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Supporting Information

Design and Virtual Screening of Donor and Non-Fullerene Acceptor for Organic Solar Cells Using Long Short-Term Memory Model

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Fig. S1 SHAP importance analysis of the 30 molecular structure descriptors used in the LSTM model for V_{OC} prediction



Fig. S2 SHAP importance analysis of the 30 molecular structure descriptors used in the LSTM model for J_{SC} prediction



Fig. S3 SHAP importance analysis of the 30 molecular structure descriptors used in the LSTM model for FF prediction

















Fig. S4 Fragments of donor molecules

























Fig. S5 Fragments of acceptor molecules

descriptors	resolve	photograph
NOCount_A	Number of nitro groups contained in the acceptor molecule	NO
NumAliphaticCarbocycl es_A	Number of alicyclic alkyl rings in the acceptor molecule	
NumAliphaticRings_A	Number of alicycles in the acceptor molecule	s s
NumAromaticCarbocycl es_A	Number of aromatic cycloalkyl rings in the acceptor molecule	
NumAromaticHeterocycl es_A	Number of aromatic heterocycles in the acceptor molecule	N N N
Number of hydrogen bond receptors (usually nitrogen, oxygen or sulfur) in the acceptor molecule		
NumHeteroatoms_A	Number of heteroatoms in the acceptor molecule	
NumRotatableBonds_A	Number of rotatable chemical bonds_A bonds in the acceptor molecule	
fr_Ar_N_A	Number of nitrogen atom groups in the aromatic ring in the acceptor	
fr_C_O_A	Number of carbon-oxygen double bonds in the acceptor	0

Table S1 Acceptor descriptors used for LSTM model

fr_NH0_A	Number of amino groups without hydrogen atoms in the acceptor	N N
fr_allylic_oxid_A	Number of allyl oxide groups in the acceptor	
fr_aryl_methyl_A	Number of arylmethyl groups in the acceptor	
fr_bicyclic_A	Number of two or more rings in the acceptor	
fr_halogen_A	Number of halogen groups in the acceptor	
fr_ketone_A	Number of ketone groups in the acceptor	0
fr_nitrile_A	Number of nitrile groups in the acceptor	
fr_thiophene_A	Number of thiophene rings in the acceptor	
fr_unbrch_alkane_A	Number of unbranched aliphatic groups in the acceptor	$\wedge \wedge$

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descriptors	resolve	photograph
NOCount_D	Number of nitro groups	
	contained in the donor molecule	
NumAliphaticCarbocycl	Number of alicyclic alkyl rings	1988 an 1974 anns an Aonaichte an
es_D	in the donor molecule	
NumAliphaticRings_D	Number of alicycles in the	
	donor molecule	

Table S2 Donor descriptors used for LSTM model

NumAromaticCarbocycl	Number of aromatic cycloalkyl				
es_D	rings in the donor molecule				
BingCount D	Number of rings contained in				
KingCount_D	the donor molecule				
	Number of carbon-oxygen				
fr_C_O_D	double bonds in the donor				
	molecule				
	Number of amino groups				
fr_NH0_D	without hydrogen atoms in the				
	donor molecule				
fr biovalia D	Number of two or more rings in				
Ir_bicyclic_D	the donor molecule				
fr. other	Number of ether groups in the	~0~			
Ir_ether	donor molecule				
fr_halogen_D	Number of halogen groups in				
	the donor molecule				
fr unbroh alkong D	Number of unbranched aliphatic				
ir_unorcn_aikane_D	groups in the donor molecule				

 Table S3 Hyperparameters of the LSTM model for PCE, V_{OC}, J_{SC}, and FF predictions

 Device Hyperparameters

PCE	hidden_size : 150 num_layers : 1 batch_size : 60 l2_foctor : 0.01 lr : 0.0001 dropout : 0.1 patience : 40 optimizer : Adam
J _{SC}	hidden_size : 90 num_layers : 1 batch_size : 310 l2_foctor : 0.1 lr : 0.01 dropout : 0.1 patience : 10 optimizer : Adam
V _{oc}	hidden_size : 220 num_layers : 1 batch_size : 10 12_foctor : 0.001 lr : 0.0001 dropout : 0.2 patience : 30 optimizer : Adam
FF	hidden_size : 90 num_layers : 1 batch_size : 220 l2_foctor : 0.1 lr : 0.01 dropout : 0.1 patience : 10 optimizer : Adam

D: A	PM6:L8-BO	PB[N][F]:	PM6:Y18	PTB7-Th:	PBDB-T:sp-
		Y6		DTC-F-F	mOEh-ITIC
Experimental V _{OC} (V)	0.88	0.85	0.84	0.84	0.87
Predictive V _{OC} (V)	0.86	0.79	0.88	0.82	0.88
V _{OC} Absolute Error (V)	0.02	0.06	0.04	0.02	0.01
Experimental J _{SC} (mA cm ⁻²)	26.49	25.00	24.91	16.01	12.13
Predictive J _{SC(} mA cm ⁻²)	23.47	24.08	24.46	12.57	11.30
J_{SC} Absolute Error (mA cm ⁻²)	3.02	0.92	0.45	3.44	0.83
Experimental FF (%)	78.92	68.10	76.40	56.00	61.00
Predictive FF (%)	68.87	68.19	67.96	54.03	61.23
FF Absolute Error (%)	10.05	0.09	8.44	1.97	0.23

Table S4 $\mathrm{V}_{\mathrm{OC}},$ J_{SC}, and FF predictions for five donor-acceptor pairs within the

database using the trained LSTM model

					PffBT4T-
	D18:L8-BO	PTQ10:	PTB7-Th:	PM6:ID-	2OD:P(4CF
D: A		ITIC-4F	Y6	C6Ph-4F	8CH-PDI-
					TT)
Experimental V _{OC} (V)	0.90	0.95	0.69	0.84	0.74
Predictive V _{OC} (V)	0.81	0.79	0.79	0.86	0.86
V _{OC} Absolute Error (V)	0.09	0.16	0.10	0.02	0.12
Experimental J _{SC} (mA cm ⁻²)	25.10	19.81	23.40	17.47	8.72
Predictive J _{SC} (mA cm ⁻²)	22.12	17.55	23.11	18.84	8.73
J _{SC} Absolute Error (mA cm ⁻²)	2.98	2.26	0.29	1.37	0.01
Experimental FF (%)	71.20	61.00	67.80	73.22	52.99
Predictive FF (%)	65.54	62.19	64.48	65.65	43.42
FF Absolute Error (%)	5.66	1.19	3.32	7.57	9.57

Table S5 $\mathrm{V}_{\mathrm{OC}},\,J_{SC},$ and FF predictions for five donor-acceptor pairs outside the

database using the trained LSTM model