MACHINE LEARNING APPLICATIONS TO PROCESS ENGINEERING

Poly-Methyl Methacrylate Reactor Unit

Ananya Bansal Lakshya Sharma

Indian Institute of Technology, Ropar May 11, 2024



CONTENTS

- **1.** Introduction to the Process
- **2.** Implementation and Model Building:
 - **i)** Feature Selection
 - **ii)** Linear Function Approximation Methods:
 - Subset Selection Method
 - Lasso Regression
 - PCA based Feature Selection
 - iii) Nonlinear Function Approximation Techniques:
 - Feed Forward Neural Network
 - LSTM model
 - iv) Just-in-time learning model/ kNN based model
- **3.** Conclusions
- **4.** Acknowledgements
- **5.** References

Introduction:

What is PMMA, what does the process encompass, and what is our aim for this model

Introduction to the process

The PMMA production process involves a free-radical polymerization of methyl methacrylate (MMA) using a Continuous Stirred-Tank Reactor (CSTR) configuration. Azo-bis-isobutyronitrile (AIBN) serves as the initiator for the polymerization reaction, while toluene functions as the solvent.

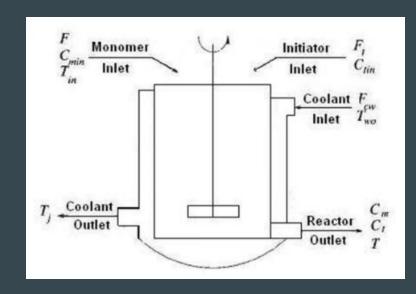


Fig 1: MMA polymerization reactor flow sheet

Process Description:

- MMA, AIBN, and toluene are continuously fed into the CSTR.
- AIBN decomposes upon heating, releasing free radicals that initiate MMA polymerization.
- MMA monomers react with the free radicals, forming PMMA chains.
- Heat generated from the exothermic reaction is controlled by a cooling jacket.
- PMMA **chains** grow in **length**, forming desired polymer product.
- Polymer solution is continuously removed from reactor for further processing.

Objectives:

- Optimizing the process of MMA to produce PMMA efficiently.
- To maintain a target monomer concentration in the reactor for consistent product quality and reaction efficiency.
- Developing predictive models for monomer concentration using multivariate regression techniques.
- To improve operation stability, efficiency, and safety and economic viability of the PMMA process.

Implementation:

Understanding and selection of features, Developing Predictive models based on various approximation methods and Testing, Presentation of Results

Implementation

The process uses 7 input variables, as can be seen from fig 1, two slides before:

- **Temperature (T):** Temperature at the reactor outlet.
- **Jacket Temperature (Tj):** Temperature of the coolant outlet.
- **Coolant water flow rate (Fcw):** Rate at which water flows through the system for temperature regulation.
- Monomer inlet Flow rate (F): Rate at which monomer is introduced into the reactor.
- Coolant inlet temperature of water (Two): Initial temperature of the coolant entering the system.
- **Inlet temperature (Tin):** Initial temperature of the input material entering the reactor.

And target variable as

• **Monomer concentration (Cm):** The concentration or amount of the monomer at the reactor outlet.

For modelling, **Fcw** and **Two** were directly dropped as they remain constant for the entire data set.

Parameters remaining, with index 0 to 4:

0: Initiator Concentration(CI)

1: Temperature (T)

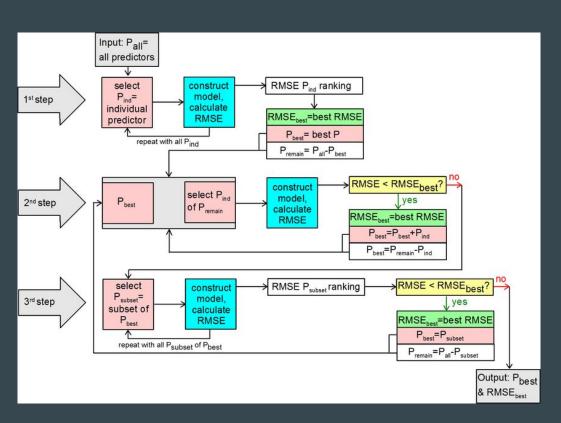
2: Jacket Temperature (Tj)

3: Monomer inlet Flowrate (F)

4: Inlet temperature of Feed (Tin)

	Initiator Concentration(CI)	Temperature (T)	Jacket Temperature (T)	Coolant water flowrate (Fcw)	Monomer inlet Flowrate (F)	Coolant inlet temp.water (Two)	Inlet temparature of Feed (Tin)	Target variable (Monomer Concentration)
count	6000.000000	6000.000000	6000.000000	6.000000e+03	6000.000000	6000.0	6000.000000	6000.000000
mean	0.025075	349.430084	331.626363	1.588000e-01	1.000060	293.2	350.000100	6.039932
std	0.000238	3.545478	3.322300	2.775789e-17	0.009995	0.0	0.079624	0.012250

Linear: Subset Selection



- → Define response variable y and matrix of predictors X .
- → Divide data into training, testing data.
- → Determine number of predictors **p** and define all possible subsets **S(p, k)**
- \rightarrow For each subset S(p, k):
 - 1. Fit linear regression model $y = X\beta S + g$
 - 2. Calculate SSE(k), R2(k), AIC(k), BIC(k) for performance evaluation.
- → Select subset S(p, k) that gives best result
- → Interpret coefficients of selected model and hence decide which features to eliminate further for optimal results

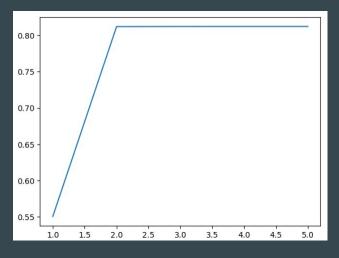
Optimal parameter subsets and their obtained R2 values:

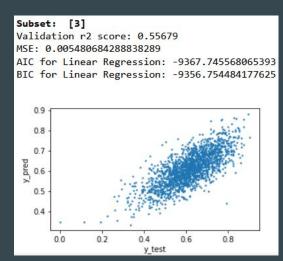
[3]: 0.55679

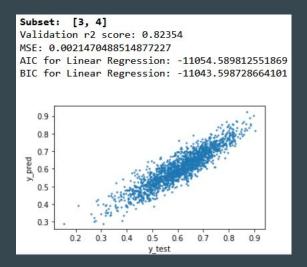
[3, 4]: 0.82354

[2, 3, 4]: 0.83216

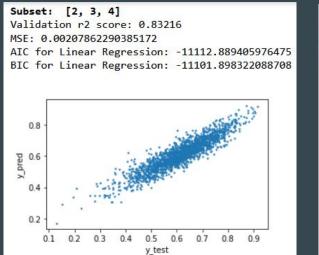
[0, 1, 3, 4]: 0.82306 [0, 1, 2, 3, 4]: 0.86836

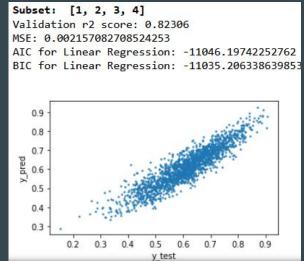


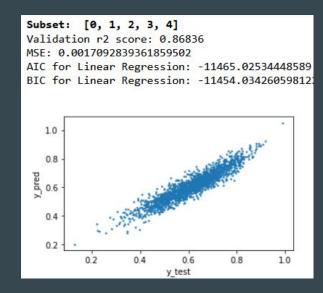




R2 vs number of features



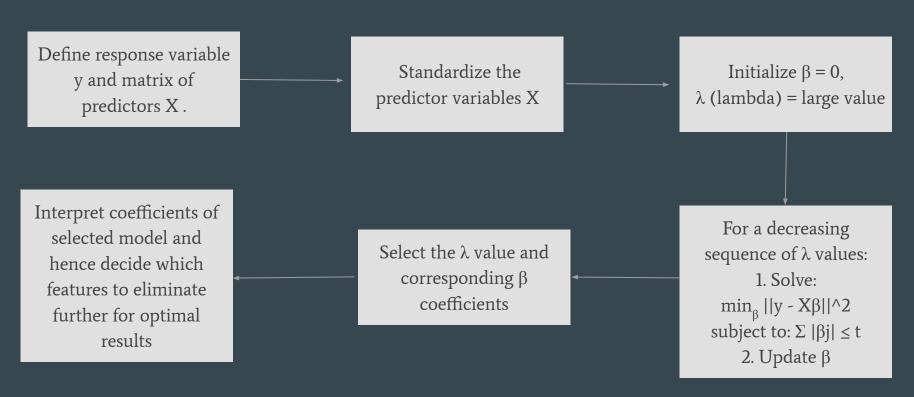




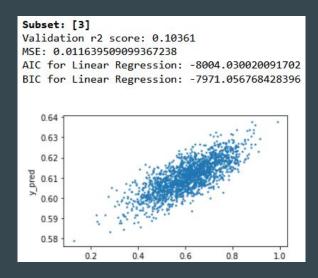
Hence, the features selected would be: [3,4] or [2,3,4] (Refer to Slide 8)

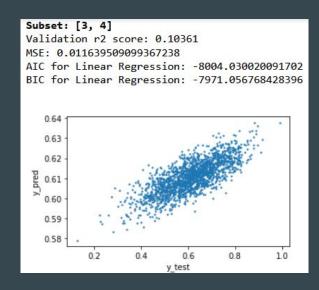
Linear: Lasso Regression

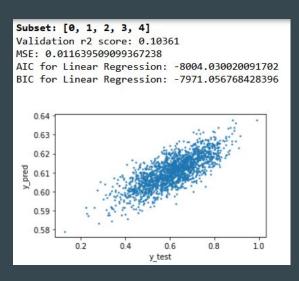
It works by introducing a bias term, ie the absolute value of the slope is added as a penalty term.



Optimal parameters and their obtained values:

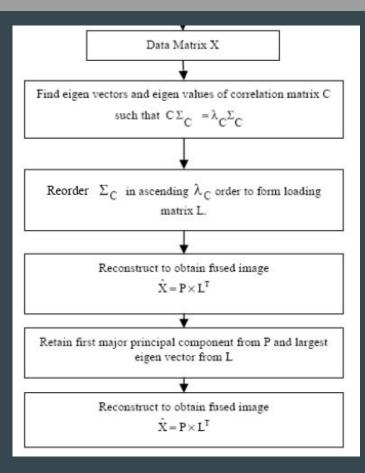






The model performs poorly for the optimal subsets if a regularisation term is introduced.

Linear: Principal Component Analysis (PCA/PCR)



Define data matrix X with n observations and p predictors

Standardize columns of

Compute covariance matrix C = X'X

Find eigenvalues λ , and eigenvectors V of the covariance matrix $CV = \lambda V$

Obtain regression coefficients q for the principal components

Construct principal component scores:

T = XVk

Regress response variable y on principal component scores T

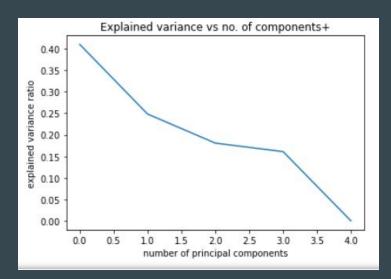
y = Tq + e

Select the number of principal components (k)to retain based on proportion of variance explained or other criteria

Sort eigenvectors V in descending order of their eigenvalues λ

Calculate coefficients for original predictors $\beta = Vk * q$ And then interpret β

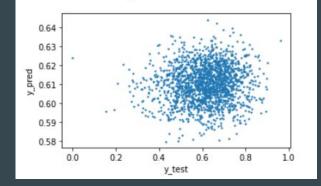
Explained variance vs Number of Components



- Elbow at 2 and then 4.
- PCR performs worse than traditional regression for n_components=2 and has comparable performance for n_components=4

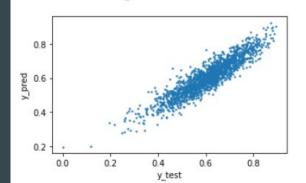
Number_of_components 2

Validation r2 score: 0.006190855951851426 Validation mse score: 0.012919676601512905 AIC for Linear Regression: -26090.022870701676 BIC for Linear Regression: -26076.623841205255



Number_of_components 4

Validation r2 score: 0.8393353611752034 Validation mse score: 0.0020590881557835455 AIC for Linear Regression: -37108.95222199654 BIC for Linear Regression: -37095.55319250012



Nonlinear: Feed Forward (Shallow Neural Network)

Define input vector x and target function f(x)

Calculate the error between y_i and f(x_i) using a loss function (e.g., mean squared error)

Update the weights and biases of the network using an optimization algorithm (e.g., gradient descent) to minimize the loss function Choose a feed-forward neural network architecture with a single hidden layer

Compute the output y_i of the network for each input x_i

Repeat the process until convergence or maximum iterations

Initialize weights and biases of the network

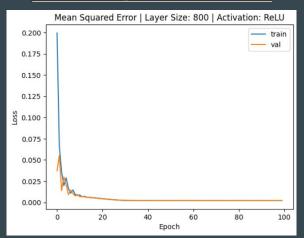
Generate a set of training data points $\{(x_i, f(x_i))\}$

The trained network approximates the target function f(x)

Keeping the number of nodes in hidden layer fixed at 800, we see

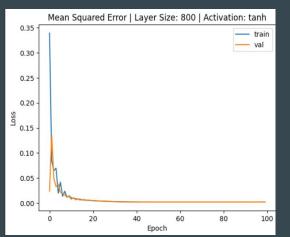
Activation Function: ReLU
Validation r2 score: 0.13207
Total no. of parameters: 644801
MSE: 0.0021650740391680784
AIC for Linear Regression: 1278558.4587042117

AIC for Linear Regression: 1278558.458704211 BIC for Linear Regression: 4822089.399662724



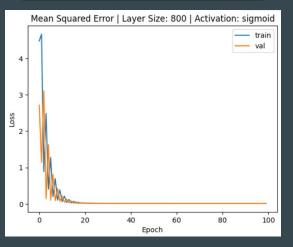
Activation Function: tanh
Validation r2 score: 0.14272
Total no. of parameters: 644801
MSE: 0.0021544549523039754

AIC for Linear Regression: 1278549.6084823934 BIC for Linear Regression: 4822080.549440905



Activation Function: sigmoid
Validation r2 score: 0.9989
Total no. of parameters: 644801
MSE: 0.012021620251940632

AIC for Linear Regression: 1281644.1125872754 BIC for Linear Regression: 4825175.053545787



TAKEAWAY: While performance of ReLU and tanh is comparable, tanh shows a better learning curve than Sigmoid.

Nonlinear: Feed Forward (Shallow NN)

| Effect of Layer Size

For tanh activation function,

Layer size: 2

Activation Function: tanh Validation r2 score: 0.83423

Total no. of parameters: 17 MSE: 0.02339032123002743

AIC for Linear Regression: -6725.779336698184 BIC for Linear Regression: -6632.355123652152

Activation Function: tanh

Validation r2 score: 0.10607 Total no. of parameters: 1841 MSF: 0.002233633071402542

Layer size: 40

Layer size: 800

AIC for Linear Regression: -7305.426510529513 BIC for Linear Regression: 2811.866208161404

Activation Function: tanh

Validation r2 score: 0.14272 Total no. of parameters: 644801

MSE: 0.0021544549523039754

AIC for Linear Regression: 1278549.6084823934 BIC for Linear Regression: 4822080.549440905

TAKEAWAYS

Performance Metric: MSE improves with layer size

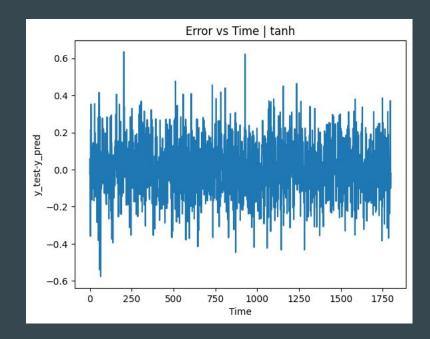
BIC being more conservative penalises layer size 40 to be worse than layer size 2 which is underfitting, While AIC considers it better.

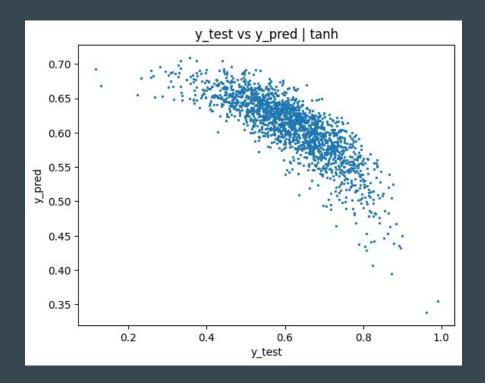
The number of trainable parameters increase exponentially with layer size

Activation Function: tanh

Validation r2 score: 0.83423 Total no. of parameters: 17 MSE: 0.02339032123002743

AIC for Linear Regression: -6725.779336698184 BIC for Linear Regression: -6632.355123652152

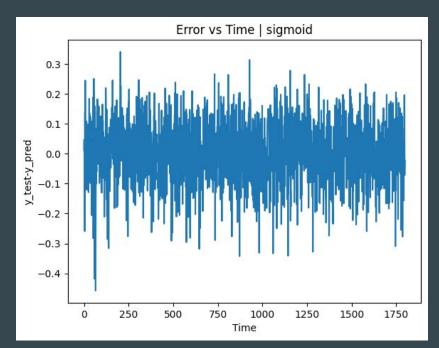


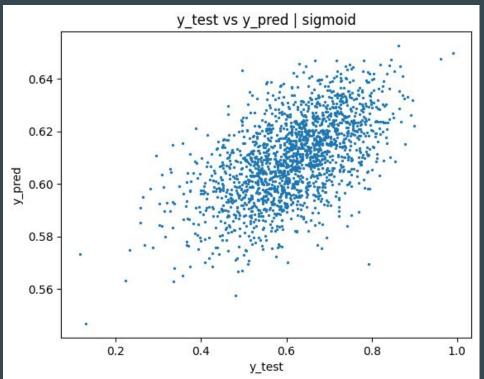


Activation Function: sigmoid Validation r2 score: 0.98098

Total no. of parameters: 17 MSE: 0.011004929270775952

AIC for Linear Regression: -8082.941583858423 BIC for Linear Regression: -7989.51737081239

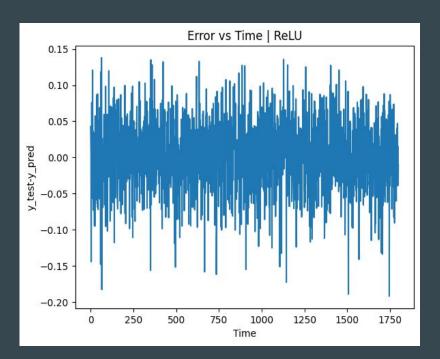


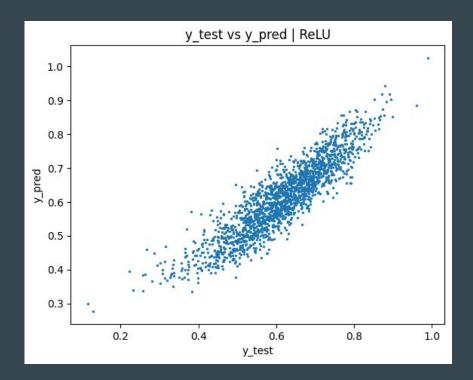


Layer Size: 40

Activation Function: ReLU Validation r2 score: 0.09309 Total no. of parameters: 1841 MSE: 0.0023713833696311732

AIC for Linear Regression: -7197.707228939935 BIC for Linear Regression: 2919.5854897509817



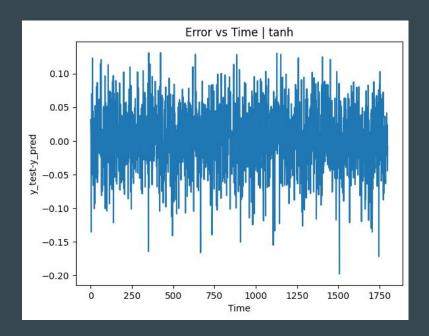


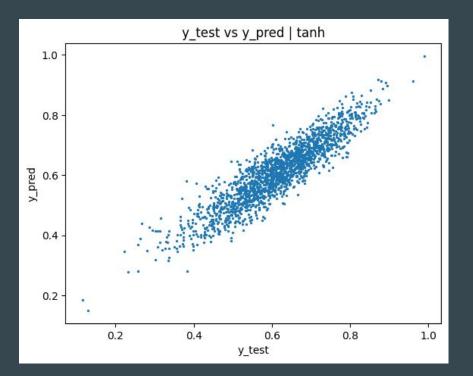
Activation Function: tanh

Validation r2 score: 0.10607 Total no. of parameters: 1841

MSE: 0.002233633071402542

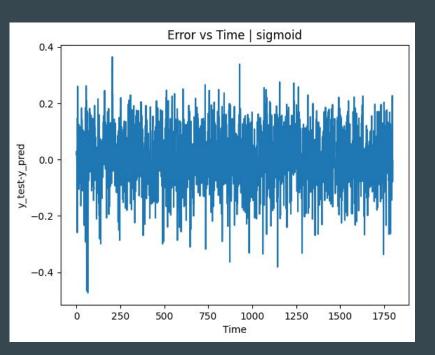
AIC for Linear Regression: -7305.426510529513 BIC for Linear Regression: 2811.866208161404

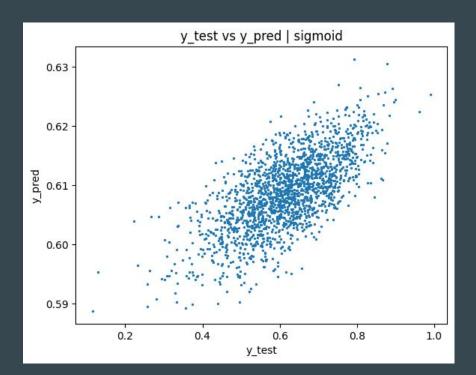




Activation Function: sigmoid Validation r2 score: 0.99685 Total no. of parameters: 1841 MSE: 0.011853492949070597

AIC for Linear Regression: -4301.238844161103 BIC for Linear Regression: 5816.053874529814





Nonlinear: Feed Forward (Deep Neural Network)

| Effect of Number of Layers

This is the same as the Shallow counterparts, other than the fact that DNN has multiple hidden layers instead of just 1.

In order to isolate effect of number of layers we take number of nodes to be same in each hidden layer. Layer Size:

15

Number of Layers = 30

Number of

Layers = 150

Activation Function: tanh Validation r2 score: 0.15807 Total no. of parameters: 6796 MSF: 0.002211925662781832

AIC for Linear Regression: 2586.994755996653 BIC for Linear Regression: 39934.69780663406

Activation Function: tanh Validation r2 score: 1.0

Total no. of parameters: 35596 MSE: 0.012794974889229245

AIC for Linear Regression: 63346.33501054779

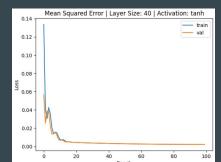
BIC for Linear Regression: 258965.64604505178

Activation Function: tanh Validation r2 score: 1.0

Number of Total no. of parameters: 83596 Layers = 350MSF: 0.13438612739476824

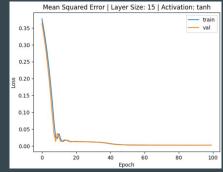
> AIC for Linear Regression: 163579.331464983 BIC for Linear Regression: 622984.6558059313

0.10 0.08



1 Hidden Layer, Parameters: 1841

- Performance deteriorates for the same amount of training if bulkier models are used for simple problems such as this.
- The learning curves for models of similar size can be very different based on number of layers.



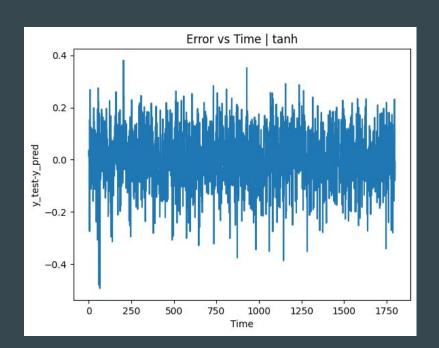
30 Hidden Layers, Parameters: 6796

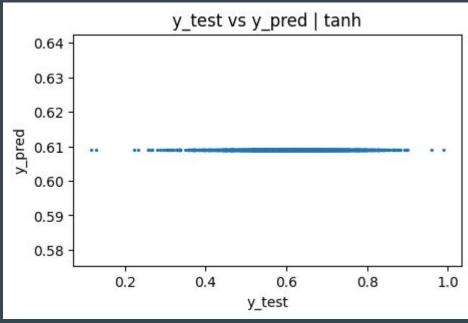
Activation Function: tanh Validation r2 score: 1.0

Total no. of parameters: 35596

MSE: 0.012794974889229245

AIC for Linear Regression: 63346.33501054779 BIC for Linear Regression: 258965.64604505178



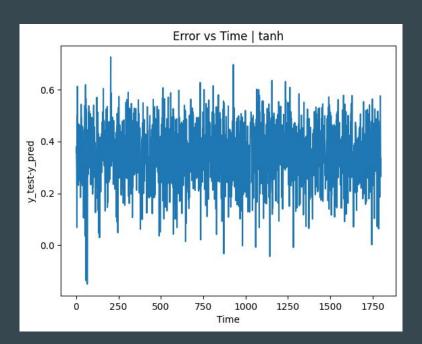


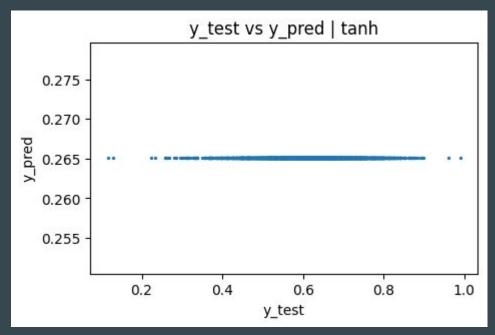
Activation Function: tanh Validation r2 score: 1.0

Total no. of parameters: 83596

MSE: 0.13438612739476824

AIC for Linear Regression: 163579.331464983 BIC for Linear Regression: 622984.6558059313





Nonlinear: LSTM model

Define the input sequence $X = (x_1, x_2, ..., x_T)$ and initialize LSTM parameters (weights and biases)

Update cell state: $c_t = f_t * c_{t-1} + i_t *$ \tilde{c}_t

Compute output gate: $o_t = \sigma(W_o[h_t-1, x_t] + b_o)$ Initialize the hidden state h_0 and cell state c_0

Compute candidate cell state: $\tilde{c}_t = \tanh(W_c[h_t-1, x_t] + b_c)$

Update hidden state: $h_t = o_t * tanh(c_t)$ For t = 1 to T:

Compute forget gate: $f_t = \sigma(W_f[h_{t-1}, x_t] + b_f)$

Next t

Obtain final hidden state h_T and/or perform further computations (e.g., feed to output layer)

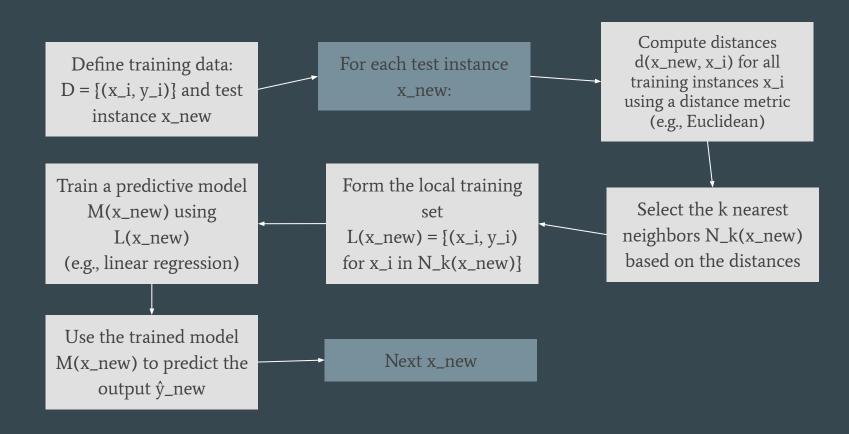
Nonlinear: LSTM model

Sequence Length: 18	Activation Function: tanh Validation r2 score: 0.98827 Total no. of parameters: 71851 MSE: 0.00014362540127040087 AIC for Linear Regression: 127775.05634973764 BIC for Linear Regression: 522635.2405597653
Sequence Length: 32	Activation Function: tanh Validation r2 score: 0.99273 Total no. of parameters: 71851 MSE: 0.00014879556106441684 AIC for Linear Regression: 127838.71291838013 BIC for Linear Regression: 522698.89712840784
Sequence Length: 56	Activation Function: tanh Validation r2 score: 0.99379 Total no. of parameters: 71851 MSE: 0.00018629797214371374 AIC for Linear Regression: 128243.30630243689

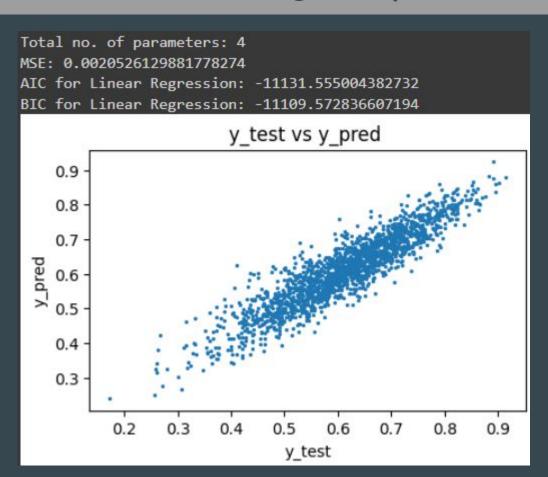
BIC for Linear Regression: 523103.49051246454

- The LSTM model shows significant improvement over DNN.
- The Sequence length was decided based on Auto-correlation of Target variable, which showed peaks on 17, 32 and 56.
- The accuracy improvement comes at significant computational cost.

Just-in-time learning based predictive model: using kNN



Just-in-time learning based predictive model: using kNN & Regression



 No significant improvement from normal regression since the data did not have multiple modes

Just-in-time learning based predictive model: using kNN & ANN

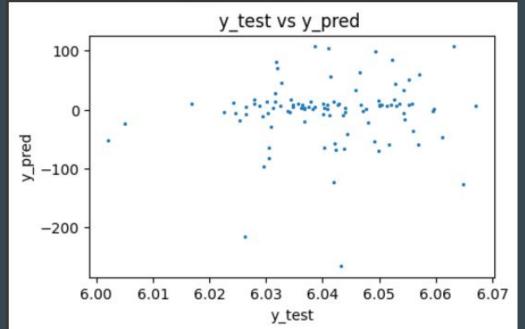
Activation Function: tanh

Validation r2 score: -21007205.14218

Total no. of parameters: 316

MSE: 3060.6601742965117

AIC for Linear Regression: 1434.6385914911355 BIC for Linear Regression: 2257.8723702633724



We did not get a satisfactory output from KNN based JITLM.

However it can be due to the following reasons:

- Coding errors
- Unoptimised KNN identification is slow.
- Tested on only 100 data points due to computation constraints.
- Not enough iterations in training

Conclusion: What we inferred from this Project

Conclusion

- We trained and developed multiple models to optimize the monomer concentration for the given PMMA data set.
- We learnt the practical implementation of all the ML concepts taught in class.
- Various regression techniques showed us that on average, the models give more consistent and better results for linear regression methods, specifically subset selection.
- The model performed poorly if the penalty term was introduced (Lasso Regression).
- PCR performs worse than traditional regression for 2 components but has a slightly better performance for 4 or 5 components.
- LSTM provides the best model in terms of MSE but has high computational cost, so in terms of scalability, linear regression still provides the best model.

Acknowledgement

We would like to thank our professor, Jayaram Valluru sir for guiding us every step of the way and introducing us to the basics of Machine Learning. Its implementation in such a conclusive way would not have been possible without his invaluable guidance and support.

References

- 1) <u>Comparison of unconstrained nonlinear state estimation techniques on a MMA polymer reactor? Arjun V Shenoy, Vinay Prasad, Sirish L Shah</u> (PMMA_Arjun)
- 2) <u>Chen, T., Morris, J., and Martin, E. (2005). Particle filters for state and parameter estimation in batch processes.</u>
- 3) Combining machine learning and process engineering physics towards enhanced accuracy and explainability of data-driven models: Bikmukhametov, Jäschke