

# Supporting Information

## Energy Differences as Descriptors for the Correlation between $J_{SC}$ and $V_{OC}$ in Nonfullerene Organic Photovoltaics

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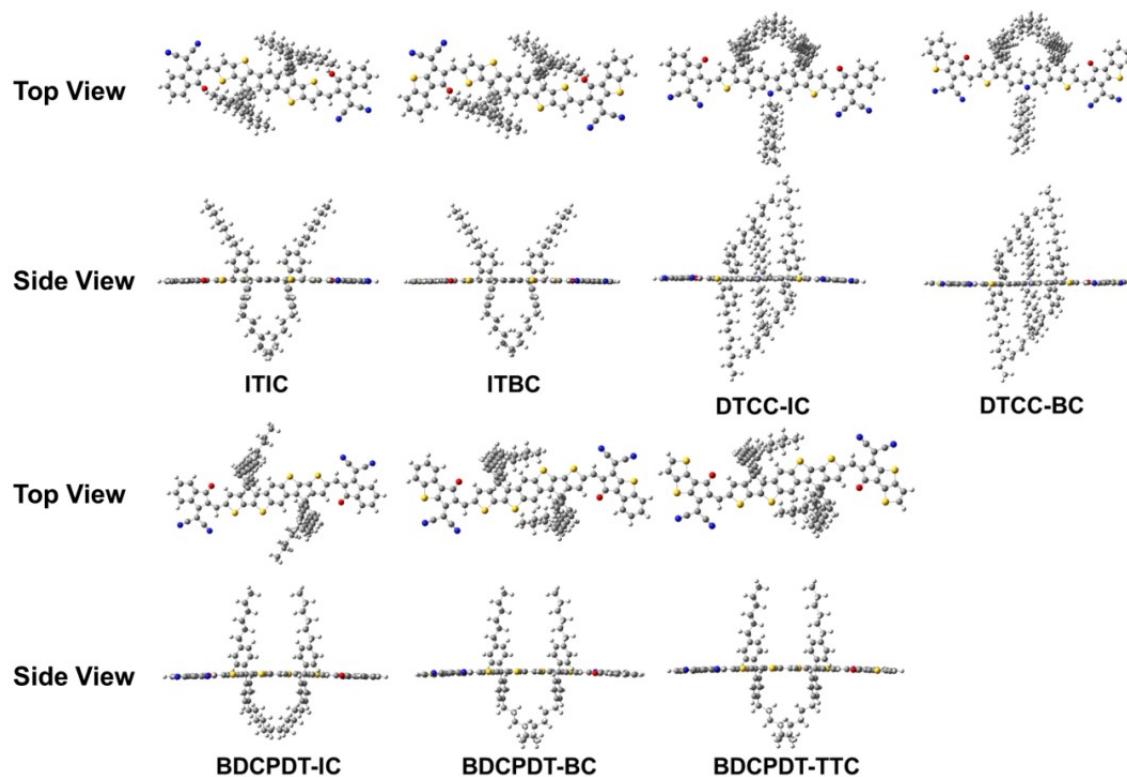
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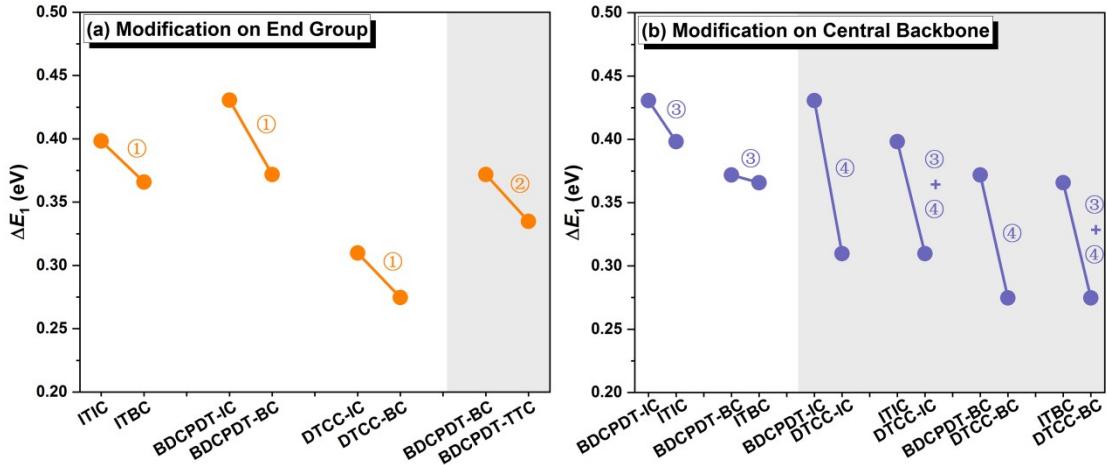
## I. Calculation Results

$$\begin{aligned}
 \Delta E_g &= (E_g)_2 - (E_g)_1 \\
 &= (E_{\text{LUMO}}^{\text{A}_2} - E_{\text{HOMO}}^{\text{A}_2}) - (E_{\text{LUMO}}^{\text{A}_1} - E_{\text{HOMO}}^{\text{A}_1}) \\
 &= (E_{\text{LUMO}}^{\text{A}_2} - E_{\text{LUMO}}^{\text{A}_1}) - (E_{\text{HOMO}}^{\text{A}_2} - E_{\text{HOMO}}^{\text{A}_1}) \\
 &= \Delta E_{\text{LUMO}} - \Delta E_{\text{HOMO}}
 \end{aligned} \tag{S1}$$

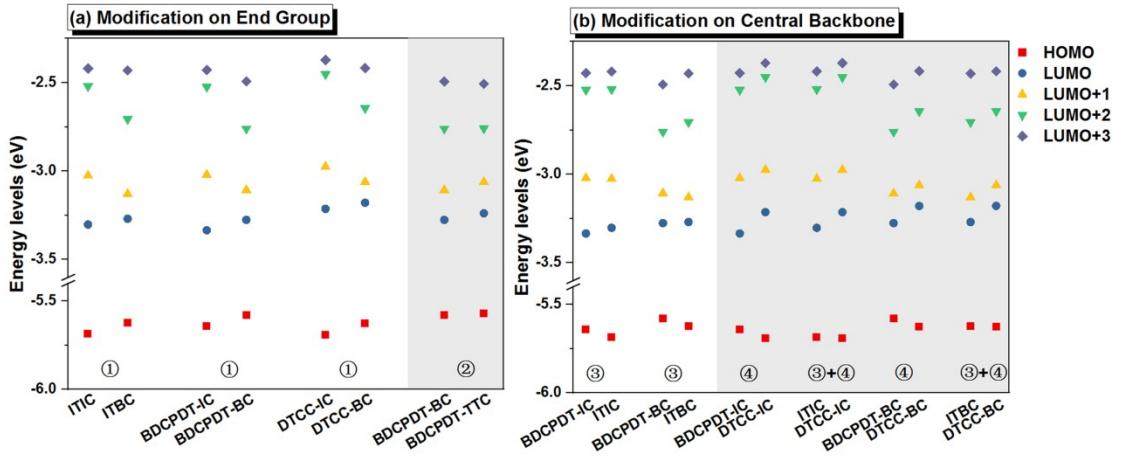
$$\begin{aligned}
 \Delta(\Delta E_1) &= (\Delta E_1)_2 - (\Delta E_1)_1 \\
 &= (E_{\text{LUMO}}^{\text{D}} - E_{\text{LUMO}}^{\text{A}_2}) - (E_{\text{LUMO}}^{\text{D}} - E_{\text{LUMO}}^{\text{A}_1}) \\
 &= E_{\text{LUMO}}^{\text{A}_1} - E_{\text{LUMO}}^{\text{A}_2} \\
 &= -\Delta E_{\text{LUMO}}
 \end{aligned}$$



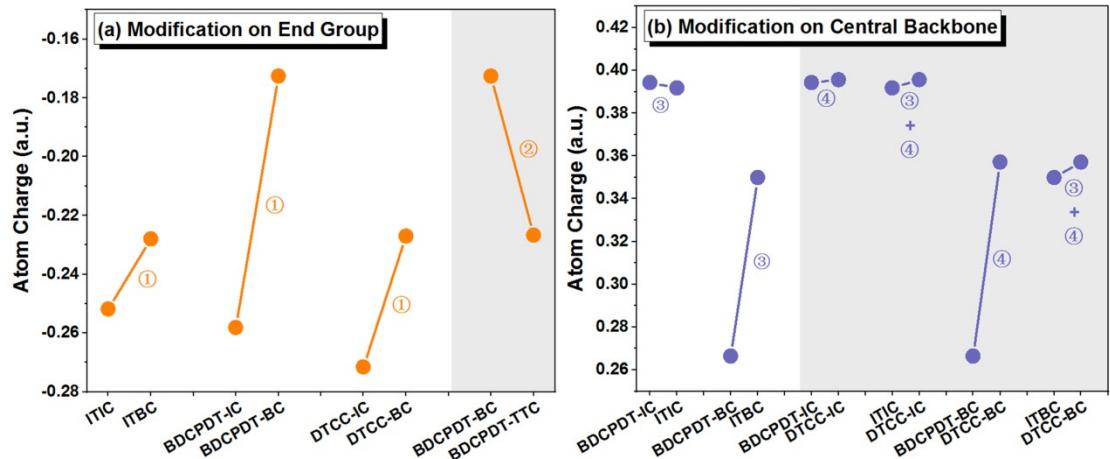
**Figure S1.** The most stable monomeric conformations of ITIC-series acceptors are optimized adopted density functional theory.



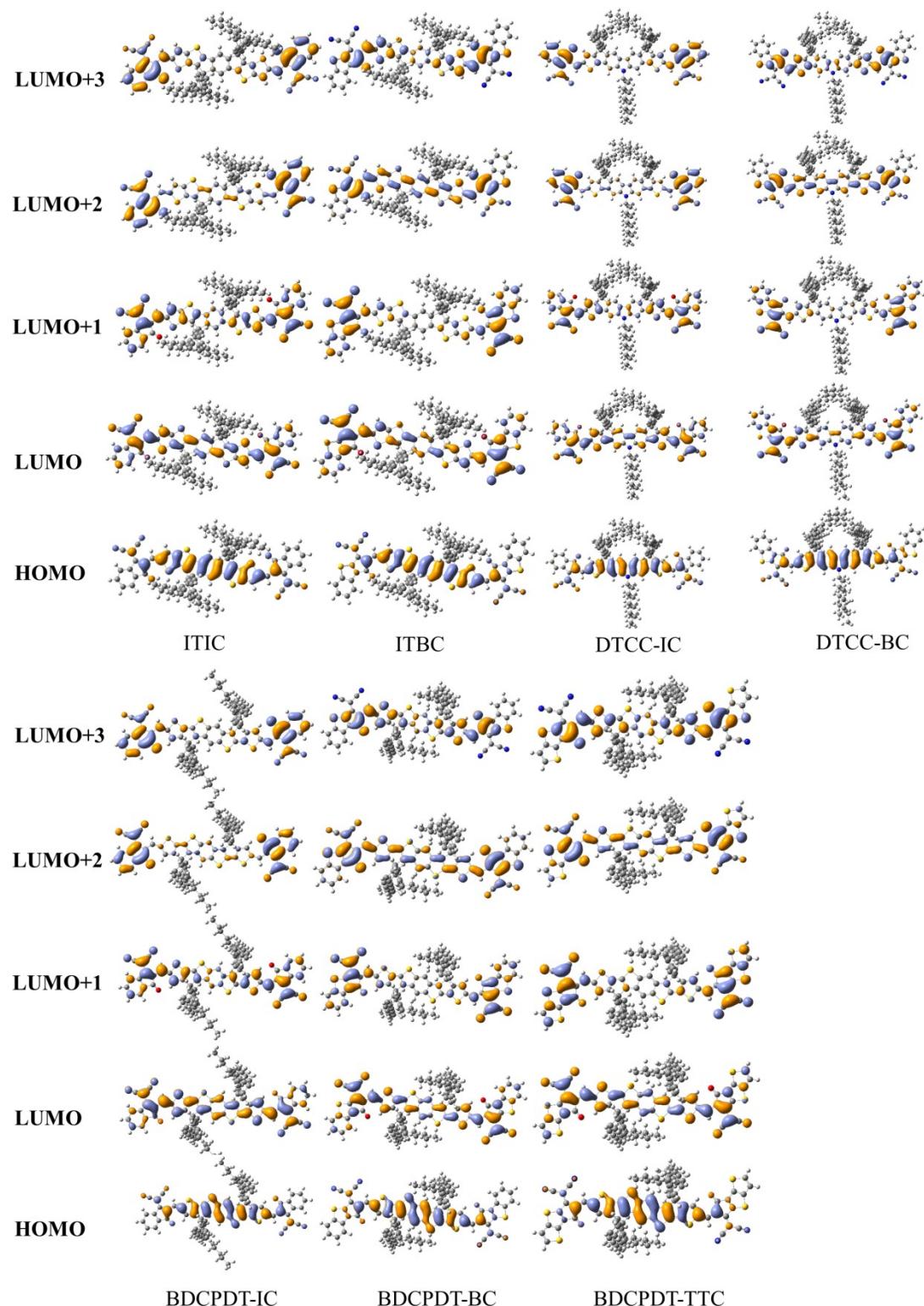
**Figure S2.** Calculated the energy driving force ( $\Delta E_1$ , eV) of 10 groups' devices with modification on (a) end group and (b) central backbone of each group, the shadow filling part represents negative correlation.



**Figure S3.** Calculated FMO energy levels (eV) of the optimized acceptors (a) modification on end group and (b) modification on central backbone at PBE0/6-31G(d) levels. The white area indicates positively correlated devices and the dark grey shaded area indicates negatively correlated devices. HOMO and LUMO, while LUMO+1 means the adjacent orbital above it and LUMO+2, LUMO+3.



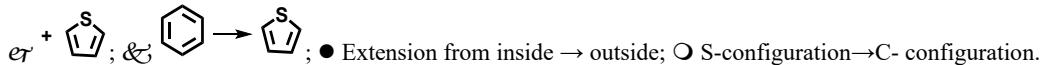
**Figure S4.** Calculated atomic charges (a.u.) of fragment between NFAs for modification on (a) end groups and (b) central backbones adopted atomic dipole moment corrected Hirshfeld population (ADCH) method at the PBE0/6-31G(d) level. When the atomic charge is positive, it indicates that the group has electron-donating property, it indicates that the electron-donating ability increases following the atomic charge increases. While the atomic charge is negative, the group has electron-withdrawing property, it means the withdrawing electron decreases with the increase of the atomic charge.



**Figure S5.** The FMOs of ITIC, ITBC, BDCPDT-IC, BDCPCT-BC, BDCPDT-TTC, DTCC-IC, and DTCC-BC at the PBE0/6-31G(d) level.

**Table S1.** Fragment modification of molecules of 10 groups with the positive and negative correlation. All acceptor were blended with same donor PBDB-T. Positive represents positive correlation between  $J_{SC}$  and  $V_{OC}$ , Negative represents negative correlation between  $J_{SC}$  and  $V_{OC}$ .

Acceptor	Modification	Location	Correlatio n	Refs.
ITIC→ITBC	$\text{er}$	end group	Positive	
BDCPDT-IC→BDCPDT-BC	$\text{er}$	end group	Positive	
DTCC-IC→DTCC-BC	$\text{er}$	end group	Positive	
BDCPDT-BC→BDCPDT-TTC	$\&$	end group	Negative	
BDCPDT-IC→DTCC-IC	○	central backbone (D')	Negative	[1,2]
ITIC→DTCC-IC	●+○	central backbone (D+D')	Negative	
BDCPDT-BC→DTCC-BC	○	central backbone (D')	Negative	
ITBC→DTCC-BC	●+○	central backbone (D+D')	Negative	
BDCPDT-IC→ITIC	●	central backbone	Positive	
BDCPDT-BC→ITBC	●	central backbone	Positive	



**Table S2.** Experiment condition of Yen-Ju Cheng's and Tang's works for ten molecules.<sup>[1,2]</sup>

	Blend ratio (wt %)	Solvent	Additive
Ref.8	1:1	chlorobenzene	0.5 vol % DIO
Ref.9	1:1	chlorobenzene	0.5 vol % DIO

**Table S3.** Photovoltaic parameters about  $J_{SC}$  ( $\text{mA cm}^{-2}$ ),  $\Delta J_{SC}$  ( $\text{mA cm}^{-2}$ ),  $V_{OC}$  (eV) and  $\Delta V_{OC}$  (eV) with the positive and negative correlation. All acceptor were blended with same donor PBDB-T.

	$J_{SC}$	$\Delta J_{SC}^{\text{a}}$	$V_{OC}$	$\Delta V_{OC}^{\text{b}}$	FF	PCE	Refs.
ITIC→ITBC	16.75→19.90	3.15	0.92→0.94	0.02	65.23→64.51	10.05→12.07	
BDCPDT-IC→BDCPDT-BC	16.56→18.55	1.99	0.86→0.92	0.06	65.52→63.41	9.33→10.82	
DTCC-IC→DTCC-BC	14.26→17.22	2.96	0.97→0.98	0.01	64.08→63.62	9.25→10.74	
BDCPDT-BC→BDCPDT-TTC	18.55→17.12	-1.43	0.92→0.92	0.00	63.41→61.19	10.82→9.64	
BDCPDT-IC→ITIC	16.56→16.75	0.19	0.86→0.92	0.06	65.52→65.23	9.33→10.05	[1,2]
BDCPDT-BC→ITBC	18.55→19.90	1.35	0.92→0.94	0.02	63.41→64.51	10.82→12.07	
BDCPDT-IC→DTCC-IC	16.56→14.26	-2.30	0.86→0.97	0.11	65.52→64.08	9.33→9.25	
ITIC→DTCC-IC	16.75→14.26	-2.49	0.92→0.97	0.05	65.23→64.08	10.05→9.25	
BDCPDT-BC→DTCC-BC	18.55→17.22	-1.33	0.92→0.98	0.06	63.41→63.62	10.82→10.74	
ITBC→DTCC-BC	19.90→17.22	-2.68	0.94→0.98	0.04	64.51→63.62	12.07→10.74	

**Table S4.** Calculated atomic charges (a.u.) of fragment between NFAs adopted atomic dipole moment corrected Hirshfeld population (ADCH) method at the PBE0/6-31G(d) level.

Molecule	Fragment charge	$\Delta$ Fragment charge
ITIC	-0.252	0.024
ITBC	-0.228	
BDCPDT-IC	-0.258	0.086
BDCPDT-BC	-0.173	
DTCC-IC	-0.272	0.044
DTCC-BC	-0.227	
BDCPDT-BC	-0.173	-0.054
BDCPDT-TTC	-0.227	
BDCPDT-IC	0.394	-0.002
ITIC	0.392	
BDCPDT-BC	0.266	0.084
ITBC	0.350	
BDCPDT-IC	0.394	0.001
DTCC-IC	0.396	
ITIC	0.392	0.004
DTCC-IC	0.396	
BDCPDT-BC	0.266	0.091
DTCC-BC	0.357	
ITBC	0.350	0.007
DTCC-BC	0.357	

**Table S5.** Calculated singlet excited energies (eV), oscillator strengths ( $f$ ) and composition of singlet excited states for ITIC-series acceptors at the TD-PBE0/6-31G(d) level. Only the  $S_2$  state of DTCC-IC and DTCC-BC with C-conformation are not forbidden transitions, and HOMO-1 and LUMO+1 participate in the electron jump, respectively. For other S-conformations, the oscillator intensity of the  $S_2$  state in which LUMO+1 participates is 0.

Acceptor	State	Energy (eV)	$f$	Composition
ITIC	$S_1$	2.018	2.590	H→L (97.9%)
	$S_2$	2.357	0	H→L+1 (96.6%)
ITBC	$S_1$	1.953	1.764	H→L (95.3%)
	$S_2$	2.112	0	H→L+1 (92.8%)
BDCPDT-IC	$S_1$	1.951	2.487	H→L (98.2%)
	$S_2$	2.314	0	H→L+1 (95.6%)
BDCPDT-BC	$S_1$	1.911	1.865	H→L (95.7%)
	$S_2$	2.089	0	H→L+1 (92.3%)
BDCPDT-TTC	$S_1$	1.936	1.945	H→L (95.7%)
	$S_2$	2.121	0	H→L+1 (91.9%)
DTCC-IC	$S_1$	2.093	2.605	H→L (97.0%)
	$S_2$	2.352	0.014	H-1→L (95.4%)

DTCC-BC	S <sub>1</sub>	2.023	1.724	H→L (93.2%)
	S <sub>2</sub>	2.162	0.005	H→L+1 (89.1%)

**Table S6.** Calculated the HOMO of donor ( $E_{\text{HOMO}}^{\text{D}}$ ), LUMO of acceptor ( $E_{\text{LUMO}}^{\text{A}}$ ), the stacking distances of donor/acceptor interface model ( $R$ ) and CT-state energies ( $E_{\text{CT}}$ ) (eV), for donor/ITIC-series acceptors complexes based on  $E_{\text{CT}} = \left| E_{\text{HOMO}}^{\text{D}} - E_{\text{LUMO}}^{\text{A}} \right| - 1/R$ .

	$E_{\text{HOMO}}^{\text{D}}$	$E_{\text{LUMO}}^{\text{A}}$	$R$	$E_{\text{CT}}$
PBDB-T:ITIC	-5.14	-3.31	3.74	2.08
PBDB-T:ITBC	-5.14	-3.27	3.64	2.15
PBDB-T:BDCPDT-IC	-5.14	-3.36	3.82	2.04
PBDB-T:BDCPDT-BC	-5.14	-3.28	3.87	2.13
PBDB-T:BDCPDT-TTC	-5.14	-3.24	3.92	2.15
PBDB-T:DTCC-IC	-5.14	-3.21	3.75	2.19
PBDB-T:DTCC-BC	-5.14	-3.18	3.74	2.23

**Table S7.** Experimental energy levels and photovoltaic parameters of devices with positive and negative correlation are used in the machine learning model.

Donor	Acceptor	HOMO	LUMO	X	$V_{\text{oc}}$	$\Delta V_{\text{oc}}$	$J_{\text{sc}}$	$\Delta J_{\text{sc}}$	PCE	$\Delta \text{PCE}$	Ref
Positive correlation											
PBDB-T	BDCPDT-BC	-5.40	-3.85	0.04	0.92	0.02	18.55	1.35	10.82	1.25	[1]
PBDB-T	ITBC	-5.49	-3.90	0.94			19.90		12.07		
PBDB-T	ITIC	-5.61	-4.01	-0.01	0.92	0.02	16.75	3.15	10.05		
PBDB-T	ITBC	-5.49	-3.9	0.94			19.90		12.07	2.02	[1]
PBDB-T	DTCC-IC	-5.47	-3.83	0.01	0.97	0.02	14.26	2.96	9.25	1.49	[1]
PBDB-T	DTCC-BC	-5.46	-3.81	0.98			17.22		10.74		
FTAZ	IDIC1	-5.51	-3.81	0.02	0.90	0.05	13.6	0.7	7.13	2.08	[3]
FTAZ	IHIC1	-5.47	-3.75	0.95			14.3		9.21		
PTB7-Th	F6IC	-5.68	-4.04	-0.2	0.61	0.07	18.15	4.1	7.00	3.7	[4]
PTB7-Th	F8IC1	-5.43	-3.99	0.68			22.25		10.70		
PM6	IT-4F	-5.74	-4.26	-0.03	0.85	0.01	20.02	1.64	12.80		
PM6	IOM-4F	-5.72	-4.27	0.86			21.66		13.41	0.61	[5]
PM6	IOM-4F	-5.72	-4.27	0.05	0.86	0.02	21.66	0.46	13.41	0.76	[5]
PM6	IM-4F	-5.69	-4.19	0.88			22.12		14.17		
PBDB-TF	BDT <sub>t</sub> IC-γCl	-5.21	-3.89	-0.04	0.81	0.08	2.26	11.51	6.9	0.71	[6]
PBDB-TF	BDT <sub>c</sub> IC-γCl	-5.29	-4.01	0.89			13.77		7.61		
PM6	SiOTC	-5.6	-3.81	-0.06	0.82	0.11	4.92	8.76	1.73	7.13	[7]
PM6	SiOTIC	-5.57	-3.84	0.93			13.68		8.86		
PM6	Z1-aa	-5.67	-3.84	-0.01	0.98	0.02	11.68	3.08	4.56	5.04	[8]

PM6	Z1-ab	-5.69	-3.87		1		14.76		9.6	
PBDB-T	BDCPDT-IC	-5.41	-3.87	0.06	0.86	0.06	16.56	0.19	9.33	0.72 [1]
PBDB-T	ITIC	-5.61	-4.01		0.92		16.75		10.05	
PM6	NOIC3	-5.83	-3.95	-0.14	0.93	0.01	12.9	3.9	7.15	2.95 [9]
PM6	NOIC4	-5.64	-3.9		0.94		16.8		10.1	
PTB7-Th	F6IC	-5.66	-4.02	-0.34	0.61	0.12	18.07	2.76	7.1	3.1 [10]
PTB7-Th	F10IC	-5.26	-3.96		0.73		20.83		10.2	
PTB7-Th	F6IC	-5.66	-4.02	-0.21	0.61	0.03	18.07	7.05	7.1	3.8 [10]
PTB7-Th	F8IC	-5.43	-4		0.64		25.12		10.9	
FTAZ	IHIC2	-5.69	-3.86	-0.2	0.775	0.13	15.7	2.8	7.45	3.75 [11]
FTAZ	IOIC2	-5.41	-3.78		0.902		18.5		11.2	
FTAZ	F5IC	-5.82	-4.05	-0.04	0.70	0.038	14.88	3.55	5.60	2.6 [12]
FTAZ	F7IC	-5.74	-4.01		0.74		18.43		8.20	
FTAZ	F7IC	-5.74	-4.01	-0.18	0.74	0.131	18.43	1.77	8.20	3.5 [12]
FTAZ	F9IC	-5.52	-3.97		0.87		20.20		11.70	
PTB7-Th	F5IC	-5.82	-4.05	-0.2	0.64	0.102	13.89	10.61	5.61	6.29 [13]
PTB7-Th	AOIC	-5.50	-3.93		0.74		24.50		11.90	
PTB7-Th	F8IC1	-5.43	-3.99	-0.05	0.68	0.04	22.25	1.19	10.70	1.6 [4]
PTB7-Th	F10IC1	-5.35	-3.96		0.72		23.44		12.30	
PM6	M8	-5.49	-3.91	0.11	0.83	0.08	8.36	15.27	4.21	11.03 [14]
PM6	M34	-5.6	-3.91		0.91		23.63		15.24	
P3HT	IDBTC	-5.25	-3.54	0.01	0.73	0.02	7.7	1.23	2.5	0.72 [15]
P3HT	IDBTFC	-5.27	-3.55		0.75		8.93		3.22	
PTB7-Th	CO5DFIC	-5.96	-4.29	-0.11	0.59	0.12	12.56	4.76	4.32	1.62 [16]
PTB7-Th	CO5DFIC-OT	-5.51	-3.95		0.71		17.32		5.94	
PTB7-Th	4TIC	-5.36	-4.11	0.10	0.7	0.04	14.58	4.64	5.26	2.87 [17]
PTB7-Th	6TIC	-5.31	-3.96		0.74		19.22		8.13	
PTB7-Th	CO5DFIC	-5.96	-4.29	-0.04	0.59	0.15	12.56	6.67	4.32	4.13 [16]
PTB7-Th	CO5DFIC-ST	-5.55	-3.92		0.74		19.23		8.45	
PTB7-Th	CO5DFIC-OT	-5.51	-3.95	0.07	0.71	0.03	17.32	1.91	5.94	2.51 [16]
PTB7-Th	CO5DFIC-ST	-5.55	-3.92		0.74		19.23		8.45	
PBDB-T	DTNIF	-5.82	-3.92	-0.38	0.9	0.02	13.26	1.23	7.15	1.58 [18]
PBDB-T	DTNSF	-5.52	-4		0.92		14.49		8.73	
PBDB-T	mOEh-ITIC-F	-5.77	-4.23	-0.02	0.68	0.01	13.74	0.45	5.79	-0.01 [19]
PBDB-T	mOEh-ITIC-C1	-5.74	-4.22		0.69		14.19		5.78	
PBDB-T	mOEh-ITIC-C1	-5.74	-4.22	0.09	0.68	0.16	12.32	0.01	4.66	
PBDB-T	mOEh-ITIC	-5.67	-4.06		0.84		12.33		6.14	1.48 [19]
J51	ArSiID-F	-5.42	-3.9	-0.01	0.78	0.1	10.4	4	3.1	4.3 [20]
J51	ArSiID	-5.37	-3.86		0.88		14.4		7.4	
J51	ArSiID-Cl	-5.45	-4.01	0.07	0.78	0.1	10	4.4	2.8	4.6 [20]
J51	ArSiID	-5.37	-3.86		0.88		14.4		7.4	
PBDB-T	NTIC-Me	-5.53	-3.73	0.03	0.963	0.002	13.03	0.49	8.3	0.31 [21]

PBDB-T	NTIC-OMe	-5.53	-3.7	0.965	13.52	8.61		
PTB7-Th	T4	-5.36	-4.01	0.16	0.61	18.57	7.01	2.81 [22]
PTB7-Th	T1	-5.42	-3.91	0.07	0.72	20.95	2.38	9.82
PTB7-Th	T3	-5.31	-4	0.07	0.61	22	0.65	9.43
PTB7-Th	T2	-5.34	-3.96	0.06	0.67	22.65	10.1	0.67 [22]
PTB7-Th	T4	-5.36	-4.01	0.03	0.61	18.57	4.07	7.01
PTB7-Th	T2	-5.34	-3.96	0.06	0.67	22.64	10.1	3.09 [22]
PBDB-T	BDCPDT-IC	-5.41	-3.87	0.06	0.86	16.56	9.33	
PBDB-T	BDCPDT-TTC	-5.38	-3.78	0.94	0.08	17.72	1.16	0.96 [2]
PBDB-T	ITIC	-5.59	-3.91	-0.03	0.87	15.45	7.8	
PBDB-T	IT-OH	-5.57	-3.92	0.21	0.89	16.71	1.26	2.6 [23]
FTAZ	ITIC-Th	-5.66	-3.93	0.01	0.92	15.84	0.5	8.90
FTAZ	ITIC-Th3	-5.67	-3.73	0.047	0.96	16.34	10.70	1.8 [24]
PBDB-T	BDCPDT-IC	-5.41	-3.87	0.01	0.86	16.56	1.99	9.33
PBDB-T	BDCPDT-BC	-5.40	-3.85	0.92	0.06	18.55	10.82	1.49 [1]
PM6	BDSeIC4Br	-5.65	-4.02	0.01	0.86	16.2	3.8	9.4
PM6	BDSeIC2Br	-5.63	-3.99	0.9	0.04	20	11.9	2.5 [25]
PBDB-T-2F	ITIC-2Br-m	-5.53	-3.9	0.01	0.87	18.01	1	10.88
PBDB-T-2F	ITIC-2Br-γ	-5.54	-3.9	0.02	0.89	19.01	12.05	1.17 [26]
PBDB-T	ITIC	-5.61	-4.02	0.01	0.9	16.8	0.64	11.22
PBDB-T	IT-M	-5.58	-3.98	0.04	0.94	17.44	12.05	0.83 [27]
FTAZ	ITIC-Th2	-5.75	-4.07	0.05	0.751	17.19	2.14	9.06
FTAZ	ITIC-Th1	-5.74	-4.01	0.098	0.849	19.33	12.1	3.04 [24]
PBDB-TF	ITIC-Cl-δ-Th	-5.31	-3.7	0.03	0.89	17.27	1.03	11.13
PBDB-TF	ITIC-Cl-γ-Th	-5.3	-3.66	0.91	0.02	18.3	12.25	1.12 [28]
J52	i-IEICO-F3	-5.34	-3.74	-0.04	0.897	16.21	4.69	7.65
J52	i-IEICO-2F	-5.29	-3.73	0.012	0.909	20.9	12.86	5.21 [29]
PBDB-T	BTTIC-0M	-5.62	-3.89	0.01	0.86	18.95	0.44	11.87
PBDB-T	BTTIC-2M	-5.60	-3.86	0.04	0.90	19.39	13.15	1.28 [30]
PBDB-T	i-mO-4Cl	-5.55	-3.83	-0.03	0.872	15.08	6.47	7.33
PBDB-T	i-mO-4F	-5.5	-3.81	0.044	0.916	21.55	13.85	6.52 [31]
PCE10	3e	-5.48	-3.65	0.00	0.92	6.72	2.38	2.52
PCE10	3d	-5.42	-3.59	0.09	1.01	9.1	4.69	2.17 [32]
PBDB-T	IDT-C8-BC	-5.39	-3.5	0.04	1.05	10.27	0.26	7.3
PBDB-T	IDT-BC	-5.5	-3.57	0.03	1.08	10.53	6.3	-1 [33]
PTB7-Th	IDTP-O-C	-5.47	-3.75	-0.01	0.77	18.08	0.7	8.61
PTB7-Th	IDTP-P-C	-5.43	-3.72	0.02	0.79	18.78	8.21	-0.4 [34]
FTAZ	ITIC1	-5.48	-3.84	-0.01	0.921	16.45	1.06	8.54
FTAZ	ITIC2	-5.43	-3.8	0.025	0.946	17.51	9.56	1.02 [35]
PBT1-C	IDTT-C6-TIC	-5.55	-3.99	0.11	0.86	15.9	1.9	9.5
PBT1-C	IDTT-C8-TIC	-5.64	-3.97	0.04	0.9	17.8	10.3	0.8 [36]
PBDB-T	m-F-ITIC	-5.69	-3.96	-0.01	0.883	15.8	2.27	8.9
				0.035				2.21 [37]

PBDB-T	o-F-ITIC	-5.66	-3.94		0.918		18.07		11.11	
PBDB-T	IDTV-ThIC	-5.43	-3.72	-0.03	0.91	0.07	2.06	2.87	0.62	1.15 [38]
PBDB-T	IDTV-PhIC	-5.52	-3.84		0.98		4.93		1.77	
PBDB-T	IDTV-PhIC	-5.52	-3.84	0.17	0.98	0.01	4.93	6.65	1.77	9.81 [38]
PBDB-T	m-IDTV-PhIC	-5.63	-3.78		0.99		11.58		11.58	
PBDB-T-2F	ClBDT-4Cl	-5.72	-3.92	-0.01	0.879	0.009	19.02	0.81	11.65	0.71 [39]
PBDB-T-2F	FBDT-4Cl	-5.7	-3.91		0.888		19.83		12.36	
PBDB-T	IDTCN-S	-5.57	-3.9	0.07	0.85	0.06	19.04	0.92	10.6	2.68 [40]
PBDB-T	IDTCN-O	-5.54	-3.8		0.91		19.96		13.28	
PM6	M38	-5.65	-3.93	-0.08	0.87	0.01	18.28	1.48	8.89	2.27 [41]
PM6	M2	-5.6	-3.96		0.88		19.76		11.16	
PM6	M38	-5.65	-3.93	-0.05	0.87	0.03	18.28	6.35	8.89	7.11 [41]
PM6	M36	-5.62	-3.95		0.9		24.63		16	
PM6	M2	-5.6	-3.96	0.03	0.88	0.02	19.76	4.87	11.16	4.84 [41]
PM6	M36	-5.62	-3.95		0.9		24.63		16	
<b>Negative correlation</b>										
PBDB-T	BDCPDT-BC	-5.40	-3.85	0.05	0.92	0.02	18.55	-0.83	10.82	-0.53 [2]
PBDB-T	BDCPDT-TTC	-5.38	-3.78		0.94		17.72		10.29	
PBDB-T	BDCPDT-IC	-5.41	-3.87	0.10	0.86	0.11	16.56	-2.3	9.33	-0.08 [1]
PBDB-T	DTCC-IC	-5.47	-3.83		0.97		14.26		9.25	
PBDB-T	BDCPDT-BC	-5.40	-3.85	0.10	0.92	0.06	18.55	-1.33	10.82	-0.08 [1]
PBDB-T	DTCC-BC	-5.46	-3.81		0.98		17.22		10.74	
PBDB-T	ITIC	-5.61	-4.01	0.04	0.92	0.05	16.75	-2.49	10.05	-0.8 [1]
PBDB-T	DTCC-IC	-5.47	-3.83		0.97		14.26		9.25	
PBDB-T	ITBC	-5.49	-3.9	0.06	0.94	0.04	19.90	-2.68	12.07	-1.33 [1]
PBDB-T	DTCC-BC	-5.46	-3.81		0.98		17.22		10.74	
PTB7-Th	NOIC	-5.76	-4.03	-0.34	0.885	-0.021	18.1	3.8	11.4	1.1 [9]
PTB7-Th	NOIC1	-5.41	-4.02		0.864		21.9		12.5	
PTB7-Th	NOIC3	-5.83	-3.95	-0.23	0.93	-0.003	12.9	7.7	7.15	6.95 [9]
PTB7-Th	NOIC2	-5.64	-3.99		0.927		20.6		14.1	
PTB7-Th	IEICO	-5.24	-3.8	-0.25	0.88	-0.16	10.7	9.8	6	4.2 [42]
PTB7-Th	IEICO-4F	-5.44	-4.25		0.72		20.5		10.2	
PTB7-Th	FBTIC	-5.7	-3.85	-0.23	0.947	-0.101	14.1	6.5	10.1	2.2 [43]
PTB7-Th	FBDIC	-5.57	-3.95		0.846		20.6		12.3	
PTB7-Th	FNIC1	-5.61	-3.92	-0.13	0.774	-0.033	19.97	3.96	10.3	2.7 [44]
PTB7-Th	FNIC2	-5.56	-4		0.741		23.93		13	
PTB7-Th	AOIC	-5.5	-3.93	-0.11	0.762	-0.018	11.03	13.48	13.7	-9.22 [13]
PTB7-Th	IUIC2	-5.32	-3.86		0.74		24.51		4.48	
PM6	IT-2F	-5.63	-4.06	-0.03	0.92	-0.06	19.3	1.5	12.7	0.9 [45]
PM6	IT-4F	-5.68	-4.14		0.86		20.8		13.6	
PM6	IT-3F	-5.67	-4.09	-0.04	0.9	-0.04	20	0.8	13.8	-0.2 [45]
PM6	IT-4F	-5.68	-4.14		0.86		20.8		13.6	

PM6	ITIC-2Cl	-5.68	-3.99	-0.03	0.92	-0.13	19.1	3.6	10.3	2.9	[46]
PM6	ITIC-4Cl	-5.75	-4.09		0.79		22.7		13.2		
PM6	ITIC-2Cl- $\beta$	-5.3	-3.71	-0.07	0.94	-0.06	18.5	0.4	11.2	1	[47]
PM6	$\alpha$ -ITIC-2Cl	-5.29	-3.77		0.88		18.9		12.2		
J71	IT-DM	-5.58	-3.82	-0.12	1.02	-0.11	16.7	1.8	12.1	1.2	[42]
J71	ITCF	-5.59	-3.95		0.91		18.5		13.3		
PBDB-T	IO-4H	-5.61	-3.65	-0.14	1.12	-0.07	5.95	6.25	2.86	5.2	[43]
PBDB-T	IO-4F	-5.65	-3.83		1.05		12.2		8.06		
PBDB-T	IDTTC	-5.4	-3.72	-0.03	1.01	0.02	18.3	-3.8	13.5	-7.04	[48]
PBDB-T	IDTTTC	-5.34	-3.69		1.03		14.5		6.46		
PM6	ITIC-Cl- $\gamma$ -Th	-5.3	-3.66	-0.07	0.91	-0.05	18.3	0.3	12.3	-0.8	[49]
PM6	ITIC-2Cl-Th	-5.31	-3.74		0.86		18.6		11.5		
PTQ10	m-ITIC-2F	-5.73	-3.95	-0.07	0.96	-0.06	19	0.8	12.5	0	[50]
PTQ10	m-ITIC-4F	-5.73	-4.02		0.9		19.8		12.5		
PM6	POIT-IC2F	-5.6	-3.98	-0.04	0.97	-0.06	18.6	2.3	12.4	1.4	[51]
PM6	POIT-IC4F	-5.65	-4.07		0.91		20.9		13.8		
PTB7-Th	C <sub>6</sub> -IDTT-T	-5.71	-3.78	0.09	1.05	0.02	14.4	-1.1	8.51	-0.99	[52]
PTB7-Th	2C <sub>6</sub> -IDTT-T	-5.74	-3.72		1.07		13.3		7.52		
PM6	C8-ITCC	-5.45	-3.85	-0.03	1.04	-0.09	16.1	1.8	10.8	1.9	[53]
PM6	C8-ITCC-Cl	-5.5	-3.93		0.95		17.9		12.7		
PBT1-C	IDTT-C8-TIC	-5.64	-3.97	0.13	0.88	0.1	20.3	-2.2	13.7	-1	[54]
PBT1-C	IDTT-C10-TIC	-5.71	-3.91		0.98		18.1		12.7		
PTB7-Th	BT-SFIC	-5.46	-4.04	-0.04	0.77	-0.04	18.5	2.8	9.52	0.58	[55]
PTB7-Th	BT-FIC	-5.48	-4.1		0.73		21.3		10.1		
PM6	HBDT-4Cl	-5.67	-3.9	0.03	0.9	-0.02	17.8	1.2	10.4	1.3	[39]
PM6	CIBDT-4Cl	-5.72	-3.92		0.88		19		11.7		
PBDB-T	BTTIC-TT	-5.54	-3.8	-0.04	0.92	0.01	19.6	-3.1	13.4	-4.26	[56]
PBDB-T	BTTIC-Ph	-5.48	-3.78		0.93		16.5		9.14		
PBDB-T	SNBDT1-F	-5.3	-3.83	0.05	0.86	0.02	21	-7.1	12.7	-6.13	[57]
PBDB-T	SNBDT2-F	-5.34	-3.82		0.88		13.9		6.57		
PBDB-T	SNBDT1-F	-5.3	-3.83	0.09	0.86	0.04	21	-8.5	12.7	-6.83	[57]
PBDB-T	SNBDT3-F	-5.36	-3.8		0.9		12.5		5.87		
PBDB-T	SNBDT2-F	-5.34	-3.82	0.04	0.88	0.02	13.9	-1.4	6.57	-0.7	[57]
PBDB-T	SNBDT3-F	-5.36	-3.8		0.9		12.5		5.87		
PBDB-T	IPT2F-Ph	-5.57	-4	0.03	0.86	-0.02	21.2	1	13.1	0.9	[58]
PBDB-T	IPT2F-TT	-5.6	-4		0.84		22.2		14		
PM6	IPTBO-4F	-5.57	-4.07	-0.03	0.92	-0.04	22.1	1.1	14.7	-0.3	[59]
PM6	IPT-4Cl	-5.58	-4.11		0.88		23.2		14.4		
PM6	IN-4F	-5.59	-3.94	-0.13	0.92	-0.09	19.5	1.7	12.5	-1.7	[60]
PM6	IPCl-4F	-5.6	-4.08		0.83		21.2		10.8		
PM6	INO-4F	-5.64	-3.93	-0.19	0.93	-0.1	20.5	0.7	13.7	-2.9	[60]
PM6	IPCl-4F	-5.6	-4.08		0.83		21.2		10.8		

P3HT	I-IDTBTRh	-5.42	-3.68	0.30	0.86	0.06	8.81	-3.81	5.38	-2.85	[61]
P3HT	a-IDTBTRh	-5.63	-3.59		0.92		5		2.53		
J61	BTA1	-5.46	-3.59	0.13	1.24	0.05	5.21	-4.37	3.02	-2.76	[62]
J61	BTA2	-5.43	-3.43		1.29		0.84		0.26		
P3HT	BTA100	-5.32	-3.23	-0.23	1.34	-0.15	1.65	3.68	1.04	2.51	[63]
P3HT	BTA101	-5.41	-3.55		1.19		5.33		3.55		
P3HT	BTA100	-5.32	-3.23	-0.36	1.34	-0.4	1.65	6.91	1.04	4.27	[63]
P3HT	BTA103	-5.37	-3.64		0.94		8.56		5.31		
P3HT	BTA101	-5.41	-3.55	-0.13	1.19	-0.25	5.33	3.23	3.55	1.76	[63]
P3HT	BTA103	-5.37	-3.64		0.94		8.56		5.31		
PM6	NT-4F	-5.88	-3.86	-0.05	0.96	-0.03	13.9	2.7	9.46	1.94	[64]
PM6	NT-4Cl	-5.89	-3.92		0.93		16.6		11.4		
PM6	T-TT-4F	-5.44	-3.51	-0.04	0.86	-0.05	18.5	0.5	10.5	-0.3	[65]
PM6	T-TT-4Cl	-5.48	-3.59		0.81		19		10.2		
PM6	IDT6CN-TM	-5.7	-3.96	-0.08	0.95	-0.09	17.4	0.9	12.4	-1.5	[66]
PM6	IDT6CN-4F	-5.78	-4.12		0.86		18.3		10.9		
PBDB-T	SN61C	-5.38	-3.93	-0.04	0.88	-0.1	16.5	6.7	9.6	3.6	[67]
PBDB-T	SN61C-4F	-5.52	-4.11		0.78		23.2		13.2		
PBDB-T	IXIC	-5.13	-3.78	-0.05	0.82	-0.09	20.9	2.7	11.3	0.9	[68]
PBDB-T	IXIC-2Cl	-5.2	-3.9		0.73		23.6		12.2		
PM6	Z1-ab	-5.69	-3.87	-0.04	1	-0.02	14.8	3.7	9.6	3.1	[8]
PM6	Z1-bb	-5.7	-3.92		0.98		18.5		12.7		
PBT1-C	TTPTTT-IC	-5.64	-3.87	-0.14	1	-0.08	12.5	4.3	7.91	3.59	[69]
PBT1-C	TTPTTT-2F	-5.67	-4.04		0.92		16.8		11.5		
PBT1-C	TTPTTT-2F	-5.67	-4.04	-0.06	0.92	-0.06	16.8	2.6	11.5	0.6	[69]
PBT1-C	TTPTTT-4F	-5.69	-4.12		0.86		19.4		12.1		
PM6	IOIC2	-5.7	-4.16	-0.09	0.97	-0.05	16.3	3.7	10.5	2.3	[70]
PM6	IOIC3	-5.64	-4.19		0.92		20		12.8		
PM6	IOIC3	-5.64	-4.19	0.08	0.92	0.04	20	-2.7	12.8	-1.7	[70]
PM6	IOIC4	-5.7	-4.17		0.96		17.3		11.1		
PM6	IOIC4	-5.7	-4.17	-0.07	0.96	-0.04	17.3	3.6	2.7		[70]
PM6	IOIC5	-5.65	-4.19		0.92		20.9		13.8		
PM6	IOIC2	-5.7	-4.16	-0.08	0.97	-0.05	16.3	4.6	10.5	3.3	[70]
PM6	IOIC5	-5.65	-4.19		0.92		20.9		13.8		
PM6	NOIC	-5.76	-4.03	-0.34	0.89	-0.03	18.1	3.8	11.4	1.1	[71]
PM6	NOIC1	-5.41	-4.02		0.86		21.9		12.5		
PM6	NOIC1	-5.41	-4.02	0.26	0.86	0.07	21.9	-1.3	12.5	1.6	[71]
PM6	NOIC2	-5.64	-3.99		0.93		20.6		14.1		
PM6	NOIC1	-5.41	-4.02	0.49	0.86	0.07	21.9	-9	12.5	-5.35	[71]
PM6	NOIC3	-5.83	-3.95		0.93		12.9		7.15		
PM6	NOIC	-5.76	-4.03	0.15	0.89	0.04	18.1	-5.2	11.4	-4.25	[71]
PM6	NOIC3	-5.83	-3.95		0.93		12.9		7.15		

PM6	NOIC1	-5.41	-4.02	0.35	0.86	0.08	21.9	-5.1	12.5	-2.4	[71]
PM6	NOIC4	-5.64	-3.9		0.94		16.8		10.1		
PM6	NOIC2	-5.64	-3.99	0.09	0.93	0.01	20.6	-3.8	14.1	-4.0	[71]
PM6	NOIC4	-5.64	-3.9		0.94		16.8		10.1		
J71	ZITI	-5.59	-3.74	0.03	0.93	-0.03	20.4	0.3	13.2	0	[72]
J71	ZITI-3F	-5.64	-3.76		0.9		20.7		13.2		
J71	ZITI-3F	-5.64	-3.76	-0.03	0.9	-0.05	20.7	0.6	13.2	0	[73]
J71	ZITI-4F	-5.66	-3.81		0.85		21.3		13.2		
PBDB-T	FTTCN	-5.54	-3.95	0.03	0.9	0.03	15.9	-0.7	10.6	-0.5	[74]
PBDB-T	FTTCN-M	-5.53	-3.91		0.93		15.2		10.1		

**Table S8.** Donor/acceptor blending ratio versus the solvent environment for 112 device preparation conditions that achieve positive correlation.

Donor	NFA	D:A ratio	Solvent
FTAZ	IDIC1	1:1.5	CHCl <sub>3</sub>
FTAZ	IHIC1		
PBDB-TF	BDT <sub>t</sub> IC- $\gamma$ Cl	1:1.2	chlorobenzene
PBDB-TF	BDT <sub>c</sub> IC- $\gamma$ Cl		
PM6	SiOTC	1:1.2	chlorobenzene
PM6	SiOTIC		
PM6	Z1-aa	1:1	chlorobenzene
PM6	Z1-ab		
PBDB-T	BDCPDT-IC	1:1	chlorobenzene
PBDB-T	ITIC		
PM6	NOIC3	1:1	chloroform
PM6	NOIC4		
PTB7-Th	F6IC	1:1.7	chloroform
PTB7-Th	F10IC		
PTB7-Th	F6IC	1:1.7	chloroform
PTB7-Th	F8IC		
FTAZ	IHIC2	1:1.5	chloroform
FTAZ	IOIC2		
FTAZ	F5IC	1:1.5	chloroform
FTAZ	F7IC		
FTAZ	F7IC	1:1.5	chloroform
FTAZ	F9IC		
PTB7-Th	F5IC	1:1.5	dichlorobenzene
PTB7-Th	AOIC		
PTB7-Th	F8IC1	1:1.2	chloroform
PTB7-Th	F10IC1		
PM6	M8	1:1	chloroform
PM6	M34		
PTB7-Th	CO5DFIC	1:1	chlorobenzene
PTB7-Th	CO5DFIC-OT		
PTB7-Th	4TIC	1:1.3	chloroform
PTB7-Th	6TIC		
PBDB-T	DTNIF	1:1	chlorobenzene
PBDB-T	DTNSF		
PBDB-T	mOEh-ITIC-F	1:1	chlorobenzene
PBDB-T	mOEh-ITIC-Cl		
PBDB-T	mOEh-ITIC-Cl	1:1	Xylene
PBDB-T	mOEh-ITIC		
J51	ArSiID-F	1:1	chloroform
J51	ArSiID		

J51	ArSiID-Cl		
J51	ArSiID	1:1	chloroform
PBDB-T	NTIC-Me		
PBDB-T	NTIC-OMe	1: 0.8	chloroform
PTB7-Th	T4		
PTB7-Th	T1	1:1.8	chlorobenzene
PTB7-Th	T3		
PTB7-Th	T2	1:1.8	chlorobenzene
PTB7-Th	T4		
PTB7-Th	T2	1:1.8	chlorobenzene
PBDB-T	ITIC		
PBDB-T	IT-OH	1:1	chlorobenzene
FTAZ	ITIC-Th		
FTAZ	ITIC-Th3	1:1.5	chloroform
PBDB-T	BDCPDT-IC		
PBDB-T	BDCPDT-BC	1:1	chlorobenzene
PM6	BDSeIC4Br		
PM6	BDSeIC2Br	1:1	chlorobenzene
PBDB-T-2F	ITIC-2Br-m		
PBDB-T-2F	ITIC-2Br- $\gamma$	1:1	chlorobenzene
PBDB-T	ITIC		
PBDB-T	IT-M	1:1	chlorobenzene
FTAZ	ITIC-Th2		
FTAZ	ITIC-Th1	1:1.5	chloroform
PBDB-TF	ITIC-Cl- $\delta$ -Th		
PBDB-TF	ITIC-Cl- $\gamma$ -Th	1:1	chlorobenzene
J52	i-IEICO-F3		
J52	i-IEICO-2F	1:1	chlorobenzene
PBDB-T	i-mO-4Cl		
PBDB-T	i-mO-4F	1:1	chlorobenzene
PBDB-T	IDT-C8-BC		
PBDB-T	IDT-BC	1:1.5	chlorobenzene
PTB7-Th	IDTP-O-C		
PTB7-Th	IDTP-P-C	1:1	chloroform
FTAZ	ITIC1		
FTAZ	ITIC2	1:1.3	chloroform
PBT1-C	IDTT-C6-TIC		
PBT1-C	IDTT-C8-TIC	1:1.3	chloroform
PBDB-T	m-F-ITIC		
PBDB-T	o-F-ITIC	1.3:1	chlorobenzene
PBDB-T	IDTV-ThIC		
PBDB-T	IDTV-PhIC	1:1	chlorobenzene
PBDB-T	IDTV-PhIC		
PBDB-T	m-IDTV-PhIC	1:1	chlorobenzene

PBDB-T-2F	CIBDT-4Cl	1:1	
PBDB-T-2F	FBDT-4Cl		chlorobenzene
PBDB-T	IDTCN-S	1:1	
PBDB-T	IDTCN-O		1,2-dichlorobenzene
PM6	M38	1:1	
PM6	M2		chloroform
PM6	M38	1:1	
PM6	M36		chloroform
PM6	M2	1:1	
PM6	M36		chloroform

## II. Calculation Methods

Since PBE0 method could provide an accurate evaluation of geometric and electronic structures for thiophene derivatives.<sup>75-76</sup> The ground geometrical configurations were optimized by density functional theory (DFT) with the PBE0 functional and 6-31G(d) basis set (**Fig. S1**). For the quantum chemical study of molecular systems, the ADCH method shows good reproducibility in terms of charge rationality, basis set dependence, *etc.*<sup>77</sup> Thereby, the atom charges of fragment were calculated adopted ADCH method in Multiwfn 3.8.<sup>78</sup> All calculations were carried out by using the Gaussian 09 package.<sup>79</sup>

## III. Machine learning model: prediction of positive and negative correlation

```

import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
from pandas import read_csv
import scipy
train_set = read_csv('train.csv', dtype = np.float32)
test_set = read_csv('test.csv', dtype = np.float32)
train_data_orig = train_set.values
test_data_orig = test_set.values
train_x = train_data_orig[:, 1:]
train_set_x = train_x .reshape(train_x .shape[0], -1).T
train_set_y = train_data_orig[:, 0]
test_x = test_data_orig[:, 1:]
test_set_x = test_x .reshape(test_x .shape[0], -1).T
test_set_y = test_data_orig[:, 0]
def sigmod(z):
    s = 1/(1+np.exp(-z))
    return s
def initialize_with_zeros(dim):
    w = np.zeros((dim, 1))
    b = 0
    return w, b
def propagate(w, b, X, Y):
    m = X.shape[1]

```

```

Z = np.dot(w.T, X)+b
A = sigmod(Z)
cost = (-(np.dot(Y, np.log(A).T)+np.dot((1-Y), np.log(1-A).T)))/m
db = np.sum(A-Y)/m
dw = np.dot(X, (A-Y).T)/m
grads = {'dw':dw, 'db':db}
return grads, cost
def optimize(w, b, X, Y, num_iterations, learning_rate):
    costs = []
    for i in range(num_iterations):
        grads, cost = propagate(w, b, X, Y)
        dw = grads['dw']
        db = grads['db']
        w = w-learning_rate*dw
        b = b-learning_rate*db
        if i%100 == 0:
            np.squeeze(cost)
            costs.append(cost)

    params = {'w':w, "b":b}

    return params, costs
def prediciton(w, b, X):
    m = X.shape[1]
    Y_pred = np.zeros((1, m))
    w = w.reshape(X.shape[0], 1)
    A = sigmod(np.dot(w.T, X)+b)
    for i in range(A.shape[1]):
        if A[0][i]<= 0.5:
            A[0][i] = 0
        else:
            A[0][i] = 1
    Y_pred = A
    return Y_pred
def model(X_train, X_test, Y_train, Y_test,num_iterations = 60000, learning_rate = 0.01):
    w, b = initialize_with_zeros(X_train.shape[0])
    parameters, costs = optimize(w, b, X_train, Y_train, num_iterations, learning_rate)
    w = parameters["w"]
    b = parameters["b"]
    Y_prediction_test = prediciton(w, b, X_test)
    Y_prediction_train = prediciton(w, b, X_train)
    print('train accuracy: {}%'.format(100-np.mean(np.abs(Y_prediction_train-Y_train))*100))
    print('test accuracy: {}%'.format(100-np.mean(np.abs(Y_prediction_test-Y_test))*100))
    d = {'costs':costs,

```

```

'Y_prediction_test':Y_prediction_test,
'Y_prediction_train':Y_prediction_train,
"w":w,
"b":b,
'learning_rate':learning_rate,
"num_iterations":num_iterations}
print("y = {}x+{}".format(d["w"], d["b"]))
return d

d = model(train_set_x, test_set_x, train_set_y, test_set_y, num_iterations = 60000, learning_rate = 0.01)
costs = np.squeeze(d['costs'])

```

#### IV. Appendix Full Name

Full Name	Abbreviation
3,9-bis(2-methylene-(3-(1,1-dicyanomethylene)-in-danone)-5,5,11,11-tetrakis(4-hexyl-phenyl)-dithieno[2,3-d:2',3'-d']-s-inndaceno[1,2-b:5,6-b'] dithiophene	ITIC
C6-olefinated2-(3-oxo-2,3-dihydro-1H-benzo[b]cyclopenta[d]thiophen-1-ylidene)-malononitrile with two 2-(3-oxo-2,3-dihydro-1H-benzo[b]cyclopenta[d] thiophen-1-ylidene)-malononitrile (BC) moieties	ITBC
Thieno[3,2-b]thiophene (BDCPDT) with two 1,1-dicyanomethylene-3-indanone (IC) moieties	BDCPDT-IC
Thieno[3,2-b]thiophene with two 2-(3-oxo-2,3-dihydro-1H-benzo[b]cyclopenta[d] thiophen-1-ylidene)-malononitrile (BC) moieties	BDCPDT-BC
Thieno[3,2-b]thiophene with two 2-(5-oxo-5,6-dihydro-7H-cyclopenta[b]thieno[2,3-d]thiophen-7-ylidene)malononitrile (TTC) moieties	BDCPDT-TTC
dithienocyclopentacarbazole (DTCC) with two 1,1-dicyanomethylene-3-indanone (IC) moieties	DTCC-IC
dithienocyclopentacarbazole (DTCC) with two 2-(3-oxo-2,3-dihydro-1H-benzo[b]cyclopenta[d] thiophen-1-ylidene)-malononitrile (BC) moieties	DTCC-BC
poly[(2,6-(4,8-bis(5-(2-ethylhexyl)thiophen-2-yl)-benzo[1,2-b:4,5-b']dithiophene))-alt-(5,5-(1',3'-di2-thienyl-5',7'-bis(2-ethylhexyl)benzo[1',2'-c:4',5'-]dithiophene-4,8-dione))])	PBDB-T

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