

**JOUL, Volume 1**

**Supplemental Information**

**Design Principles and Top Non-Fullerene**

**Acceptor Candidates**

**for Organic Photovoltaics**

**Steven A. Lopez, Benjamin Sanchez-Lengeling, Julio de Goes Soares, and Alán Aspuru-Guzik**

## SUPPORTING INFORMATION

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#### Supplementary Tables.

**Table S1.** Summary table of references values utilized in the calibration set.

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#### Supplementary Datafiles

**Datafile S1. 'Molecules.csv'** – Comma separated value text file containing information on the entire dataset of acceptor molecules, 51280 in total. Data included is: InChI key, smiles string, HOMO energy, LUMO energy, Open Circuit Voltage (Voc), Short circuit current (Jsc), Fill Factor (FF) and power conversion efficiency (PCE). GP Calibrated values (`_calib`) and their uncertainty (`_calib_std`) are included for HOMO, LUMO and PCE values. The PCE calibrated values are `PCE_calib_plus`.

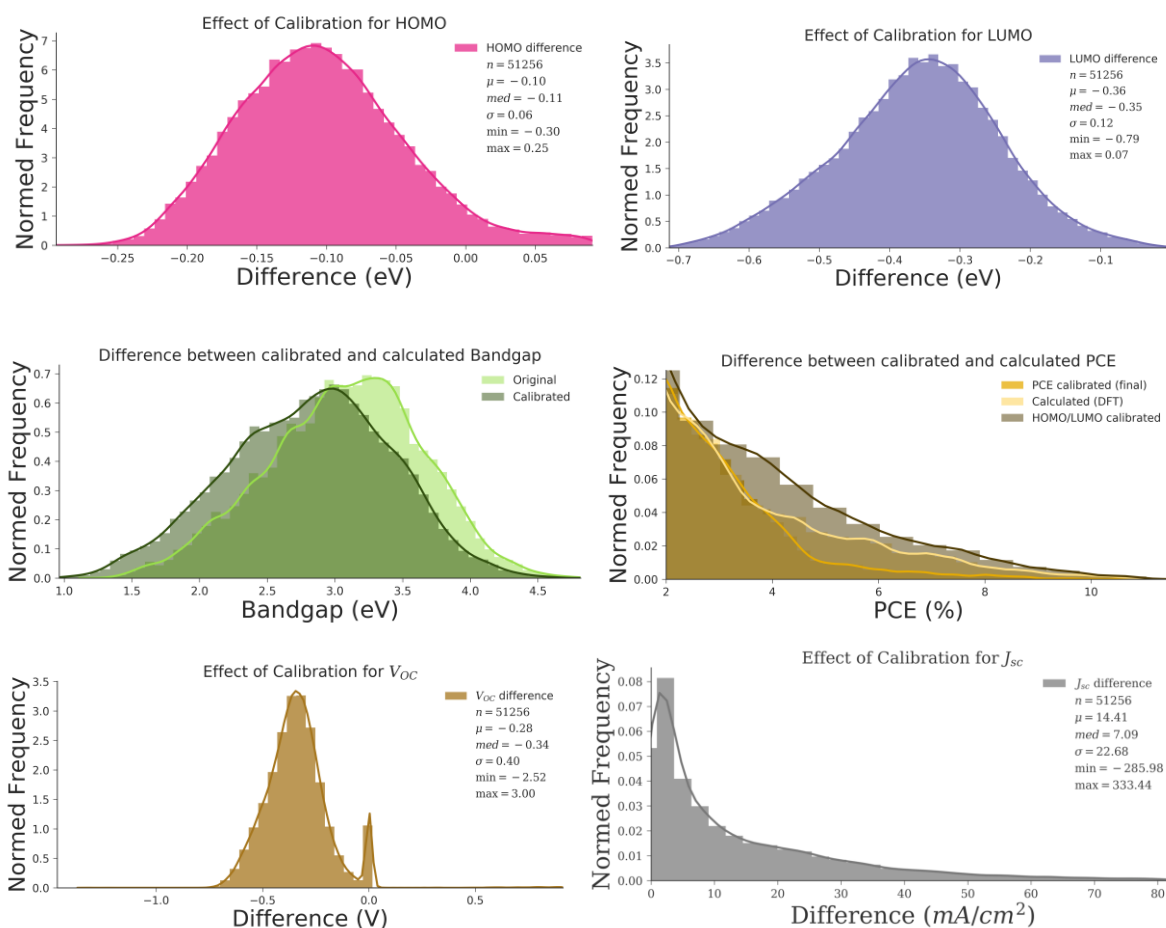
**Datafile S2. 'Fragments.csv'** – Comma separated value text file containing information on the 112 fragments used to build the library of molecules. Data included is: Group classification for fragment (group), label, smiles string, representative PCE statistics (mean, SD, median, etc.) in python dictionary format for molecules that contain that fragment, PCE mean and SD, count of molecules containing a fragment for three categories: global, PCE < 8 and PCE > 8 (`global_pop` and `global_pop_percent`, etc.) and Z-score for that particular fragment.

**Datafile S3. 'Bits.csv'** – Comma separated value text file containing information on fingerprint analysis utilizing 8192 bits with a connectivity radius of 4. Data included is: Bit number (Bit), count (`_count`) and percentage of molecules ( `Pop %`) with that particular activated bit percentage for two categories: entire data set (global) and top candidates (top), representative PCE statistics (mean, SD, median, etc.) in python dictionary format for molecules that have a activated bit, PCE mean and SD, mean and std of radius of the fingerprint for all molecules and Z-score for a particular activated bit.

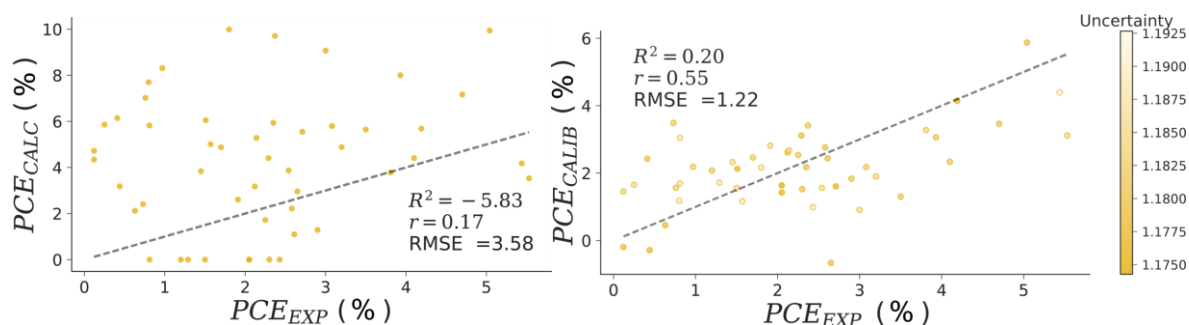
**Datafile S4. 'HOMO-LUMO\_calibration.csv'** – Comma separated value text file containing information on molecules used for HOMO-LUMO GP calibration. Contains experimental values and values calculated with TDDFT.

**Datafile S5. 'PCE\_calibration.csv'** – Comma separated value text file containing information on molecules used for PCE GP calibration. Contains experimental values, values calculated with TDDFT, scharber model values and values obtained after calibration.

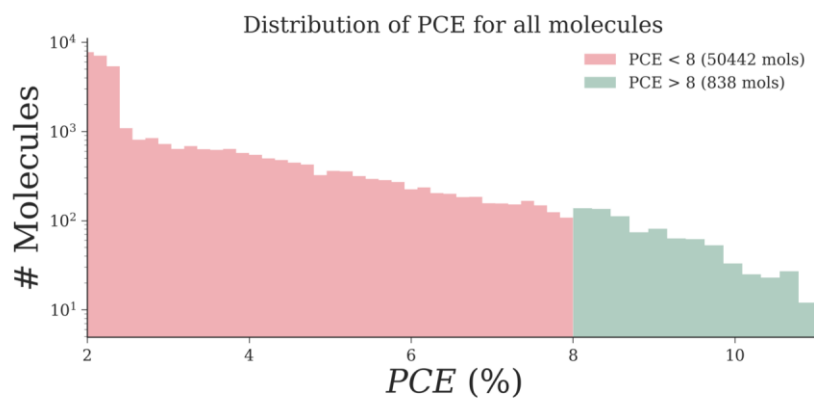
**Figure S1. Histograms of the distributions of difference between calibrated and calculated values.** *Related to Figure 2.* For some properties we also show the statistical measures for these distributions.



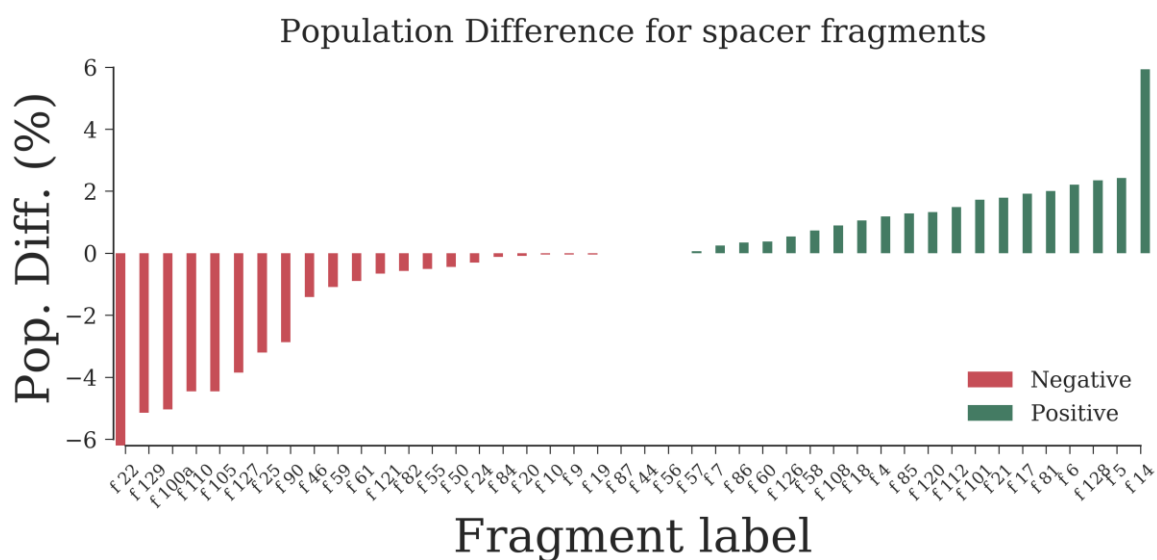
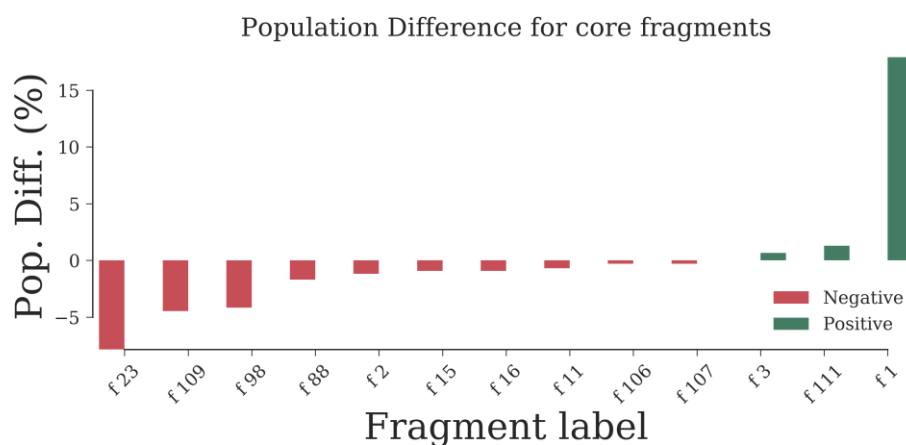
**Figure S2. GP calibration of PCE values.** Related to section *Machine learning: Gaussian Processes regression*. Experimental (EXP), calculated (DFT/Scharber model) and GP calibrated LOO-CV predictions (CALIB) are compared. Uncertainty increases with lightly shaded data points. Dotted line represents the ideal fit.

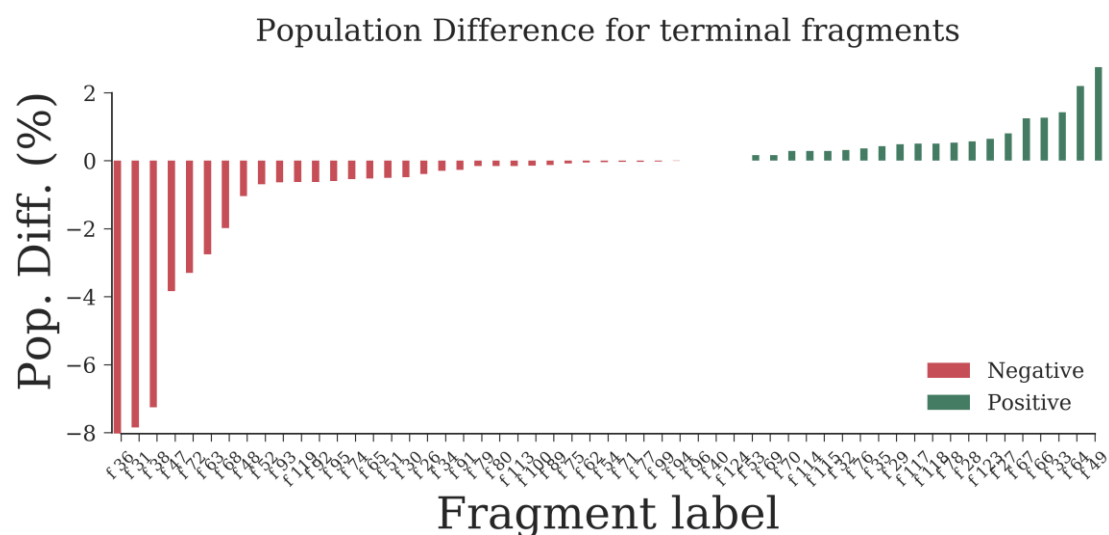


**Figure S3. Histogram of distribution of calibrated PCE values.** Related to section *Scharber Model: PCE calculations*. Top (green) and non-top (red) molecules are color coded, y-axis is using a log scale. Statistical measures of the global distribution are included.

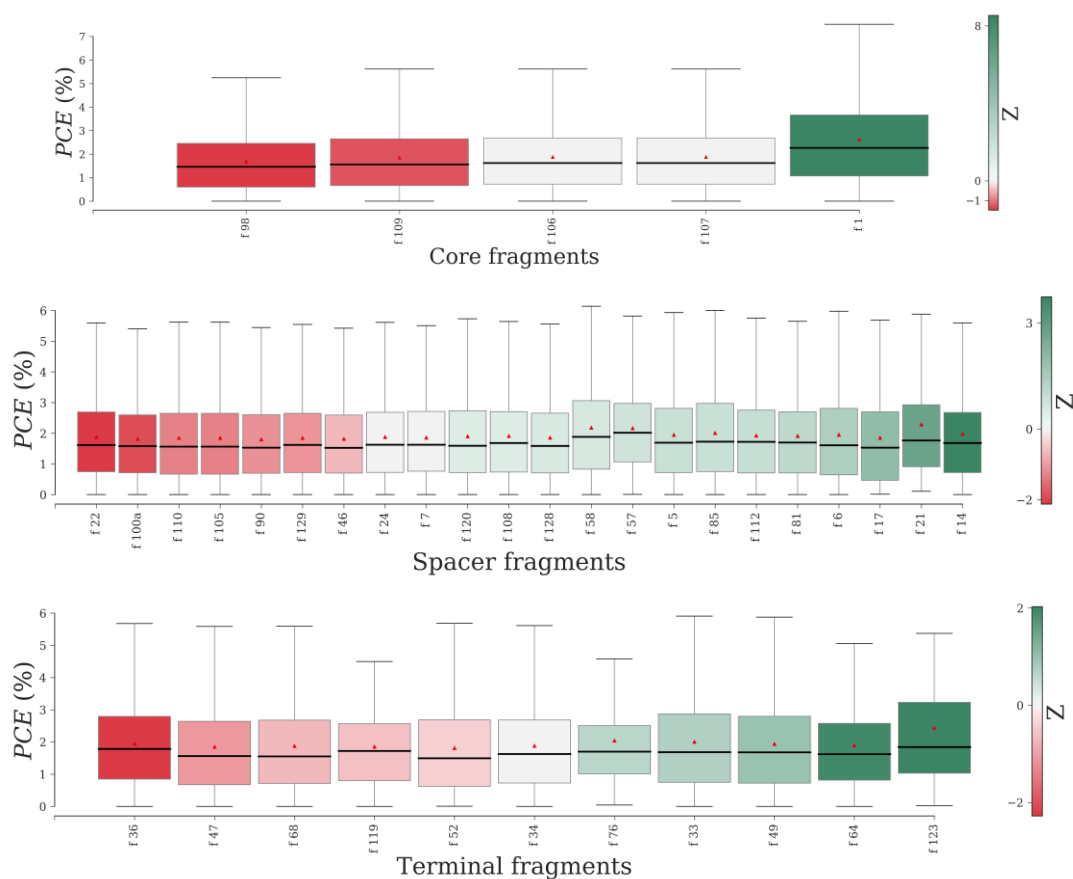


**Figure S4. Barplots of population difference between top and non-top molecules, grouped by fragment group.** Related to section *Scharber Model: PCE calculations*. Color-coded based on positive or negative representation. Positive differences indicate a fragment is more present in top candidate than non-top candidates.

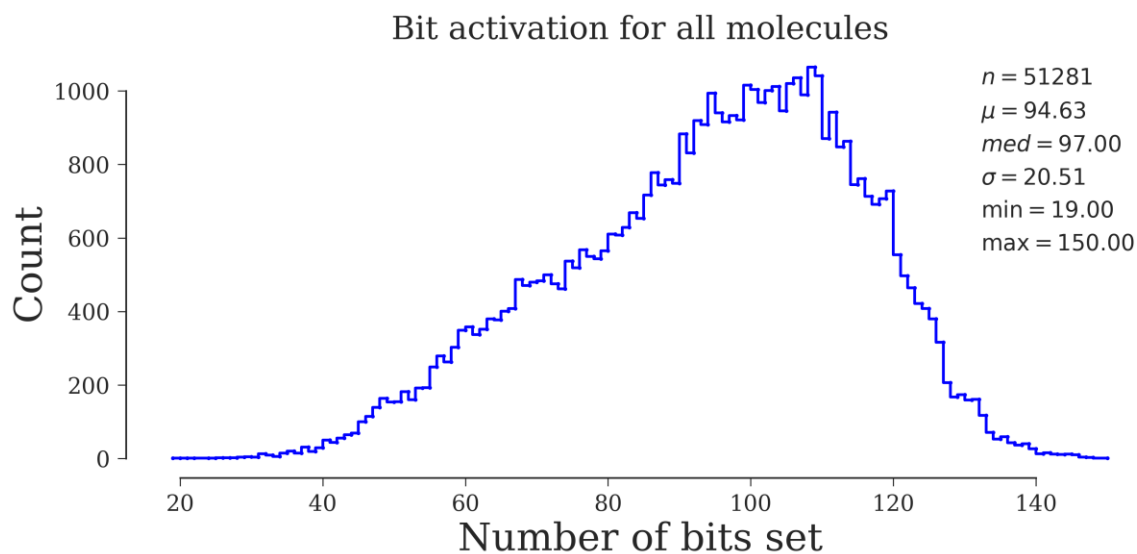




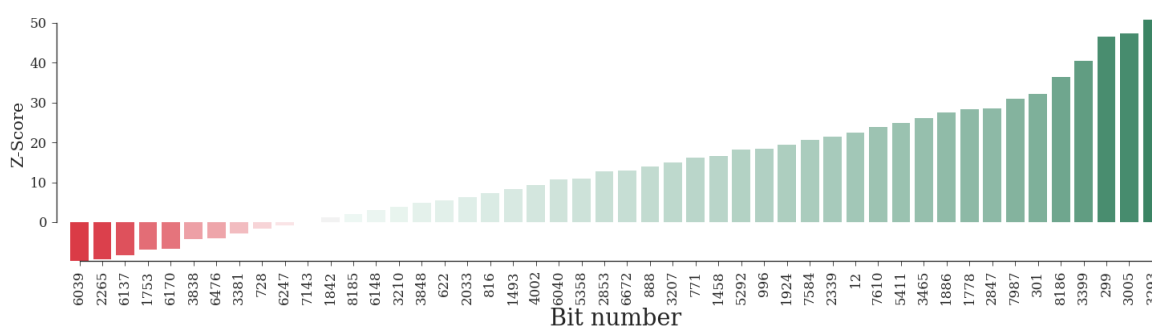
**Figure S5. Boxplot of PCE values for groups of molecules containing a particular fragment.** Related to section *Scharber Model: PCE calculations*. Colored based on Z-score and grouped by fragment group. Mean values are indicated as black lines, median as red dots. Each box represents the 25% to 75% quartile of the values with min and max values as whiskers.



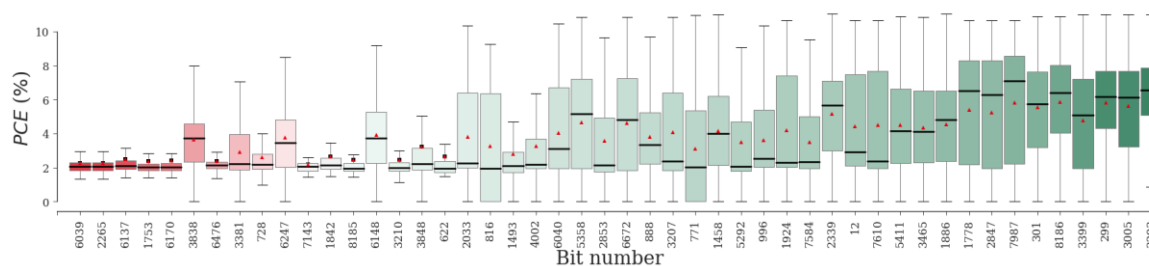
**Figure S6. Distribution of counts of activated bits in all molecules.** Related to section *Fingerprint statistics*.



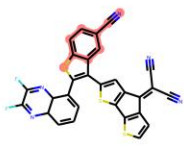
**Figure S7. Z-scores for a representative sample of 48 bits.** *Related to section Fingerprint statistics.* Representative sample was created using a histogram on the Z-scores and picking random bits from each bin. Binning was using the 'doane' estimator which works better with non-normal datasets and takes in to account outliers and data variability.



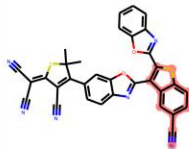
**Figure S8. Boxplots for PCE values of molecules with activated bits.** *Related to section Fingerprint statistics.* Using a representative sample of 48 bits. Boxes are colored based on their z-scores.



**Figure S9. Grid of example molecules of the representative sample of 48 bits.** *Related to section Fingerprint statistics.* Structure encoded in bit is highlighted in red.



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Bit 2265 (-9.33)



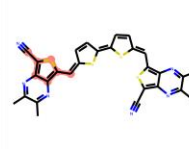
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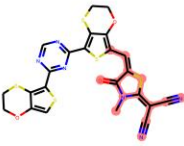
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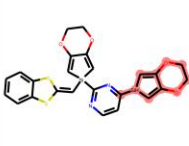
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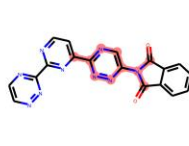
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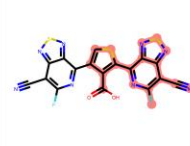
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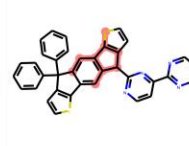
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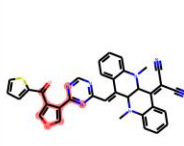
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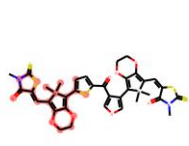
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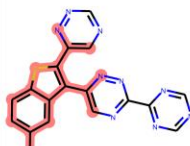
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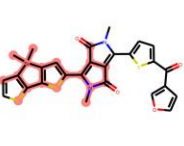
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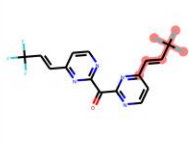
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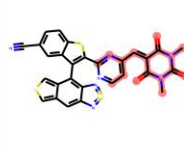
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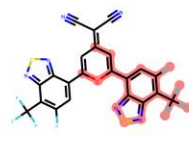
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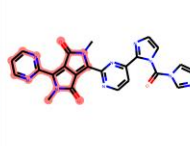
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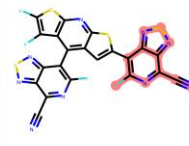
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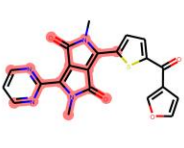
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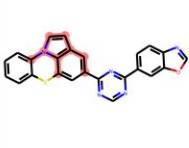
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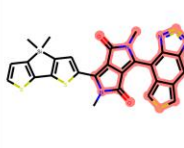
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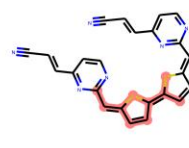
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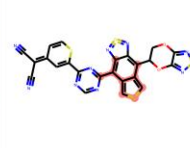
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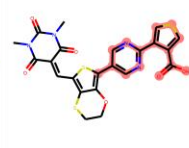
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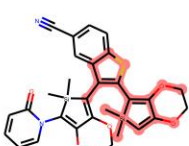
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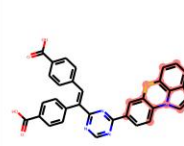
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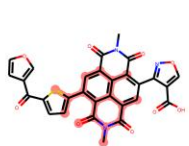
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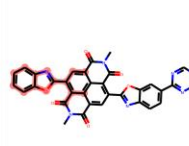
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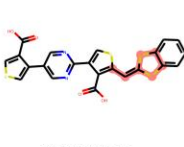
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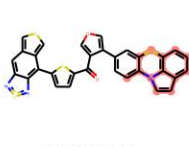
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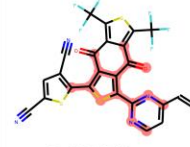
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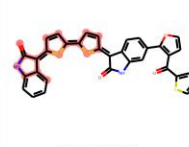
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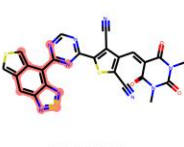
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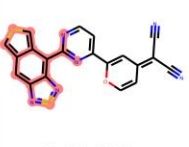
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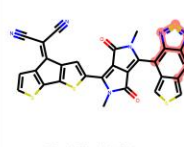
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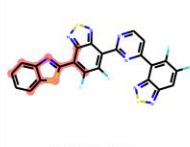
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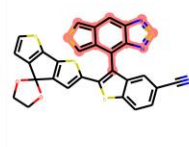
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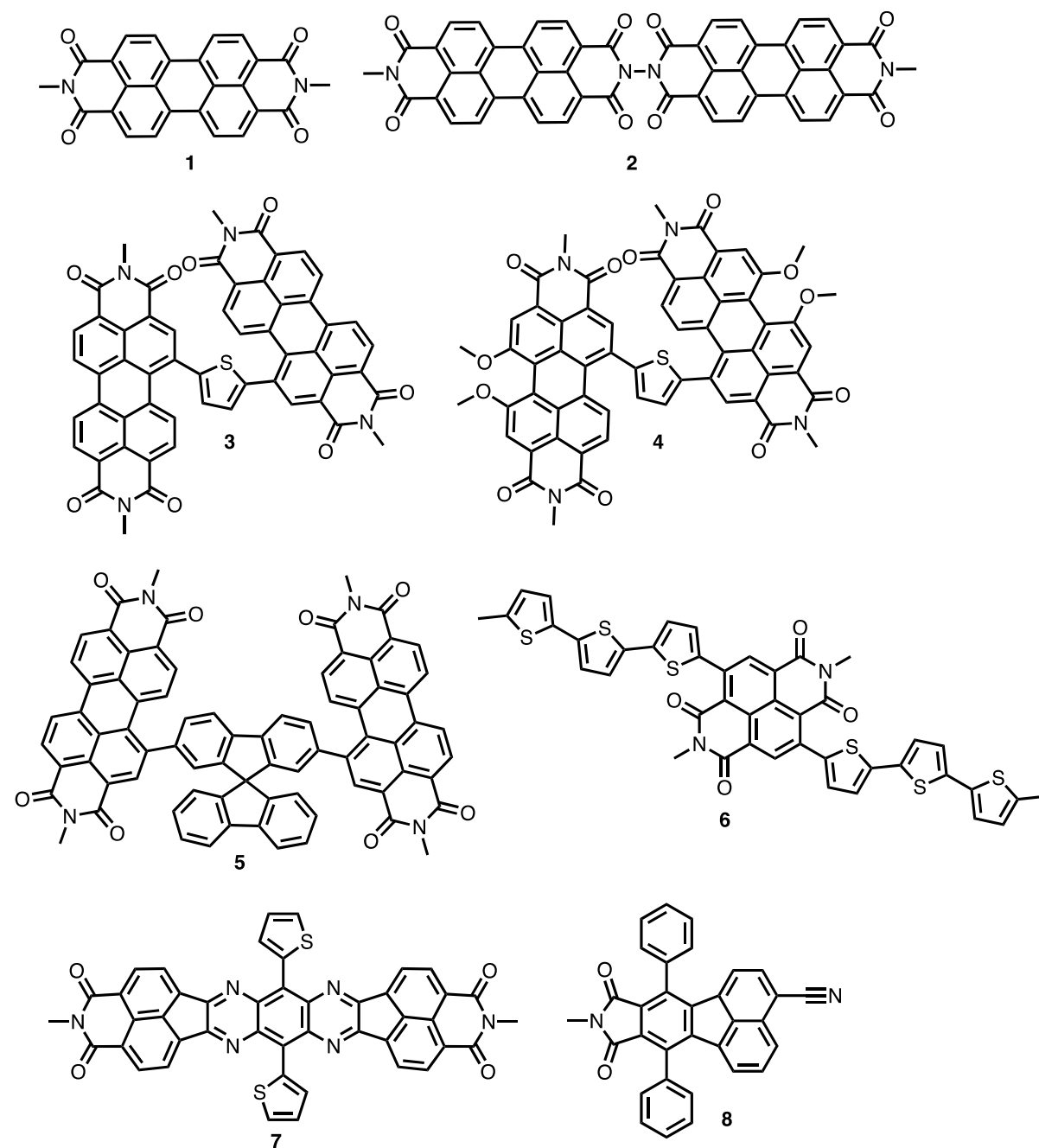
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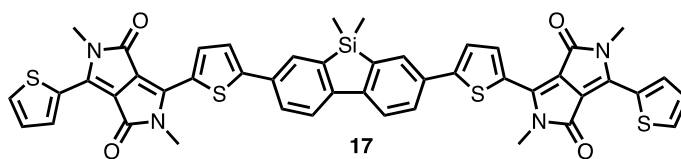
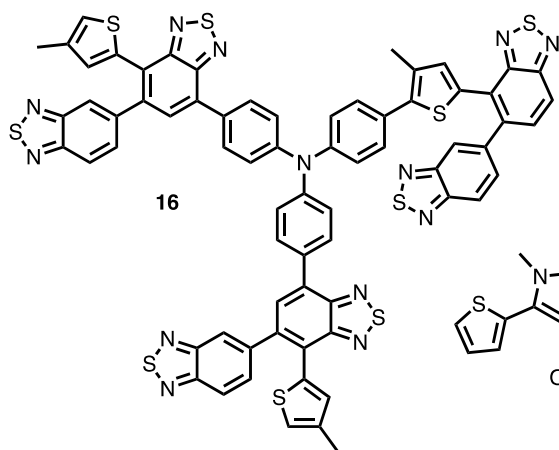
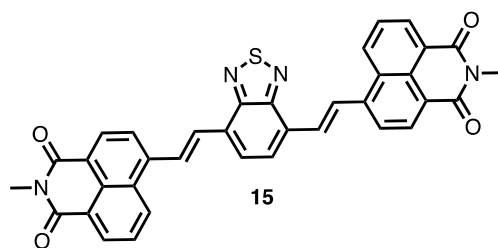
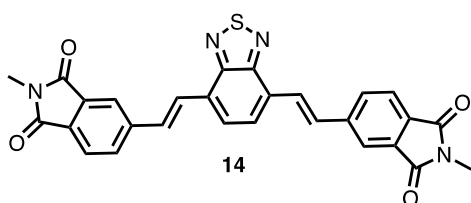
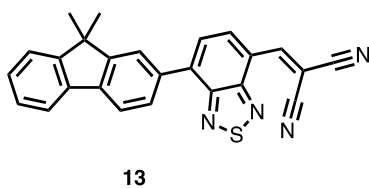
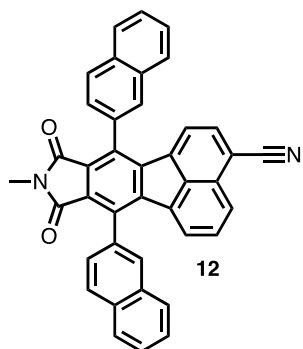
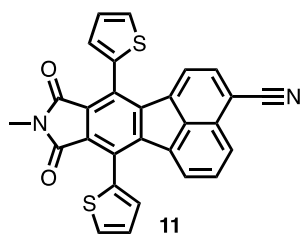
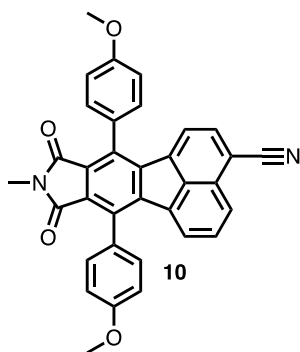
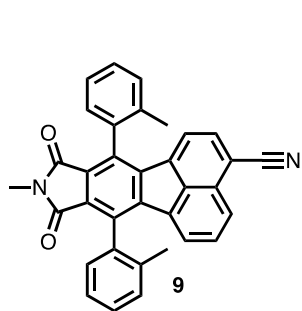
**Table S1. Summary table of references values utilized in the calibration set.** *Related to section Molecular generation and DFT calculations.* Experimental and computed frontier molecular orbitals used for training in the GP calibration. Includes source for reference.

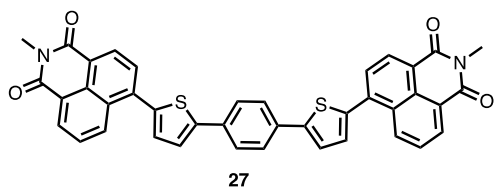
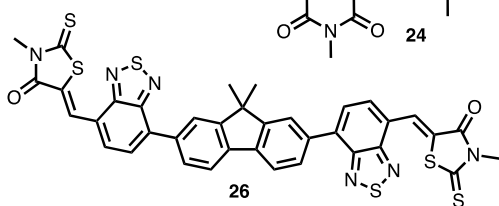
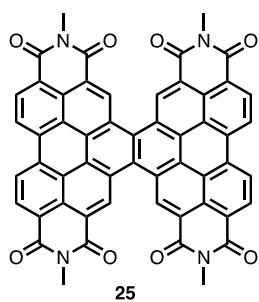
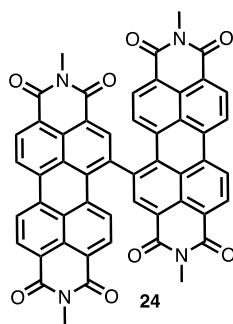
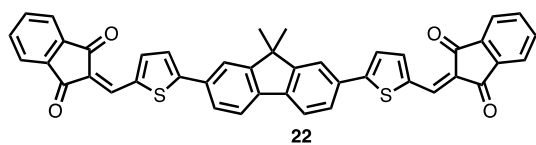
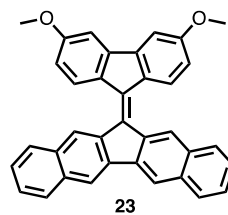
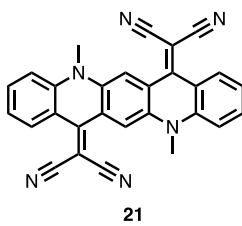
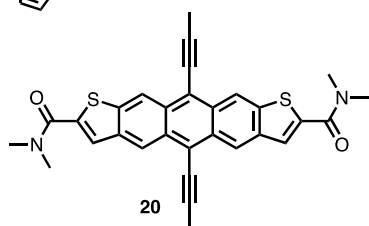
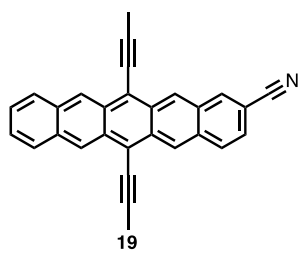
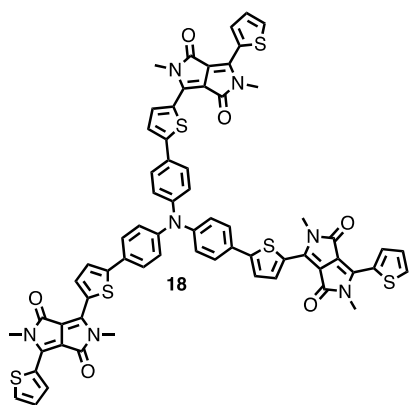
SI Code	Reference	Experimental		Calculated		Calibrated	
		HOMO (eV)	LUMO (eV)	HOMO (eV)	LUMO (eV)	HOMO (eV)	LUMO (eV)
1	3	-5.48	-3.84	-5.50	-3.05	-5.57	-3.76
2	4	-5.71	-3.71	-5.65	-3.06	-5.74	-3.64
3	5	-5.94	-3.84	-5.71	-3.30	-5.85	-3.86
4	6	-5.81	-3.61	-5.76	-3.07	-5.83	-3.59
5	7	-5.26	-3.52	-5.27	-3.22	-5.37	-3.57
6	8	-5.90	-4.09	-5.71	-3.63	-5.84	-4.11
7	9	-6.12	-3.43	-6.24	-3.18	-6.03	-3.394
8	10	-5.57	-3.95	-5.32	-3.61	-5.44	-3.90
10	11	-6.02	-4.01	-6.07	-3.61	-5.95	-3.92
11	12	-5.65	-3.84	-6.04	-3.86	-5.8	-4.03
14	13	-6.04	-3.77	-6.25	-3.91	-5.98	-3.93
19	14	-5.71	-3.71	-5.35	-3.55	-5.65	-3.76
20	15	-5.90	-4.10	-6.00	-3.35	-5.95	-4.01
21	9	-6.27	-3.44	-6.17	-3.21	-6.28	-3.53
22	16	-5.80	-3.60	-6.11	-3.59	-5.97	-3.70
23	17	-6.00	-3.60	-6.09	-3.34	-5.99	-3.53
24	16	-5.80	-3.80	-5.59	-3.67	-5.55	-3.77
27	18	-5.48	-3.83	-5.66	-3.61	-5.59	-3.89
28	6	-5.81	-3.61	-6.53	-3.67	-6.12	-3.77
30	19	-5.18	-3.24	-5.03	-2.63	-5.19	-3.17
31	20	-5.69	-3.71	-5.41	-3.08	-5.61	-3.65
32	21	-5.53	-4.05	-5.89	-3.45	-5.89	-3.98
33	22	-5.88	-3.80	-6.05	-3.58	-5.97	-3.79
34	22	-5.46	-3.78	-5.25	-3.51	-5.38	-3.94
35	23	-6.74	-4.35	-6.72	-3.93	-6.60	-4.21
48	25	-6.28	-4.23	-6.17	-3.18	-6.27	-4.07
50	14	-6.40	-4.10	-6.17	-3.18	-6.38	-3.95

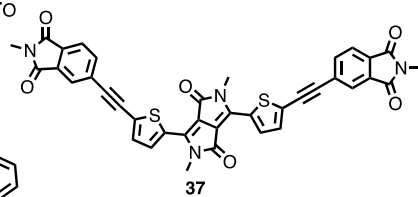
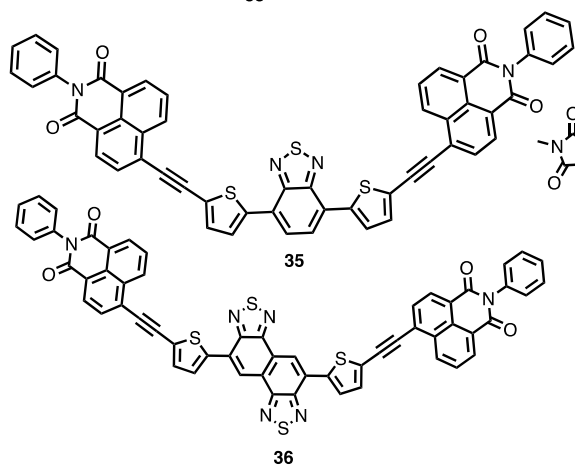
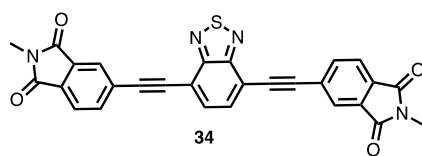
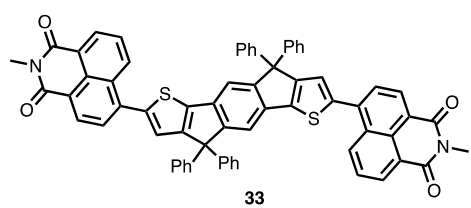
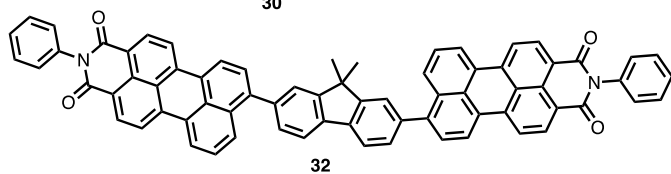
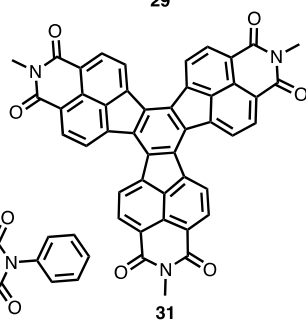
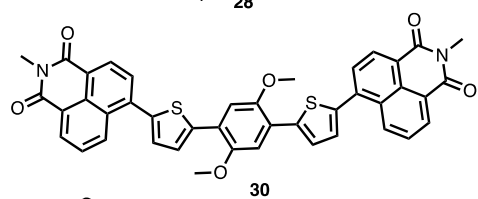
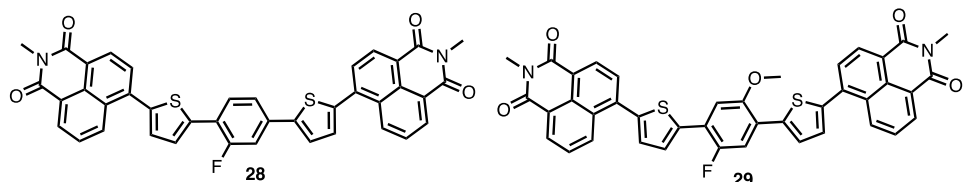


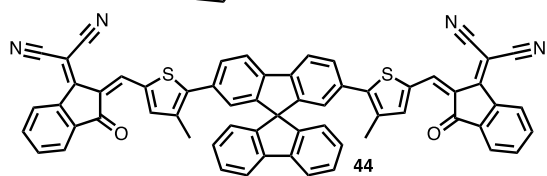
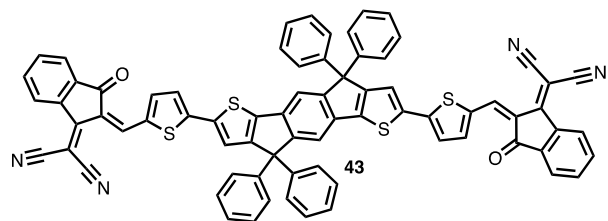
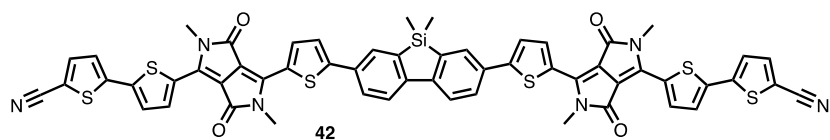
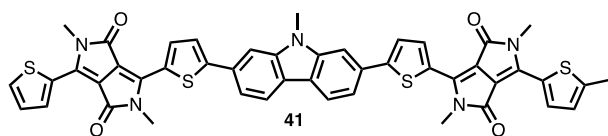
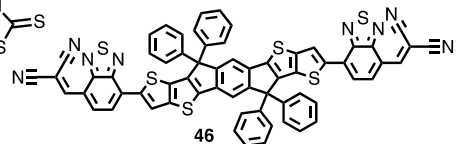
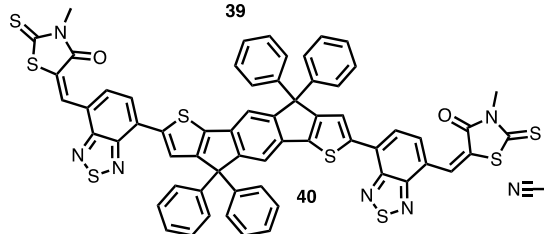
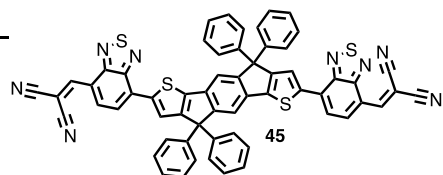
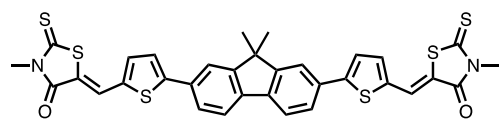
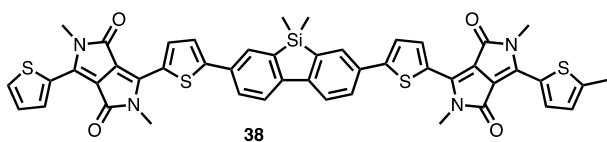
**Table S2. Structures of *n*-type materials in table S1.** Related to section *Molecular generation and DFT calculations*.

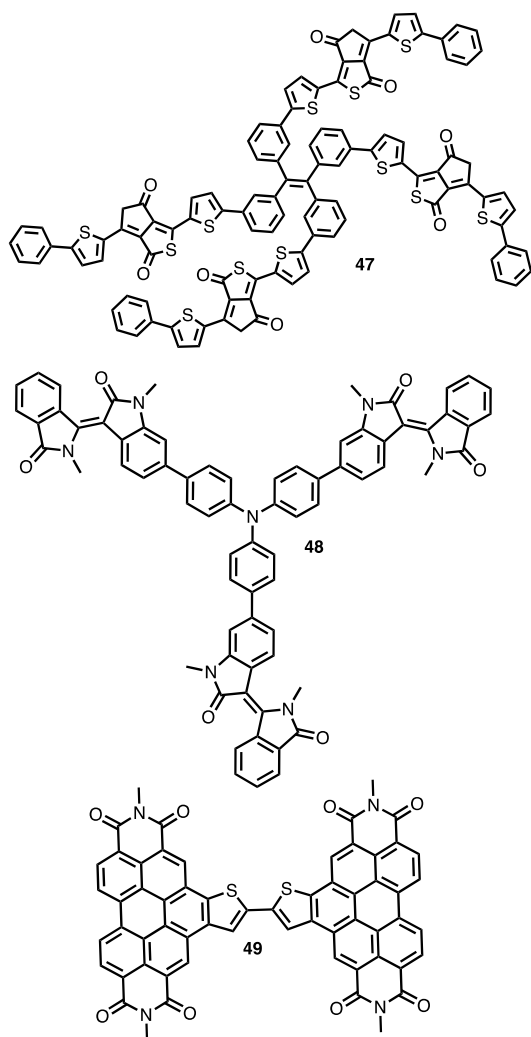












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