# COMP0078 Assignment 2

Student Numbers: 21168615 & 19004608

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# 1 PART I

# 1.1 Kernel Perceptron

#### 1.1.1 Experimental Results

1. To generalise the kernel perceptron to *K* classes:

```
Algorithm 1 An training algorithm for multi-class kernel perceptron
```

```
for k \in \{1, 2, ..., K\} do w_k^{1,1} \leftarrow 0 end for for i \in \{1, 2, ..., M\} do: \Rightarrow Number of Epochs for j \in \{2, ..., N\} do \Rightarrow Number of training points (skipping the first point) for k \in \{1, 2, ..., K\} do \Rightarrow Number of classes \hat{y}_k^{i,j} \leftarrow \sum_{i'=1}^{i-1} \sum_{j'=1}^{j-1} (w_k^{i',j'} \cdot k(\mathbf{x}_{j'}, \mathbf{x}_{j})) \Rightarrow Prediction for class k with kernel k(\cdot, \cdot) if sign(\hat{y}_k^{i,j}) \neq sign(y_k^{i,j}) then w_k^{i,j} \leftarrow y_k^{i,j} else w_k^{i,j} \leftarrow 0 end if end for end for end for
```

The number of epochs was determined by training on the mini training data set until the performance of the mini testing data set no longer changed. This was considered a reasonable number of epochs to use to train the actual data set.

For prediction, the k with the maximum value of  $\hat{y}_k$  (the argmax of  $\hat{y}$ ) is considered the label prediction by the model.

### Basic Results:

	Number of Training Epochs	Train Error	Test Error
degree=1.0e+00	5	8.59%±1.10%	10.63%±1.21%
degree=2.0e+00	2	2.88%±0.58%	5.69%±0.62%
degree=3.0e+00	1	2.78%±0.72%	5.29%±0.80%
degree=4.0e+00	3	0.32%±0.14%	3.27%±0.44%
degree=5.0e+00	1	1.86%±0.38%	4.70%±0.59%
degree=6.0e+00	2	0.32%±0.07%	3.24%±0.36%
degree=7.0e+00	2	0.27%±0.08%	3.32%±0.45%

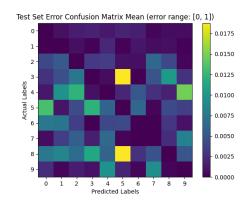
Figure 1: 20 Runs with a Polynomial Kernel

### 2. Cross Validation Results:

	Optimal degree	Number of Training Epochs	Train Error	Test Error
Run				
1	4.0e+00	3	0.31%	3.28%
2	4.0e+00	3	0.38%	3.28%
3	6.0e+00	2	0.03%	3.28%
4	7.0e+00	2	0.09%	2.80%
5	4.0e+00	3	0.13%	2.96%
6	4.0e+00	3	0.38%	2.96%
7	6.0e+00	2	0.11%	2.58%
8	4.0e+00	3	0.30%	3.28%
9	4.0e+00	3	0.31%	3.01%
10	4.0e+00	3	0.20%	3.60%
11	4.0e+00	3	0.26%	3.87%
12	6.0e+00	2	0.12%	2.90%
13	6.0e+00	2	0.16%	3.17%
14	6.0e+00	2	0.19%	2.53%
15	4.0e+00	3	0.22%	2.47%
16	4.0e+00	3	0.26%	4.03%
17	6.0e+00	2	0.07%	2.69%
18	4.0e+00	3	0.35%	3.82%
19	4.0e+00	3	0.38%	2.26%
20	7.0e+00	2	0.05%	3.82%
Across Runs	4.9e+00±1.1e+00	2.60±0.49	0.21%±0.11%	3.13%±0.50%

Figure 2: 5-fold Cross-Validation for 20 Runs with a Polynomial Kernel

#### 3. Confusion Matrix:



Test Set Error Confusion Matrix St. Dev. (error range: [0, 1])

0 - 0.014
- 0.012
- 0.010
- 0.008
- 0.006
- 7 - 0.004
- 0.002
- 0.000
- 0.000
- 0.0002
- 0.0002
- 0.0000
- 0.0000
- 0.0000

Figure 3: Polynomial Kernel Mean

Figure 4: Polynomial Kernel St. Dev.

		Predicted Lab	eis								
		0	1	2	3	4	5	6	7	8	9
Actual Labels	0	0.0%±0.0%	0.08%±0.18%	0.14%±0.19%	0.17%±0.25%	0.13%±0.16%	0.19%±0.27%	0.16%±0.16%	0.04%±0.11%	0.06%±0.16%	0.06%±0.13%
	1	0.0%±0.0%	0.0%±0.0%	0.08%±0.16%	0.02%±0.08%	0.21%±0.28%	0.02%±0.09%	0.12%±0.18%	0.06%±0.14%	0.08%±0.16%	0.02%±0.08%
	2	0.4%±0.34%	0.52%±0.54%	0.0%±0.0%	0.31%±0.44%	0.34%±0.42%	0.08%±0.2%	0.08%±0.18%	0.63%±0.58%	0.41%±0.51%	0.03%±0.12%
	3	0.27%±0.4%	0.46%±0.62%	0.74%±0.75%	0.0%±0.0%	0.06%±0.18%	1.87%±1.46%	0.0%±0.0%	0.48%±0.66%	1.02%±0.55%	0.28%±0.31%
	4	0.12%±0.32%	0.96%±0.79%	1.22%±1.18%	0.09%±0.21%	0.0%±0.0%	0.2%±0.33%	0.51%±0.62%	0.26%±0.34%	0.17%±0.32%	1.48%±1.15%
	5	1.33%±1.13%	0.11%±0.25%	0.43%±0.62%	1.21%±0.99%	0.6%±0.58%	0.0%±0.0%	0.62%±0.61%	0.03%±0.15%	0.42%±0.46%	0.35%±0.7%
	6	0.74%±1.04%	0.73%±0.64%	0.35%±0.43%	0.0%±0.0%	0.41%±0.41%	0.41%±0.39%	0.0%±0.0%	0.0%±0.0%	0.29%±0.4%	0.09%±0.22%
	7	0.06%±0.19%	0.34%±0.42%	0.57%±0.57%	0.15%±0.27%	0.65%±0.71%	0.03%±0.14%	0.0%±0.0%	0.0%±0.0%	0.25%±0.36%	0.82%±0.57%
	8	0.75%±0.8%	0.86%±0.79%	0.67%±0.75%	1.12%±0.89%	0.6%±0.72%	1.88%±1.27%	0.26%±0.55%	0.47%±0.53%	0.0%±0.0%	0.5%±0.76%
	9	0.24%±0.36%	0.03%±0.14%	0.15%±0.33%	0.09%±0.22%	0.96%±0.81%	0.19%±0.44%	0.03%±0.12%	0.92%±0.95%	0.03%±0.14%	0.0%±0.0%

Figure 5: Polynomial Kernel Confusion Matrix Error Rate Table

4. The images with the most errors made by the 20 models trained in part 2 were considered the hardest to predict and visualised. It is not surprising that these are hard to predict. We can see that in all five cases, the images do not look like their labels, in fact, they mostly look like vertical or horizontal bars. Thus it would be unreasonable to expect a model to predict them correctly.

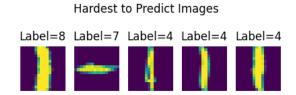


Figure 6: Images with the Most Errors from the Trained Models

5. To select a good range of values for the kernel width sigma for the Gaussian kernel, the mini training and testing sets was used to quickly gauge the test error for different hyper-parameters. The hyper-parameter values above and below the best performing hyper-parameter (with respect to test set error) was chosen as the hyper-parameter range to define *S*, the set of values for cross-validation for repeating parts 1 and 2 with the Gaussian Kernel.

	Number of Training Epochs	Train Error	Test Error
sigma=1.0e-05	6	22.44%±4.69%	22.74%±6.04%
sigma=2.1e-05	4	11.24%±2.05%	11.94%±2.08%
sigma=4.3e-05	5	8.54%±1.79%	9.24%±1.89%
sigma=8.9e-05	11	2.65%±0.96%	4.20%±1.63%
sigma=1.8e-04	5	2.78%±0.84%	4.11%±1.54%
sigma=3.8e-04	10	2.62%±1.66%	4.04%±2.14%
sigma=7.8e-04	3	2.04%±0.95%	2.83%±1.55%
sigma=1.6e-03	5	0.56%±0.34%	2.36%±1.37%
sigma=3.4e-03	3	0.28%±0.33%	2.23%±1.00%
sigma=7.0e-03	2	0.17%±0.20%	1.91%±0.99%
sigma=1.4e-02	2	0.10%±0.15%	1.72%±0.99%
sigma=3.0e-02	2	0.03%±0.06%	1.59%±1.02%
sigma=6.2e-02	1	1.44%±0.43%	3.41%±1.36%
sigma=1.3e-01	2	0.06%±0.11%	3.60%±1.63%
sigma=2.6e-01	2	0.12%±0.14%	3.73%±1.48%
sigma=5.5e-01	1	0.62%±0.31%	6.40%±1.22%
sigma=1.1e+00	1	0.09%±0.08%	36.11%±3.68%
sigma=2.3e+00	1	0.15%±0.03%	57.32%±3.65%
sigma=4.8e+00	1	0.15%±0.03%	60.54%±3.61%
sigma=1.0e+01	1	0.15%±0.03%	60.70%±3.52%

Figure 7: Performance for mini data set (to narrow down a hyper-parameter range)

### Basic Results:

	Number of Training Epochs	Train Error	Test Error
sigma=1.4e-02	2	0.31%±0.10%	3.37%±0.37%
sigma=2.2e-02	3	0.04%±0.02%	2.96%±0.31%
sigma=3.0e-02	2	0.18%±0.06%	3.28%±0.27%
sigma=3.8e-02	1	1.38%±0.17%	4.68%±0.45%
sigma=4.6e-02	3	0.05%±0.04%	3.75%±0.35%
sigma=5.4e-02	3	0.05%±0.03%	4.06%±0.44%
sigma=6.2e-02	1	1.77%±0.15%	5.61%±0.45%

Figure 8: 20 Runs with a Gaussian Kernel

#### Cross Validation Results:

	Optimal sigma	Number of Training Epochs	Train Error	Test Error
Run				
1	2.2e-02	3	0.05%	2.69%
2	2.2e-02	3	0.07%	2.63%
3	2.2e-02	3	0.04%	3.66%
4	2.2e-02	3	0.01%	2.69%
5	2.2e-02	3	0.04%	2.10%
6	2.2e-02	3	0.04%	2.37%
7	2.2e-02	3	0.07%	2.85%
8	2.2e-02	3	0.07%	2.69%
9	2.2e-02	3	0.05%	2.42%
10	2.2e-02	3	0.01%	2.47%
11	2.2e-02	3	0.00%	2.31%
12	2.2e-02	3	0.04%	2.90%
13	2.2e-02	3	0.08%	2.63%
14	2.2e-02	3	0.05%	2.90%
15	2.2e-02	3	0.05%	1.88%
16	2.2e-02	3	0.01%	2.63%
17	2.2e-02	3	0.08%	1.99%
18	2.2e-02	3	0.04%	2.85%
19	2.2e-02	3	0.04%	2.42%
20	2.2e-02	3	0.13%	3.49%
Across Runs	2.2e-02±3.5e-18	3.00±0.00	0.05%±0.03%	2.63%±0.42%

Figure 9: 5-fold Cross-Validation for 20 Runs with a Gaussian Kernel

6. An alternative method to generalise the kernel perceptron to k-classes:

#### Algorithm 2 Another training algorithm for multi-class kernel perceptron

```
for k \in \{1, 2, ..., K\} do
                                                                               ▶ Initialise weights for first training example
     w_k^{1,1} \leftarrow 0
end for
for i \in \{1, 2, ..., M\} do:
                                                                                                                ▶ Number of Epochs
     for j \in \{2, ..., N\} do
                                                                                                      ▶ Number of training points
         \hat{y}_k^{i,j} \leftarrow \sum_{i'=1}^{i-1} \sum_{j'=1}^{j-1} (w_k^{i',j} \cdot k(\mathbf{x}_{j'},\mathbf{x}_j)) end for
                                                                                                                 ▶ Number of classes
                                                                                   ▶ Prediction for class k with kernel k(\cdot, \cdot)
          \hat{c} \leftarrow argmax_k \hat{y}_k^{i,j}
                                                                                                                      ▶ Predicted label
          c \leftarrow argmax_k y_k^{i,j}
                                                                                                                          > Actual label
          if \hat{c} \neq c then
                                                                                         ▶ Compare predicted and actual label
                                                                 ▶ Non-zero weights only for actual and predicted label
                     \leftarrow 0 \forall k \in \{1,2,...,K\}, k \neq c, k \neq \hat{c}
                                                                 ▶ All zero weights if actual and predicted labels match
               w_k^{i,j} \leftarrow 0 \forall k \in \{1, 2, ..., K\}
     end for
end for
```

### Basic Results:

	Number of Training Epochs	Train Error	Test Error
degree=1.0e+00	5	9.47%±2.42%	11.66%±2.36%
degree=2.0e+00	3	3.46%±1.15%	6.44%±1.29%
degree=3.0e+00	2	2.33%±0.66%	5.35%±0.69%
degree=4.0e+00	2	1.52%±0.45%	4.56%±0.71%
degree=5.0e+00	2	0.98%±0.34%	4.10%±0.39%
degree=6.0e+00	2	0.62%±0.14%	3.68%±0.44%
degree=7.0e+00	2	0.59%±0.20%	3.79%±0.49%

Figure 10: 20 Runs with a Polynomial Kernel (Alternative Training Method)

### Cross Validation Results:

	Optimal degree	Number of Training Epochs	Train Error	Test Error
Run				
1	7.0e+00	2	0.69%	3.60%
2	6.0e+00	2	0.62%	3.66%
3	6.0e+00	2	0.75%	4.19%
4	5.0e+00	2	0.99%	4.30%
5	7.0e+00	2	0.77%	3.87%
6	7.0e+00	2	0.77%	4.09%
7	5.0e+00	2	0.95%	4.03%
8	7.0e+00	2	0.55%	3.87%
9	7.0e+00	2	0.50%	3.98%
10	6.0e+00	2	0.93%	3.92%
11	6.0e+00	2	0.59%	3.71%
12	7.0e+00	2	0.62%	4.19%
13	6.0e+00	2	0.67%	4.14%
14	7.0e+00	2	0.63%	3.71%
15	6.0e+00	2	0.67%	3.60%
16	6.0e+00	2	0.95%	3.82%
17	7.0e+00	2	0.65%	3.17%
18	7.0e+00	2	0.66%	4.57%
19	5.0e+00	2	0.78%	3.92%
20	6.0e+00	2	1.26%	5.05%
Across Runs	6.3e+00±7.1e-01	2.00±0.00	0.75%±0.18%	3.97%±0.39%

Figure 11: 5-fold Cross-Validation for 20 Runs with a Polynomial Kernel (Alternative Training Method)

#### 1.1.2 Discussions

For these experiments, the number of folds, number of runs, and train/test split were parameters that were arbitrarily chosen, not chosen through cross-validation. Moreover, the kernel parameters were crossvalidated to find optimal values, however the range for which cross validation was performed over (i.e. degree 1 to 7 for the polynomial kernel) was chosen relatively arbitrarily. For Gaussian kernels, this was done in a two step process, where a very wide range for sigma was initially used, although this wide range was still chosen arbitrarily. Increasing the number of folds would provide a better estimate of the out of sample performance (i.e. in the limit where we have N folds, each training on N-1 points and testing on the single excluded data point). However, this can be quite computationally heavy and a compromise must be drawn. This a similar situation for the number of runs. By increasing the number of runs, we gain a better understanding of average performance, however this is at a computational cost of performing more runs. The train/test split used for these experiments was 80% training data and 20% testing data. This follows general rule of thumb for train/test splits. Given a finite data set size, increasing the training percentage can provide inaccurate out of sample performance with the test set. Conversly, a smaller training percentage can poorly represent the model's performance due to limited training examples. Thus a balance must be drawn between the two. The range of hyper-parameters to search for cross validation requires past experience working with kernels and knowledge about the given data set. However these ranges can be quite general (as shown for the Gaussian kernel where we search a space from 1e - 05 to 1e + 01) and we can iteratively narrow our search space.

Comparing the two methodologies for generalising the perceptron to k-class classifiers, we can see that the main implementation difference is with the training process, where Algorithm 1 can update an arbitrary number of its weights to be non-zero while Algorithm 2 will update only two weights if the prediction is incorrect. In this sense, Algorithm 1 is like training k independent binary classifiers, where updates are made for each sub-classifier independent of other the prediction of the others. Algorithm 2 introduces dependence between sub-classifiers during the training phase. This is because whether a sub-classifier is selected to be updated depends on if it is the label predicting classifier (i.e. the argmax) or the classifier for the actual label. Otherwise, it is not updated. In some sense, Algorithm 1 is able to pass "correction" signal to all of its classifiers at each step, while Algorithm 2 is constrained in its "speed" of training its classifiers to a maximum of two at a time. Comparing Figure ?? and Figure ??, we can see that Algorithm 1 performs better than Algorithm 2 and this is likely due to this ability of Algorithm 1 to perform corrections for all its classifiers even if they are not the actual label or wrongly predicted label for a training example.

Comparing Figure ?? and Figure ??, we can see that the Gaussian Kernel performs better than the Polynomial Kernel. This is expected because the Gaussian Kernel provides a more expressive (infinite) feature space to model the data compared to a Polynomial kernel.

These experiments were implemented by vectorising and simulating the training behaviour of this online algorithm for faster computation. However, the results exactly mimic the behaviour of the algorithms depicted. The vectorisation starts with initialising a zeros weight matrix  $\mathbf{w} = \mathbf{0} \in \mathbb{R}^{KxN}$ . During the  $j^{th}$  step of training, we only update the  $j^{th}$  column of weights in  $\mathbf{w}$ . Thus, during prediction for the  $j^{th}$  training example, we know that all weights for training examples  $\geq j$  will be zero and we will be simulating an online algorithm that only predicts using the < j examples. Moreover during prediction, the operation  $\sum_{i'=1}^{i-1} \sum_{j'=1}^{j-1} (w_k^{i',j'} \cdot k(\mathbf{x}_{j'},\mathbf{x}_j))$  can be factored as  $\sum_{j'=1}^{j-1} (\sum_{i'=1}^{i-1} w_k^{i',j'}) \cdot k(\mathbf{x}_{j'},\mathbf{x}_j)$ . Thus, instead of storing a weight matrix of size  $\mathbb{R}^{Kx(N\cdot M)}$ , for M epochs, we can sum the weights corresponding to the sample training example over multiple epochs and have a weight matrix of constant size  $\mathbb{R}^{KxN}$ . The kernel evaluations were also precomputed to simulate the online model and speed up computation. Precomputing the gram matrix for a kernel helps vectorise the operation. Thus during the simulated online model, the gram matrix

is simply indexed at the appropriate training example to perform the weight updates. Additionally, across different hyper-parameters for a given kernels, a "pre-gram matrix" is pre-computed and used across the different hyper-parameters of a kernel. For example with a polynomial kernel, the gram matrix of inner products is computed as a "pre-gram matrix". Then for each polynomial experiment, the elements of the matrix is simply raised to the  $d^{th}$  degree. This also significantly improves the speed of the experiments because a single "pre-gram matrix" can be shared across experiments during a hyper-parameter search. Although the perceptron training algorithm is sequential in nature (i.e. the new weight update depends on the weight updates of the previous step), each hyper-parameter and each run is independent. This invites the opportunity to run these in parallel, performing weight updates for multiple runs in parallel. However we ran into issues of memory constraints when trying to run this on our machine with the given data set, so our implementation did not take advantage of this.

### 2 PART II

#### 2.1 Semi-supervised Learning via Laplacian Interpolation

Experimental report for the laplacian interpolation approach:



Figure 12:

And for the laplacian kernel method:



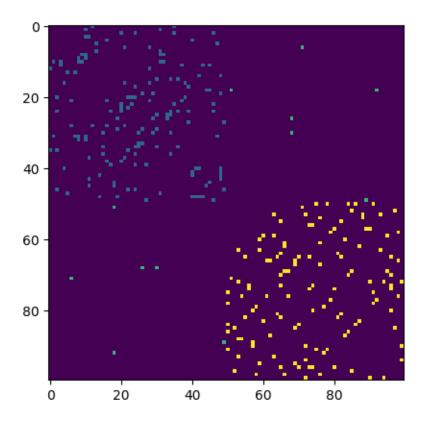
Figure 13:

Some observations to make here are:

- a) Both models exhibit high accuracy in all low-data settings.
- b) Increasing the number of labelled datapoints generally increases the accuracy in both models.
- c) Increasing the total number of datapoints generally decreases the error variance in the seminorm interpolation model, but has a limited impact on the kernel interpolation model.
- d) The laplacian kernel method outperforms the vanilla inerpolation approach considerably, in accuracy, and less so for variance.

The main reason for the success of these models is the high degree of cluster separation observed in the dataset.

For illustration, here is a diagram representing the graph adjacency matrix for 50 labels per class, with colours showing the labelling of an edge. Here blue edges are where both labels are -1, teal when both are different, and yellow when both are +1.



We see here that the vast majority of datapoints are connected only to points of the same labelling, and that the graph is highly separated into 2 clusters. Hence with minimal training information, both methods are able to predict with a high degree of accuracy. For the laplacian seminorm interpolation method, this may be observed as its solution in the objective-minimisation form essentially describes a graph cut of minimal size, respecting the true labels. We should note that in both of our models, the only pints that will be classified incorrectly with a reasonably high probability are these teal points in the diagram above.

Note that error variance decreases as a function of the number of known labels, but has a more significant impact for the vanilla interpolation method. A reasonable explanation for this is that our

We see that the kernel approach consistently outperforms the simple laplacian interpolation method. A reasonable explanation for this is that the laplacian interpolation approach utilises local information to 'diffuse' label information through the graph. This means that only datapoints close to labelled data receive direct information from the labels themselves, and accuracy is likely reduced for datapoints far from the labels. In contrast, the kernel interpolation method takes global information from all the labelled datapoints and weights them according to their proximity and connectedness to eachother to predict labels.

# 3 PART III

# 3.1 Questions

a.

b.

c.

D. We note that since for any  $X_t$ ,  $y_t = X_{1,t}$ , we have the following expression representing the linear separability of our dataset:

$$(v \cdot x_t)y_t \ge 1$$

Where  $v = (1, 0, ..., 0)^T$ . Note that ||v|| = 1.

Hence, in our online mistake bound for the perceptron, we have that  $\gamma = 1$ . Further note that  $\forall t$ ,  $x_t \in \{-1, 1\}^n \implies ||x_t||^2 = n$ . This gives us that  $R = \max_t ||x_t|| = \sqrt{n}$ . Hence our mistake bound for the perceptron algorithm is given by:

.

Using the theorem on page 60 of the online learning notes, we arrive at:

$$Prob(\mathcal{A}_{\mathcal{S}}(x') \neq y') \leq \frac{n}{m}$$

E. From our experimental observations we may expect the sample complexity of 1-NN to be lower bounded by some exponential function. We formalise this in the following proposition:

#### **Proposition:**

Following the data-generating distribution described above, we have that our sample complexity given by:

$$m(n) = \Omega(2^n)$$

#### **Proof:**

We note the following result on generalisation error.

$$L_D(1-NN) = \mathbf{E}_{(x,y)\sim\mathcal{D},S\sim\mathcal{D}^m}(\mathbf{1}_{[\mathcal{A}_S(x)\neq y]})$$

Note that since  $\mathbf{1}_{[\mathcal{A}_S(x)\neq f(x)]}$  is a bernoulli variable, its expectation is given by:  $P_{x\sim\mathcal{D},S\sim\mathcal{D}^m}(\mathcal{A}_S(x)\neq y)$ , which is our new expression for the generalisation error.

Suppose we sample a training set  $S = \{(x_1, y_1), \dots, (x_n, y_n)\}$ , x sampled uniformly from the set  $\{-1, 1\}^n$ , with associated labels defined by the rule  $y|x = x_1$ , and use this training set for inference on an arbitrary test point  $(x, x_1)$ , sampled uniformly from the same data generating distribution. We note the following observation:

#### Obs:

if  $x \in S$ , then  $\mathcal{A}_S(x)$  = y, where y is the true label for x.

To justify this, observe that our dataset represents a realiseable learning problem, and as such, if (x, y), (x', y') are datapoints sampled from our distribution,  $x = x' \implies y = y'$ . Trivially, x is the 1-nearest neighbour to itself, so the algorithm makes a correct prediction for any datapoint present in our training set.

Hence,  $\mathcal{A}_S$  makes an error on  $x \implies x \notin S$ .

Hence, 
$$\{S \subseteq \{-1, 1\}^n : \mathcal{A}_S(x) \neq y\} \subseteq \{S \subseteq \{-1, 1\}^n : x \notin S\}$$

Hence, for a given x,  $P_S(\mathcal{A}_S(x) \neq y) \leq P_S(x \notin S)$ .

Since S is a collection of points sampled iid from our data generating distribution,

$$P_S(x \notin S) = \prod_{i=1}^m P(x_i \neq x) = \prod_i (1 - P(x_i = x)) = (1 - 2^{-n})^m$$

We note that our choice of x was arbitrary, and since we sampled x uniformly, we arrive at the following generalisation error bound:

$$\mathbb{E}_{S \sim \mathcal{D}^m, x \sim \mathcal{D}} [\mathcal{L}_S(x)] \le (1 - 2^{-n})^m$$

Using the identity  $1 - x \le e^{-x}$  provided frequently in the notes, we simplify this bound to give:

$$\mathbb{E}_{S \sim \mathcal{D}^m, x \sim \mathcal{D}}[\mathcal{L}_S(x)] \leq exp(-2^{-n}m)$$

Suppose we seek m such that our generalisation error is less than some fixed  $\epsilon$ .

Hence we require  $\mathbb{E}_{S \sim \mathcal{D}^m, x \sim \mathcal{D}}[\mathcal{L}_S(x)] \leq exp(-2^{-n}m) \leq \epsilon$ 

$$\implies -m2^{-n} \le log(\epsilon)$$

Hence,  $m \ge -log(\epsilon)2^n = log(\frac{1}{\epsilon})2^n$ 

$$\implies m = \Omega(2^n)$$