

## Extracting the parameters of two-energy-level defects in silicon wafers using machine learning

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With the alarming issues of climate change and energy shortage, the importance of sustainable energy technologies has significantly increased [1] [2]. Photovoltaic (PV) energy is one of the most promising renewable technologies [3], whose cost has dramatically decreased over the past thirty years [4]. According to the International Energy Agency (IEA), PV-generated electricity is the cheapest form of energy in human history [5]. Nevertheless, the price of PV systems should be further reduced to improve the accessibility of PV energy in developing countries [6] and to obtain the full potential of this technology. One of the key methods to do this is by improving the power-conversion efficiency of solar cells [7]. A key limitation towards improving cell efficiencies is the recombination-active defects in the silicon bulk [8]. To minimise the detrimental effects of these defects, it is essential to predict their energy levels ( $E_1$ ) and their electron and hole capture cross-sections ( $\sigma_0$  and  $\sigma_p$ , respectively) [9] [10].

Commonly, bulk defects are classified as single-energy-level [9] or multi-energy-level [11]. This study will focus on two-level defects, which are a subset of multi-energy-level defects. Temperature- and injection-dependent lifetime spectroscopy (TIDLS) [12] is often used to extract the parameters of the defect [10] [13] [12]. In the case of single-energy-level defects, a few analysis methods are commonly employed. The most common method is the defect parameter solution surface (DPSS) method [12]. Other methods include the linearised DPSS method [14], the Newton Raphson method [10], and the machine learning (ML)-based approach [15]. However, in the case of two-level defects, there is only the pioneering study by Zhu *et al.* [13]. Here, we present a novel ML-based approach to extracting the defect parameters of two-level defects.

To train the developed ML algorithms, a dataset of 800,000 defects and the corresponding lifetime curves with different temperatures was generated using the Sah-Shockley equation [11] with random selections of the defect parameters ( $E_{t1}$ ,  $E_{t2}$ ,  $\sigma_{n1}$ ,  $\sigma_{p1}$ ,  $\sigma_{n2}$ ,  $\sigma_{p2}$ , and  $N_t$ ; the subscript '1' or '2' indicates the level of the defect):

$$\tau_{\text{two-level}} = \frac{1 + \frac{\sigma_{\text{n1}} v_{\text{n}} n_{1} + \sigma_{\text{p1}} v_{\text{p}} p}{\sigma_{\text{p1}} v_{\text{p}} p_{1} + \sigma_{\text{n1}} v_{\text{n}} n} + \frac{\sigma_{\text{p2}} v_{\text{p}} p_{2} + \sigma_{\text{n2}} v_{\text{n}} n}{\sigma_{\text{n2}} v_{\text{n}} n_{2} + \sigma_{\text{p2}} v_{\text{p}} p}}{N_{\text{t}} (n_{0} + p_{0} + \Delta n) \left[ \left( \frac{\sigma_{\text{n1}} \sigma_{\text{p1}} v_{\text{n}} v_{\text{p}}}{\sigma_{\text{p1}} v_{\text{p}} p_{1} + \sigma_{\text{n1}} v_{\text{n}} n} \right) + \left( \frac{\sigma_{\text{n2}} \sigma_{\text{p2}} v_{\text{p}} v_{\text{p}}}{\sigma_{\text{n2}} v_{\text{n}} n_{2} + \sigma_{\text{p2}} v_{\text{p}} p} \right) \right]},$$
Eq. 1

 $n_1$ ,  $n_2$ ,  $p_1$ , and  $p_2$  are defined by Eqs. 2-5:

$$n_1 = n_i e^{\frac{E_{t1} - E_i}{k_b T}}$$
, Eq. 2

$$n_2 = n_{\rm i} e^{\frac{E_{\rm t2} - E_{\rm i}}{k_{\rm b}T}}$$
, Eq. 3

$$p_1 = n_i e^{\frac{-E_{t_1} + E_i}{k_b T}},$$
 Eq. 4

$$p_2 = n_{\rm i} e^{\frac{-E_{\rm t2} + E_{\rm i}}{k_{\rm b}T}}$$
. Eq. 5

In Eq. 1,  $r_{\text{two-level}}$  refers to the recombination lifetime of the two-level defect,  $\sigma_{\text{n1}}$  ( $\sigma_{\text{p1}}$ ) is the electron (hole) capture cross-section of the first energy level,  $\sigma_{\text{n2}}$  ( $\sigma_{\text{p2}}$ ) is the electron (hole) capture cross-section of the second energy level, and  $v_{\text{n}}$  ( $v_{\text{p}}$ ) is the electron (hole) thermal velocity calculated using the model of Green *et al.* [16]. Furthermore, n (p) is the electron (hole) density,  $N_{\text{t}}$  is the total defect concentration,  $n_{\text{0}}$  ( $p_{\text{0}}$ ) is the thermal equilibrium electron (hole) density, and  $\Delta n$  is the excess carrier density. In Eqs. 2 to 5,  $n_{\text{i}}$  is the intrinsic carrier density calculated using the model of Couderc *et al.* [17] with bandgap narrowing by the model of Yan *et al.* [18],  $E_{\text{t1}}$  and  $E_{\text{t2}}$  are the defect energy levels,  $E_{\text{t1}}$  is the transition energy between the most positively charged state and the middle charge state



while  $E_{12}$  is the transition energy between the most negatively charged state and the middle charge state [13] [19]. Once the dataset was created, it was randomly split into a training set (90% of the simulated data) and a validation set (10%).  $E_i$  is the intrinsic fermi energy of silicon,  $k_b$  is the Boltzmann's constant, and T is the bulk temperature.

Two solutions for  $E_t$  are usually obtained in one-level defect parameter extraction methods such as DPSS [12], one in each bandgap half (higher or lower than  $E_i$ ). As a result, these two cases require separate training instances for the ML-based one-level defect parameter extraction [15]. Similarly, for a two-level defect, the DPSS residual map using TIDLS data proposed by Yan [13] identifies multiple possible solutions in different bandgap halves as well. Since two-level-defects have two energy levels, this study will use four cases: both energy levels above  $E_i$  (named here Set 11); both below  $E_i$  (Set 00);  $E_{t1} > E_i$  and  $E_{t2} < E_i$  (Set 10);  $E_{t1} < E_i$  and  $E_{t2} > E_i$  (Set 01).

The proposed method is divided into two steps as shown in Fig. 1. In the first step, the defect parameters are extracted by regression for the four cases listed above. In the second step, the ML algorithm is trained to classify the most likely solution out of the four cases.

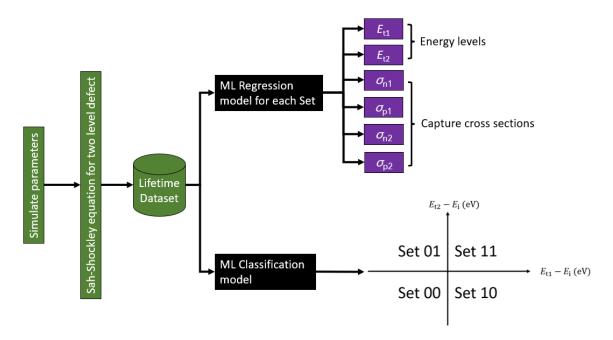


Figure 1: A flow chart for defect parameter extraction using machine learning.

To evaluate the performance of the developed ML models, two scoring methods are used. For the regression task, the coefficient of determination ( $R^2$  score) is used as defined by Eq. 6 [20]:

$$R^{2} = 1 - \frac{\sum_{i=1}^{N} \left(y_{i}^{\text{pred}} - y_{i}^{\text{true}}\right)^{2}}{\sum_{i=1}^{N} \left(y_{i}^{\text{true}} - y_{\text{mean}}^{\text{true}}\right)^{2}},$$
 Eq. 6

where the label 'pred' and 'true' refers to the predicted and true values of the test dataset respectively and  $R^2$  ranges from zero (very poor prediction) to unity (perfect prediction). The accuracy matrix is utilised for the classification task, which is the ratio of the number of correct predictions to the total number of predictions [21].

For the regression task, a random forest model [22] was trained. The predicted vs simulated values of the validation set for a p-type wafer (bulk doping of  $10^{15}$  cm<sup>-3</sup>, Set 11) are presented in Fig. 2, the red line represents the ideal prediction while the green dots are the actual prediction. The defect parameters of the first level ( $E_{t1}$ ,  $\sigma_{n1}$ ,  $\sigma_{p1}$ ) are extracted with a high level of accuracy ( $R^2 > 0.9$ ). The predictions of  $E_{t2}$ ,  $\sigma_{n2}$ ,  $\sigma_{p2}$ , are reasonable, but less accurate ( $R^2 > 0.6$ ). The difference in the prediction qualities could be as the Sah-Shockley recombination lifetime is more sensitive to changes in the first-level defect parameters in p-type wafers.

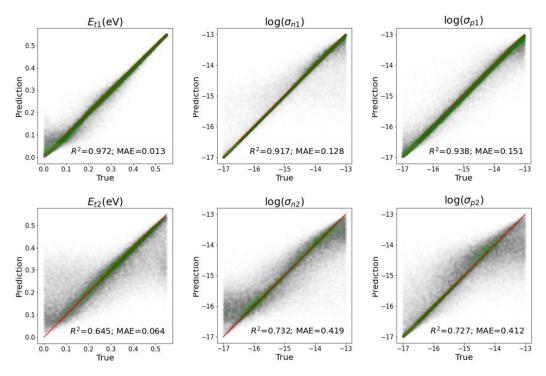


Figure 2: The predicted vs simulated parameters for a p-type wafer in Set 11 (bulk doping density of  $10^{15}$  cm<sup>-3</sup>).

The correlation coefficients of the different sets are summarised in Table 1. The reason for the use of the logarithm of the capture cross sections is that it ranges over several orders of magnitude (typically  $10^{-13}$  cm<sup>-3</sup> to  $10^{-17}$  cm<sup>-3</sup> [23]). It can be seen that for *p*-type doping when the set number begins with the subscript "1", meaning  $E_{t1} > E_{i}$ , the prediction for  $E_{t1}$ ,  $\sigma_{n1}$ , and  $\sigma_{p1}$  are more accurate, and vice versa for  $E_{t2}$ . The reason for this pattern is still under investigation.

Table 1: The coefficient of correlation ( $R^2$  score) for the four sets (a p-type wafer)

|        | $E_{t1}$ | $\log(\sigma_{\rm n1})$ | $\log(\sigma_{\rm p1})$ | $E_{t2}$ | $\log(\sigma_{\rm n2})$ | $\log(\sigma_{\mathrm{p}2})$ |
|--------|----------|-------------------------|-------------------------|----------|-------------------------|------------------------------|
| Set 11 | 0.972    | 0.919                   | 0.939                   | 0.643    | 0.726                   | 0.726                        |
| Set 10 | 0.936    | 0.987                   | 0.941                   | 0.821    | 0.807                   | 0.703                        |
| Set 01 | 0.813    | 0.916                   | 0.726                   | 0.693    | 0.689                   | 0.783                        |
| Set 00 | 0.827    | 0.882                   | 0.739                   | 0.934    | 0.849                   | 0.766                        |

The confusion matrix of the classification model is shown in Table 2, the rows correspond to the true class while the columns represent the predicted class; the number represents the percentage of this prediction compared to the dataset size. The correct predictions are shown in the diagonal. At this stage, the average accuracy for the four sets is about 67%. Furthermore, the specificity of each set is shown on the rightest column, which is the ratio between the correct classification over the actual population of each set. It seems that the ML model has difficulties identifying Set 11 correctly.

Table 2: The confusion matrix of the classification model.

| Prediction<br>True | Set 11 | Set 10 | Set 01 | Set 00 | Specificity |
|--------------------|--------|--------|--------|--------|-------------|
| Set 11             | 13.1%  | 5.9%   | 0.8%   | 5.3%   | 52.2%       |
| Set 10             | 4.5%   | 17.9%  | 1.6%   | 1.2%   | 71.0%       |
| Set 01             | 0.6%   | 1.7%   | 18.4%  | 4.5%   | 73.0%       |
| Set 00             | 2.9%   | 0.3%   | 3.3%   | 18.2%  | 73.7%       |



To summarise, this research presented a novel ML-based approach to solving a challenging problem – the extraction of the defect parameters of a two-level defect from TILDS data. The defect parameters of the four possible cases are first extracted. A classification model then predicts the most likely solution. We note a difference between the prediction accuracy of  $E_{t1}$ ,  $\sigma_{n1}$ ,  $\sigma_{p1}$  and  $E_{t2}$ ,  $\sigma_{n2}$ ,  $\sigma_{p2}$ . This is due to the difference in the sensitivity of the lifetime with different parameters for a given set of doping densities and/or temperatures. However, this is not a limitation when the same two-level defect is present in both p- and n-type wafers. Future work will focus on improving prediction accuracy and uncovering the reason for better predictions of some parameters than others.

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