# A Python code for a neural network machine-learning approach for the interpretation of TDS experiments

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#### Abstract

Documentation that accompanies the Python code for the implementation of the neural network machine-learning approach for the interpretation of TDS experiments, available from here. This documentation explains the usage of the machine-learning framework and highlights the main files. Special attention is paid to the parts of the code that implement the model building (e.g., hyperparameter selection) and the model training algorithm. The visualisation of results (i.e., the reconstruction of the TDS spectrum with the predicted trapping parameters and the comparison with experimental data) is also discussed through an example test case.

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## Contents

1	Intr	oducti	on	2
	1.1	Requir	red external software libraries	2
	1.2	Basic	usage	3
2	Sun	nmary	of included files	3
	2.1	Main.	py	3
		2.1.1	Parameters and setup	3
		2.1.2	Workflow	4
		2.1.3	Additional considerations	Ē
2.2 TDS simulation files		TDS s	imulation files	5
		2.2.1	TDS_Sim.py	6
		2.2.2	TDS_Material.py	7
	2.3	ML m	odel files	8

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		2.3.1	ModelEnsemble.py	8
		2.3.2	ClassificationModel.py	9
		2.3.3	RegressionModel.py	10
		2.3.4	Model_Parameters.py	11
	2.4	Exper	imental data files	12
		2.4.1	ExpDataParameters.py	12
		2.4.2	ExpDataProcessing.py	13
3	Spe	ecifics:	NN building and training	14
	_		NN building and training  High-density low-energy trap	14 16
	_	ecifics:		
	Spe	ecifics:	High-density low-energy trap	16
4	<b>Spe</b> 4.1 4.2	ecifics:	High-density low-energy trap  Generation Modifications	<b>16</b>

## 1. Introduction

Understanding hydrogen-material interactions, such as diffusion and trapping, provides insight into the susceptibility of a material to hydrogen embrittlement. Thermal desorption spectroscopy (TDS) is a widely used bulk experimental technique for quantifying the key trapping characteristics of metallic alloys, specifically the trapping capacity, including trap binding energy and density, of different microstructural features (e.g., dislocations, grain boundaries and precipitates). Interpreting the output of TDS experiments, i.e., extracting the number of trap sites, and the binding energy and density associated with each trap, from the TDS spectrum, is far from straightforward. To overcome the limitations of current TDS analysis techniques, we developed a machine learning-based approach, comprising a multi-neural network (NN) machine learning (ML) model trained exclusively on synthetic data, to predict the trapping parameters directly from experimental data. Here, we present a Python implementation of the proposed ML approach. In the remainder of this documentation, we outline the synthetic training data generation, NN model building and training, and required inputs for each test case (e.g., material parameters, TDS test setup, and model hyperparameters). The experimental data preprocessing and model output visualisation will also be discussed. An example test case is provided in Section 5.

## 1.1. Required external software libraries

In addition to standard Python libraries, the following external software libraries are also required to conduct the analysis:

• TensorFlow (recommended version 2.19.0)

• Keras (recommended version 3.10.0)

Note: Python 3.11.0 is recommended.

1.2. Basic usage

Running "Main.py" performs all actions required for conducting the analysis: It defines the test case-specific inputs and model hyperparameters, checks if the training data and ML models for the given inputs exist and loads them, if they don't exist, it generates the training data and trains the model, performs model validation (i.e., generates validation dataset of 500 data points and plots confusion matrix and predicted vs actual scatter plots for trap density and binding energy), fits the experimental data and plots comparison plots between the model predictions and the experimental data (including the individual trap contributions).

A parameter sweep to compare multiple trained models, differing in the number of training data points, can be carried out by running "TrainMultipleDataSets.py".

2. Summary of included files

The code is set up in an object-oriented manner, defining Python classes for each sub-component and providing accompanying methods. As such, a clear distinction is made between different components, and each can be used and altered with limited/no impact on other components. Here, the different files (and corresponding classes) are described.

2.1. Main.py

This is the main file, from which all classes are constructed, and the actual analysis is performed. Within it, all properties used within the required classes are defined as inputs.

2.1.1. Parameters and setup

• NumTraining (int): Number of samples used for training

• NumVerification (int): Number of samples used for model verification

• Regenerate\_Data (bool): Flag to regenerate synthetic data

• Regenerate\_Training (bool): Flag to retrain models from scratch

• n\_cpu\_cores (int): Number of CPU cores allocated for parallel processing

• Traps (str): Trap configuration type ("Random" in this case)

• Concentrations (str): Concentration assignment method ("Random")

• MaxTraps (int): Maximum number of traps to predict.

3

- HD\_Trap\_params (dict) or None: Controls the configuration of a high-density low-energy trap (utilised in test cases 2 and 3 of the manuscript). When provided as a dictionary, it must contain the following three keys:
  - High\_Density\_Trap (bool): Specifies whether a high-density trap is present in the sample
  - HDT\_NRange (list): Defines the allowable density range bounds for the trap
  - HDT\_ERange (list): Defines the allowable binding energy range bounds for the trap

When set to "None", the following default values are applied:

```
    High_Density_Trap = False (i.e., no high-density low-energy trap is present)
    HDT_NRange = None
```

- HDT\_ERange = None
- ExpName (str): Experimental dataset identifier
- trap\_model (str): Trap model selection ("McNabb" or "Oriani")
- Material: Material object initialised with experiment and trap model
- HyperParameters: Model hyperparameters loaded from predefined optimised set

**Note**: An overview of how the presence of a high-density low-energy trap is accounted for in the data generation and model training phases is provided in Section 4.

## 2.1.2. Workflow

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Once all parameters have been defined, the ModelEnsemble class, which automates the training data generation as well as the model setup, training, and inference in a unified interface, is initialised with the required inputs:

Subsequently, model validation is carried out. Synthetic verification data is generated using TDS\_Sim.SimDataSet (discussed in Section 2.2.1) and the trained model used to predict traps, concentrations, and energies from the verification data. Plot comparisons between predicted and actual values for traps, concentrations, and energies are provided.

```
# Verification

TDS_Curves, Actual_Traps, Actual_Concentrations, Actual_Energies, TDS_Temp =

TDS_Sim.SimDataSet(Material, NumVerification, MaxTraps, Traps, Concentrations, n_cpu_cores)

Predicted_Traps, Predicted_Concentrations, Predicted_Energies = Model.predict(TDS_Curves)

Model.PlotComparisonTraps(Predicted_Traps, Actual_Traps, TDS_Curves, TDS_Temp)

Model.PlotComparisonConcentrations(Predicted_Concentrations, Actual_Concentrations)

Model.PlotComparisonEnergies(Predicted_Energies, Actual_Energies)
```

Finally, the experimental TDS data of the corresponding test case ("ExpName") is loaded and processed using the ExpDataProcessing class and passed to the trained model as input for the parameter inference.

```
# Experimental data fit

FileName = 'TDSData/Novak_200_Experimental.xlsx'

Exp_Processed_Data = ExpDataProcessing.ExpDataProcessing(FileName, Material, HyperParameters)

Exp_Temp = Exp_Processed_Data.Temperature

Exp_Flux = Exp_Processed_Data.Flux

Exp_TDS_Curve = Exp_Processed_Data.TDS_Curve

Exp_Predicted_Traps, Exp_Predicted_Concentrations, Exp_Predicted_Energies =

Model.predict(Exp_TDS_Curve)
```

The visualisation of the experimental data fit and predicted trap parameter reconstructed curve comparison is performed by the following line:

```
# Plot predictions

Model.PlotComparisonExpData(Exp_Temp, Exp_Flux, Exp_Predicted_Concentrations,

Exp_Predicted_Energies)
```

#### 2.1.3. Additional considerations

- The script uses parallel CPU cores to speed up data generation and training
- Model training and data generation can be toggled with Regenerate\_Training and Regenerate\_Data
- Energy labelling in the output adjusts according to the selected trap model

## 2.2. TDS simulation files

The TDS simulation is carried out by two classes: TDS\_Sim.py (see Section 2.2.1) and TDS\_Material (see Section 2.2.2). The TDS\_Sim.py class is responsible for conducting the FEM-based TDS simulation

and for generating the full training dataset. The TDS\_Material class defines the material properties and test parameters used in TDS\_Sim.py. It serves as the data storage for material constants, experimental conditions, and numerical setup.

# 2.2.1. TDS\_Sim.py

Initialization:. An instance of this class requires a fully defined TDS\_Material object, which contains all material properties, numerical settings, and TDS test parameters. The material object is typically initialised from user-defined inputs (via ExpDataParameters.py), allowing flexible switching between materials or test setups. Once the material is defined, the simulation functions in TDS\_Sim.py can be called to perform individual or batch TDS experiments.

The simulation workflow is primarily handled by two functions:

Function: GenerateDataPoint(). Generates a single data point by simulating the TDS process under a given set of predefined parameters. It either sets the specified trap energies and concentrations or randomly assigns trapping energies and concentrations, initialises a sample, and performs the three major stages of TDS:

- 1. Charging: Fills the traps with hydrogen (Sample.Charge())
- 2. Resting: Keeps the sample at room temperature for a specified period (Sample.Rest())
- 3. TDS Measurement: Heats the sample and records the desorption signal (Sample.TDS())

# Function definition:

```
def GenerateDataPoint(i :int,

Material :TDS_Material.TDS_Material,

MaxTraps :int,

NumTraps :Union[int, str],

Concentration :Union[float, str]):
```

# Core TDS simulation steps:

```
Sample = TDS_Sample(Material, N_traps, E_traps, PlotWhileSolving) #initializes material

Sample.Charge() #performs charging

Sample.Rest() #leave at atmospheric pressure

[T,J] = Sample.TDS() #perform TDS
```

Function: SimDataSet(). Generates a complete dataset by repeatedly invoking GenerateDataPoint() in parallel, utilising multiple CPU cores.

## Function definition:

```
def SimDataSet (Material
711
                                          :TDS_Material.TDS_Material,
                       NumVerification
                                         :int,
712
                       MaxTraps
                                          :int,
713
                       NumTraps
                                          :Union[int, str],
714
                                          :Union[float, str],
                       Concentrations
                       n_cpu_cores
                                          :int
                       ):
717
```

This function leverages joblib.Parallel to distribute the data point simulations across available CPU cores, enabling efficient generation of large datasets.

```
Res = joblib.Parallel(n_jobs=n_cpu_cores)(joblib.delayed(GenerateDataPoint)(i, Material,

MaxTraps, NumTraps, Concentrations) for i in range(0, NumVerification))
```

# 2.2.2. TDS\_Material.py

This class centralises and organises all input parameters for TDS simulations, including material properties, experimental setup (e.g., heating rates, charging conditions), and numerical parameters.

The class takes the following as inputs:

- ExpName (str): A user-defined label or identifier for the experimental dataset. This is used to load and register the correct parameters via ExpDataParameters
- material\_param (dict): Material properties for the specified experimental dataset
- test\_param (dict): TDS test configuration parameters for the specified experimental dataset
- numerical\_param (dict): Numerical parameters for the specified experimental dataset
- HD\_Trap\_param (dict): Controls the configuration of a high-density low-energy trap
- trap\_model (str): Specifies the trapping model to use. Options include:
  - "McNabb" McNabb-Foster model
  - "Oriani" Oriani's equilibrium trapping model

The initialisation behaviour depends on whether parameter dictionaries are provided:

- When parameter dictionaries are not provided (all None): The class checks whether ExpName corresponds to a registered experimental dataset in the ExpDataParameters class or matches a predefined experiment (such as the test cases presented in the manuscript, e.g., "Novak\_200"). If found, the class retrieves the material\_param, test\_param, numerical\_param and HD\_Trap\_param from ExpDataParameters. If ExpName is not recognised, a KeyError is raised.
- When parameter dictionaries are provided: The class uses these user-supplied parameters and registers the experiment in ExpDataParameters for future use.

Further information on the ExpDataParameters class functionalities and additional methods can be found in Section 2.4.1.

## 2.3. ML model files

The implementation of the ML models is carried out by three classes: ModelEnsemble (see Section 2.3.1), ClassificationModel (see Section 2.3.2) and RegressionModel (see Section 2.3.3).

Both models leverage TensorFlow's Keras API for model training and inference. This is discussed in Section 3.

## 2.3.1. ModelEnsemble.py

The ModelEnsemble class acts as a wrapper for both the classification and regression models. Its primary function is to predict the number of active hydrogen traps in a material and, for the identified number of traps, estimate the corresponding trap energies and concentrations. It automates model setup, training, and inference in a unified interface.

The ensemble structure enables a two-step prediction process:

- 1. A classification model determines the number of distinct trap types active in a given TDS curve
- 2. A corresponding regression model (one per possible number of traps) predicts the associated trap energies and concentrations

*Initialization:*. An instance of ModelEnsemble requires material properties, training settings, and control flags for regenerating data and retraining models. Depending on whether a fixed number of traps is specified, it either:

- Loads a single regression model (for fixed-trap problems), or
- Loads one regression model per possible trap count and a classification model (for variable-trap problems)

Function: predict(). This function accepts a list of processed TDS curves and performs full model inference. It returns:

- The predicted number of traps for each curve
- Predicted trap energies and concentrations

*Plotting functions:*. The class includes several helper functions for visual comparison between predicted and actual outputs:

- PlotComparisonExpData(): Overlays model predictions on experimental TDS curves
- PlotComparisonTraps(): Visualizes classification performance and confusion matrix
- PlotComparisonEnergies(), PlotComparisonConcentrations(): Compare predicted vs. true regression outputs

These plots are saved automatically.

## 2.3.2. ClassificationModel.py

The ClassificationModel class defines a neural network classifier that predicts the number of traps present in a material based on its TDS response.

**Initialisation:** The constructor requires simulation parameters and optionally pre-trained regression models. If no saved model is found, or if retraining is explicitly requested, the network is trained from scratch.

```
def __init__(self, Material, MaxTraps, Concentrations, HyperParameters, NumTraining,

→ RegModels, Regenerate_Training, n_cpu_cores):
```

```
if ((os.path.isfile(self.TrainedModelName) == False) or (Regenerate_Training)):
    self.Train(RegModels)

self.TrainedModel = saving.load_model(self.TrainedModelName)
```

## Main methods:

GetData(RegModels) Retrieves TDS curves and trap count labels from synthetic regression model data. Returns training features and one-hot encoded labels.

Train(RegModels) Handles data preprocessing (thresholding, optional log transformation, noise injection), builds a neural network classifier, trains it, and saves the final model.

predict\_traps(TDS\_Curve) Processes a new TDS curve and returns the predicted number of traps.

PlotConfMatrix(y\_test, y\_pred) Generates a normalised confusion matrix to evaluate classification accuracy across trap classes.

# Example usage:

```
n_traps = model.predict_traps(new_TDS_curve)
```

# 2.3.3. RegressionModel.py

The RegressionModel class implements a neural network for predicting binding energies and site concentrations for a known number of traps. The outputs depend on the trapping model: Oriani (binding energies) or McNabb-Foster (de-trapping energies).

**Initialisation:** The constructor accepts all necessary parameters for training and simulation. If a trained model is not found or retraining is requested, a new model is trained using synthetic data.

```
def __init__(self, Material, MaxTraps, NumTraps, Concentrations, HyperParameters,

→ NumTraining, Regenerate_Data, Regenerate_Training, n_cpu_cores):
```

```
if ((os.path.isfile(self.DataName) == False) or (Regenerate_Data)):
    #Data needs to be generated
    self.GenerateData()

if ((os.path.isfile(self.TrainedModelName) == False) or (Regenerate_Training)):
    #Train model
    self.Train()
```

# Main methods:

GetData(RegModels) Loads synthetic TDS curves and corresponding continuous output labels (trap energies and concentrations). Returns scaled inputs and targets.

Train(RegModels) Applies preprocessing steps (normalisation, optional noise), constructs and trains a deep neural network regression model, and saves the trained weights and output scalers.

predict\_parameters(TDS\_Curve) Accepts a new TDS curve and returns predicted trap parameters (energy and concentration vectors).

PlotPredictionResults(y\_true, y\_pred) Visualises model performance by plotting predicted versus actual trap parameter values.

# Example usage:

```
trap_params = model.predict_parameters(new_TDS_curve)
```

# 2.3.4. Model\_Parameters.py

The Model\_Parameters class encapsulates all hyperparameters used for training both regression and classification models. It supports two modes of initialisation: "optimised" (uses predefined, tuned hyperparameters) and "user\_defined" (allows user input via a parameter dictionary).

## **Initialisation:**

- ParameterSet (str): Specifies the mode ("optimised" or "user\_defined").
- UserDefinedParameters (dict, optional): Dictionary of custom hyperparameters (currently a placeholder for future extension).

## Core attributes:

- flux\_threshold (float): Minimum flux value for numerical stability (default: 1e-12)
- log\_transformation (bool): Whether to apply a base-10 log transformation to input data (default: True)
- noise\_std\_dev (float): Standard deviation of Gaussian noise added during data augmentation (default: 0.05)
- epoch\_coeff\_reg (int): Multiplier for the number of training epochs in regression models (default: 200)
- epoch\_coeff\_class (int): Multiplier for training epochs in classification models (default: 100)
- batch\_size (int): Number of samples per training batch (default: 32)

# Optimised hyperparameters:

# • Regression model:

- Number of layers: 5

- Nodes per layer: [64, 64, 32, 16, 8]

- Learning rate: 1e-3

- Weight decay: 1e-3

## • Classification model:

- Number of layers: 4

- Nodes per layer: [256, 128, 64, 32]

- Learning rate: 1e-3

- Weight decay: 1e-3

# 2.4. Experimental data files

The experimental data files comprise two classes: ExpDataParameters, which stores all parameters for each experimental dataset, and ExpDataProcessing, which processes raw experimental data and prepares it for model input.

# 2.4.1. ExpDataParameters.py

The ExpDataParameters class serves as a central repository for all material properties, experimental conditions and numerical settings associated with different experimental datasets. The class manages two types of experiments: predefined test cases and user-registered experiments.

Predefined experiments correspond to the test cases presented in the manuscript (i.e., "Novak\_200", "Novak\_100", "Novak\_50", "Wei\_Tsuzaki", "Depover"). These can be accessed by initialising the class with the corresponding ExpName:

```
exp_params = ExpDataParameters(ExpName)
exp_material = exp_params.material
exp_test = exp_params.test
exp_numerical = exp_params.numerical
```

User-registered experiments are datasets added during runtime and can be accessed using the get\_experiment method:

```
exp_material, exp_test, exp_numerical = ExpDataParameters.get_experiment(ExpName)
```

Registration of new experiments is accomplished using the register\_experiment method, which stores the experiment parameters for future use:

```
ExpDataParameters.register_experiment(ExpName, material_param, test_param,
numerical_param)
```

Additional methods include:

- list\_experiments(): Returns a list of all registered experiments
- experiment\_exists(ExpName): Checks whether a specified experiment exists in the registry

## 2.4.2. ExpDataProcessing.py

The ExpDataProcessing class handles preprocessing of raw experimental TDS data to generate uniformly sampled, smoothed datasets compatible with the ML model inputs.

Initialisation parameters:

- file\_name (str): Path to the Excel file containing raw experimental data
- material: A fully defined TDS\_Material object containing all required parameters
- hyperparameters: A Model\_Parameters object containing preprocessing settings (including ntp for temperature grid points)

Upon instantiation, the class automatically executes the data loading and preprocessing pipeline through sequential calls to \_load\_data() and \_process\_data().

\_load\_data(). Loads raw experimental data from an Excel file. The expected format requires temperature values (in K) in the first column and corresponding desorption rates (in wppm/s) in the second column.

\_process\_data(). Processes the raw data through the following pipeline:

- 1. Scales desorption rates using material thickness and density to obtain physically meaningful flux units
- 2. Applies Savitzky-Golay filtering to smooth experimental noise
- 3. Interpolates the smoothed data onto a uniform temperature grid with ntp points using cubic interpolation
- 4. Clips desorption values above the maximum temperature (TMax) to a predefined flux threshold to handle extrapolation artefacts
- 5. Stores the processed arrays as class attributes for model input

The processed data are accessible through the following attributes:

- self.Temperature (numpy.ndarray): Uniformly spaced temperature values (K), excluding the final temperature point
- self.Flux (numpy.ndarray): Corresponding desorption flux rates after preprocessing
- self.TDS\_Curve (tensorflow.Tensor): Processed data formatted as a TensorFlow tensor for direct model input

# 3. Specifics: NN building and training

In both ClassificationModel and RegressionModel classes, the model construction and training are handled by the Train() function. This function is only called if the model has not been previously trained (i.e., no saved model file is found) or if training is explicitly requested using the Regenerate\_Training flag. This is done for the regression model as:

```
if ((os.path.isfile(self.TrainedModelName) == False) or (Regenerate_Training)):

#Train model
self.Train()
```

And for the classification model as:

```
if ((os.path.isfile(self.TrainedModelName) == False) or (Regenerate_Training)):
self.Train(RegModels)
```

The classification model's Train() function takes the regression models as input in order to access the training data. Specifically, the regression model defines the filenames for the input data, which are retrieved via the GetData() function.

Within both classes, the training process follows the same four stages:

- 1. Preprocessing the training data
- 2. Train-test splitting
- 3. Model building
- 4. Model training

A detailed breakdown of each stage (for the classification model) is given below.

- I. Preprocessing training data. This stage involves three key preprocessing steps:
  - Flux Thresholding: Small values are clipped for numerical stability.

```
All_TDS = np.where(All_TDS < self.hp.flux_threshold, self.hp.flux_threshold,

All_TDS)
```

• Log Transformation: Optionally applies a base-10 logarithmic transformation.

```
if self.hp.log_transformation:

All_TDS = np.log10(All_TDS)
```

• Noise Augmentation: Adds independent Gaussian noise to each feature for robustness.

```
All_TDS_noise = All_TDS.copy()

selected_features = range(All_TDS.shape[1])

for feature_idx in selected_features:

noise = np.random.normal(0, self.hp.noise_std_dev, All_TDS.shape[0])

All_TDS_noise[:,feature_idx] += noise
```

II. Train-test splitting. The preprocessed data is split into training and test sets. The test set size is the minimum of 200 samples or 20% of the dataset, scaled by the number of traps:

```
TestsetSize = min(200, math.floor(0.2*self.NumTraining))*self.MaxTraps

X_train, X_test, y_train, y_test = train_test_split(All_TDS_noise, Labels,

→ test_size=TestsetSize)
```

III. Model building. The model architecture consists of an input normalisation layer, a series of hidden layers, and a softmax output layer. The number of hidden layers and their sizes are determined by hyperparameters.

```
# Build model
145
               NormLayer = layers.Normalization()
146
               NormLayer.adapt(X_train)
147
               model = models.Sequential()
148
               model.add(layers.Input(shape=(X_train.shape[1],)))
149
               model.add(NormLayer)
150
151
               # Add Dense layers based on num_layers
152
               for i in range(0,self.hp.num_layers_class):
                   model.add(layers.Dense(self.hp.nodes_class[i]*self.MaxTraps,
154

    kernel_initializer=initialiser))

                   model.add(layers.ReLU())
155
156
               # Add Output layer
               model.add(layers.Dense(self.MaxTraps, activation="softmax"))
158
159
               # Compile model
160
```

```
model.compile(optimizer=tf.keras.optimizers.Adamax(learning_rate=self.hp.lr_class,

→ weight_decay=self.hp.wd_class),

loss='categorical_crossentropy')
```

IV. Model training. The model is trained using model.fit(). Training hyperparameters are defined in the Model\_Parameters class and are passed during class initialisation.

```
# Train model

model.fit(x=X_train, y=y_train,

validation_data=(X_test, y_test),

batch_size=self.hp.batch_size,

epochs=self.hp.epoch_coeff_class*self.MaxTraps,

verbose=1)
```

The regression model follows the same general pipeline, with one key distinction: it applies separate normalisation (i.e., scalers) to the energy and density outputs using two independent MinMaxScaler instances.

```
SolScalerEnergy = MinMaxScaler(feature_range=(0,1))

SolScalerConcentration = MinMaxScaler(feature_range=(0,1))
```

Both scalers are then saved separately.

# 4. Specifics: High-density low-energy trap

To account for the presence of a high-density low-energy trap in the sample, modifications are required in both the data generation and model training processes.

## 4.1. Data Generation Modifications

During data generation, the first trap is generated using the specified high-density trap parameters (density and binding energy bounds), as shown in the code below:

```
if Material.High_Density_Trap:

# Define parameters for high density trap

# E1_abs = Material.E_Diff #set equal to lattice activation energy
```

```
E1_b = random.uniform(Material.HDT_ERange[0], Material.HDT_ERange[1])

E1_des = E1_b + E1_abs

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643

if (isinstance(Concentration, str)):

644

N1 = random.uniform(Material.HDT_NRange[0], Material.HDT_NRange[1])

645

else:

646

N1 = Concentration

647

648

traps.append([E1_abs, E1_des, N1])
```

The remaining traps are then generated using the standard procedure described previously.

## 4.2. Model Training Adjustments

The regression model training requires adjustment to handle the large magnitude difference between the high-density trap densities and those of standard traps. To address this scale disparity, a  $\log_{10}$  transformation is applied to the high-density trap density value, bringing it to a similar scale as the other traps:

```
if (isinstance(self.Concentrations, str)):
182
                       if self.Material.High_Density_Trap:
                           for j in range(0,self.NumTraps):
184
                                if j == 0:
185
                                    sol.append(np.log10(concentration_trapping_sites[i][j]))
186
                                else:
                                    sol.append(concentration_trapping_sites[i][j])
188
                       else:
189
                           for j in range(0,self.NumTraps):
190
                                sol.append(concentration_trapping_sites[i][j])
191
```

Following the model prediction, the log transformation of this trap is inverted to report the actual trap density values rather than their logarithmic equivalents.

## 5. Sample results

In its current setup, running Main.py will conduct the analysis for the experimental data obtained by Novak et al. [1], corresponding to a high-strength AISI tempered martensitic steel ( $\phi = 200^{\circ}\text{C/h}$ ), i.e., test case 1 of the manuscript.

The test case-specific parameters (i.e., material parameters, TDS setup configuration and numerical parameters) have been included in the ExpDataParameters class under "Novak\_200". The optimised hyper-

parameters (as described in the manuscript) have been included in the Model\_Parameters class and can be selected by setting the ParameterSet variable as "optimised".

First, all parameters required for the analysis are specified, and the material and hyperparameters (all other classes) are initialised. Subsequently, the analysis is carried out as described in Section 2.1.

**Parameter definition:.** The test case is set as "Novak\_200", and the McNabb-Foster framework is specified as the trap model. The material can then be defined/initialised (see Main.py)

**Data generation and model training:**. The data generation, model building and training, model evaluation and experimental data pre-processing and fit are carried out as described in Section 2.1.

Results and figures: Several figures are produced and saved to the Figures folder in the folder specific to the test case, i.e., Novak\_200 folder. Six figures are saved: the confusion matrix, scatter plot predicted vs actual for trap density, scatter plot predicted vs actual for binding energy, the wrongfully predicted TDS curves, plot of the experimental data vs the reconstructed TDS spectrum based on the model predictions, and the reconstructed TDS spectrum with the trap contributions (i.e., individual peak corresponding to each trap). The first four figures correspond to the model validation. The latter two figures are the experimental data fit. These are shown in Fig. 1.

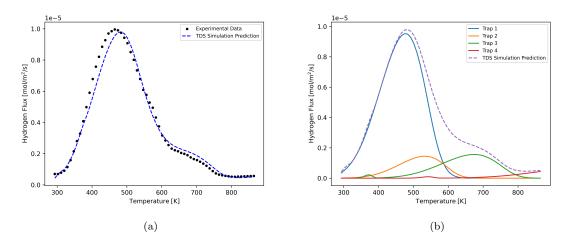


Figure 1: Results produced from running "Main.py", for the high-strength AISI tempered martensitic steel ( $\phi = 200^{\circ}\text{C/h}$ ), corresponding to test case 1 of the manuscript or "Novak\_200" in "ExpDataParameters.py". A comparison between the reconstructed TDS spectrum based on the ML model predictions and the experimental data is provided in (a). The individual trap contributions are depicted in (b).

Additionally, the values of binding/de-trapping energy and the trap density of each trap are printed in the terminal. Note: the energies are reported as absolute values in ascending order.

If the McNabb-Foster framework is used, the values appear as:

Number of traps: 4

```
Trap 1: De-trapping energy = 58290.01 J/mol, Trap density = 9.26 mol/m<sup>3</sup>
```

Conversely, if the Oriani framework is used, the values appear:

```
Number of traps: 4
Trap 1: Binding energy = 52600.01 J/mol, Trap density = 9.26 mol/m<sup>3</sup>
```

## References

[1] P. Novak, R. Yuan, B.P. Somerday, P. Sofronis, and R.O. Ritchie. A statistical, physical-based, micromechanical model of hydrogen-induced intergranular fracture in steel. *Journal of the Mechanics and Physics of Solids*, 58(2):206–226, February 2010. Publisher: Elsevier BV.