

Cite this: *Digital Discovery*, 2025, 4, 1306

# Predicting the excited-state properties of crystalline organic semiconductors using GW+BSE and machine learning†

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Excited-state properties of crystalline organic semiconductors are key to organic electronic device applications. Machine learning (ML) models capable of predicting these properties could significantly accelerate materials discovery. We use the sure-independence-screening-and-sparsifying-operator (SISSO) ML algorithm to generate models to predict the first singlet excitation energy, which corresponds to the optical gap, the first triplet excitation energy, the singlet–triplet gap, and the singlet exciton binding energy of organic molecular crystals. To train the models we use the “PAH101” dataset of many-body perturbation theory calculations within the GW approximation and Bethe–Salpeter equation (GW+BSE) for 101 crystals of polycyclic aromatic hydrocarbons (PAHs). The best performing SISSO models yield predictions within about 0.2 eV of the GW+BSE reference values. SISSO models are selected based on considerations of accuracy and computational cost to construct materials screening workflows for each property. The screening targets are chosen to demonstrate typical use-cases relevant for organic electronic devices. We show that the workflows based on SISSO models can effectively screen out most of the materials that are not of interest and significantly reduce the number of candidates selected for further evaluation using computationally expensive excited-state theory.

Received 16th December 2024  
Accepted 7th April 2025

DOI: 10.1039/d4dd00396a

rsc.li/digitaldiscovery

## Introduction

In analogy with their inorganic counterparts, crystalline organic semiconductors have the advantages of uniform well-defined electronic and optical properties, as well as improved charge carrier mobility, owing to band-like transport and fewer traps, compared to amorphous films.<sup>1–11</sup> Advances in processing techniques have led to better prospects and increasing interest in organic devices based on crystalline materials, including organic field effect transistors (OFETs),<sup>6,7,12–15</sup> organic light emitting diodes (OLEDs),<sup>16–22</sup> organic photovoltaics (OPV),<sup>23–25</sup> photodetectors,<sup>26–28</sup> and scintillators.<sup>29–32</sup> The properties of crystalline organic semiconductors can be tuned in many ways, including chemical substitution and side-group functionalization, which can modify the molecular properties as well as the crystal packing,<sup>7,33–35</sup> promoting the crystallization of polymorphs with different properties,<sup>32,36–38</sup> co-crystallization,<sup>27,39–44</sup> and doping.<sup>16,20–22,29,45,46</sup> This opens up a vast design space with

endless possibilities for devices based on crystalline organic semiconductors.

Key parameters for device performance are derived from excited-state properties of crystalline organic semiconductors. The optical gap, which corresponds to the lowest singlet exciton energy, is of fundamental importance for any device based on absorption or emission of light. The triplet exciton energy is important for devices that involve conversion between singlet and triplet states. To increase the efficiency of OLEDs, it is desirable to convert electrically generated triplet excitons into emissive singlet excitons *via* thermally activated delayed fluorescence (TADF).<sup>47–53</sup> In OPV, the thermalization loss may be reduced by converting high-energy singlet excitons into two triplet excitons, *via* singlet fission (SF).<sup>54–62</sup> Conversely, the transmission loss of photons with energies below the optical gap of the absorber may be reduced by up-converting two low-energy triplet excitons into one singlet exciton *via* triplet–triplet annihilation (TTA).<sup>63–71</sup> TTA also plays a crucial role in the detection of X-rays, gamma rays, and neutrons in scintillators.<sup>30</sup> Another important parameter for organic devices is the exciton binding energy, which is the electrostatic attraction between the electron and hole that must be overcome in order to separate an exciton into free charge carriers.<sup>72</sup> The band lineup *e.g.*, between the donor and acceptor in organic solar cells, is often engineered to overcome the exciton binding energy and provide the driving force for charge separation.

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† Electronic supplementary information (ESI) available. See DOI: <https://doi.org/10.1039/d4dd00396a>

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Different applications impose different requirements on the excited-state energies of molecular crystals. For example, solar cells require materials with broad absorption in the visible range, whereas OLEDs require materials with sharp emission in specific colors. Other applications may require absorption in the infrared (IR) or ultra-violet (UV) ranges. SF requires the triplet excitation energy to be slightly smaller than half the singlet excitation energy, whereas TTA requires the triplet energy to be slightly larger than half the singlet energy. TADF requires the difference between the singlet excitation energy and the triplet excitation energy, also known as the singlet–triplet gap, to be as small as possible or even negative. In addition, compatibility with other device components may require singlet and/or triplet energies in a specific range and/or specific band edge positions. Although the structures of over a million organic molecular crystals are known,<sup>73,74</sup> the electronic and optical properties of most of them are unknown. A material originally produced for one purpose may turn out to be useful for another.<sup>75</sup> Even materials that formed as reaction byproducts may turn out to have useful properties.<sup>76</sup> Therefore, the ability to predict the excited-state properties of molecular crystals could lead to materials discovery and advances in organic semiconductor devices.

On present day supercomputers, it is possible to screen thousands of materials in a “high-throughput” manner using density functional theory (DFT) with computationally efficient semi-local exchange–correlation functionals.<sup>77,78</sup> However, DFT is a ground-state theory, which inherently cannot describe the excited-state properties required for organic semiconductor devices. The excited-state properties of isolated molecules may be calculated relatively efficiently with time dependent DFT (TDDFT). Indeed, TDDFT has been employed for high-throughput screening efforts to discover organic chromophores.<sup>79–86</sup> The properties of crystalline organic semiconductors depend not only on the molecular properties, but also on the crystal packing and the resulting electronic interactions between molecules. There is a vast body of literature reporting how changes in the crystal packing affect device relevant properties of organic semiconductors from excited-state energies to charge carrier mobility.<sup>32,36–38,40,44,87–90</sup> Therefore, for applications based on crystalline organic semiconductors it is important for computational materials discovery efforts to focus on predicting the excited-state properties of crystals as opposed to isolated molecules.

The excited-state properties of molecular crystals can be calculated within the framework of Green’s function based many-body perturbation theory. The GW approximation, where  $G$  represents the one-particle Green’s function and  $W$  represents the screened Coulomb interaction, can be used to calculate properties associated with charged excitations, such as the fundamental band gap. Subsequently, the GW quasiparticle energies are fed into the Bethe–Salpeter equation (BSE) to calculate properties associated with neutral excitations, such as the singlet and triplet excitation energies.<sup>91–95</sup> The high computational cost of GW+BSE is prohibitive for large-scale materials screening. Machine learning (ML) can be used to perform preliminary screening and reduce the need for expensive simulations to a smaller number of promising

candidates.<sup>96–107</sup> However, training ML models such as neural networks typically requires a large amount of data,<sup>108,109</sup> which is difficult to acquire with GW+BSE. Compared to DFT and even TDDFT datasets, GW+BSE datasets are scarce and contain a relatively small amount of data for isolated molecules<sup>77,110,111</sup> or for inorganic crystals with a small number of atoms in the unit cell.<sup>112</sup> As a consequence, efforts to use ML to predict the outcomes of GW+BSE calculations have also been restricted to small organic molecules and small inorganic systems.<sup>112–116</sup> This has limited the ability to train ML models to predict the excited state properties of molecular crystals.

Recently, we have published a first of its kind dataset of GW+BSE calculations for 101 molecular crystals of polycyclic aromatic hydrocarbons (PAHs) with up to ~500 atoms in the unit cell, known as PAH101.<sup>117</sup> We have chosen to focus on PAHs because they are the fundamental building blocks of materials for organic electronics.<sup>7,88,118–126</sup> Fig. 1a and b shows the size distribution of the molecules and the unit cells in the PAH101 set. The data records contain the GW+BSE singlet and triplet excitation energies, whose distributions are shown in Fig. 1c and d as well as the GW fundamental band gaps. From these quantities we may also calculate the singlet–triplet gap and the singlet exciton binding energy, whose distributions are shown in Fig. 1e and f. Detailed technical validation of the PAH101 dataset is provided in ref. 117, including comparison of the relaxed geometries with experimental data, convergence tests of the GW+BSE calculations, and comparison of the optical gaps and absorption spectra to available experimental data.

The PAH101 dataset was originally generated for the purpose of SF materials discovery.<sup>127</sup> To address the challenge of constructing transferable ML models with a small amount of training data, we used the sure-independence-screening-and-sparsifying-operator (SISSO)<sup>128,129</sup> algorithm. SISSO takes advantage of physical/chemical knowledge to train ML models based on “small data”. The input of SISSO is a set of scalar primary features, which are physical/chemical descriptors thought to be related to the target property. The primary features used in ref. 127 are also provided in the PAH101 data records.<sup>117</sup> SISSO generates a huge feature space by repeatedly combining the primary features using a set of linear and nonlinear algebraic operations with rules to avoid unphysical combinations. Subsequently, linear regression is performed to identify the most predictive models. In the last few years SISSO has been applied increasingly broadly for diverse classes of materials and target properties and continues to perform well with relatively small amounts of data.<sup>130–149</sup> In ref. 127 SISSO successfully produced several models that were able to predict the GW+BSE values of the SF driving force, which corresponds to the difference between the singlet exciton energy and twice the triplet exciton energy, with a root mean square error (RMSE) below 0.2 eV. Based on considerations of model accuracy and primary feature computational cost, two of the SISSO models were selected to build a classifier for materials screening. Later, the simpler model of the two was found to deliver robust performance outside of the PAH101 set, whereas the more complex model was found to be overfitted.<sup>54</sup>

Here, we use the PAH101 dataset to train SISSO models for the singlet exciton energy, which corresponds to the optical gap, the



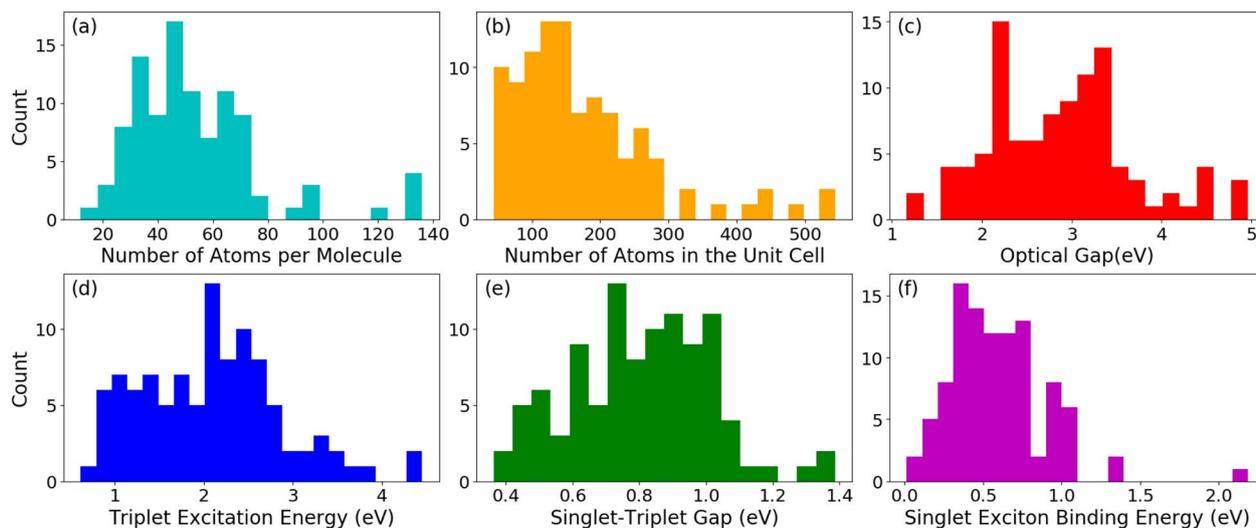


Fig. 1 Distributions of (a) the number of atoms per molecule, (b) the number of atoms in the unit cell, (c) the singlet excitation energy (optical gap), (d) the triplet excitation energy, (e) the singlet–triplet gap, and (f) the singlet exciton binding energy in the PAH101 dataset.

triplet exciton energy, the singlet–triplet gap, and the singlet exciton binding energy. For all four properties, the best performing SISO models yield predictions within about 0.2 eV of the GW+BSE reference values. We then select SISO models based on considerations of accuracy and computational cost to construct materials screening workflows for each property. The screening targets are chosen to demonstrate typical use-cases relevant for organic electronic devices. We demonstrate screening for materials with an optical gap in a particular color; materials meeting the requirements for up-conversion of infrared light to the visible range *via* TTA, namely, a triplet excitation energy in the infrared, optical gap in the visible range, and a higher TTA driving force than rubrene; materials with a small singlet–triplet gap, which is desirable for OLEDs; and materials with a small singlet exciton binding energy to facilitate the separation of excitons into free charge carriers. We show that the workflows based on SISO models can effectively screen out most of the materials that are not of interest and significantly reduce the number of candidates selected for further evaluation using computationally expensive excited-state theory.

## Methods

SISO models were trained following the same procedure used in ref. 127. The same primary features were used, summarized in Table 1 (also provided in the PAH101 dataset<sup>117</sup>). The primary features include both single molecule and crystal properties. DFT primary features were calculated with the FHI-aims<sup>150,151</sup> code using the Perdew–Burke–Ernzerhof (PBE)<sup>152,153</sup> exchange–correlation functional, as described in ref. 127. The DFT estimates for the SF driving force of the single molecule and crystal ( $DF_s$  and  $DF_c$ ), which were used as primary features in ref. 127, were excluded because they are not relevant to the properties studied here. For the singlet–triplet gap model, two additional primary features were included: the DFT estimates for the single molecule and crystal singlet–triplet gap ( $\Delta E_{ST}^S$  and  $\Delta E_{ST}^C$ ). These were evaluated by subtracting the molecule/crystal

triplet formation energy, as an approximation for the triplet excitation energy, from the molecule/crystal PBE gap, as an approximation for the singlet excitation energy (*i.e.*,  $\Delta E_{ST}^S = \text{Gap}^S - E_T^S$  and  $\Delta E_{ST}^C = \text{Gap}^C - E_T^C$ ). The total number of SISO primary features was thus 14 for the singlet exciton energy, triplet exciton energy, and singlet exciton binding energy models and 16 for the singlet–triplet gap models.

In ref. 127, A subset of 10 PAH crystals of different sizes with a range of SF driving force values were completely left out of the SISO training to serve as the test set of unseen data. The same 10 structures were withheld here as an unseen validation set and kept completely separate from the remaining 91 structures, which were used for model training. An additional set of 9 hydrocarbon crystals, not included in the PAH101 set, was used to test the performance of the best SISO-generated models outside of the PAH101 set. This set, referred to as “Test 2”, includes: terylene (CSD reference code AZOXOF), benzo [e] dinaphtho[2,3-*a*; 10,20,30,40-*ghi*]fluoranthene (CSD reference code ZERXED), 7,14-diphenylnaphtho[1,2,3,4-*cde*]bisanthene (CSD reference code ZERXIH), 9,18-diphenyldibenzo[*a,o*]naphtho[1,2,3,4-*ghi*]perylene (CSD reference code ZERXON), 8,9-bis(4-methylphenyl)-10-phenylpentalen[1,2-*a*]naphthalene (CSD reference code BEGJOO), 3,6-bis-(diphenylmethylene)-1,4-cyclohexadiene (CSD reference code DUPRIP), heptafulvalene (CSD reference code HEPFUL10), 3,12-Di-*t*-butyl-2,2,13,13-tetramethyl-tetradeca-3,5,7,9,11-pentaene (CSD reference code GAFDUO), and (*E*)-1-cycloocta-tetraenyl-2-phenylethene (CSD reference code GIWHUP). GW+BSE results were published in ref. 154 for terylene, in ref. 76 for ZERXED, ZERXIH, and ZERXON, and in ref. 54 for the remaining materials. Some of the materials in the second test set are structurally very different from the materials in PAH101. HEPFUL10 and GIWHUP contain 7- and 8-membered carbon rings, respectively, and GAFDUO comprises a long polyene chain. Therefore, they can provide an estimate for the performance of the SISO models outside of the PAH101 set.



**Table 1** Description and relative cost of the primary features used to construct SISO models. Single molecule features are denoted by an "S" superscript and crystal features are denoted by a "C" superscript. The cost of each feature was evaluated as a multiple of the cost of a PBE calculation for a single molecule in the ground state, averaged over the 10 materials in the validation set

Feature	Cost	Description
Gap <sup>S</sup>	1	Single molecule gap, <i>i.e.</i> , the energy difference between the highest occupied molecular orbital (HOMO) and the lowest unoccupied molecular orbital (LUMO)
IP <sup>S</sup>	2	Single molecule ionization potential, evaluated as the total energy difference between a cation and neutral molecule
PolarTensor <sup>S</sup>	2	Trace of the molecular polarization tensor, calculated using PBE with the many-body dispersion (MBD) method <sup>155</sup>
$E_{\text{T}}^{\text{S}}$	3	Single molecule triplet formation energy, corresponding to the total energy difference between the ground-state and triplet-state molecule
EA <sup>S</sup>	3	Single molecule electron affinity, evaluated as the total energy difference between an anion and neutral molecule
$\Delta E_{\text{ST}}^{\text{S}}$	3	Estimated single molecule singlet-triplet gap, evaluated as $\text{Gap}^{\text{S}} - E_{\text{T}}^{\text{S}}$
Gap <sup>C</sup>	33	The crystal band gap, extracted from the band structure
VB <sub>disp</sub> <sup>C</sup>	33	Valence band dispersion, corresponding to the energy range of the HOMO-derived band
CB <sub>disp</sub> <sup>C</sup>	33	Conduction band dispersion, corresponding to the energy range of the LUMO-derived band
$\epsilon^{\text{C}}$	42	Dielectric constant calculated using the PBE + MBD@rSCS polarizability in the Clausius-Mossotti equation <sup>156</sup>
$H_{\text{ab}}$	93	The highest value of the transfer integral obtained out of all the unique molecular dimers extracted from the crystal, calculated with fragment orbital DFT <sup>157</sup>
$E_{\text{T}}^{\text{C}}$	148	Crystal triplet formation energy, corresponding to the total energy difference between the ground-state and triplet-state crystal
$\Delta E_{\text{ST}}^{\text{C}}$	181	Estimated crystal singlet-triplet gap, evaluated as $\text{Gap}^{\text{C}} - E_{\text{T}}^{\text{C}}$
MolWt <sup>S</sup>	0	Molecular weight in atomic mass units (amu)
$\rho^{\text{C}}$	0	Crystal density in $\text{amu} \text{ \AA}^{-3}$
AtomNum <sup>C</sup>	0	Number of atoms in the crystal unit cell

To generate the feature space, features were constructed with a maximum rung (the number of times primary features are combined) of 3 and a maximum dimension (Dim) of 4. Features were combined using the operator set  $H = \{+, -, \times, \div, \exp, \log, ()^{-1}, ()^2, ()^3, \sqrt{\cdot}, \sqrt[3]{\cdot}, \lfloor \cdot \rfloor\}$ . The maximum complexity, *i.e.*, the maximum number of operators in one combined model, was set to 10. For the singlet exciton energy, triplet exciton energy, and singlet exciton binding energy a total of roughly  $5 \times 10^2$ ,  $4 \times 10^5$ , and  $6 \times 10^{10}$  features were generated by SISO with a rung of 1, 2, and 3, respectively. For the singlet-triplet gap, a total of roughly  $6.5 \times 10^2$ ,  $7.5 \times 10^5$ , and  $2 \times 10^{11}$  features were generated by SISO with a Rung of 1, 2, and 3, respectively.

After feature generation, SISO performs linear regression to yield the model prediction, where each model is the scalar product of the SISO-generated feature with a vector of fitted coefficients. Then, the models are ranked according to their prediction performance. Sure independence screening (SIS) is used to select optimal subspaces from the huge feature space. The number of features saved after SIS was set to 20. SISO then uses  $\ell_0$ -norm minimization as a sparsifying operator (SO) to determine the sparse solution for each such subspace. For each combination of dimension and rung, 40 rounds of leave-10-out cross validation (LCV) were performed. In each round, 10 data points (out of the 91 points used for model training) were randomly selected and held out as an unseen validation set. The model with the lowest RMSE for the validation set was selected in each round. Finally, the model with the lowest RMSE for the combined training and validation data was selected out of the 40 models. This model is denoted as  $M_{\text{Dim, Rung}}$ . A full account of the SISO models for each target property is provided in the SI. For each model, the training RMSE was calculated for the whole training set with 91 structures and the test RMSE was calculated for the 10 withheld materials, which were excluded from the LCV.

The computational cost of SISO-generated models varies depending on the number and type of primary features they contain. The relative cost of each primary feature was evaluated in a similar manner to ref. 127, relative to the calculation of the single molecule gap, Gap<sup>S</sup>, whose cost is the lowest of all the primary features. The computer time required to calculate Gap<sup>S</sup> was assigned a value of 1 cost unit and the cost of other features is given in Table 1 as multiples of that unit. The cost of all the primary features has been updated to account for new developments in the latest version of FHI-aims. In particular, the many-body dispersion (MBD)<sup>155</sup> calculation has become significantly more efficient than in older versions of the code. In addition, we averaged the cost over the 10 structures in the validation set, rather than picking one system of average size, as in ref. 127. The cost of each SISO model was evaluated by summing over the costs of all the primary features included in it. The cost of features that appear in the model more than once was counted only once. The cost of features that do not require additional calculations was also counted only once. For example, the crystal band gap, Gap<sup>C</sup>, valence band dispersion, VB<sub>disp</sub><sup>C</sup>, and conduction band dispersion, CB<sub>disp</sub><sup>C</sup>, are extracted from the same band structure calculation. Therefore, if two or more of them are included in a SISO model the cost is counted only once.

A comparison of the SISO models to baseline models is provided in the ESI.† Linear regression (LR) and Gaussian process regression (GPR) models for each property were trained based on the SISO primary features and the many-body tensor representation (MBTR).<sup>158</sup> The LR and GPR models were trained using the scikit-learn<sup>159</sup> and GPyTorch<sup>160</sup> Python packages, respectively. For the singlet excitation energy, the triplet excitation energy, and the singlet-triplet gap, we also provide a comparison with a linear fit based only on the corresponding DFT-level approximations, namely the single molecule HOMO-LUMO gap and crystal band gap (Gap<sup>S</sup>, Gap<sup>C</sup>) for the singlet excitation energy, the single molecule and crystal triplet formation energy ( $E_{\text{T}}^{\text{S}}$ ,  $E_{\text{T}}^{\text{C}}$ ) for the triplet excitation energy, and



the DFT estimate for the single molecule and crystal singlet-triplet gap ( $\Delta E_{\text{ST}}^{\text{S}}, \Delta E_{\text{ST}}^{\text{C}}$ ), evaluated by calculating the difference between the aforementioned features. For the singlet exciton binding energy there is no corresponding DFT feature. In all cases, the SISO models provide better prediction performance with less over-fitting than the baseline models (see detailed discussion in the ESI†).

## Results and discussion

SISO models were trained to predict the singlet excitation energy, which corresponds to the optical gap, the triplet excitation energy, the singlet-triplet gap, and the singlet exciton binding energy. The equations of all the models, as well as tabulated values of their cost and accuracy are provided in the ESI.† We note that although in some instances models produced by symbolic regression algorithms have rediscovered or reproduced known physical relations,<sup>161</sup> this is not generally the case. Therefore, we refrain from ascribing physical meaning to the models produced by SISO.

In Fig. 2, the models generated by SISO for each property are evaluated based on considerations of cost vs. accuracy, represented by Pareto plots. For accuracy, we consider both the cross validation “train” RMSE, obtained for the training set of 91 materials, and the “test” RMSE, obtained for the unseen validation set of 10 materials. In general, the computational cost tends to increase with the model complexity because the more complex models contain more primary features. The training set RMSE generally tends to decrease with the model complexity. The validation set RMSE is generally higher than the training set RMSE. For some of the more complex models, the validation set RMSE is significantly higher than the training set RMSE, indicating over-fitting to the training data.

A detailed discussion of the results for each property is provided in the corresponding sub-sections below. For each property, we select the three models that deliver the most robust performance out of each group of models with a similar computational cost, based on the Pareto plots shown in Fig. 2. For these models, we perform a more detailed analysis of the correlation between the model predictions and the GW+BSE reference data. We examine the most significant outliers and evaluate whether the models perform robustly for the additional materials not included in the PAH101 set. The models that perform robustly for both the PAH101 set and the additional materials are selected to construct hierarchical screening workflows, as suggested in ref. 127, in which a decreasing number of candidate materials are evaluated with increasingly accurate and more expensive models. The performance of the workflows is evaluated in terms of the number of true positives and false positives that pass each filtering stage out of the entire set of 110 materials (PAH101 and the additional test set). In the Conclusion section, we remark on the overall performance of SISO for predicting the excited-state properties of crystalline organic semiconductors and provide a comparative assessment of the performance for different properties.

### Singlet excitation energy (optical gap)

Fig. 2a shows a Pareto plot of the accuracy vs. the computational cost of the SISO-generated models for the singlet excitation energy (optical gap). The models are clustered in three groups of lower, intermediate, and higher computational cost. In this case, even the models in the lower-cost group have a relatively high cost because all models contain crystal primary features. There is no model based only on single molecule primary features. This is consistent with the observation that singlet

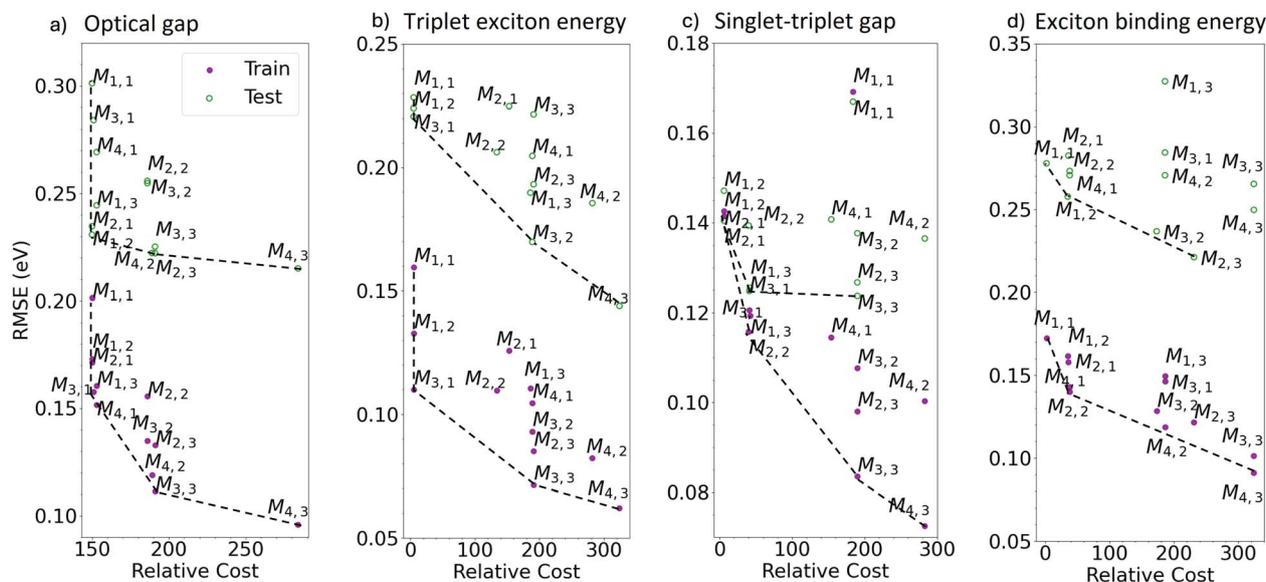


Fig. 2 Pareto charts displaying the accuracy vs. computational cost of the SISO models for (a) the singlet excitation energy (optical gap), (b) the triplet excitation energy, (c) the singlet-triplet gap, (d) the singlet excitation binding energy. The accuracy is represented by the training set cross-validation RMSE (purple filled circles) and the unseen test set RMSE (green open circles). The cost is given in multiples of the computer time required for a PBE calculation of a single molecule in the ground state. The dashed lines indicate the Pareto front.



excitons in molecular crystals are often delocalized over many molecules,<sup>54</sup> leading to a significant dependence of the singlet exciton energy on the crystal packing. The relatively expensive evaluation of the crystal triplet formation energy,  $E_T^C$ , is the main contribution to the computational cost of the lower-cost models. The models in the intermediate-cost group are more complex and contain a larger number of primary features, whose evaluation contributes to their computational cost. The higher-cost group contains only one model,  $M_{4,3}$ , which is the most complex model, whose computational cost is high owing to the need to evaluate multiple primary features, including  $H_{ab}$ , which is relatively expensive to evaluate.

We select models out of each cost group that are close to the Pareto front for both the train and test RMSE. In the lower-cost group  $M_{4,1}$  and  $M_{3,1}$  yield the lowest RMSE of 0.15 eV and 0.16 eV, respectively, for the training set. However, their performance for the test set is significantly worse with RMSEs of 0.27 eV and 0.28 eV, respectively. In comparison,  $M_{1,3}$  yields a similar train RMSE of 0.16 eV and a lower test RMSE of 0.24 eV. For this reason, we select  $M_{1,3}$  out of the lower-cost group:

$$M_{1,3} = -7.32 \times \frac{E_T^C \times \ln(\text{Gap}^S / \rho^C)}{(E_T^S - \text{IP}^S) \times E_T^S} + 1.38 \quad (1)$$

In the intermediate-cost group,  $M_{3,3}$  is on the Pareto front with a training set RMSE of 0.11 eV, significantly lower than that of the lower-cost models, and a validation set RMSE of 0.23 eV, which is somewhat lower than that of  $M_{1,3}$ :

$$M_{3,3} = -7.33 \times \frac{E_T^C \times \ln(\text{Gap}^S / \rho^C)}{(E_T^S - \text{IP}^S) \times E_T^S} - 0.939 \\ \times \frac{(EA^S + CB_{\text{disp}}^C) \times \text{AtomNum}^C}{\text{PolarTensor}^S \times (CB_{\text{disp}}^C - EA^S - \text{Gap}^S + \text{Gap}^C)} - 0.0641 \\ \times \frac{1}{\text{MolWt}^S \times |E_T^S - \text{Gap}^C| \times |(E_T^C)^2 - \text{Gap}^C \times \text{Gap}^S|} + 1.25 \quad (2)$$

The train RMSE of the high-cost model  $M_{4,3}$ , 0.10 eV, is slightly lower than the intermediate-cost models  $M_{3,3}$  and its test RMSE of 0.22 eV is similar to  $M_{3,3}$ . The need to evaluate  $H_{ab}$  is the main reason for the higher cost of  $M_{4,3}$  compared to  $M_{3,3}$ .

$$M_{4,3} = -7.57 \times \frac{E_T^C \times \ln(\text{Gap}^S / \rho^C)}{(E_T^C - \text{IP}^S) \times E_T^S} \\ - 0.000409 \times \frac{H_{ab} + CB_{\text{disp}}^C}{|E_T^S - \text{Gap}^C| \times |E_T^C / \text{Gap}^C - \text{Gap}^S / E_T^C|} \\ - 0.959 \times \frac{(EA^S + CB_{\text{disp}}^C) \times \text{AtomNum}^C}{\text{PolarTensor}^S \times (CB_{\text{disp}}^C - EA^S - \text{Gap}^S + \text{Gap}^C)} \\ - 0.0000398 \times \frac{|E_T^C - \text{Gap}^C| \times \text{MolWt}^S}{E_T^S \times |E_T^C + CB_{\text{disp}}^C - \text{Gap}^S + VB_{\text{disp}}^C|} + 1.22 \quad (3)$$

Fig. 3 shows the predictions of the SISSO models selected based on considerations of cost and accuracy as a function of the GW+BSE reference values of the singlet exciton energy (optical gap). Parity plots for all other SISSO models are

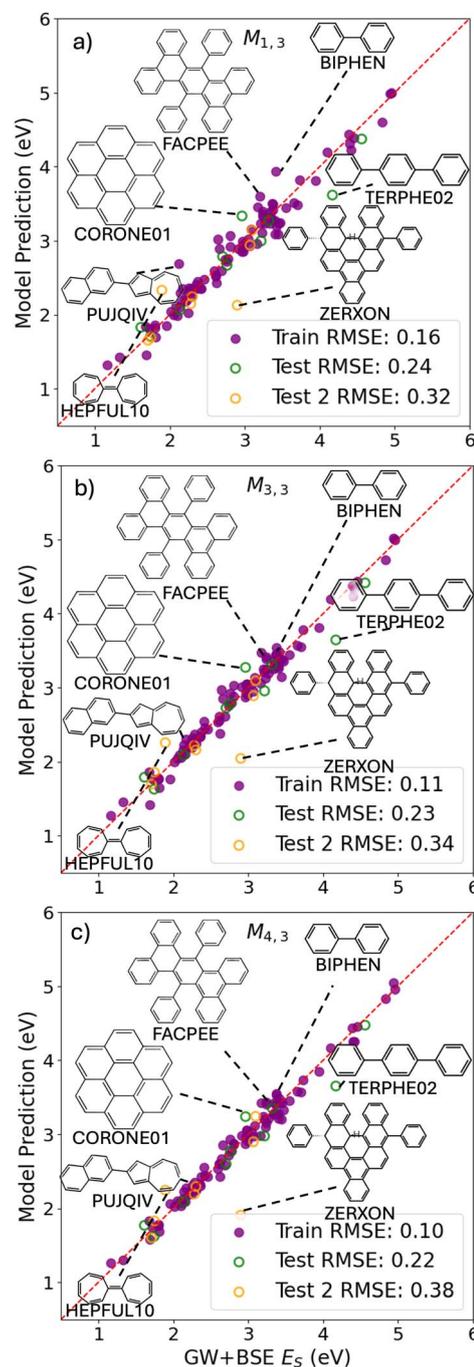


Fig. 3 SISSO model predictions as a function of the GW+BSE reference data for the singlet excitation energy (optical gap). (a)  $M_{1,3}$ , (b)  $M_{3,3}$ , and (c)  $M_{4,3}$  were selected based on considerations of cost vs. accuracy. The filled purple circles represent the training set, the open green circles represent the test set, and the open orange circles represent the additional test set not included in PAH101. Molecular structures and CSD reference codes of some of the outliers are also shown.



provided in the ESI.† Overall the model predictions are quite close to the reference values. The low-cost model  $M_{1,3}$  has a few outliers with larger deviations from the reference values. Some of the outliers are chemically and/or structurally distinct from the majority of systems in the dataset.<sup>54,127</sup> For example, 2-(naphthalen-2-yl)azulene (CSD reference code PUJQIV) contains a 7-membered ring fused with a 5-membered ring and heptafulvalene (HEPFUL10) contains two 7-membered rings. Biphenyl (CSD reference code BIPHEN) and terphenyl (CSD reference code TERPHE02) comprise benzene rings connected by single bonds, rather than the large aromatic systems of fused rings characteristic of most of the PAH101 set. For biphenyl in particular, the optical gap value of 3.41 eV included in the PAH101 dataset<sup>117</sup> is a significant underestimation compared to the experimental values of 4.1–4.18 eV.<sup>162–165</sup> Therefore, it could be argued that this data point is less reliable and the model prediction is in fact closer to experiment. Coronene (CSD reference code CORONE01), 9,18-diphenyltetrabenz(*a,c,h,j*)anthracene (CSD reference code FACPEE), and 9,18-diphenyldibenzo[*a,o*]naphtho[1,2,3,4-*ghi*]perylene (CSD reference code ZERXON) do not appear chemically or structurally distinct from most of the materials in the PAH101 set. Coronene and FACPEE have unusually high values for the crystal triplet formation energies,  $E_T^C$ , which may explain why their optical gaps are overestimated by  $M_{1,3}$ . Conversely, ZERXON has a particularly low  $E_T^C$ , which causes its optical gap to be significantly underestimated. For many of the  $M_{1,3}$  outliers out of the PAH101 set, the predictions of the intermediate-cost  $M_{3,3}$  and higher-cost  $M_{4,3}$  models are closer to the reference values, with the exception of terphenyl, which remains an outlier. For the additional test set, which was not included in PAH101, the RMSE deteriorates with the model complexity, which indicates over-fitting.

The PAH101 set contains materials with a wide range of optical gaps, as shown in Fig. 1c. Crystalline quaterrylene (QUATER10) and hexacene (ZZZDKE01) have the smallest optical gaps of 1.33 eV and 1.17 eV, respectively. Fig. 4 demonstrates a two-stage screening workflow constructed based on the SISO models selected for the optical gap. The first stage is  $M_{1,3}$  and the second stage is  $M_{3,3}$ . We have decided not to use  $M_{4,3}$  because of its markedly worse performance for the Test 2 set. For demonstration purposes, we screen for materials with an optical gap in the range of 2.2–2.5 eV, corresponding to green color. There are 17 such materials in total in the

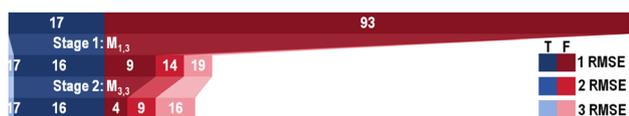


Fig. 4 A two-stage screening workflow for materials with an optical gap in the green range of 2.2–2.5 eV. The first stage is  $M_{1,3}$  and the second stage is  $M_{3,3}$ . The number of true positives (shades of blue) and the number of false positives (shades of red/pink) that pass each stage of screening is shown when the thresholds are set to one, two, and three times the training set RMSE of each model. In each case,  $n/2 \times$  RMSE is applied on either end of the target energy range.

combined set of PAH101 and the additional test set. The screening thresholds for each stage of the workflow are set to  $2.2 \text{ eV} - 0.5 n \times \text{RMSE}$  to  $2.5 \text{ eV} + 0.5 n \times \text{RMSE}$ , where  $n = 1, 2, 3$  and RMSE refers to the training set RMSE of each model.

With  $n = 1$ , the first-stage model,  $M_{1,3}$ , screens out most of the materials, whose optical gap is not within the target range, leaving 9 false positives. 16 of the 17 materials, whose gaps are in the target range are successfully identified. Further screening by the second-stage model,  $M_{3,3}$ , significantly reduces the number of false positives from 9 to 4, while retaining the 16 true positives. Setting  $n = 2$  does not improve the performance of the workflow because the final outcome is the same 16 true positives only with a higher number of false positives. With  $n = 3$  all 17 true positives pass the screening along with 19 and 16 false positives after the first and second stage of screening, respectively. Using the same workflow to screen for materials with optical gaps in the red range and blue range produces very similar results, as shown in the ESI.† Based on this, we would suggest a two-stage workflow with thresholds defined depending on one's tolerance for false positives. We also note that the optical gaps obtained from GW+BSE are typically within 0.2 eV from experiment.<sup>117</sup> The training RMSE values of  $M_{1,3}$  and  $M_{3,3}$  are 0.16 eV and 0.11 eV, respectively. Therefore, it may be reasonable to set  $n = 2$  to account for the errors of GW+BSE on top of the errors of the ML models.

### Triplet excitation energy

Fig. 2b shows a Pareto plot of the accuracy vs. the computational cost of the SISO-generated models for the triplet excitation energy. Similar to the results for the optical gap, the triplet exciton energy models are clustered in three groups of lower, intermediate, and higher computational cost. In contrast to the optical gap models, in this case, the models in the lower-cost group only require the calculation of single molecule primary features, which makes them very cheap to evaluate. This is consistent with the observation that triplet excitons in molecular crystals are typically localized on one molecule (in contrast to singlet excitons),<sup>54</sup> leading to a weaker dependence on the crystal packing. The models in the intermediate-cost group are more complex and contain various crystal primary features. The higher-cost group contains the higher complexity fourth rung models,  $M_{4,2}$  and  $M_{4,3}$ . These models contain both of the high-cost features  $E_T^C$  and  $H_{ab}$ , as well as multiple other primary features, which contribute to their high cost.

We select models out of each cost group that are close to the Pareto front for both the train and test RMSE. Of the lower-cost group  $M_{3,1}$  yields the lowest RMSEs of 0.11 eV and 0.22 eV, respectively, for the training set and the validation set.

$$M_{3,1} = 6.5 \times (E_T^S / \text{IP}^S) + 0.00051 \times (E_T^S \times \text{AtomNum}^C) + 0.0091 \times (e^{E_T^S}) - 0.052 \quad (4)$$

None of the models in the intermediate-cost group is on the Pareto front for both the training set and the validation set.  $M_{3,3}$  yields the lowest RMSE of 0.07 eV for the training set, however



its performance deteriorates considerably for the validation set with an RMSE of 0.22 eV. The next best model for the training set,  $M_{2,3}$ , with an RMSE of 0.09 eV also performs significantly worse for the validation set with an RMSE of 0.19 eV. We therefore choose the third model,  $M_{3,2}$ , because its RMSE of 0.09 eV for the training set is similar to  $M_{2,3}$ , but its performance for the validation set is markedly better with an RMSE of 0.17 eV.

$$M_{3,2} = 0.079 \times ((E_T^C + E_T^S) \times \ln(\text{PolarTensor}^S)) + 55 \times \frac{(\text{CB}_{\text{disp}}^C)^3}{\text{EA}^S \times (\text{AtomNum}^C)} - 0.0000053 \times ((\text{Gap}^C - \text{Gap}^S) \times (\text{AtomNum}^C)^2) - 0.035 \quad (5)$$

Of the high-cost group,  $M_{4,3}$  is on the Pareto front for both the training set and the validation set with RMSEs of 0.06 eV and 0.14 eV, respectively.

$$M_{4,3} = 0.093 \times ((E_T^C + \text{CB}_{\text{disp}}^C) \times \ln(\text{PolarTensor}^S) - (\text{CB}_{\text{disp}}^C - E_T^S) \times \ln(\text{AtomNum}^C)) + 83 \times \frac{(\text{CB}_{\text{disp}}^C \times \varepsilon^C) / (\text{AtomNum}^C)^2}{(\text{EA}^S)^2 - (H_{\text{ab}} \times \text{Gap}^C)} + 0.0000039 \times \frac{(\text{EA}^S - \text{Gap}^S) \times (\text{VB}_{\text{disp}}^C \times \text{PolarTensor}^S)}{|(E_T^C + \text{CB}_{\text{disp}}^C) - (\text{Gap}^S - \text{VB}_{\text{disp}}^C)|} + 0.080 \times \left| \frac{E_T^S \times \text{CB}_{\text{disp}}^C}{\text{Gap}^S - E_T^C} - ((H_{\text{ab}} + \text{CB}_{\text{disp}}^C) \times \ln(\text{PolarTensor}^S)) \right| - 0.082 \quad (6)$$

Fig. 5 shows the predictions of the SISSO models selected based on considerations of cost and accuracy as a function of the GW+BSE reference values of the triplet exciton energy. Parity plots for all other SISSO models are provided in the ESI.† Similar to the optical gap, the SISSO model predictions are quite close to the reference values. The low-cost model  $M_{3,1}$  has a few outliers with larger deviations from the reference values. Biphenyl and terphenyl are significant outliers, similar to the optical gap models. 1,2,3,4-Tetraphenylbenzene (CSD reference code FOVVOB) also comprises benzene rings connected by single bonds. Biphenyl, 9,9'-bianthracenyl (CSD reference code KUBWAF01), hexabeno(bc,ef,hi,kl,no,qr)coronene (CSD reference code HBZCOR), and 1,12-benzoperylene (CSD reference code BNPERY) have unusually high single molecule triplet formation energy,  $E_T^S$ , values exceeding the GW+BSE reference values of their crystal triplet excitation energies, which causes the  $M_{3,1}$  predictions to be overestimated. 7,14-Diphenylnaphtho[1,2,3,4-cde]bisanthene (CSD reference code ZERXIH) has an unusually large value for the trace of the molecular polarization tensor  $\text{PolarTensor}^S$ , which causes it to become a significant outlier for  $M_{3,2}$ . Interestingly, the lowest-cost model  $M_{3,1}$  has

a lower RMSE for the Test 2 set than for the PAH101 held out validation set. In contrast, the performance of the more complex models  $M_{3,2}$  and  $M_{4,3}$  deteriorates very significantly for the additional test set, indicating over-fitting.

The PAH101 set contains materials with a wide range of triplet excitation energies, as shown in Fig. 1d. Fig. 6 demonstrates a one-stage screening workflow for the triplet exciton energy based on the  $M_{3,1}$  SISSO model, which was selected because it retains robust performance for the additional test set, unlike the more complex models. Because of the recent interest in TTA up-conversion of infrared (IR) light to the visible range,<sup>71</sup> we screen for materials, whose triplet excitation energy is in the IR, below 1.6 eV. There are 37 such materials in total in the combined set of PAH101 and the additional test set. The screening thresholds are set to 1.6 eV +  $n \times \text{RMSE}$ , where  $n = 1, 2, 3$  and RMSE refers to the training set RMSE of  $M_{3,1}$ , 0.11 eV. With  $n = 1$ , the  $M_{3,1}$  model correctly classifies all 37 materials with triplet excitation energy in the IR with only 3 false positives. Setting  $n$  to 2 or 3 is not beneficial because it only

increases the number of false positives. In this case, screening only with  $M_{3,1}$  delivers robust performance.

Next, we assess whether the 37 materials in the combined set of PAH101 and the additional test set with triplet excitation energies in the IR are likely to undergo TTA. The thermodynamic driving force for TTA is given by the difference between twice the triplet excitation energy and the singlet excitation energy (the opposite of the SF driving force). As discussed in ref. 54,63 and 127, GW+BSE systematically underestimates the SF driving force and therefore overestimates the TTA driving force. Hence, we assess the likelihood of a given material to undergo SF/TTA relative to other known SF/TTA materials. Only 8 out of the 37 materials have a higher TTA driving force than rubrene and perylene: pyracylene (KEGHEJ01), 11-phenylbenzo[*a*]naphtho[2,1,8-*cde*]perylene (KAGFUV), the two polymorphs of diindeno[1,2,3-*cd*:1',2',3'-*lm*]perylene (POBPIG and POBPIG06), anthra(2,1,9,8-*hijkl*)benzo(*de*)naphtho(2,1,8,7-*stuv*)pentacene (BOXGAW), dibenzo(*def,i*)naphtho(1,8,7-*v,w,x*)pyranthrene (DUPHAX), benzo[*e*]dinaphtho[2,3-*a*; 10,20,30,40-*ghi*]fluoranthene (ZERXED), and heptafulvalene (HEPFUL10). All of these have GW+BSE triplet excitation energies in the near-IR



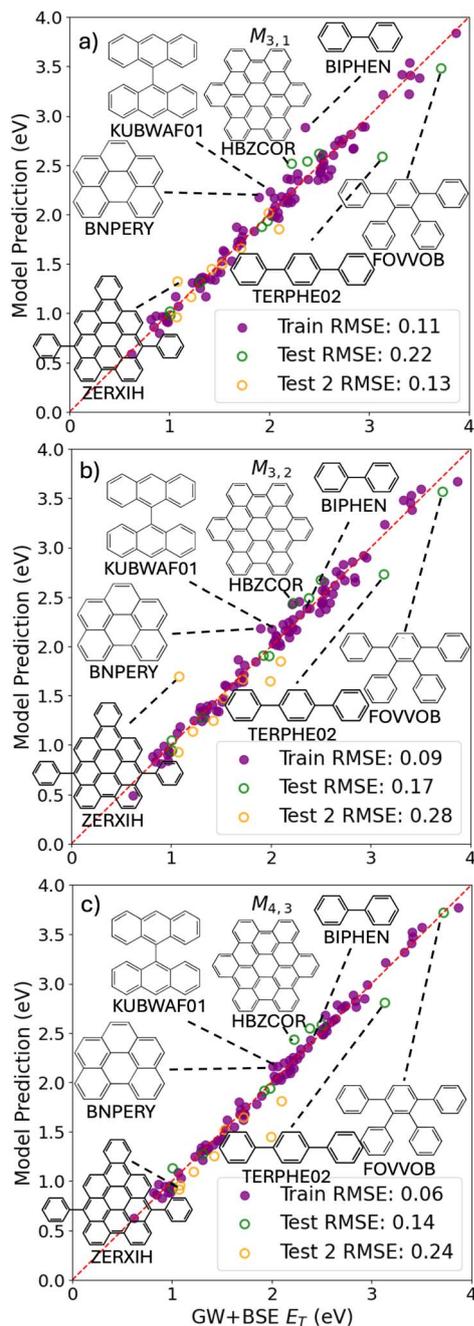


Fig. 5 SISSO model predictions as a function of the GW+BSE reference data for the triplet excitation energy. (a)  $M_{3,1}$ , (b)  $M_{3,2}$ , and (c)  $M_{4,3}$  were selected based on considerations of cost vs. accuracy. The filled purple circles represent the training set, the open green circles represent the test set, and the open orange circles represent the additional test set not included in PAH101. Molecular structures and CSD reference codes of some of the outliers are also shown.

and their GW+BSE optical gaps range from 1.89 eV for crystal-line heptafulvalene to 2.41 eV for KAGFUV. Materials in the PAH101 set with lower triplet excitation energies tend to be more likely to undergo SF than TTA. Five of these compounds have been previously assessed in ref. 63 as isolated molecules rather than crystals, using a different GW+BSE implementation.



Fig. 6 A one-stage screening workflow for materials with a triplet excitation energy in the infrared, below 1.6 eV, based on the  $M_{3,1}$  model. The number of true positives (shades of blue) and the number of false positives (shades of red/pink) that pass the screening is shown when the thresholds are set to one, two, and three times the training set RMSE of the model.

Therein, the likelihood of the competing process of conversion of two excitons in the first triplet state to one exciton in a higher triplet state instead of a singlet exciton was also considered. Based on these energetic considerations, pyracylene (labeled as compound C1 in ref. 63) appeared promising, however it is not a good TTA chromophore because of its short triplet state lifetime and the rapid non-radiative decay of its lowest singlet state.<sup>166</sup> POBPIG is a perylene derivative (labeled as compound D16 in ref. 63), for which the competing conversion to a higher triplet state was found to be too energetically favorable for high-yield TTA. KAGFUV, BOXGAW, and DUPHAX are large aromatic compounds comprising several fused six-membered rings that can be classified as graphene flakes. Of these, KAGFUV (labeled as compound E2 in ref. 63) and DUPHAX (labeled as compound E3 in ref. 63) were identified as TTA candidates based on energetic considerations. ZERXED has been evaluated based on the same criteria in ref. 76 (therein it was labeled “Compound I”) and found to be a promising TTA candidate. Heptafulvalene is reported here for the first time as a potential TTA candidate and could indicate an interesting new direction of exploring compounds with 7-membered rings as TTA/SF candidates.

The workflow presented here demonstrates how ML models can be used to identify new potential TTA candidates with triplet exciton energies in the IR, optical gaps in the visible range, and favorable TTA energetics. We note that the optical gap SISSO models presented above and the SF driving force SISSO models from ref. 127 could be used to perform further fast screening of materials that pass the triplet exciton energy screening. Thus, the number of candidates selected for computationally expensive excited-state calculations can be significantly reduced.

### Singlet triplet gap

Fig. 2c shows a Pareto plot of the accuracy vs. the computational cost of the SISSO-generated models for the singlet–triplet gap. The singlet–triplet gap models are clustered in four groups of very-low, low, intermediate, and higher computational cost. The very-low cost group contains models, whose evaluation only requires the calculation of single molecule primary features. The low-cost group contains models, whose evaluation only requires calculating the crystal band gap,  $\text{Gap}^C$ , in addition to single molecule features. Models in the intermediate cost group require the calculation of the crystal triplet formation energy,  $E_{\text{T}}^C$ , and/or the estimated crystal singlet–triplet gap,  $\Delta E_{\text{ST}}^C$ . The models in the high-cost group also contain  $H_{\text{ab}}$ .



We select models out of each cost group that are close to the Pareto front for both the train and test RMSE. In the very-low-cost group  $M_{1,2}$  and  $M_{2,1}$  contain the same primary features and have a similar performance with train RMSE of 0.14 eV and test RMSEs of 0.15 eV and 0.14 eV, respectively. Therefore we use an ensemble of the two models:

$$\begin{aligned} M_{1,2} \& M_{2,1} &= \frac{1}{2} \times M_{1,2} + \frac{1}{2} \times M_{2,1} \\ &= \frac{1}{2} \times \left( -0.41 \times \left( (\rho^C)^2 \times (EA^S + E_T^S) \right) + 1.6 \right) \\ &+ \frac{1}{2} \times \left( -3.6 \times \left( \sqrt{\rho^C} \right) - 0.26 \times (EA^S + E_T^S) + 4.83 \right) \end{aligned} \quad (7)$$

In the low-cost group  $M_{2,2}$  yields the lowest RMSE of 0.12 eV for the training set. The RMSE of 0.14 eV for validation set is similar to the lowest cost models.

$$\begin{aligned} M_{2,2} &= -0.52 \times (\rho^C \times (EA^S + \text{Gap}^S)) \\ &- 0.036 \times \frac{\text{Gap}^S / \Delta E_{ST}^S}{(\text{Gap}^C)^2} + 2.3 \end{aligned} \quad (8)$$

In the intermediate-cost group  $M_{3,3}$  is on the Pareto front for both the training set and the validation set with RMSEs of 0.08 eV and 0.12 eV, respectively.

$$\begin{aligned} M_{3,3} &= -121 \times \frac{\sqrt{\text{Gap}^C} / e^{\rho^C}}{(\Delta E_{ST}^S - E_T^S) - (EA^S + E_T^S)} \\ &- 0.0021 \frac{(EA^S - \text{Gap}^C) / (\Delta E_{ST}^C + \Delta E_{ST}^S)}{(E_T^S - \text{Gap}^C) / (E_T^C - E_T^S)} \\ &+ 0.071 \frac{(EA^S \times \Delta E_{ST}^C) / (\Delta E_{ST}^C + \text{CB}_{\text{disp}}^C)}{(E_T^C / EA^S) + (EA^S / \Delta E_{ST}^C)} - 0.73 \end{aligned} \quad (9)$$

In the high-cost group  $M_{4,3}$  has the lowest RMSE of 0.07 eV for the training set, however its performance deteriorates very significantly to 0.3 eV for the validation set, indicating overfitting.

Fig. 7 shows the predictions of the SISSO models selected based on considerations of cost and accuracy as a function of the GW+BSE reference values of the singlet–triplet gap. Parity plots for all other SISSO models are provided in the ESI.† We note that the energy range of GW+BSE reference data for the singlet–triplet gap is significantly smaller than the energy range of the singlet and triplet exciton energies (see the distributions in Fig. 1). Therefore, the deviations of the SISSO model predictions appear larger in comparison to the reference data even though they are of comparable magnitude or even smaller than the deviations of the SISSO models for the singlet and triplet exciton energies. In the discussion of the outliers we focus on materials with relatively small singlet–triplet gaps because these would be of most interest for OLEDs. The very-low-cost model  $M_{1,2}$  &  $M_{2,1}$  has several significant outliers in the range of singlet–triplet gaps below 0.6 eV. These include materials that are also among the outliers for the singlet and triplet exciton energy models, PUJQIV, FACPEE, HEPFUL10, and

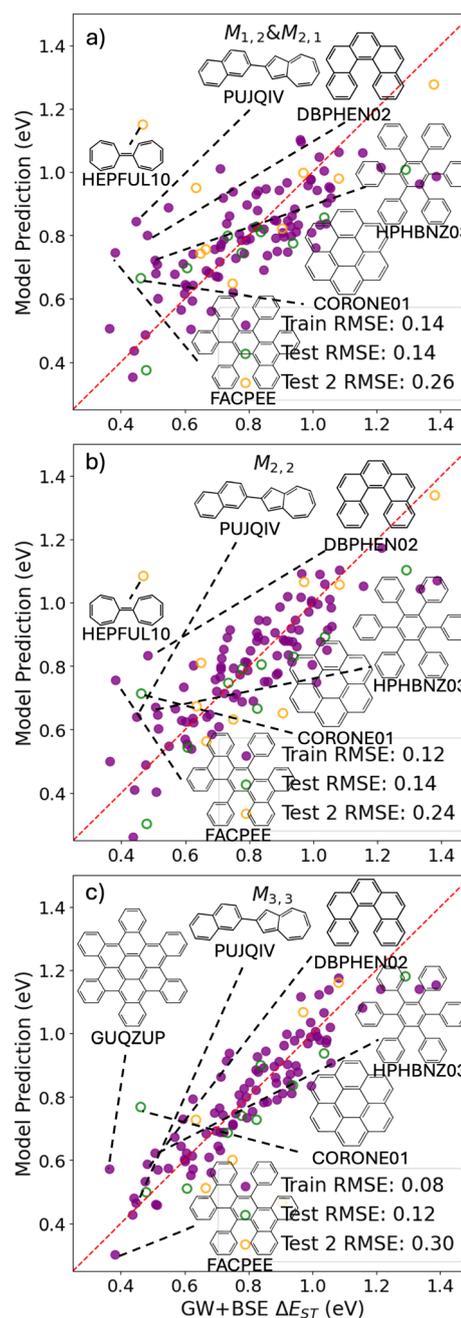


Fig. 7 SISSO model predictions as a function of the GW+BSE reference data for the singlet–triplet gap. (a) Ensemble model of  $M_{1,2}$  &  $M_{2,1}$ , (b)  $M_{2,2}$ , and (c)  $M_{3,3}$  were selected based on considerations of cost vs. accuracy. The filled purple circles represent the training set, the open green circles represent the test set, and the open orange circles represent the additional test set not included in PAH101. Molecular structures and CSD reference codes of some of the outliers are also shown.

coronene. Hexaphenylbenzene (CSD reference code HPHBNZ03) comprises phenyl rings connected by single bonds like some of the outliers for the singlet and triplet excitation energies.

Trinaphtho[1,2,3,4-*fg*h:1',2',3',4'-*pqr*:1'',2'',3'',4''-*za*\_1\_b\_1\_]trinaphthylene (CSD reference code GUQZUP) and (5)helicene (CSD reference code DBPHEN02) are not structurally



or chemically distinct from most of the materials in the PAH101 set. In addition, they do not have any primary features with unusual values. It is possible that the relatively large number of outliers, especially in the low singlet–triplet gap range, is a reflection of the difficulty of training reliable models to predict very small target values, in particular considering that there are few materials with a singlet–triplet gap below 0.5 eV in the training data (See Fig. 1e). The performance of all three models for the Test 2 set is worse than for the PAH101 validation set. In particular, for the more complex model,  $M_{3,3}$ , the Test 2 RMSE is significantly higher than the lower cost models. For HEPFUL10,  $M_{3,3}$  even predicts a negative value. The poor performance of  $M_{3,3}$  for the Test 2 set indicates overfitting.

Fig. 1e shows the distribution of singlet–triplet gaps in the PAH101 dataset. Small singlet–triplet gaps are rare among this class of materials. The materials with lowest singlet–triplet gaps (in parentheses) are: trinaphtho[1,2,3,4-fgh:1',2',3',4'-pqr:1'',2'',3'',4''-za\_1\_b\_1\_]trinaphthylene (GUQZUP; 0.36 eV), 9,18-diphenyltetrabenz[*a,c,h,i*]anthracene (FACPEE; 0.38 eV), acenaphtho[3,2,1,8-fghij]tetrabenz[*a,c,m,o*]picene (VUFHUA; 0.435 eV), benzo(1,2,3-*bc*:4,5,6 *b',c'*)dicononene (YOFCUR; 0.44 eV), and 2-(naphthalen-2-yl)azulene (PUJQIV; 0.45 eV). Even the lowest singlet–triplet gaps in the PAH101 set would be considered marginal or too high for TADF. It is interesting to note that with the exception of PUJQIV (shown in Fig. 3 and 7), the materials with the smallest singlet–triplet gaps in the PAH101 set bear no resemblance to the donor–acceptor compounds typically used for TADF.<sup>47,167</sup> Rather, they are large PAHs with extended  $\pi$  systems. FACPEE (shown in Fig. 3 and 7), VUFHUA, and YOFCUR have segments that could lead to charge-transfer-like intramolecular excitations. GUQZUP can be described as a graphene flake with no obvious segments. The twisted conformation it adopts in the crystal structure may contribute to orbital localization and charge-transfer-like excitations. The effect of crystal packing and intermolecular vs. intramolecular charge-transfer excitations on singlet–triplet gaps is not well-understood and should be further investigated in relation to TADF in crystalline materials.<sup>168,169</sup>

Fig. 8 demonstrates a two-stage screening workflow constructed based on the SISO models selected for the singlet–triplet gap. The  $M_{1,2}$  &  $M_{2,1}$  and  $M_{2,2}$  models were selected for the first and second stages of the workflow, respectively, based on their performance for the Test 2 set. For demonstration purposes, we screen for materials with a singlet–triplet gap below 0.5 eV. There are 10 such materials in total in the

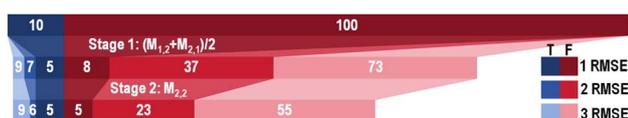


Fig. 8 A two-stage screening workflow for materials with a singlet–triplet gap below 0.5 eV. The first stage is an ensemble model of  $M_{1,2}$  &  $M_{2,1}$  and the second stage is  $M_{2,2}$ . The number of true positives (shades of blue) and the number of false positives (shades of red/pink) that pass each stage of screening is shown when the thresholds are set to one, two, and three times the training set RMSE of each model.

combined set of PAH101 and the additional test set. The screening thresholds for each stage of the workflow are set to  $0.5 \text{ eV} + n \times \text{RMSE}$ , where  $n = 1, 2, 3$  and RMSE refers to the training set RMSE of each model. With  $n = 1$ , the first-stage ensemble model of  $M_{1,2}$  &  $M_{2,1}$  screens out most of the materials, whose singlet–triplet gap is above 0.5 eV, leaving only 8 false positives. However, 5 of the 10 materials whose singlet–triplet gaps are below 0.5 eV are also screened out, leaving only 5 true positives. Further screening by the second-stage model,  $M_{2,2}$  reduces the number of false positives to 5 without losing any additional true positives. Setting  $n = 2$  results in 7 and 6 true positives passing the first and second stage, respectively, with a significantly higher number (23) of false positives passing the screening. With  $n = 3$ , 9 true positives pass the screening along with 55 false positives. One may define the screening thresholds depending on their tolerance for false positives. With the thresholds set to one RMSE, the workflow presented here effectively eliminates most of the materials that are not of interest, which significantly reduces the number of candidates that need to be evaluated using more accurate and computationally expensive methods. Training better models for predicting the singlet–triplet gap would require acquiring more GW+BSE data for materials with small singlet–triplet gaps. To this end, the present models could be used to select materials for data acquisition.

### Singlet exciton binding energy

Fig. 2d shows a Pareto plot of the accuracy vs. the computational cost of the SISO-generated models for the singlet exciton binding energy. We note that terphenyl (TERPHE02) is a major outlier for all the exciton binding energy models (see parity plots provided in the ESI†). Therefore, we also consider the validation RMSE without terphenyl. The SISO models are clustered in four cost groups. The very-low-cost model  $M_{1,1}$  only requires a calculation of the molecular ionization potential,  $\text{IP}^S$ . The models in the low-cost group require a crystal band structure calculation in addition to single molecule features. Most of the models in the intermediate cost group include the crystal triplet formation energy,  $E_T^C$ , except for  $M_{3,2}$ , which includes  $H_{ab}$  and the crystal dielectric constant,  $\epsilon^C$ . The models in the high-cost group contain  $E_T^C$  and  $H_{ab}$ .

We select models out of each cost group that are close to the Pareto front for both the train and test RMSE. The very-low-cost model  $M_{1,1}$  yields RMSEs of 0.17 eV for the training set and 0.28 eV for the validation set (0.19 eV without terphenyl).

$$M_{1,1} = 0.24 \times (\text{IP}^S/\rho^C) - 1.35 \quad (10)$$

Of the low-cost group,  $M_{4,1}$  and  $M_{2,2}$  yield the same RMSEs of 0.14 eV for the training set and 0.27 eV for the test set.  $M_{2,2}$  delivers a slightly better performance for the validation set without terphenyl with an RMSE of 0.18 eV, compared to 0.20 eV for  $M_{4,1}$ . Therefore, we select  $M_{2,2}$ :

$$M_{2,2} = 0.012 \times (E_T^S + \text{IP}^S) \times (\text{IP}^S/\rho^C) + 0.020 \\ \times ((E_T^S - \text{Gap}^S) \times (\text{CB}_{\text{disp}}^C \times \text{AtomNum}^C)) - 0.17 \quad (11)$$



Of the intermediate-cost group  $M_{4,2}$  yields the lowest RMSE of 0.12 eV for the training set, however its performance deteriorates significantly for the validation set with an RMSE of 0.27 eV (0.18 eV without terphenyl), indicating over-fitting. The next model  $M_{2,3}$  has a more robust performance with RMSEs of 0.12 eV for the training set and 0.22 eV for the validation set (0.13 eV without terphenyl).

$$M_{2,3} = 5.2 \times \frac{(\text{Gap}^S)^2 - (\text{IP}^S \times \text{CB}_{\text{disp}}^C)}{\sqrt[3]{\text{MolWt}^S \times (\text{Gap}^S \times \epsilon^C)}} + 0.033 \times \frac{(E_{\text{T}}^C - E_{\text{T}}^S)/(E_{\text{T}}^C - \text{Gap}^S)}{\ln\left(\frac{\text{EA}^S - \text{CB}_{\text{disp}}^C}{\text{EA}^S}\right)} + 0.24 \quad (12)$$

Both models in the high-cost group are over-fitted, as indicated by the large difference between their performance for the training set vs. the validation set. Therefore, these overly complex models are not useful.

Fig. 9 shows the predictions of the SISSO models selected based on considerations of cost and accuracy as a function of the GW+BSE reference values of the singlet exciton binding energy. Parity plots for all other SISSO models are provided in the ESI.† We note that, similar to the singlet-triplet gap, the energy range of the GW+BSE reference data for the singlet exciton binding energy is significantly smaller than the energy range of the singlet and triplet exciton energies (see the distributions in Fig. 1). Therefore, the deviations of the SISSO model predictions appear larger in comparison to the reference data even though they are of comparable magnitude or even smaller than the deviations of the SISSO models for the singlet and triplet exciton energies. Terphenyl is a major outlier to the point that it noticeably skews the test RMSE. This is perhaps not surprising because terphenyl is an outlier for most other properties considered here. Some of the outliers encountered for other properties are also outliers for the singlet exciton binding energy models, including PUJQIV and FACPEE (not shown), as well as HEPFUL10. Some of the outliers, such as  $\beta$ -tribenzopyrene (CSD reference code TBZPYR) and dinaphtho(1,2-a:1',2'-h)anthracene (CSD reference code DNAPAN), do not appear structurally or chemically distinct from most of the materials in the PAH101 set. For  $M_{1,1}$ , a relatively high value of the single molecule ionization potential,  $\text{IP}^S$ , combined with a low crystal density,  $\rho^C$  can lead to significant overestimation (e.g., for terphenyl, DNAPAN, and GAFDUO). Conversely, a low  $\text{IP}^S$  value combined with a high  $\rho^C$  can lead to significant underestimation (e.g., for TBZPYR, BEGJOO and HEPFUL10). The relatively large number of persistent outliers for the singlet exciton binding energy models could be a reflection of the difficulty of training reliable models to predict very small target values. The simpler models  $M_{1,1}$  and  $M_{2,2}$  have a lower RMSE for the Test 2 set than for the PAH101 validation set, whereas the more complex model  $M_{2,2}$  has a significantly higher RMSE for the Test 2 set, indicating over-fitting.

Fig. 1f shows the distribution of singlet exciton binding energies in the PAH101 dataset. In most organic materials the exciton binding energy is significant compared to inorganic materials because the dielectric screening of charges is not as strong. However, some materials in the PAH101 set have low exciton binding energies (in parentheses), including: anthra(2,1,9,8-*hijkl*)benzo(2,1,8,7-*stuv*)pentacene (BOXGAW; 0.013 eV), dinaphtho(1,2-*a:1',2'-h*)anthracene (DNAPAN; 0.071 eV), tetrabenzo(2,1,8,7-*stuv*)heptacene (TBZHCE; 0.130 eV), benzo[*lm*]chryseno[1,12,11,10-*opqrab*]

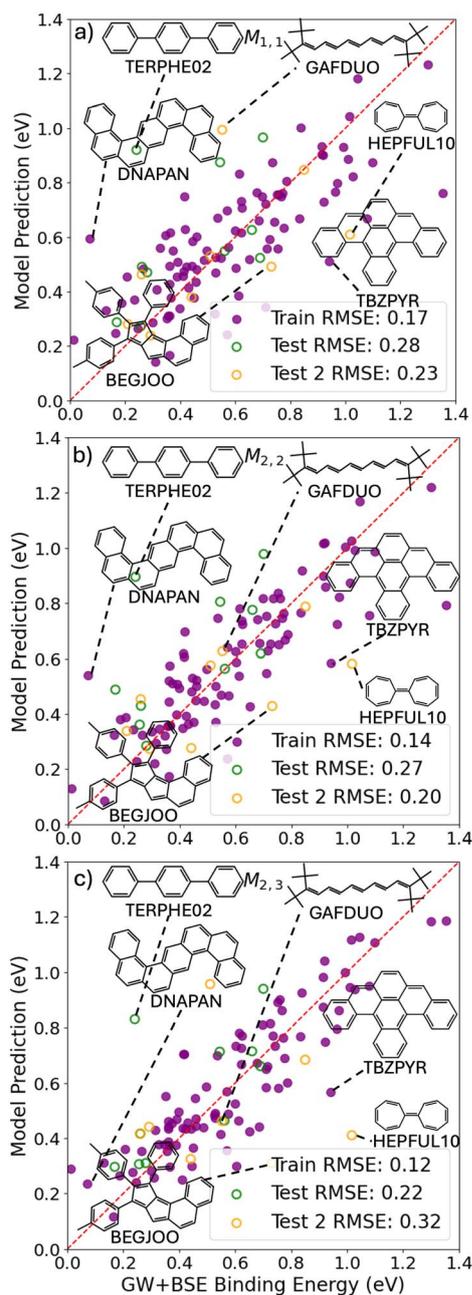


Fig. 9 SISSO model predictions as a function of the GW+BSE reference data for the singlet exciton binding energy. (a)  $M_{1,1}$ , (b)  $M_{2,2}$ , and (c)  $M_{2,3}$  were selected based on considerations of cost vs. accuracy. The filled purple circles represent the training set, the open green circles represent the test set, and the open orange circles represent the additional test set not included in PAH101. Molecular structures and CSD reference codes of some of the outliers are also shown.





Fig. 10 A two-stage screening workflow for materials with a singlet exciton binding energy below 0.3 eV. The first stage is  $M_{1,1}$  and the second stage is  $M_{2,2}$ . The number of true positives (shades of blue) and the number of false positives (shades of red/pink) that pass each stage of screening is shown when the thresholds are set to one, two, and three times the training set RMSE of each model.

perylene (YUNYAJ; 0.165 eV), and hexabenzobenzene (*bc,ef,hi,kl,no,qr*) coronene (HBZCOR; 0.169 eV). All of these compounds are characterized by very extended and/or elongated  $\pi$  systems, which likely lead to an already low molecular exciton binding energy (not calculated here), further reduced by dielectric screening in the solid form.

Fig. 10 demonstrates a two-stage screening workflow constructed based on the SISO models selected for the singlet exciton binding energy. The  $M_{1,1}$  and  $M_{2,2}$  models have been selected for the first and second stage, respectively, based on their robust performance for the Test 2 set. A small exciton binding energy means it is easier to separate excitons into free charge carriers. It is also often associated with strong dielectric screening and better charge transport. For demonstration purposes, we screen for materials with a singlet exciton binding energy below 0.3 eV. There are 17 such materials in total in the combined set of PAH101 and the additional test set. The screening thresholds for each stage of the workflow are set to  $0.3 \text{ eV} + n \times \text{RMSE}$ , where  $n = 1, 2, 3$  and RMSE refers to the training set RMSE of each model. With  $n = 1$ , the first-stage model,  $M_{1,1}$ , screens out many of the materials, whose exciton binding energy is above 0.3 eV, leaving 21 false positives. However, 4 of the 17 materials with exciton binding energy below 0.3 eV are also screened out, leaving 13 true positives. Further screening by the second-stage model,  $M_{2,2}$  reduces the number of false positives to 15 with two additional true positives lost. Setting  $n = 2$  results in 16 out of 17 true positives (all except terphenyl) passing the screening, but with a high number (41) of false positives. Setting  $n = 3$  is not beneficial because the same 16 true positives pass the screening (terphenyl is still misclassified) along with 57 false positives. One may define the screening thresholds depending on their tolerance for false positives. With the thresholds set to one RMSE, the workflow presented here effectively eliminates most of the materials that are not of interest, which significantly reduces the number of candidates that need to be evaluated using more accurate and computationally expensive methods. Training better models for predicting the singlet exciton binding energy would require acquiring more GW+BSE data for materials with small exciton binding energies. To this end, the present models could be used to select materials for data acquisition.

## Conclusion

In summary, we used the PAH101 dataset of GW+BSE calculations to train SISO models to predict the first singlet excitation energy, which corresponds to the optical gap, the first triplet

excitation energy, the singlet–triplet gap, and the singlet exciton binding energy of organic molecular crystals. SISO models were selected based on considerations of accuracy and computational cost to design materials screening workflows for each property. The screening targets were chosen to demonstrate typical use-cases relevant for organic electronic devices. We demonstrated screening for materials with an optical gap in a particular color; materials meeting the requirements for up-conversion of infrared light to the visible range *via* TTA, namely, a triplet excitation energy in the infrared, optical gap in the visible range, and a higher TTA driving force than rubrene; materials with a small singlet–triplet gap, which is desirable for OLEDs; and materials with a small singlet exciton binding energy to facilitate the separation of excitons into free charge carriers.

For all four properties, the workflows based on SISO models can effectively screen out most of the materials that are not of interest. However, the classification performance varies across properties and screening targets in terms of the number of false positives that pass the workflow and the number of true positives missed. The workflows for the optical gap and triplet excitation energy yield a more robust performance than the workflows for the singlet–triplet gap and exciton binding energy. The SISO models for the triplet exciton energy performed particularly well. The lowest cost model, which only requires evaluating single molecule DFT features, successfully classified all the materials with triplet exciton energies in the IR with a very small number of false positives. This analysis helped identify compounds with 7-membered rings, such as heptafulvalene, as a potential new direction to be explored for TTA/SF.

The overall narrow energy range of the singlet–triplet gap and exciton binding energy in the PAH101 dataset, as well as the small number of samples with the desirable low values, made it more challenging to train reliable models for these properties. Improving the models would require additional data acquisition. To this end, the present models could be used to select materials likely to have low singlet–triplet gaps or low exciton binding energy for additional GW+BSE calculations. We also note that ML models trained on PAH101 are not guaranteed to perform well for materials that are significantly different chemically or structurally. This has been demonstrated here and in ref. 54 by the worse performance of some of the more complex SISO models for an additional test set of 9 hydrocarbon crystals. Therefore, we recommend carefully validating the performance of these models for materials outside of the PAH101 set before using them for large scale screening.

In conclusion, we have demonstrated that the SISO algorithm can generate ML models to predict the excited-state properties of molecular crystals using only a small amount of GW+BSE training data by incorporating physical/chemical knowledge into the selection of primary features. The resulting ML models, which require only relatively low-cost DFT calculations to evaluate, can provide estimates for excited-state properties of molecular crystals including the optical gap, triplet exciton energy, singlet–triplet gap, and exciton binding energy. These properties would not otherwise be accessible *via* DFT calculations. Using ML models in the early stages of materials screening workflows can effectively narrow down the number of candidates selected for further



evaluation using computationally expensive excited-state theory. ML models can thus significantly accelerate the discovery of crystalline organic semiconductors with desirable properties for applications in optoelectronic devices.

## Data and code availability

The PAH101 dataset<sup>117</sup> is available *via* the NOvel Materials Discovery (NOMAD) repository<sup>170</sup> and can be accessed at DOI: <https://doi.org/10.17172/NOMAD/2024.12.05-1>. The SISSO code,<sup>128</sup> used to perform sure independent screening and sparsifying operator model training, is available in the GitHub repository SISSO. SISSO version 3.3 dated July 2023 was used here. The SISSO primary features were calculated using version 18.06.07 of the FHI-aims<sup>150</sup> electronic structure package available *via* the FHI-aims website. Scripts used to calculate the primary features are available in the GitHub repository MLfeat\_FHI-aims, DOI: <https://doi.org/10.5281/zenodo.15093306>. Scripts for preparing the input for SISSO, running the training and model evaluation, analyzing the SISSO output, and making Pareto plots and correlation plots between the SISSO model predictions and the true labels are provided in the GitHub repository SISSOonPAH, DOI: <https://doi.org/10.5281/zenodo.15093308>.

## Author contributions

S. G., Y. L., and X. L. performed the calculations and data analysis. N. M., S. G., Y. L., and X. L. wrote the manuscript. N. M. conceived and led the project.

## Conflicts of interest

There are no conflicts of interest to declare.

## Acknowledgements

This work was supported by the National Science Foundation (NSF) Designing Materials to Revolutionize and Engineer our Future (DMREF) program under award DMR-2323749. This research used resources of the Argonne Leadership Computing Facility (ALCF), which is a DOE Office of Science User Facility supported under Contract DE-AC02-06CH11357.

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