |
| BPDM3            | HOMO                                | 9.5                                                          | 28.7                                                       | 61.8                                                        |
|                  | LUMO                                | 3.0                                                          | 81.4                                                       | 15.6                                                        |
| BPDM4            | HOMO                                | 9.9                                                          | 28.4                                                       | 61.7                                                        |
|                  | LUMO                                | 3.0                                                          | 80.7                                                       | 16.2                                                        |
| BPDM5            | HOMO                                | 10.5                                                         | 26.6                                                       | 62.9                                                        |
|                  | LUMO                                | 8.9                                                          | 54.2                                                       | 37.0                                                        |
| BPDM6            | HOMO                                | 10.3                                                         | 27.0                                                       | 62.7                                                        |
|                  | LUMO                                | 6.5                                                          | 63.3                                                       | 30.3                                                        |
| BPDM7            | HOMO                                | 9.7                                                          | 29.5                                                       | 60.8                                                        |
|                  | LUMO                                | 9.6                                                          | 51.9                                                       | 38.5                                                        |

Table S5 Ionization potential (IP), electron affinity (EA), chemical hardness ( $\eta$ ), and global softness (S) for the R and designed molecules (BPDM1–BPDM7) calculated at the B3LYP/6-31G(d,p) level

| <b>Molecules</b> | <b>IP<br/>(eV)</b> | <b>EA<br/>(eV)</b> | <b><math>\eta</math><br/>(eV)</b> | <b>S<br/>(eV)</b> |
|------------------|--------------------|--------------------|-----------------------------------|-------------------|
| R                | 5.66               | 3.46               | 1.10                              | 0.450             |
| BPDM1            | 5.48               | 3.33               | 1.075                             | 0.465             |
| BPDM2            | 5.58               | 3.45               | 1.065                             | 0.469             |
| BPDM3            | 5.36               | 3.51               | 0.925                             | 0.541             |
| BPDM4            | 5.49               | 3.61               | 0.94                              | 0.532             |
| BPDM5            | 5.66               | 3.48               | 1.09                              | 0.459             |
| BPDM6            | 5.68               | 3.61               | 1.035                             | 0.483             |
| BPDM7            | 5.47               | 3.30               | 1.085                             | 0.461             |

Table S6 The reorganization energies of holes ( $\lambda_h$ ) and electrons ( $\lambda_e$ ) for the R and designed molecules (BPDM1–BPDM7)

| <b>Molecules</b> | <b><math>\lambda_e</math><br/>(eV)</b> | <b><math>\lambda_h</math><br/>(eV)</b> |
|------------------|----------------------------------------|----------------------------------------|
| R                | 0.007425                               | 0.007731                               |
| BPDM1            | 0.006702                               | 0.103544                               |
| BPDM2            | 0.016091                               | 0.015454                               |
| BPDM3            | 0.004301                               | 0.006650                               |
| BDPM4            | 0.004053                               | 0.006267                               |
| BDPM5            | 0.007098                               | 0.458086                               |
| BPDM6            | 0.105526                               | 0.463267                               |
| BPDM7            | 0.007235                               | 0.228532                               |

Table S7 Calculated fill factor (FF), open-circuit voltage ( $V_{oc}$ ), and PCE for the R and the designed molecules (BPDM1–BPDM7)

| <b>Molecules</b> | <b>Fill factor<br/>(FF)</b> | <b><math>V_{oc}</math></b> | <b>PCE</b> |
|------------------|-----------------------------|----------------------------|------------|
| R                | 92.43%                      | 1.76                       | 25.85%     |
| BPDM1            | 92.82%                      | 1.88                       | 27.83%     |
| BPDM2            | 92.43%                      | 1.76                       | 25.85%     |
| BPDM3            | 92.22%                      | 1.70                       | 24.75%     |
| BPDM4            | 91.87%                      | 1.61                       | 23.29%     |
| BPMD5            | 92.36%                      | 1.74                       | 25.54%     |
| BPDM6            | 91.87%                      | 1.61                       | 23.29%     |
| BPDM7            | 92.96%                      | 1.93                       | 28.72%     |