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1 **Evaluation of Machine Learning Algorithms to Predict Internal Concentration Polarization**  
2 **in Forward Osmosis**

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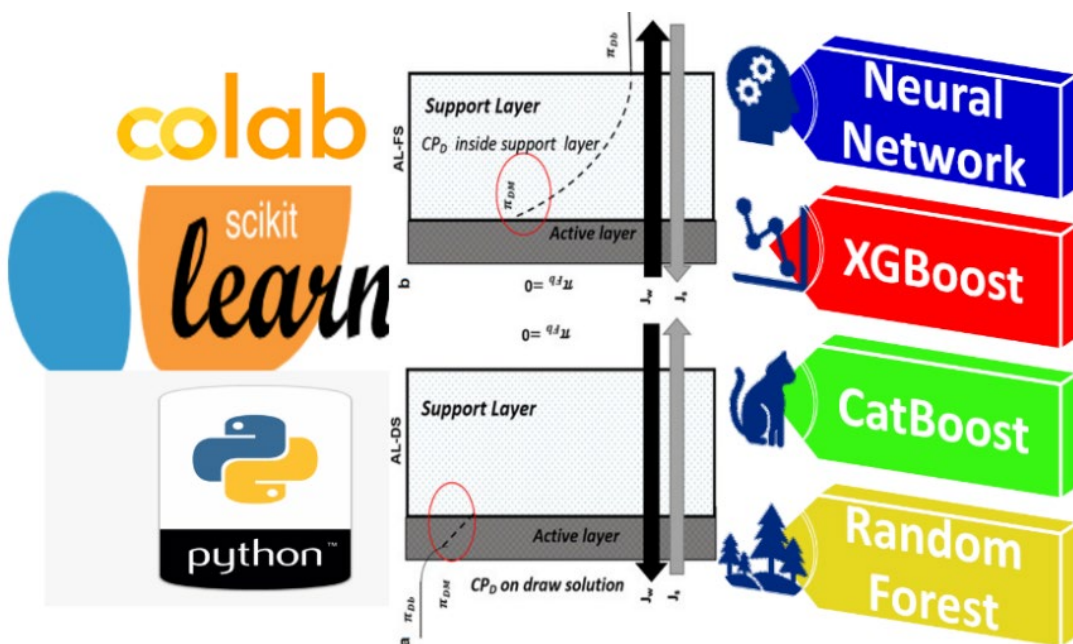
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12 **Graphical Abstract**



## 14 **Abstract**

15 Internal concentration polarization (ICP) is currently a major bottleneck in the forward  
16 osmosis process. Proper modelling of the internal concentration polarisation is therefore vital  
17 for improving the process performance and efficiency. This study assessed the feasibility of  
18 several machine learning methods for internal concentration polarisation prediction,  
19 including artificial neural networks, extreme gradient boosting (XGBoost), Categorical  
20 boosting (CatBoost), Random forest, and linear regression. Among the many algorithms  
21 evaluated, the CatBoost regression outperformed other methods in terms of coefficient of  
22 determination ( $R^2$ ) and the mean square error. The CatBoost algorithm's prediction power  
23 was then evaluated using non-training (user-provided) data and compared to solution  
24 diffusion models. The results indicated that the machine learning algorithms could predict ICP  
25 in the process with high accuracy for the provided dataset and excellent generalizability for  
26 future testing data. Furthermore, machine learning algorithms may offer insights into the  
27 input features that majorly affect ICP modelling in the forward osmosis process.

28 **Keywords:** Forward osmosis (FO), internal concentration polarization (ICP), machine learning  
29 modelling, artificial neural network, and wastewater treatment

## 30 **1. Introduction**

31 Forward osmosis (FO) membrane-based desalination has attracted tremendous attention due  
32 to its numerous advantages over pressure-driven membrane processes, particularly for  
33 treating complex wastewaters [1, 2]. The FO process advantages include low membrane  
34 fouling and a high-water recovery rate [3, 4]. However, the process being driven by osmosis  
35 or concentration differences between the feed and the draw solutions, is hindered by the  
36 problem of concentration polarization (CP) [5-7]. In the FO process, two types of CP co-occur,

37 viz. external concentration polarization (ECP) and internal concentration polarization (ICP) [5].  
38 The ECP has been reported in all membrane processes, whereas the ICP is unique to the  
39 forward osmosis process. Both ICP and ECP lead to a decrease in the concentration at the  
40 boundary layer near the membrane surface compared to the actual concentration in the bulk  
41 solutions, leading to a decrease in the net osmotic driving force in the FO process [5].  
42 However, ECP is not a severe issue compared to ICP in the FO process since ECP can be  
43 mitigated by varying hydrodynamic conditions.

44 Earlier studies on the FO process revealed that ICP accounts for about 80% of flux reduction  
45 in the FO process [7-9]. Since CP is such a significant issue in the forward osmosis process,  
46 modelling water flux behaviour in the presence of the CP is of utmost importance to  
47 understand the process performance and possible optimisation. Several models for CP  
48 measurements and predictions have been reported in the literature [10-15]. The first FO  
49 water flux model incorporating concentration polarization was proposed by Lee et al. [16].  
50 The model has been revised and witnessed a lot of improvements by several researchers over  
51 time. This model was later updated by McCutcheon and Elimelech [10] by considering  
52 external and internal CPs on both sides of the FO membrane. This model was widely used in  
53 the FO literature; however, the impact of reverse salt flux (RSF) from the draw solution to the  
54 feed solution, osmotic swelling of the FO membrane, as well as the mass transfer resistance  
55 provided by the support layer was neglected. Yip et al. [13] presented an updated model for  
56 FO transport in the presence of external and internal CPs along with the impact of the RSF.  
57 Details equation of these models is presented in the methodology section. Several other  
58 models are also presented in the literature, such as Bui et al. [15] model, resistance in the  
59 series model by Nagy [14], empirical model by [17] and several new models based on CFD

60 (computational fluid dynamics)[18], and two dimensional-FEM (finite element analysis)  
61 models [19-21].

62 Each model given in the literature has limitations and was developed based on certain  
63 assumptions or ideal circumstances. The measurement, prediction and estimation of  
64 concentration polarization, water flux and RSF are based on iterative procedures in these  
65 models. Furthermore, these models are based on exacting methodologies, are  
66 mathematically sophisticated and are time-consuming. Despite these drawbacks, these  
67 models are still widely employed in FO studies; however, there is a lack of universality since  
68 each model is limited to a certain range of parameters. For instance, if the draw solution is  
69 changed from NaCl (sodium chloride) to a mixture (e.g. NaCl + MgSO<sub>4</sub> (magnesium sulphate)  
70 + MgCl<sub>2</sub> (magnesium chloride)), most FO flux models would be inaccurate because the  
71 diffusion coefficient for the mixture would have to be determined separately through a  
72 lengthy series of exhaustive experiments. Recently, there has been a surge in ML modelling  
73 due to their high precision and accuracy [22-24], and ML algorithms have recently been used  
74 for FO's performance in several studies [23, 25]. ML is the study and interpretation of patterns  
75 and structures in data to automate learning, reasoning, and prediction [23]. Further, ML  
76 enables users to feed massive amounts of data to a computer algorithm, then analyses and  
77 makes data-driven recommendations and predictions based entirely on the input data.  
78 Moreover, unlike solution diffusion models, ML models can model complex and non-linear  
79 systems with high accuracy [26, 27].

80 Very few studies have investigated ML techniques to model forward osmosis performance.  
81 For instance, Jawad et al. [23] employed artificial neural networks to predict forward osmosis  
82 permeate flux. K et al. [28] employed artificial neural networks and ANFIS (Adaptive-Neuro

83 Fuzzy Inference System) to model forward osmosis treatment of textile wastewater.  
84 Hosseinzadeh et al. [29] employed ANN and the ANFIS to model water flux in an osmotic  
85 membrane bioreactor. The prediction of all of these studies is limited to water flux prediction,  
86 which otherwise can also be predicted accurately using solution diffusion or empirical models.  
87 Fundamentally, the problem is with the prediction of internal concentration polarisation,  
88 where solution diffusion models or empirical models have limitations. As a result, the use of  
89 ML techniques can be a viable tool to predict the ICP in the forward osmosis process.

90 This research aims to assess and compare various ML models for ICP prediction and compare  
91 the prediction to solution diffusion (SD) models. ML predictions can be superior to SD models  
92 since ML algorithms can learn from the input data, find hidden patterns and predict the  
93 output with high precision and accuracy. On the other hand, using solution diffusion-based  
94 models, for example, if  $(\text{NH}_4)\text{HCO}_3$  (ammonium bicarbonate) or  $\text{Ca}(\text{NO}_3)_2$  (calcium nitrate)  
95 is used as a draw solution, the diffusion coefficient of these compounds is not accessible in  
96 the literature, and the solution diffusion model would fail to estimate the severity of the ICP.  
97 Furthermore, when draw solutions mixture is employed, calculating the diffusion coefficient  
98 becomes difficult. As a result, we propose and analyse several ML models for ICP prediction  
99 based on various inputs and FO process parameters for the first time. By applying different  
100 ML algorithms to the dataset, two critical questions will be addressed; which ML algorithm is  
101 the most effective for the prediction of ICP modulus, and what is the potential of ML models,  
102 in general, to give insights into the most important features that impact ICP on the FO process.  
103 For assessing the modulus of ICP, the findings are compared to the SD model.

## 104 **2. Theory and methods**

## 105 **2.1. Study design**

106 Initial data were collected from published papers that reported values for K (solute resistance  
107 to diffusion), "S," predicted water flux using SD models, and experimental water flux. The rest  
108 of the data for input parameters were gathered from the experimental design and results of  
109 each study after the published studies were chosen (**Appendix Figure A.3 and A.4, heatmaps**)  
110 [5, 8, 11-15, 17, 30-33]. ML modelling was done in Python using Google Collaboratory or  
111 Google Collab on the Google research platform, a free Jupyter notebook in the Google cloud.  
112 Modelling was done following each ML algorithm-specific methodology without major  
113 feature engineering.

## 114 **2.2. Data collection and pre-processing**

115 The data collected for the ML models included a total of 438 instances and 19 features,  
116 including output features for CICP (concentrative internal concentration polarization) and  
117 DICP (dilutive internal concentration polarization) [10, 17, 34-36]. The total number of data  
118 points for DICP were 351 instances and 19 features (Heatmap is shown in Figure A.3, Appendix  
119 A. On the other hand, the total number of data points of CICP was 84 instances and 19  
120 features (Heatmap shown in Figure A.4, Appendix A). To the best of our knowledge, no study  
121 recommends a minimum quantity of data for the ANN and other ML algorithms to perform  
122 well; however, some studies have reported good performance even with datasets ranging  
123 from 20-80 data points [37, 38]. The data from images were extracted using an online  
124 software known as "WebPlotDigitizer 4.5". The data included all the input parameters that  
125 impact internal concentration polarization in the FO process, except reverse salt flux (RSF)  
126 since RSF data was scarce in the published studies. The forward osmosis system used  
127 throughout all the studies was operated in the FO mode, also termed the AL-FS mode, and  
128 the PRO mode, also known as the AL-DS mode (**Table 1**). The crossflow velocities ranged from

129 0.0192 m.sec<sup>-1</sup> to a maximum of 1.22 m.sec<sup>-1</sup>. Recent studies have revealed that flow  
 130 arrangement impacts the ICP [15]; thus, it was also considered an input parameter. Most of  
 131 the data were collected using commercial CTA, TFC and NaNoH<sub>2</sub>O membranes provided by  
 132 different suppliers of FO membranes. All of the studies' pure water permeability A value, salt  
 133 rejection R-value, and salt permeability B value were collected. Later, rejection rate data were  
 134 not considered in the input since most studies reported a similar rejection for the FO  
 135 membrane. The osmotic pressure data calculated by OLI software was taken directly from the  
 136 studies or calculated by Van't Hoff equation when it was not directly available. The data was  
 137 normalized in Python by utilising the data's minimum and maximum values.

$$138 \quad X_{sc} = \frac{x-x_{min}}{x_{max}-x_{min}} \quad (1)$$

139 **Table 1:** Input and output parameters for the collected data

Input/output Parameters	Mean	Maximum value	Minimum value	Standard deviation
Flow arrangement (Cocurrent or counter current)	N/A	N/A	N/A	N/A
Membrane orientation AL-FS	N/A	N/A	N/A	N/A
Membrane orientation AL-DS	N/A	N/A	N/A	N/A
FS cross-flow velocity (CFV)	0.21	122.22	0.56	26.64
DS CFV	0.21	122.22	1.92	26.26
FS molarity	0.13			0.21
FS Osmotic Pressure	6.59	52.02	0	10.76
DS molarity	1.36 M	9	0.04	1.03



DS osmotic pressure	71.98	1.96	483.14	54.71
FS type	N/A	N/A	N/A	N/A
DS type	N/A	N/A	N/A	N/A
FS temperature	300.98 K	323	273	6.61
DS temperature	300.98 K	323	273	6.23
DS Molecular weight	67.47 g	119.02	47.60	12.02
Osmotic pressure difference	71.98 atm	0	472	52.54
Membrane pure water permeability (A) value	3.01E-07 m/sec.atm	7.12E-07	1.1E-07	1.24E-07
Membrane salt permeability (B) value	6.24E-07 m/s	9.8E-08	5E-08	4.18E-07
Water flux	3.92E-06 m/s	2.1E-05	2.7E-10	3.41E-06
DICP (output)	0.38	1	0	0.33
CICP (output)	5.72	1	147.10	19.26
Mass transfer coefficient ( $k$ )	2.81E-05	6.3E-05	1.7E-05	1.93E-05
Solute resistance to diffusion ( $K$ )	2.6E05	3.5E05	2.5E05	3.1E05

140 \*RSF data was not available for most studies in the literature and CP models, therefore was excluded.

#### 141 **Data for solution diffusion and empirical model**

142 The ICP modulus calculation requires the value solute resistance to diffusion. The  $k$  value  
143 (mass transfer coefficient of the FO channel) and  $K$  (solute resistance to diffusion) must be  
144 calculated for the SD model. This was either taken directly from the studies where it was  
145 available or calculated accordingly to the two models of [10] and [13]. According to [10],  
146 water flux in the FO process operating in the PRO mode (active membrane layer faces the  
147 draw solution) is given by Equation (2). For the FO membrane operating in the FO mode  
148 (active membrane layer faces the feed solution), water flux is presented by Equation (3).

149  $J_w^{PRO} = A \left[ \pi_{Db} \exp\left(\frac{-J_w}{k}\right) - \pi_{Fb} \exp(J_w K) \right]$  (2)

150  $J_w^{FO} = A \left[ \pi_{Db} \exp(J_w K) - \pi_{Fb} \exp\left(-\frac{J_w}{k}\right) \right]$  (3)

151 Where  $J_w$  is the experimental flux,  $\pi_{Db}$  and  $\pi_{Fb}$  are the bulk feed concentration of draw and  
 152 feed solution, respectively,  $k$  is the convective mass transfer coefficient, and  $K$  is the salt  
 153 resistivity. Unfortunately, the proposed model neglects the effects of salt transport and the  
 154 external mass transfer resistance on the support layer [14]. Yip et al. [13] presented a  
 155 modified mathematical model in Equations (4) and (5) for the FO and the PRO orientation,  
 156 respectively.

157  $J_w^{FO} = A \left[ \frac{\pi_{Db} \exp(-J_w K) - \pi_{Fb} \exp\left(\frac{J_w}{k}\right)}{1 + \frac{B}{J_w} \left\{ \exp\left(\frac{J_w}{k}\right) - \exp(-J_w K) \right\}} \right]$  (4)

158  $J_w^{PRO} = A \left[ \frac{\pi_{D,b} \exp\left(-\frac{J_w}{k}\right) - \pi_{F,b} \exp(J_w K)}{1 + \frac{B}{J_w} \left\{ \exp(J_w K) - \exp\left(-\frac{J_w}{k}\right) \right\}} \right]$  (5)

159 The mass transfer resistance at the porous support layer, on the other hand, was not  
 160 considered in equations (4) and (5), but it's by far the most widely used model for the  
 161 calculation of  $K$  value. The experimental flux was taken as reported directly from the studies.  
 162 The modelling studies considered the initial water flux (30 minutes), and constant draw  
 163 solution concentration is maintained, or the dilution of draw solution is ignored and not  
 164 considered.

### 165 2.3. Design settings and performance evaluation metrics

166 Two different models were designed and evaluated for DICP and CICIP, respectively (**Table 2**).  
 167 The former CP occurs in the AL-FS mode and the latter CP in the AL-DS mode, and both do not  
 168 occur simultaneously in the FO process, so one was made redundant when modelling for the  
 169 other parameter. The efficacy of the constructed ML models can be determined using the  
 170 correlation coefficient ( $R^2$ ), which is defined as:

171 
$$R^2 = \frac{\sum_{i=1}^n (P_i - \bar{y}_i)^2}{\sum_{i=1}^n (y_i - \bar{y}_i)^2} \quad (6)$$

172 Where  $P_i$  denotes the predicted output of the ML model,  $\bar{y}_i$  denotes the mean of the sample  
 173 data, and  $y_i$  represent the actual output. In general, R-square values range from 0 to 1, with  
 174 a higher R-squared value indicating a better model. The MSE (mean square error) for each  
 175 model was calculated using equation (7).

176 
$$MSE = \frac{1}{n} \sum (y_i - P_i)^2 \quad (7)$$

177 The MSE indicates how close the regression model line (or fit) is to the data. The MSE of the  
 178 training data in supervised ML shows how good is the model in detecting the anomaly in the  
 179 dataset. However, it is the MSE of the test data, which indicates the performance of the ML  
 180 model. In this study, the MSE for the test data was considered the performance indicator of  
 181 the predictive power of the ML models. In general, the higher the  $R^2$  value and lower the MSE,  
 182 the better the model's predictive power. The different models were built and run on the  
 183 Google Colaboratory platform using Python (Python 3.9) using the built-in libraries. Python  
 184 offers numerous advantages over MATLAB for ML as Python is free, open-source and has  
 185 various inbuilt libraries for ML such as NumPy, Matplotlib, Keras and Pandas. Further, the  
 186 back-end libraries in Python, such as TensorFlow, chose the best and efficient way to define  
 187 and train an ML model with a few lines of code. The model's input data comprised the FO's  
 188 independent variables, and the output layer consisted of either CICP or DICP based on the  
 189 membrane orientation.

190 **Table 2:** Different ML approaches used in this study

Model Name	Output	Validation method	Python Library
ANN	CICP/DICP	R square/MSE	TensorFlow

Extreme Gradient boosting (XGBoost)	CICP/DICP	R square/MSE	XGBoost
CatBoostRegressor	CICP/DICP	R square/MSE	CatBoost
RandomForestRegressor	CICP/DICP	R square/MSE	Sklearn.ensemble
Linear regression	CICP/DICP	R square/MSE	Scikit-Learn

191

192 **2.4. Artificial neural network**

193 This study designed a neural network in Python, using the Keras (A Python library for neural  
 194 networks). After loading the dataset, the model was one-hot encoded to create dummies  
 195 from all the categorical variables such as flow arrangement (countercurrent or co-current  
 196 flow), membrane orientation (AL-FS or AL-DS), types of FS and DS. This method of encoding  
 197 was chosen as there was no direct relationship between the categorical input variables.  
 198 Customized models were trained for both datasets of DICP and CICP. In the datasets, various  
 199 categories of input variables were processed into dense layers after the sequential model was  
 200 implemented.

201 **2.5. Gradient Boosting Models Theory**

202 Three gradient tree boosting models were employed in this study, known as extreme gradient  
 203 boosting or XGBoost, Categorical boosting or CatBoost and RandomForest. The most common  
 204 functions (predictive learners) used in the Gradient Boosting framework are Decision Trees  
 205 [39]. Let's say we have a set of input variables  $x = \{x_1, x_2, x_3 \dots \dots \dots x_p\}$ . The Decision Trees  
 206 are built in a greedy manner, with the best split points chosen based on purity scores such as  
 207 Gini or to minimise the loss. In this model, we have all the parameters which have an impact  
 208 on ICP as input variables. Our output variable  $y$  is either CICP or DICP. In Gradient Boosting

209 models, a single terminal tree is used to create an initial basis function  $F_0(x)$  given by  
210 equation (9) [40, 41].

$$211 \quad F_0(x) = \operatorname{argmin}_{\beta} \sum_{i=1}^n L(y_i, \beta) \quad (9)$$

212 In equation (6)  $L(y_i, \beta)$  represents the loss function,  $\beta$  represents the set of split points for  
213 the tree's internal nodes, and  $n$  represents the number of input and output variables. Using  
214 some samples from a training dataset, the goal is to find the values of  $\beta$  that minimize the  
215 loss function to the minimum. Detailed methodology and equation derivation can be found  
216 in [40].

### 217 **3. Results and Discussions**

218 In the FO mode or the AL-FS orientation, the DS is diluted inside the support layer leading to  
219 DICP output for the ML model. On the other hand, when the membrane orientation is the AL-  
220 DS mode, the model's output is the CICIP. The modulus of ICP is always less than 1, whereas  
221 the modulus of CICIP is always greater than 1. The "K" value reported for SD models is usually  
222 higher in the AL-DS mode than the AL-FS mode.

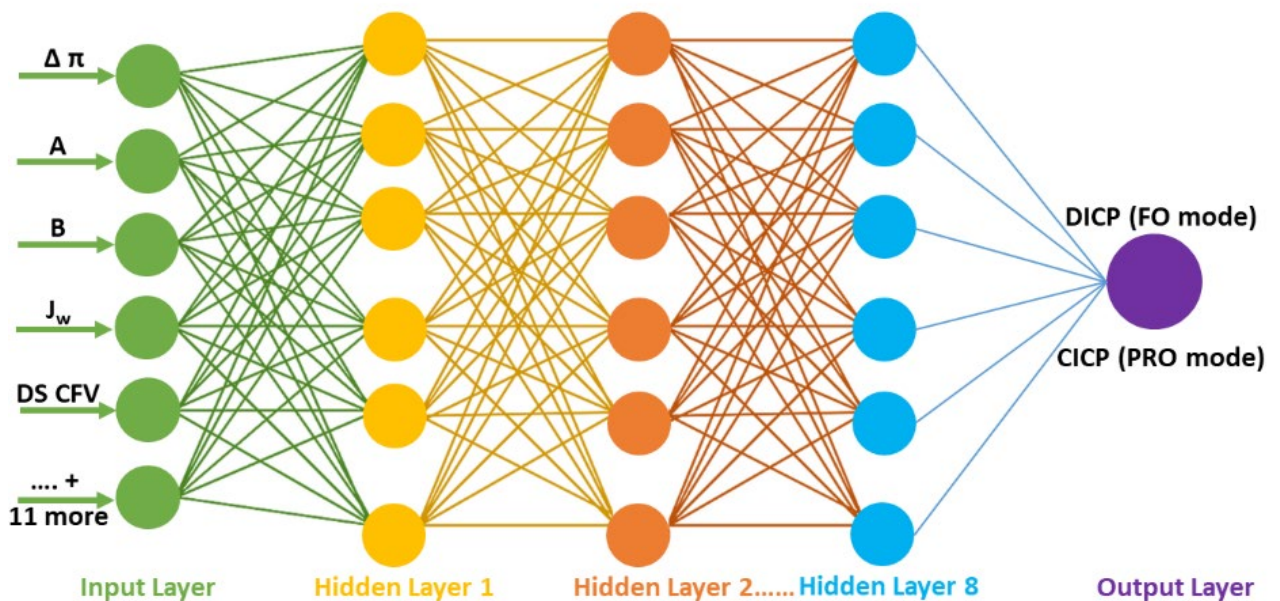
#### 223 **3.1. Prediction of DICP and CICIP using ANN**

224 A neural network using a deep learning approach was designed in Python to predict DICP and  
225 CICIP, respectively. Since there is no linear relationship between the input and output  
226 parameters, to create an efficient neural network, we must carefully choose the appropriate  
227 or the optimum number of layers, the learning rate of the algorithm, and the number of  
228 training epochs. For instance, if the hidden layer contains insufficient neurons, the ANN will  
229 not reflect nonlinearity in the training data and lead to inaccurate predictions. Initially, setting  
230 the hidden layers to one, the optimal number of hidden layers was determined through trial

231 and error. An initial neural network was designed by considering all the input parameters  
232 listed in Table 1 without hyperparameter optimisation. The designed neural network had an  
233 input layer, three hidden layers, and an output layer (for CICIP or DICIP, depending on the  
234 membrane orientation). The neural network exhibited good predictive power for the CICIP;  
235 however, it was found that this neural network was prone to overfitting and didn't exhibit  
236 good predictive power for the DICIP dataset (Fig.A.1 for DICIP and Fig.A.2 for CICIP (Appendix  
237 A.1)).

238 Further optimisation was performed by selecting appropriate inputs to the ANN model based  
239 on the correlation matrix indicated in Fig. A.3 and Fig.A.4 for the DICIP (blue) and CICIP (pink),  
240 respectively (Appendix A.1). It was found from the heatmap that several parameters have  
241 correlated to each other with a correlation factor of 1, such as the correlation of FS CFV with  
242 DS CFV and FS molarity with FS osmotic pressure. It meant one of those should not be  
243 considered as input and should be removed from the input dataset, which may increase the  
244 ANN model performance. As a result, FS CFV and FS molarity were removed from the main  
245 dataset. In general, DS CFV has more impact on the ICP than FS CFV, and FS osmotic pressure  
246 is generally considered in all ICP equations rather than molarity. The hyperparameters tuning  
247 of the ANN model was further performed through Grid search (GridSearchCV in Python),  
248 leading to selecting the best parameters for the model performance. Additionally, batch  
249 normalisation was added after the activation functions for the normalisation of the output of  
250 each layer. The resulting optimised neural network had an input layer, eight hidden layers,  
251 and an output layer (for CICIP or DICIP, depending on the membrane orientation). It should be  
252 noted that the optimal number of hidden layers was determined through trial and error.  
253 Using Python Scikit learn, 80 % of the input data was used as the training set for the neural  
254 network, 10% was used to validate the data, and the rest of the 10 % was utilized for testing

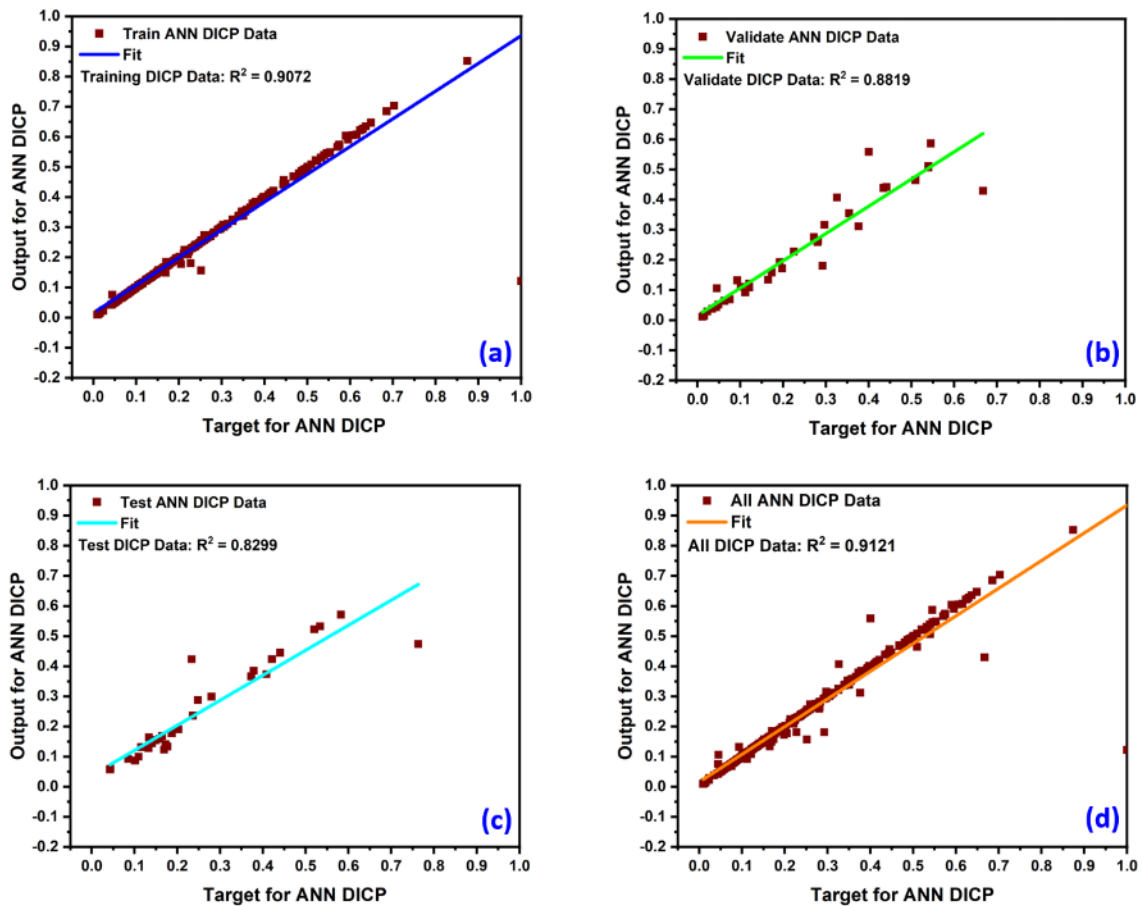
255 the ANN for the prediction of DICP and CICIP. The ANN was implemented using the TensorFlow  
 256 Sequential API, and the Relu (rectified linear activation function) and LeakyRelu activated  
 257 function was chosen for the DICP and the CICIP models. The categorical data include  
 258 membrane orientation (AL-FS or AL-DS), flow arrangement (countercurrent or cocurrent), FS  
 259 types such as NaCl, DI water, KCl, DS types and membranes in the dataset was transformed  
 260 into numerical values using Python "LabelEncoder". To reduce the learning rate and avoid  
 261 overfitting of the model, "early stopping" was implemented. After some trial and error with  
 262 the number of epochs based on the validation R square and mean square error, the neural  
 263 network with the best possible results was chosen and presented here in this study (Fig.1).



264

265 **Figure 1:** ANN network for prediction of DICP and CICIP

266 For each phase of the simulation, the  $R^2$  value was calculated for the training, validation,  
 267 testing, and all data. For the DICP model, The  $R^2$  values for training and test data are 0.90 and  
 268 0.88, respectively (Fig 2a and 2c). The testing data  $R^2$  (0.82) shows a good performance of the  
 269 optimised ANN compared to the initial one (reported in Appendix A.1).



270

271 **Figure 2:** Predictive power of the ANN using  $R^2$  square metric for the prediction of DICP, **a)**

272 For training data, **b)** For validating the data, **c)** for testing the data, **d)** for all data.

273 Since the  $R^2$  value for the testing set is rational, the neural network shows good

274 generalization overall for the prediction of DICP using the given dataset (**Fig 2c**). The MSE

275 for the testing data was calculated to be 0.075, which shows the model good performance for

276 the DICP prediction for the testing data. The MSE of the training data for the DICP model was

277 0.0029. Thus, the test MSE reveals that the model has good generalisation ability and can

278 predict output variables of the test data with good accuracy.

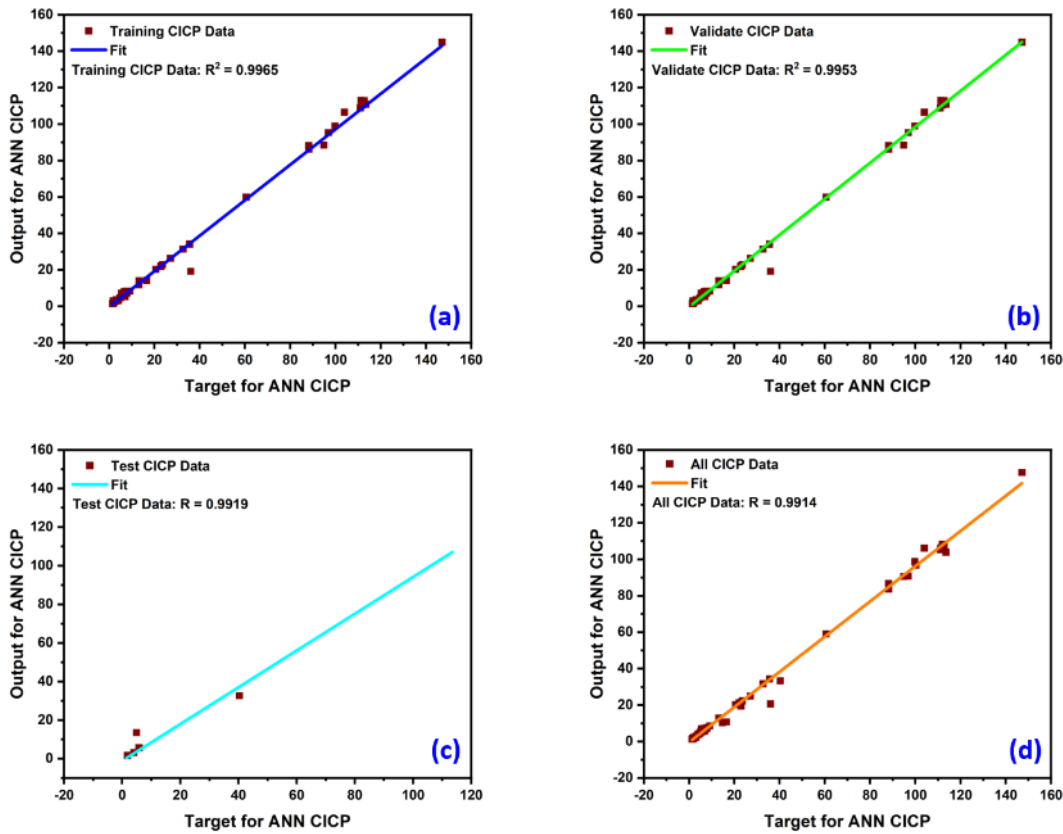
279 Compared to the DICP, the neural network for CICIP gave a high  $R^2$  value for the training,

280 validation and testing data (**Fig. 3a, 3b, 3c**). Overall, the  $R^2$  value for all the data was quite

281 high (0.99), as presented in **Fig. 3d**. The  $R^2$  value for the training data was almost closer to 1;



282 on the other hand, the  $R^2$  value of the testing data (0.9730) was higher than that of the DICP  
283 prediction for the neural network. The MSE for the CICIP test data was evaluated to be 5.07.



284

285 **Figure 3:** Predictive power of the ANN using  $R^2$  square metric for the prediction of CICIP, **a)** For  
286 training data, **b)** For validating the data, **c)** for testing the data, **d)** for all data.

287 On the other hand, the MSE of the training data for the CICIP model was 2.74. The higher MSE  
288 for the CICIP may be due to the smaller data points for the CICIP dataset than the DICP. Since  
289 CICIP is more prevalent in the PRO mode [42], ANN can be a viable tool to predict the CICIP in  
290 the FO process with great accuracy In the PRO mode. Further optimisation can improve the  
291 accuracy of the ANN by minimising the number of inputs further to the neural network.  
292 However, further optimisation was not performed in this study. Although ANN can be a  
293 powerful tool and can predict CICIP with high accuracy, it should be noted that the ANN

294 network consists of ten thousand neurons with hidden relationships between the input and  
295 output functions. With such modelling, users can get no insights into which feature of the  
296 data has a high or low impact on shaping the neural network's output. Further, a major issue  
297 in the ANN is the overfitting of the model [43]. Overfitting occurs when the model's training  
298 data shows good performance but has poor predictive power for the test data, especially for  
299 larger datasets. It should also be emphasised that the modulus of the ICP in the FO process  
300 does not give much information alone unless we know what input parameters to the FO  
301 process affect its severity. Therefore, it is important to evaluate other advanced models to  
302 better predict the ICP in the FO mode and the CICP in the PRO mode.

### 303 **3.2. Prediction of DICP and CICP using Gradient tree Boosting models**

304 Different tree-based models were also evaluated in this study for the prediction of DICP and  
305 CICP data. In tree-based machine learning models, the residuals are utilised to correct the  
306 previous prediction at each iteration of gradient boosting, allowing the stated loss function  
307 to be optimised [44]. Three different tree-based models, XGBoost, CatBoost and Random  
308 Forest, were evaluated to predict DICP and CICP modulus, respectively. The initial model was  
309 implemented using Extreme Gradient boosting, also known as XGBoost, to predict DICP and  
310 CICP and compared to the optimised ANN. The same dataset was used for the tree models  
311 like the one used for optimised ANN (removing FS CFV and FS molarity). XGBoost is an  
312 ensemble learning algorithm that uses boosted trees, and by weighting, the learning rate aids  
313 in shrinking the boosting process, making fitting more conservative [45]. The main advantage  
314 of using this algorithm is that XGBoost can provide insights into the importance of various  
315 inputs that impact the ICP in the FO process. The XGBoost algorithm also offers several other  
316 advantages in ML, such as cross-validation, high flexibility and speed of operation compared  
317 to the neural network approach. For XGBoost, 80 % of the data was used for training, and the

318 remaining 10 % was used for validation and testing (10% for validation, 10 % for testing). The  
319 models were trained on the training set and the parameters were adjusted based on their  
320 performance on the validation set. The predictive power was evaluated based on the metrics  
321 of validation and test data  $R^2$  and MSE. The results from the XGBoost simulation  
322 demonstrated that the model using this dataset is rational for predicting DICP with an  $R^2$  value  
323 of 0.94 for the training data, 0.92 for the validation data and 0.81 for the test data (**Fig.4a**).  
324 The MSE for DICP prediction on the test data was 0.076, confirming the model's efficient  
325 predictive power for the DICP data. On the other hand, the  $R^2$  value for the prediction of CICIP  
326 was 0.92 for the validation data and 0.88 for the test data (**Fig.4b**). For the prediction of the  
327 DICP data, the XGBoost shows almost similar predictive power as the ANN in terms of  $R^2$  and  
328 MSE; however, the test data  $R^2$  for this model for the CICIP model was lower than the neural  
329 network modelling for the CICIP, and hence it was not optimised any further. It should be  
330 noted that XGBoost performs better than ANN generally in terms of computational time.  
331 Furthermore, XGBoost requires less coding and hyperparameter tuning compared to the ANN  
332 program implemented in this study.

333 A slightly different approach to XGBoost is the categorical boosting, also known as the  
334 CatBoost algorithm, which was evaluated to predict DICP and CICIP in this study. The main  
335 advantage of Categorical boosting or CatBoost over other models is that it can learn from its  
336 mistakes during the learning phase of the simulation. Furthermore, it is easy to implement,  
337 has good stability, has less computational requirements and workload [46], and can  
338 sometimes outperform XGBoost in terms of accuracy. Furthermore, we do not have to pre-  
339 process any categorical variables such as membrane orientation, types of draw and feed  
340 solutions, flow arrangement and membrane types. The basic idea behind boosting is to  
341 combine numerous weak models sequentially and create a strong competitive predictive

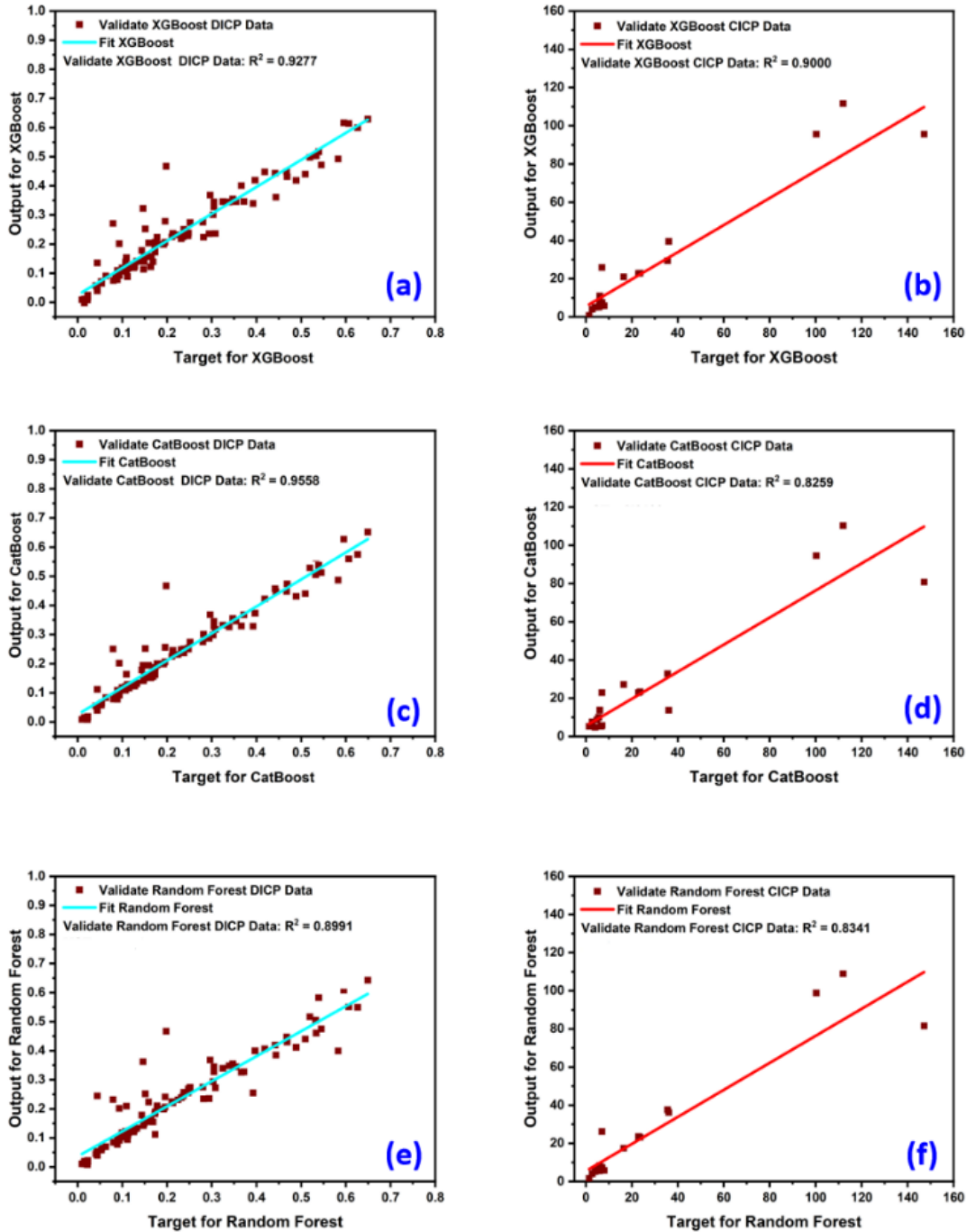
342 model via greedy search. Unlike other models, this ML method does not require large data to  
343 learn from and perform efficiently, even with a smaller dataset. Similar to the XGBoost  
344 simulation, the data was split into 80 % training data, and the remaining 10 % was used to  
345 validate and test the model on a dataset not seen by the model. It should be noted that the  
346 model simulated here was done using minimal feature engineering. **Fig. 4c** and **Fig.4d** show  
347 the model performance for DICP and CICIP, respectively. For both the CICIP and DICP, the  
348 learning rates of the models were set to default. For CICIP, the learning rate obtained from the  
349 program was 0.17065, whereas for the DICP model, the learning rate was 0.2142. The best  
350 performance was obtained for DICP predictions with an  $R^2$  value of 0.99 for the training data,  
351 0.95 for validation data, and 0.92 for the test data. The MSE for the DICP model was 0.049,  
352 which shows the model superiority over the Xgboost algorithm and the ANN. The  $R^2$  value of  
353 CICIP was also superior to both ANN and XGBoost ( $R^2=0.99$  for training data and 0.99 for the  
354 test data). It should be noted that the  $R^2$  can be boosted further by doing some feature  
355 engineering, later discussed in the next section. For both the DICP prediction and CICIP, the  
356 CatBoost provides the highest  $R^2$  and lower MSE compared to the ANN and Xgboost.

357 To overcome the limitations of boosting ML algorithms for the prediction of ICP, the study  
358 checked the validity of the Random Forest regressor, which uses multiple decision trees (that  
359 is why called forest) and can perform both regression and classification tasks using a  
360 technique known as bagging. Instead of relying on individual decision trees, the basic idea of  
361 Random Forest ML is to combine multiple decision trees to determine the final output, in this  
362 case, CICIP or DICP. Random Forests, in general, produce better results, perform well on large  
363 amounts of data, and work with missing data by generating estimates for it. Further, it avoids  
364 the overfitting of the model since overfitting is one of the most serious problems in machine  
365 learning. Still, it is rarely a problem with the random forest method. If the forest contains a

366 sufficient number of trees, the classifier will avoid overfitting the model. The RandomForest  
367 regressor was used in Python for fitting the Random Forest regression to the forward osmosis  
368 data. The “. fit ()” function fits the training input and output data to the RandomForest  
369 regressor. The hyperparameter of the model (n\_estimators) was set to 100. This is simply the  
370 number of trees the algorithm constructs prior to performing maximum voting or calculating  
371 prediction averages. In general, increasing the number of trees improves performance and  
372 increases the stability of the predictions. **Fig. 4e** and **4f** show the  $R^2$  value for the DICP and  
373 CICIP for the Random Forest Regressor with 0.81  $R^2$  for the DICP test data and 0.93 for the  
374 CICIP test data. The MSE for the DICP and CICIP calculated on the test data for this model was  
375 0.076 and 12.16, respectively. Overall, the Random Forest regression provides an acceptable  
376  $R^2$  and MSE score for CICIP and DICP; however, the DICP  $R^2$  score obtained is less than the  
377 CatBoost model. On the other hand, the MSE for the DICP of this model is also slightly higher  
378 than the CatBoost. While the model simulation performance was almost comparable with the  
379 Xgboost and ANN, it should be noted that if there are many trees in a random forest model,  
380 the model's predictive power can become very slow and inefficient, hindering the model  
381 application in the real world. For this reason, the model was not further optimised. Table 3  
382 lists the tree-based algorithm training data  $R^2$ , testing data  $R^2$  and MSE of the testing data.

383

384



385

386 **Figure 4:** Predictive power of the different gradient tree models using R<sup>2</sup> square metric for  
 387 validation data **a)** XGBoost for validating the data of DICP, **b)** XGBoost for validating the data  
 388 of CICIP, **c)** CatBoost for validating the data of DICP, **d)** CatBoost for validating the data of CICIP,  
 389 **e)** Random Forest for validating the data of DICP, **f)** Random Forest for validating the data of  
 390 CICIP.

391 Table 3: Tree-based algorithms R<sup>2</sup> for training, validation and testing data and MSE for testing  
 392 data

	Training Data R <sup>2</sup>	Testing Data R <sup>2</sup>	MSE (Test Data)
XGBoost DICP	0.94	0.81	0.0771
XGBoost CICIP	0.99	0.88	13.32
CatBoost DICP	0.99	0.92	0.049
CatBoost CICIP	0.99	0.99	3.61
Random Forest DICP	0.96	0.81	0.076
Random Forest CICIP	0.98	0.90	12.16

393

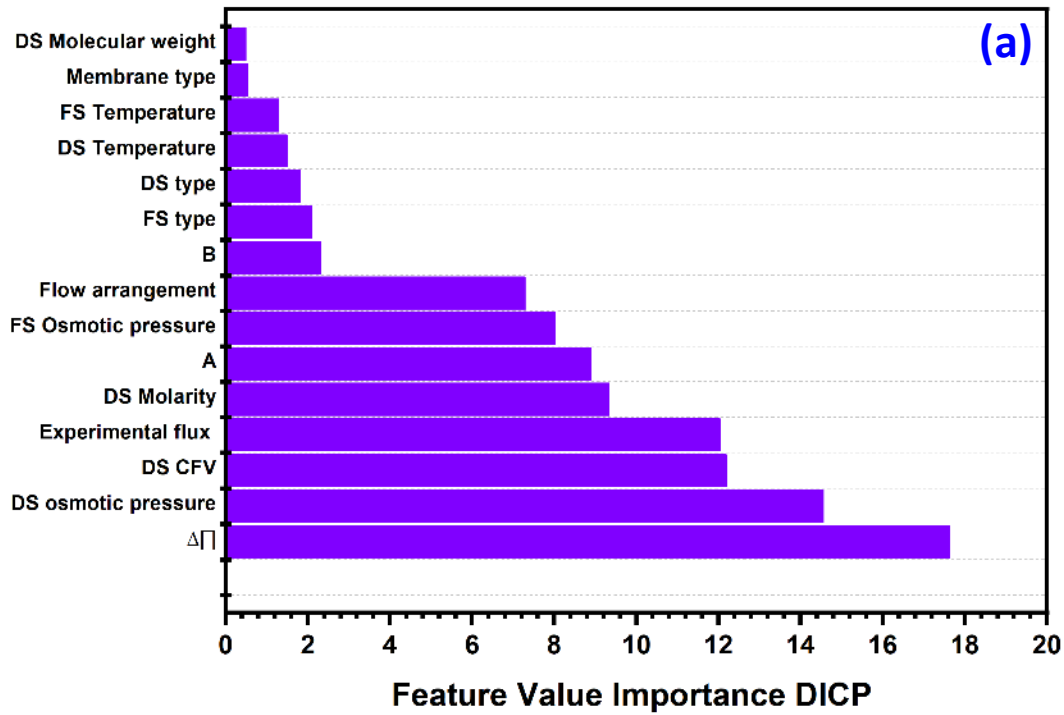
### 394 3.3. Feature importance Scores for the Catboost model

395 Tree-based algorithms such as Xgboost, CatBoost and Random Forest can provide further  
 396 insights about the importance of the input parameters or the input features using a metric  
 397 called “Feature Importance” or FI score. Since there are many complex relationships between  
 398 the input parameters to the FO process and the ICP, the importance of various input  
 399 parameters should be considered when building a model for ICP. This will allow us to improve  
 400 the model further for accurate predictions and only consider important input parameters.  
 401 Feature importance was evaluated for the best performing model, CatBoost. **Fig. 5a** and **Fig.**  
 402 **5b** present the FI for the training data of CatBoost for DICP and CICIP, respectively. For DICP,  
 403 it can be observed that the highest score is for the osmotic pressure difference (FI score of  
 404 17.64) followed by the DS osmotic pressure (FI score of 14.55). The result implies that the  
 405 osmotic pressure difference and DS osmotic pressure values have a significant impact on the  
 406 model, whereas the features with the lowest FI score values have a small impact on the  
 407 modelling performance, such as DS molecular weight and membrane type. Generally, the  
 408 highest the osmotic pressure difference between the FS and the DS, the higher the dilution

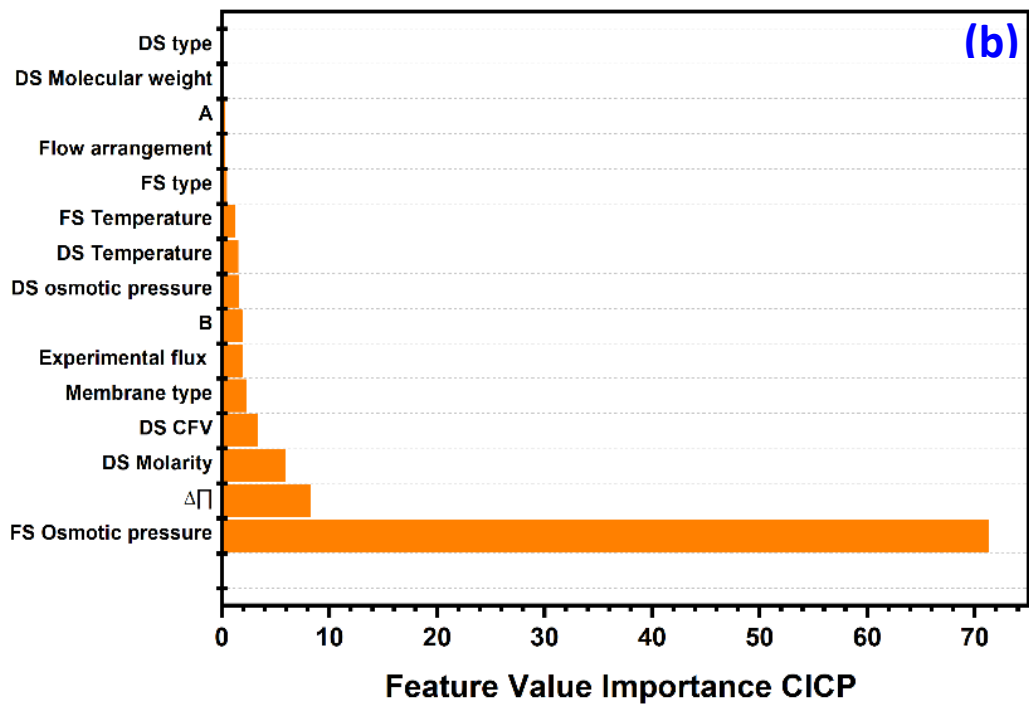
409 and thus the higher the DICP value. On the other hand, the higher the DS osmotic pressure,  
410 the higher the permeation drag of the DS enabling more water transport across the FO  
411 membrane, and therefore, more dilution of the DS, leading to higher values of the DICP. DS  
412 CFV and experimental flux have also a significant impact on the DICP. For instance, higher DS  
413 CFV will result in increased flux in the forward osmosis process, and thus reducing the impacts  
414 of DICP. Recent studies have also confirmed that increasing CFV will result in minimizing the  
415 impacts of DICP as well as CACP [15]. For CACP (**Fig.5b**), the highest score is for the osmotic  
416 pressure of the FS and is significantly higher than all the other features. Generally, when the  
417 FS is DI water (FS osmotic pressure=0), a higher water flux is obtained in the PRO mode, as  
418 there is no CACP on the FS of the membrane. On the other hand, when the FS in the PRO mode  
419 is FS with high osmotic pressure, such as wastewater, the flux in the PRO mode is less than  
420 the FO mode due to the severity of the CACP. The lowest FI score for the CACP CatBoost was  
421 obtained for the DS type and the DS molecular weight. The lowest FI score for the CACP  
422 CatBoost model were obtained for the DS type and the DS molecular weight. As CACP occurs  
423 on the FS of the FO membrane, the DS type and the DS molecular weight may have a negligible  
424 effect on the CACP. DS weight and the type of DS shows to have no significant impact on the  
425 modelling of the CACP process, and hence, these can be ignored when modelling for the CACP  
426 process in the PRO mode. This will allow for more accurate predictions of CACP in the FO  
427 process in the PRO mode.

428





429



430

431 **Figure 5:** a) Feature importance score of the CatBoost model for DICP, b) Feature importance  
 432 score of the CatBoost model for CICP.

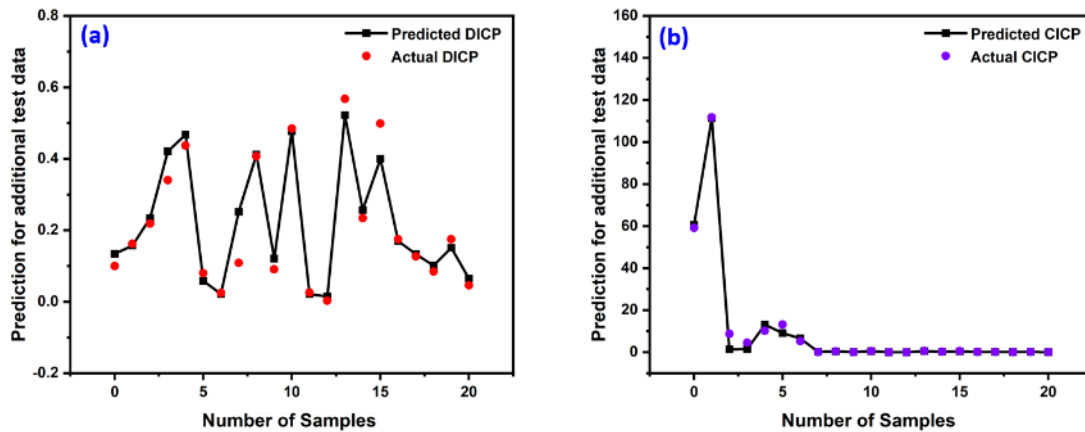
433 These can be made redundant when modelling the effects of CICP on the FO process. As CICP  
434 occurs on the FS of the FO membrane, the DS type and the DS molecular weight may have a  
435 negligible effect on the CICP.

436 The CatBoost FI score optimises the model by including the highest scores and neglecting  
437 those with the lowest scores. This enables the ML algorithm to predict the output more  
438 accurately. Features that are unimportant or relevant to the mode introduce noise in the  
439 modelling process, decreasing accuracy [47]. Since ML-based predictive models do not  
440 perform well when exposed to noisy data, particularly in the training phase. Furthermore, the  
441 model built on noisy data tends not to generalise well to data outside the training set, possibly  
442 leading to over-fitting the model. It is interesting to see that the feature importance of the  
443 CatBoost model for the test data is similar to the one obtained for the training data. **Fig. A.4**  
444 and **Fig.A.5 (Appendix A.1)** present the feature importance for DICP and CICP, respectively,  
445 for the CatBoost model, respectively. The feature importance is an important metric in the  
446 CatBoost model, and it tells us how much the prediction in CICP or DICP would change if any  
447 of the feature's change. Even though we use all of the features of the given dataset based on  
448 self-collected user data, not all of them contribute equally to the prediction of the CICP and  
449 the DICP. Modelling the most important features of the predictive model can lead us to a high  
450  $R^2$  value and lower MSE for the CatBoost algorithm. The model for the DICP or CICP can be  
451 further optimised by cleaning the data and removing the features with a very low score, such  
452 as type of membrane and DS molecular weight.

### 453 **3.4. Prediction of additional test samples for CatBoost**

454 **Fig.6a** and **Fig.6b** present additional test samples tested on the CatBoost model for DICP and  
455 optimised CatBoost model for CICP. The model shows very good agreement with the actual

456 values. It should be noted here that the test data here was outside of the training dataset (not  
457 used for training data or unseen data), so overall; the model exhibits very good generalisation.

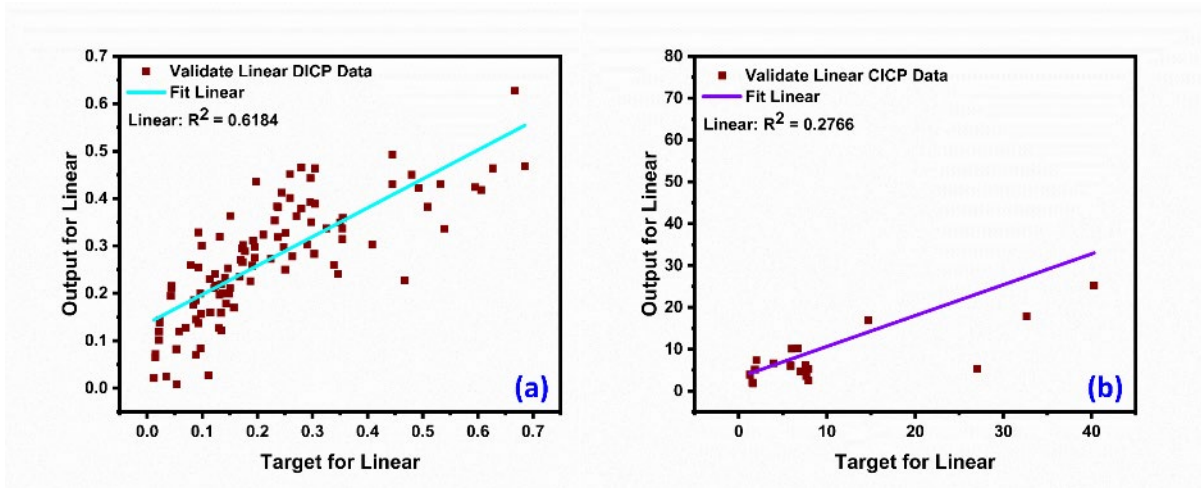


458

459 **Figure 6:** The predictive power of the CatBoost model using  $R^2$  square metric **a)** For validating  
460 the data of DICP, **b)** For validating the data of CICP.

### 461 3.5. Linear Regression

462 Linear regression was also evaluated in Python to test the prediction of the ICP. The  $R^2$  value  
463 for the DICP model for linear regression was 0.48, and the lowest  $R^2$  value amongst all the  
464 models of 0.27 was for the CICP prediction (**Fig.7a** and **Fig.7b**). This implies that the predictive  
465 power of linear regression is very bad for the CICP prediction and is lower than CatBoost for  
466 ICP predictions. This could be due to the linear regression model's inability to predict the ICP  
467 with different inputs with no linear relationships. As a result, the  $R^2$  is low, resulting in  
468 inaccurate results.



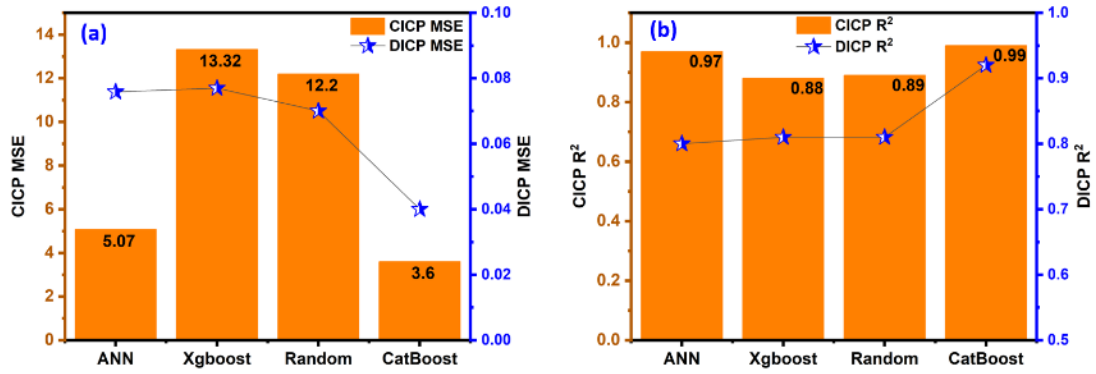
469

470 **Figure 7:** Predictive power of the linear regression model using  $R^2$  square metric, **a)** For  
 471 validating the data of DICP, **b)** For validating the data of CICP

472 **3.6. Summary of Comparison for ML models**

473 Amongst all the models, the best performance was obtained by CatBoost, followed by ANN.

474 **Fig.8a** and **8b** show the comparison of the MSE and  $R^2$  value for each model except linear  
 475 regression, which was excluded since the performance was very poor. It should be  
 476 emphasised that ANN requires data normalisation compared to CatBoost, which can handle  
 477 categorical values such as membrane orientation, DS type, FS type, flow direction and type of  
 478 membrane used in the FO study. CatBoost also has the additional ability to handle missing  
 479 values or information in the data by using estimates. Finally, in terms of execution and design,  
 480 CatBoost is much efficient and requires less computational and processing time.



481

482 **Figure 8:** Comparison of mean square error (MSE) and R<sup>2</sup> value for different models for CICP  
 483 and DICP, **a)** MSE for CICP and DICP, **b)** R<sup>2</sup> value comparison for different models for CICP and  
 484 DICP.

### 485 3.7. Comparison of CatBoost vs prediction by literature models

486 The prediction of the CatBoost was also compared with the solution diffusion model to show  
 487 the applicability of the ML models for CP prediction. As discussed before, the CatBoost model  
 488 for CICP was optimised further to improve the accuracy. The initial CatBoost model had a  
 489 good score for the prediction of DICP and the CICP in terms of R<sup>2</sup> and MSE. Although the  
 490 evaluation metrics can be further optimised with hyperparameter tuning and data cleaning,  
 491 the initial CatBoost model was not optimised further as the performance was efficient  
 492 compared to other models. Furthermore, major feature engineering is avoided here so that  
 493 environmental engineers do not have to apply complex codes, which otherwise would require  
 494 advanced programming skills in Python. The results are presented in **Table 4** with the  
 495 absolute error between the values obtained for the solution diffusion model and the CatBoost  
 496 algorithm.

497

498

499 **Table 4:** Additional test data for comparison against SD models.

<b>AL-DS Mode, Temperature of FS and DS =25, CFV of FS and DS = 0.36 Concentrative ICP Modulus</b>						
DS Osmotic Pressure atm	FS Osmotic pressure atm	Water flux m/s	K (experimental) s/m	K (average) s/m	CICP SD	CICP CatBoost
70.04	2.33	8.82E-06	2.02E+05	2.2 E05	5.93	7.2
70.04	4.67	7.35E-06	2.12 E+05	2.2 E05	4.75	5.18
70.04	23.35	3.11E-06	2.36 E+05	2.2 E05	2.08	2.00
70.04	46.70	1.14E-06	2.47 E+05	2.2 E05	1.32	1.29
<b>AL-FS Mode, Temperature of FS and DS = 25, CFV of FS and DS = 45.3 cm/s</b>					<b>DICP SD</b>	<b>DICP CatBoost</b>
70.04	0.00	5.05E-06	2.83E05	2.7E05	0.25	0.25
70.04	3.06	4.70E-06	2.80E05	2.7E05	0.28	0.28
70.04	6.00	4.36E-06	2.81E05	2.7E05	0.30	0.30
70.04	27.11	2.59E-06	2.56E05	2.7E05	0.49	0.47
70.04	49.63	1.06E-06	2.52E05	2.7E05	0.75	0.71

500

501 It is evident that the ML method used here can predict the internal concentration polarization  
 502 based on the process input parameters. A user can simulate and predict the ICP in the FO  
 503 process, as long as the membrane's operating parameters and the A and B are known.  
 504 Changes in the operating parameters would also reflect the corresponding changes in ICP and  
 505 the process performance. This will allow for improvement in membrane design without long

506 experimentation. On the other hand, with the solution diffusion model, modelling and  
507 prediction are done using a constant value of solute resistance to diffusion “ $K$ ”.

#### 508 **4. Conclusion**

509 Accurate measurement and mitigation of internal concentration polarization are vital for the  
510 FO process to compete against the existing state of the art RO and other technologies. ML  
511 models can accurately predict ICP in the FO process based on the input variables or process  
512 conditions. The following are the main findings from this study.

- 513 a) ML models can predict both dilutive and concentrative ICP in the FO process without the  
514 need for structural or solute resistance to diffusion inputs.
- 515 b) The models can also be applicable where the diffusivity of the draw solution is unknown.
- 516 c) Data captured from ML models can be used to optimize the FO process at the pilot scale.
- 517 d) The CatBoost algorithm best predicts dilutive and concentrative internal polarization in  
518 the FO process among the different ML models used in this study.
- 519 e) The feature importance score in the ML models can help select the best input variables  
520 for modelling and redesign the model based on the score.
- 521 f) The CatBoost model can predict ICP with an  $R^2$  value of 0.92 and has more predictive  
522 power than the ANN.

523 The model designed here is solely reliant on the input or trained data collected from the  
524 literature. The accuracy of the data is directly proportional to the accuracy of the model. Any  
525 discrepancies in the methodologies of the user data from the literature studies might lead to  
526 inaccurate predictions in the DICP or CICIP occurring in the FO process. ML models also  
527 perform well when there is a large amount of data available.

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## Appendix A

665 **Evaluation of Machine Learning Algorithms to Predict Internal Concentration Polarization**

666

### **(ICP) in Forward Osmosis (FO)**

667 Ibrar Ibrar<sup>1</sup>, Sudesh Yadav<sup>1</sup>, Ali Braytee<sup>2</sup>, Ali Altaee<sup>1, \*</sup>, Ahmad HosseinZadeh<sup>1</sup>, Akshaya K.

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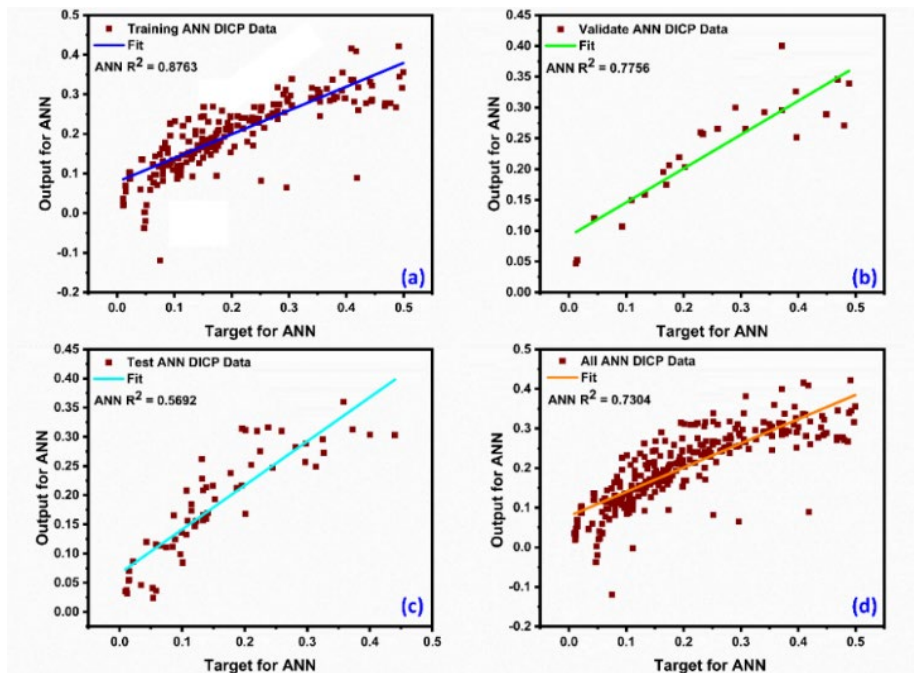
#### 676 **A.1. Prediction of DICP and CICIP with ANN without optimisation**

677 Initially, all inputs were considered without any consideration, and the ANN performance for

678 the training, validation data, testing data and all the data is presented here in Fig.A.1 and

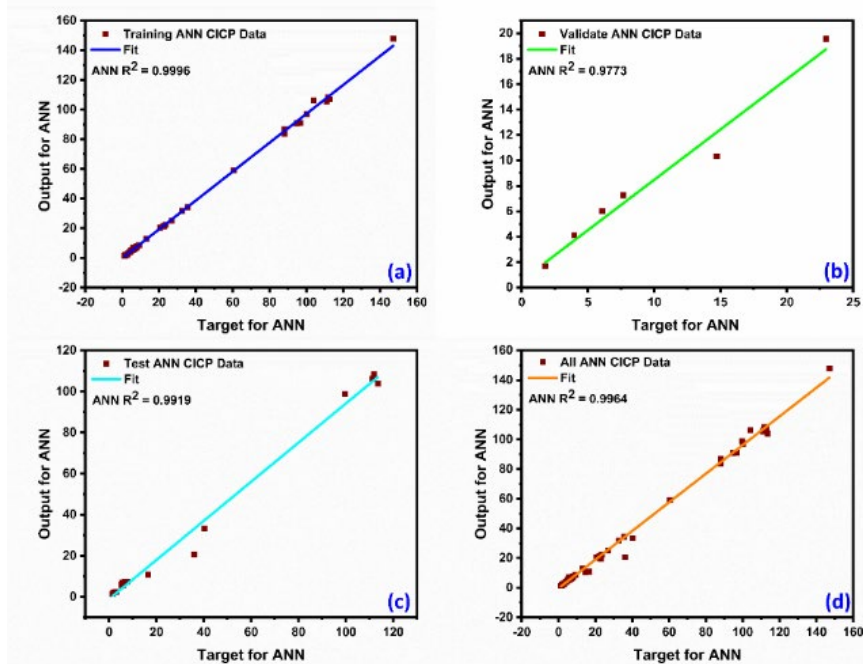
679 Fig.A.2. For the DICP data, the neural network was prone to overfitting. However, the

680 performance was acceptable for the CICIP data.



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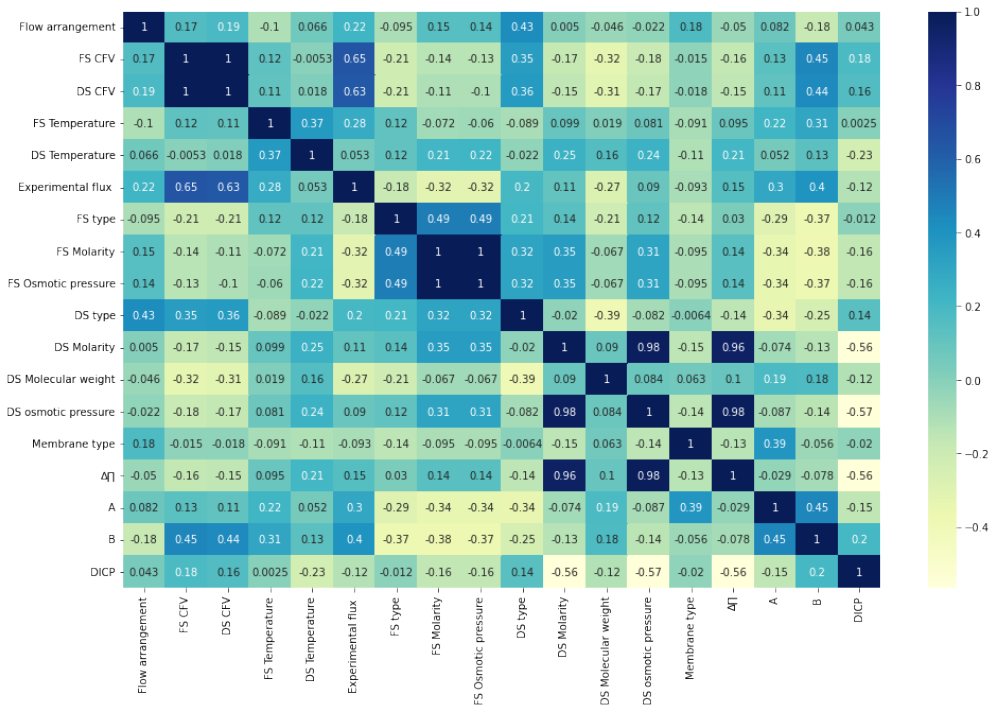
682 **Figure A.1.:** Predictive power of the ANN (no optimisation) using  $R^2$  square metric for the  
 683 prediction of DICP, **a)** For training data, **b)** For validating the data, **c)** for testing the data, **d)**  
 684 for all data.



685

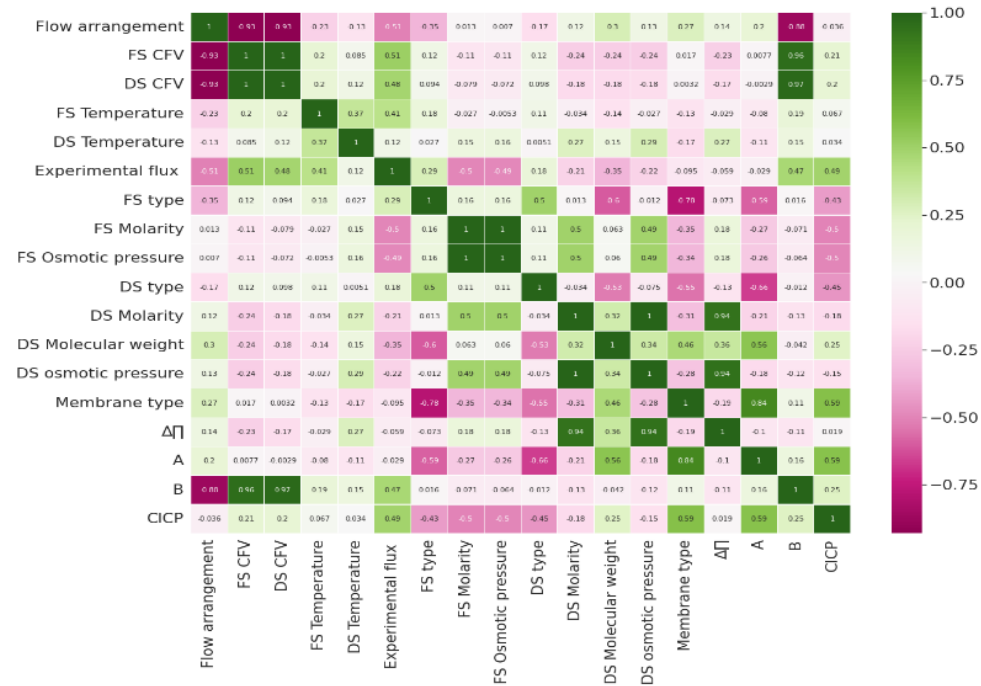
686 **Figure A.2.:** Predictive power of the ANN (no optimisation) using  $R^2$  square metric for the  
 687 prediction of CICP, **a)** For training data, **b)** For validating the data, **c)** for testing the data, **d)**  
 688 for all data.

689



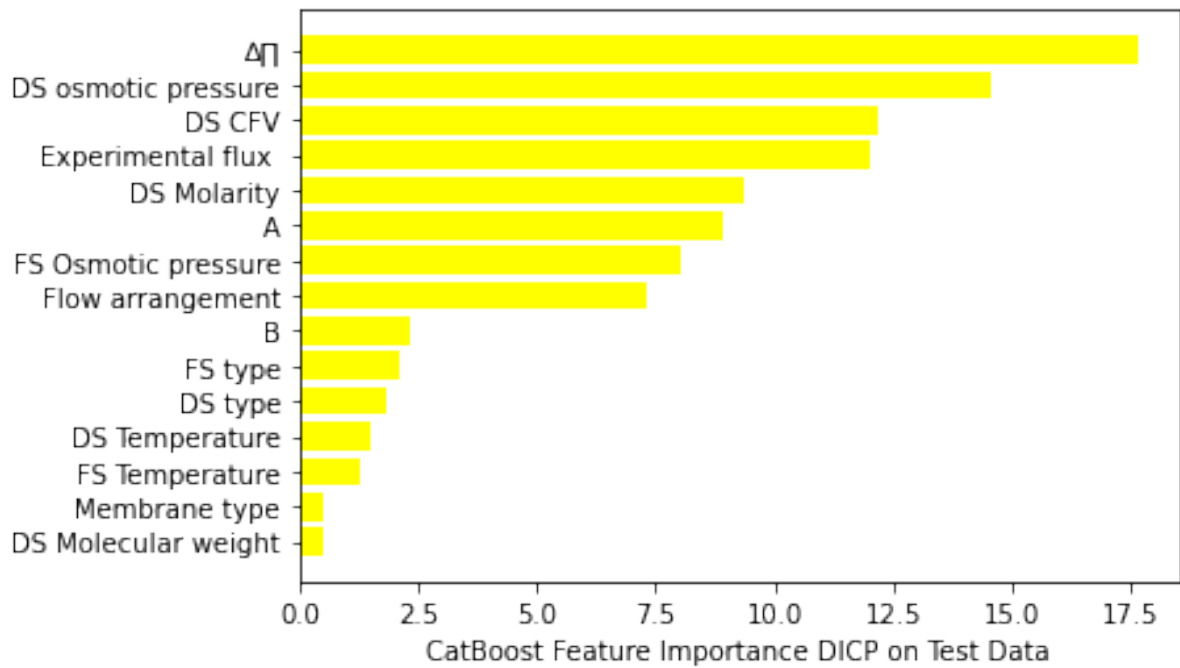
690

691 **Figure A.3:** Heat map of the input parameters to the FO process for DICP model.



692

693 **Figure A.4:** Heat map of the input parameters to the FO process for C1CP model.

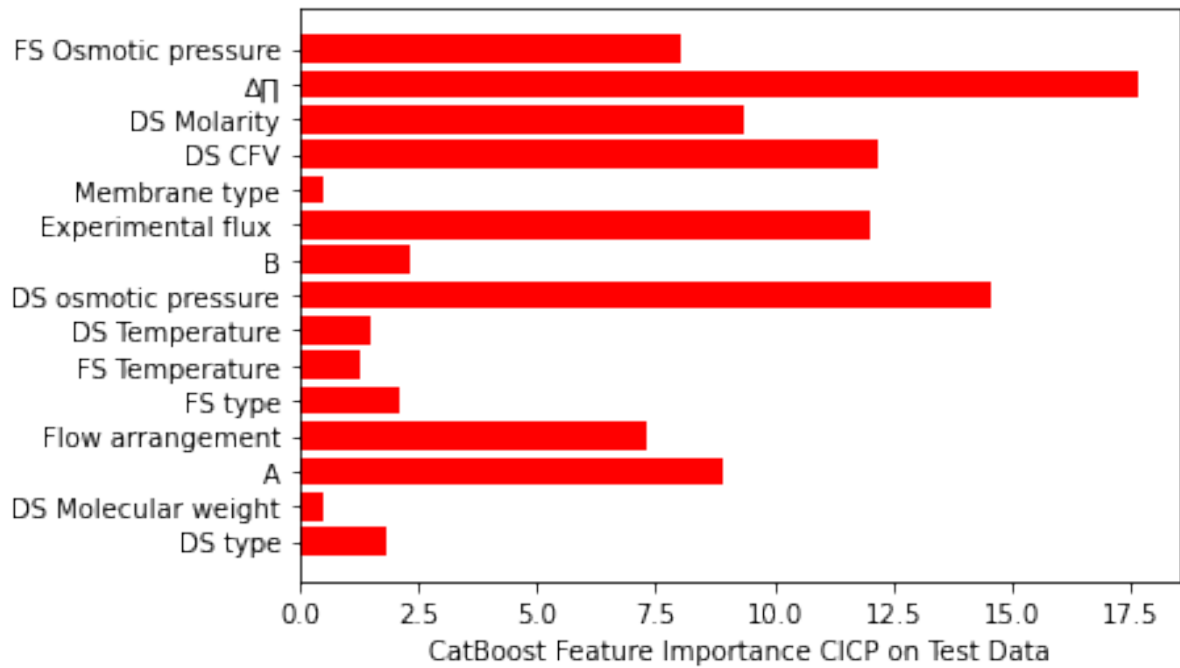


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**Figure. A.5. Catboost Feature Importance based on the test dataset for the DICP predictions, (Figure directly exported from the program)**

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**Figure. A.6. Catboost Feature importance score based on the test data for the CICP predictions (Figure directly exported from the program).**

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