## **Report**

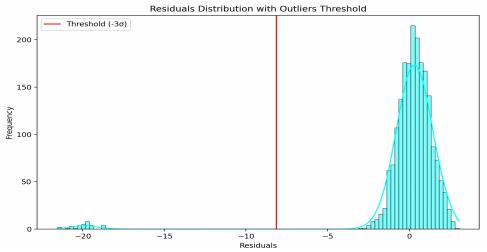
## Homework – 1

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## Task 1 - Linear Regression

- 1) Packages Used:
  - a) Pandas
  - b) Sklearn
- 2) Steps followed to find mislabeled data.
  - a) Loaded the linear\_regression CSV file.
  - b) Got the description of the dataset using 'describe'.
  - c) Checked for outliers.
  - d) Created two data frames X and y, one is for training and other one for testing.
  - e) Used Linear Regression model to fit the training dataset.
  - f) Using the trained model, predicted the target value for the entire dataset.
  - g) Found the **residual** between y and y\_predict
  - h) Checked for minimum residual value and the mean residual value
  - i) The mean was around 3.44 whereas the minimum was -21 which meant there are mislabeled samples

j) After checking the graph and I created a threshold for filtering the data based on standard deviation of the residual values

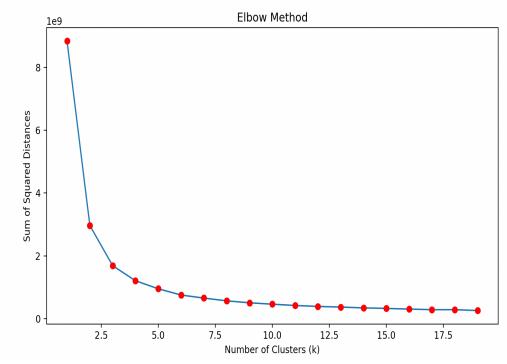


- k) I found 32 mislabeled samples after filtering the dataset
- l) Labelled them as 1 and stored it in new column Outliers

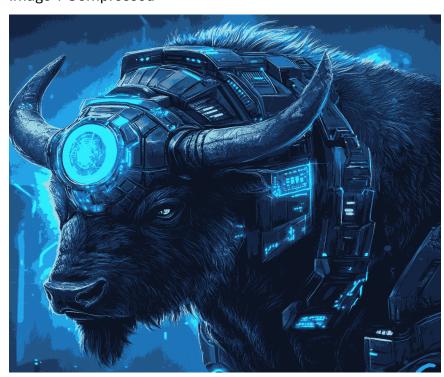
## Task 2 – Image Editing using k-means and kNN

- 1) Packages Used
  - a) PIL
  - b) Numpy
  - c) Sklearn
  - d) Matplotlib
  - e) Tqdm
- 2) Steps followed to create the compressed image 1 and 2
  - a) Used Image function from package PIL to open image
  - b) Converted the image to numpy array
  - c) Used k values ranging from 2 to 20

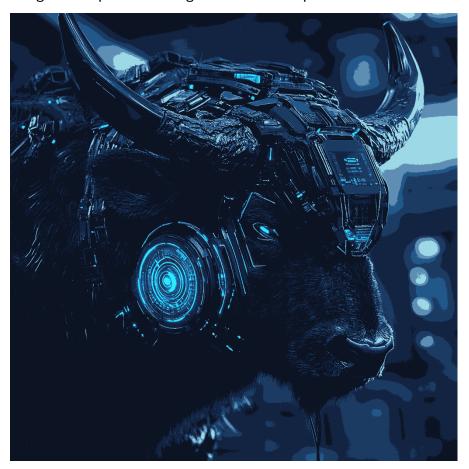
d) Using Elbow method, I found out that  $\mathbf{k} = \mathbf{9}$  had the minimum sum of squared distance and beyond that point the difference was minor.



- e) Used K\_Means to fit the image1
- f) Using kNN, k-value and the centroid from K\_Means, I fitted the model
- g) Compressed the image2 and here is the output
- h) Image 1 Compressed



i) Image 2 compressed using the same color palate



- j) If we had used a higher k value (for instance  $k \ge 60$ ), we could have got a vibrant image
- k) But using elbow method we determine that increasing k yields will have diminishing returns in terms of SSE reduction, hence adding more k (cluster value) does not significantly improve the quality of the clusters thus helping to avoid overfitting and unnecessary complexity.