- 1. Classification
 - (a) K-Nearest Neighbor
 - (b) Support Vector Machines
 - (c) Adaboost
 - (d) Iterative Dichotomiser 3
 - (e) C4.5
 - (f) Naive Bayes
 - (g) Bagging
 - (h) Random Forest
- 2. Neural Networks
 - (a) Perceptron
 - (b) Back-Propagation
 - (c) Learning Vector Quantization
 - (d) Self Organizing Map
- 3. Clustering Algorithms
 - (a) Hierarchial Agglomerative Clustering
 - (b) Hierarchial Division Clustering
- 4. Regression Algorithms
 - (a) Lasso Regression
 - (b) Logistic Regression
- 5. Deep Learning Algorithms
 - (a) Deep Q-Learning
- 6. Other Methods
 - (a) Gradient Descent
 - (b) Gaussian Process

Algorithm 1 k-Nearest Neighbor [Tay et al., 2014] link:36

Input: X: training data, Y:Class labels of X, x: unknown sample

Output: Class label of unknown sample

- 1: **function** CLASSIFY(X, Y, x)
- 2: **for** i = 1 to m **do**
- 3: Compute distance $d(X_i, x)$
- 4: end for
- 5: Compute set I containing indices for the k smallest distances $d(X_i, x)$
- 6: Return majority label $\{Y_i \text{ where } i \in I\}$
- 7: end function

```
Algorithm 2 Adaboost
                       Schapire, 2014
```

```
Input:
```

Training data $\{(x_i, y_i)_{i=1}^N \text{ where } x_i \in \mathbb{R}^k \text{ and } y_i \in \{-1, 1\}\}$ Large number of classifiers denoted by $f_m(x) \in \{-1, 1\}$ 0-1 loss function I defined as

$$I(f_m(x,y)) = \begin{cases} 0, & \text{if } f_m(x_i) = y_i \\ 1, & \text{if } f_m(x_i) \neq y_i \end{cases}$$
 (1)

Output: The final classifier 1: **for** i = 1 to N **do** for i = 1 to M do 2: Fit weak classifier m to minimize the objective function: 3: $\epsilon_m = \frac{\sum_{i=1}^N w_i^m I(f_m(x_i)) \neq y_i}{x^2 + 2x + 1}$ where $I(f_m(x_i) \neq y_i) = 1$ if $f_m(x_i) \neq y_i$ and 0 otherwise 4: 5: $\alpha_m = \ln \frac{1 - \epsilon_m}{\epsilon_m}$ 6: end for 7: $\begin{aligned} & \mathbf{for} \ \text{all} \ i \ \ \mathbf{do} \\ & w_i^{m+1} = w_i^{(m)} e^{\alpha_{mI(f_m(x_i) \neq y_i)}} \end{aligned}$ 8: 9:

11: end for

Algorithm 3 Adaboost [Hertzmann et al., 2015]

Input:

10:

Training data $\{(x_i, y_i)_{i=1}^N \text{ where } x_i \in \mathbb{R}^k \text{ and } y_i \in \{-1, 1\}\}$ Output: Weighted sum that represents the final output of the boosted classifier 1: Given Training data $\{(x_i, y_i) \text{ where } y_i \in \{-1, 1\}\}$

2: initialize D_1 = uniform distribution on training examples 3: for t = 1 to T do

4:

end for

Train weak classifier h_t on D_t choose $\alpha_t > 0$ 5:

compute new distribution D_{t+1} : 6:

7: for all i do multiply $D_t(x)$ by 8:

$$\begin{cases}
e^{-\alpha_t}, & (<1) \text{ if } y_i = h_t(x_i) \\
e^{\alpha_t}, & (>1) \text{ if } y_i \neq h_t(x_i)
\end{cases} \tag{3}$$

9: renormalize 10: end for

output final classifier $H_{final(x)} = sign(\sum \alpha_t h_t(x))$ 11:

12: end for

Algorithm 4 Adaboost4 [Hertzmann et al., 2015] Link:23,35,93,95

Input:

Training data $\{(x_i, y_i)_{i=1}^N \text{ where } x_i \in \mathbb{R}^k \text{ and } y_i \in \{-1, 1\}\}$

Output: Weighted sum that represents the final output of the boosted classifier

1: Set uniform example weights.

2: for each base learner do do

Train base learner with weighted sample. 3:

4: Test base learner on all data.

Set learner weight with weighted error. 5:

Set example weights based on ensemble predictions.

7: end for

Algorithm 5 Random forest [Bernstein, 2016] Link:42

```
Input: S: training set, F:Features and number of trees in forest B
Output: Constructed tree
 1: function RANDOMFOREST(S, F)
        H \leftarrow \emptyset
       for i \in 1, ....B do
 3:
           S^{(i)} \leftarrow A bootstrap sample from S
 4:
           h_i \leftarrow RANDOMIZEDTREELEARN(S^i, F)
 5:
           H \leftarrow H \bigcup \{h_i\}
 6:
        end for
 7:
       return H
 8:
 9: end function
10: function RANDOMIZEDTREELEARN(S, F)
       At each node:
11:
        f \leftarrow \text{a very small subset of } F
12:
       Split on best feature in f
13:
       return The learned tree
14:
15: end function
```

Algorithm 6 Iterative Dichotomiser 3 [., 2015a] Link:40

```
Input: D: Training Data, X: Set of Input Attributes
Output: A decision tree
 1: function ID3(D,X)
        Let T be a new tree
        if all instances in D have the same class c then
 3:
 4:
            Label (T) = c; Return T
        end if
 5:
        if X = \emptyset or no attribute has positive information gain then
 6:
            Label (T) = most common class in D; Return T
 7:
 8:
        end if
        X \leftarrow attribute with highest information gain
 9:
        Label(T) = X
10:
        {\bf for} \ {\rm each} \ {\rm value} \ x \ {\rm of} \ X \ \ {\bf do}
11:
12:
            D_x \leftarrow \text{ instances in } D \text{ with } X = x
            if D_x is empty then
13:
               Let T_x be a new tree
14:
               Label(T_x) = most common class in D
15:
16:
               T_x = ID3(D_x, X - \{x\})
17:
18:
        end if
        Add a branch from T to T_x labeled by x
19:
20: end for
21: return T
end function
```

Algorithm 7 Perceptron [Brownlee, 2015d] Link:65

```
Input: ProblemSize, InputPatterns, iterations_max, learn_rate
Output: Weights

1: for i = 1 to iterations_{max} do

2: Pattern_i \leftarrow SelectInputPattern(InputPatterns)

3: Activation_i \leftarrow ActivateNetwork(Pattern_i, Weights)

4: Output_i \leftarrow TransferActivation(Activation_i)

5: UpdateWeights(Pattern_i, Output_i, learn_{rate})

6: end for

7: Return Weights
```

```
Algorithm 8 Back-propagation [Brownlee, 2015a] Link:17,42
```

```
Input: ProblemSize, InputPatterns, iterations_{max}, learn_{rate}

Output: Network

1: Network \leftarrow ConstructNetworkLayers()

2: Network_{weights} \leftarrow InitializeWeights(Network, ProblemSize)

3: \mathbf{for} \ i = 1 \ to \ iterations_{max} \ \mathbf{do}

4: Pattern_i \leftarrow SelectInputPattern(InputPatterns)

5: Output_i \leftarrow ForwardPropagate(Pattern_i, Network)

6: BackwardPropagateError(Pattern_i, Output_i, Network)

7: UpdateWeights(Pattern_i, Output_i, Network, learn_{rate})

8: \mathbf{end} \ \mathbf{for}

9: Return \ Network
```

Algorithm 9 Back-propagation2 [Ng,

```
Input:
    Training Set x^{(1)}, y^{(1)}, \dots, (x^{(m)}, y^{(m)})

Output:
    Gradient of the cost function

1: \Delta_{ij}^{(l)} = 0 (for all l, i, j)

2: for i = 1 to m do

3: Set a^{(1)} = x^{(i)}

4: Perform forward propagation to compute a^{(l)} for l = 2, 3, \dots L

5: Using y^{(i)}, compute \delta^{(L)} = a^{(L)} - y^{(i)}

6: Compute \delta^{(L-1)}, \delta^{(L-2)}, \dots \delta^{(2)}

7: \Delta_{ij}^{(l)} := \Delta_{ij}^{(l)} + a_j^{(l)} \delta_i^{(l+1)}

8: end for

9: D_{ij}^{(l)} := \frac{1}{m} \Delta_{ij} + \lambda \theta_{ij}^{(l)} if j \neq 0

10: D_{ij}^{(l)} := \frac{1}{m} \Delta_{ij} if j = 0

11: \frac{\partial}{\partial \theta_{ij}^{(l)}} J(\theta) = D_{ij}^{(l)} l
```

```
Algorithm 10 Learning Vector Quantization [Brownlee, 2015c] Link: 50 and 58
```

```
Input: Problem Size, Input Patterns, iterations_{max}, Codebook Vectors_{num}, learn_{rate}
Output: CodebookVectors
 1: CodebookVectors \leftarrow InitializeCodebookVectors(CodebookVectors_{num}, ProblemSize)
 2: for i = 1 to iterations_{max} do
        Pattern_i \leftarrow SelectInputPattern(InputPatterns)
 3:
        Bmu_i \leftarrow SelectBestMatchingUnit(Pattern_i, CodebookVectors)
 4:
        for Bmu_i^{attribute} \in Bmu_i do
 5:
            if Bmu_i^{class} \equiv Pattern_i^{class} then
 6:
                 Bmu_i^{attribute} \leftarrow Bmu_i^{attribute} + learn_{rate} \times (Pattern_i^{attribute} - Bmu_i^{attribute})
 7:
 8:
                 Bmu_i^{attribute} \leftarrow Bmu_i^{attribute} - learn_{rate} \times (Pattern_i^{attribute} - Bmu_i^{attribute})
 9:
10:
        end if
11: end for
end for
Return CodebookVectors
```

```
Algorithm 11 Self Organizing Map [Brownlee, 2015b] Link:45
```

```
Input: InputPatterns, iterations_{max}, learn_{rate}, Grid_width, Grid_height
Output: CodebookVectors
 1: \ Codebook Vectors \leftarrow Initialize Codebook Vectors (Grid_{width}, Grid_{height}, Input Patterns)
 2: for i = 1 to iterations_{max} do
        Learn_{rate}^{i} \leftarrow CalculateLearningRate(i, learn_{rate}^{init})
 3:
        neighborhood_{size}^{i} \leftarrow CalculateNeighborhoodSize(i, neighborhood_{init}^{size})
 4:
 5:
        Pattern_i \leftarrow SelectInputPattern(InputPatterns)
        Bmu_i \leftarrow SelectBestMatchingUnit(Pattern_i, CodebookVectors)
 6:
        Neighborhood \leftarrow Bmu_i
 7:
        Neighborhood \leftarrow SelectNeighbors(Bmu_i, CodebookVectors, neighborhood_{size}^i)
 8:
        for Vector_i \in Neighborhood do
 9:
            for Vector_i^{attribute} \in Vector_i do
10:
                 Vector_i^{attribute} \leftarrow Vector_i^{attribute} + learn_{rate} \times (Pattern_i^{attribute} - Vector_i^{attribute})
11:
            end for
12:
        end for
13:
14: end for
15: Return CodebookVectors
```

Algorithm 12 Hierarchial Agglomerative Algorithm [Stein, 2016a] Link:24,54

```
Input:
```

```
\langle V, E, w \rangle. Weighted graph d_c. Distance measure for two clusters Output: \langle V_T, E_T \rangle. Cluster hierarchy or dendogram
```

```
1: C = \{\{v \mid v \in V\}\} \triangleright Initial Clustering \ge V_t = \{v_C \mid C \in C\}, E_T = \emptyset \Rightarrow Initial Dendogram 3: while |C| > 1 do 4: update\_distance\_matrix(C, G, d_c) \ge (C, C') = \underset{\{C_i, C_j\} \in C: C_i \neq C_j \\ \{C_i, C_j\} \in C: C_i \neq C_j\}}{\{C_i, C_j\} \in C: C_i \neq C_j\}} \Rightarrow Merging 7: V_T = V_T \cup \{v_{C,C'}\}, E_T = E_T \cup \{\{v_{C,C'}, v_C\}, \{v_{C,C'}, v_C\}\}\} \Rightarrow Dendogram 8: end while 9: Return T
```

Algorithm 13 Hierarchial Agglomerative Algorithm 2 [?] Link:28,54

Input:

 $\langle V, E, w \rangle$. Weighted graph

 d_c . Distance measure for two clusters

Output: $\langle V_T, E_T \rangle$. Cluster hierarchy or dendogram

- 1: while More than one cluster remains do
- 2: Compute the proximity graph if necessary
- 3: repeat
- 4: Merge the closest two clusters.
- 5: Update the proximity matrix to reflect the proximity between the new cluster and the original clusters.
- 6: end while

```
Algorithm 14 Hierarchial Divisive Algorithm
                                                           [Stein, 2016b]
Input:
     \langle V, E, w \rangle. Weighted graph
     d_c. Distance measure for two clusters
Output: \langle V_T, E_T \rangle. Cluster hierarchy or dendogram
 1: C = \{V\}
                                                                                                                 ▶ Initial Clustering
 2: V_t = \{v_C \mid C \in C\}, E_T = \emptyset
                                                                                                               ▶ Initial Dendogram
 3: while \exists C_x : (C_x \in C \land |C| > 1) do
         update\_distance\_matrix(C, G, d_c)
        \{C, C'\} = \underset{\{C_i, C_j\}: C_i \cup C_j = C_x \land C_i \cap C_j = \emptyset}{argmax} d_c(C_i, C_j)
C = (C \setminus \{C, C'\}) \cup \{C \cup C'\}
                                                                                                                            ▶ Merging
         V_T = V_T \cup \{v_{C,C'}\}, E_T = E_T \cup \{\{v_{C,C'}, v_C\}, \{v_{C,C'}, v_C\}\}\
                                                                                                                       ▶ Dendogram
 8: end while
 9: Return T
Algorithm 15 Hierarchial Divisive Algorithm [Stein, 2016b]
Input:
     \langle V, E, w \rangle. Weighted graph
     d_c. Distance measure for two clusters
Output: \langle V_T, E_T \rangle. Cluster hierarchy or dendogram
 1: while More than one cluster remains do
         Create a new cluster by breaking the link corresponding to the largest distance (smallest similarity).
 3: end while
Algorithm 16 C4.5 [Dai and Ji, 2014] Link :22,55
Input:
    T: Training dataset
     S: Attributes
Output: decision tree Tree
 1: function C4.5(T)
        if T is NULL then
 2:
 3:
             return failure
        end if
 4:
        if S is NULL then
 5:
 6:
             return Tree as a single node with most frequent class label in T
        end if
 7:
        if |S| = 1 then
 8:
             return Tree as a single node S
 9:
10:
         end if
11:
        set Tree = \{\}
         for a \in S do
12:
             set Info(a,T) = 0 and SplitInfo(a,T) = 0
13:
             compute Entropy(a)
             for v \in values(a, T) do
15:
                 set T_{a,v} as the subset of T with attribute a = v
16:
                 Info(a,T) + = \frac{|T_{a,v}|}{|T_a|} Entropy(a)
17:
                 SplitInfo(a,T) + = -\frac{|T_{a,v}|}{|T_a|} \log \frac{|T_{a,v}|}{|T_c|}
18:
19:
             \begin{aligned} Gain(a,T) &= Entropy(a) - Info(a,T) \\ Gain(a,T) &= \frac{Gain(a,T)}{SplitInfo(a,T)} \end{aligned}
20:
21:
         end for
22:
23:
        set a_{best} = argmax\{GainRatio(a, T)\}
        a_{best}into Tree
24:
        for v \in values(a_{best}, T) do call C4.5(T_{a,v})
25:
```

26:

27:

end for return Tree

28: end function

Algorithm 17 Gradient Descent [., 2015b] Link: 59

```
Input:
    f
    starting value x_1
    termination tolerances
Output: x_{maxIters}
 1: for i = 1 to maxIters do
        Compute the search direction d_t = -\delta f(x_t)
        if |d_T| < \epsilon_q then
 3:
            return "Converged to critical point", output x_t
 4:
            Find \alpha_t so that f(x_t + \alpha_t d_t) < f(x_t)
 5:
        end if
 6:
 7:
        if |\alpha_t d_T| < \epsilon_x then
            return "Converged in x", output x_t
 8:
            Find \alpha_t so that f(x_t + \alpha_t d_t) < f(x_t)
 9:
        end if
10:
        Let x_{t+1} = x_t + \alpha_t d_t
11:
12: end for
13: Return "Max number of iterations reached", output x_{maxIters}
```

Algorithm 18 Naive Bayes [., d] Link: 9, 56

```
Input:
    C: A  fixed set of classes
    D: Documents
Output: Category(Class) of the Documents
 1: function TrainMultinomialNB(C, D)
        V \leftarrow EXTRACTVOCABULARY(D)
       N \leftarrow COUNTDOCS(D)
 3:
 4:
       for each c \in C do
           N_c \leftarrow COUNTDOCSINCLASS(D, c)
 5:
           prior|c| \leftarrow N_c/N
 6:
           text_c \leftarrow CONCATENATETEXTOFALLDOCSINCLASS(D,C)
 7:
 8:
           for each t \in V do
              condprob|t||c| \leftarrow \frac{T_{ct}+1}{\sum_{t'}(T_{ct'+1})}
 9:
           end for
10:
       end for
11:
       return V, prior, condprob
12:
13: end function
    \mathbf{function} \ \mathbf{ApplyMultinomialNB}(C, D, prior, condprob, d)
       W \leftarrow EXTRACTTOKENSFROMDOC(V, d)
15:
       for each c \in C do
16:
17:
           score|c| \leftarrow \log \ prior|c|
           for each t \in W do
18:
               score|c| + = \log condprob|t||c|
19:
20:
           end for
       end for
21:
       return arg\ max_{c \in C} score|c|
23: end function
```

Algorithm 19 Lasso Regression

```
Input:
     ipy: Inner product vector, ipy_i = \langle y, X_{\cdot i} \rangle
     ipx: Inner product matrix, ipx_{ij} = \langle X_{\cdot i}, X_{\cdot j} \rangle
     \lambda: Penalty parameter
     N: Number of samples
Output: beta: Regression parameter vector
 1: function FastLasso(ipy, ipx, \lambda, N)
         stop_-thr
                                                                                                   ▶ Threshold for stopping iteration
         p \leftarrow length(ipy)
 3:
 4:
         beta \leftarrow 0 with length p
 5:
         gc \leftarrow 0 with length p
         while difBeta_{max} \geq \text{stop\_thr do}
 6:
              difBeta_{max} \leftarrow 0
 7:
 8:
              for j = 1 \leftarrow p do
                  z \leftarrow (ipy|j| - gc|j|)/N + beta|j|
 9:
                  \mathtt{beta\_tmp} \leftarrow \max(0, z - \lambda) - \max(0, -z - \lambda)
10:
                  difBeta \leftarrow \mathtt{beta\_tmp} - beta|j|
11:
                  difabs \leftarrow abs(difBeta)
12:
                  if difabs > 0 then
13:
                       beta|j| \leftarrow \mathtt{beta\_tmp}
14:
                       gc \leftarrow gc + ipx|j| \times difBeta
15:
                       difBeta_{max} = max(difBeta_{max}, difabs)
16:
                  end if
17:
              end for
18:
         end while
19:
20: end function
```

Algorithm 20 Bagging [., a] Link:57 Input:

```
B: the number of bags or base hypotheses
    L: Base Learning Algorithm
Output: New Training Sets
 1: function BAGGING(examples, B, L)
       for i = 1 to B do
 2:
 3:
           examples_i \leftarrow a bootstrap sample of examples
       end for
 4:
       Compute set I containing indices for the k smallest distances d(X_i, x)
 5:
       h_i \leftarrow \text{apply}Ltoexamples_i
 6:
 7:
       Return h_1, h_2, ...h_B
 8: end function
```

```
Algorithm 21 Deep Q-Learning with Experience Replay [Mnih et al., 2013]
Input:
     D: data set
     Q: Action-Value Function
Output: New Training Sets
 1: for i = 1 to M do
          Initialise sequence s_1 = \{x_1\} and preprocessed sequenced \phi = \phi(s_1)
 2:
 3:
          for i = 1 to T do
               With probability \epsilon select a random action a_t otherwise select a_t = max_aQ * (\phi(s_t).a : \theta)
 4:
               Execute action a_t in emulator and observe reward r and image x_{t+1}
 5:
               Set s_{t+1} = s_t, a_t, x_{t+1} and preprocess \phi_{t+1} = \phi(s_{t+1})
 6:
 7:
               Store transition (\phi_t, a_t, r_t, \phi_{t+1}) in D
               Set y_i =
 8:
                                             \begin{cases} r_j, & \text{for terminal } \phi_{j+1} \\ r_j + \gamma \max_{a'} Q(\phi_{j+1,a';\theta}), & \text{for terminal } \phi_{j+1} \end{cases} 
                                                                                                                                                      (5)
                                                                                                                                                      (6)
               Perform a gradient descent step on (y_j - Q(\phi_j, a_j; \theta))^2 according to the following equation
 9:
10:
                                  \Delta_{\theta}L_{i}(\theta_{i}) = \mathbb{E}_{s,a \sim \rho(.);s' \sim \epsilon[(r + \gamma max_{a'}Q(s',a';\theta_{i-1}) - Q(s,a;\theta_{i}))\Delta_{\theta_{i}}Q(s,a;\theta_{i})]}
          end for
11:
```

Algorithm 22 PageRank [?] Link:24,64,98,99

12: end for

```
Input:
     G: inlink file
     iteration: Number of iteration
Output: PageRank
 1: function PageRank(G, iteration)
         d \leftarrow 0.85
                                                                                                                  ⊳ damping factor: 0.85
         oh \leftarrow G
                                                                                                              \triangleright get outlink hash from G
 3:
         ih \leftarrow G
 4:
                                                                                                                ⊳ get inlink hash from G
         N \leftarrow G
                                                                                                         ⊳ get number of pages from G
 5:
         for all p in the graph do
 6:
              opg[p] \leftarrow \frac{1}{N}
 7:
         end for
 8:
 9:
         while iteration > 0 do
              dp \leftarrow 0
10:
              for all p that has no out-links do
11:
                  dp \leftarrow dp + d * \frac{opg[p]}{N}
12:
              end for
13:
              for all p in the graph do
14:
                  npg[p] \leftarrow dp + \frac{[1-d]}{N} for all ip in ih[p] do
15:
16:
                       npg[p] \leftarrow dp + \frac{d*opg[ip]}{oh[ip]}
17:
                  end for
18:
19:
              end for
              opg \leftarrow npg
20:
              iteration \leftarrow iteration - 1
21:
         end while
22:
23: end function
```

Algorithm 23 K means Link:25,29,40,104

```
Input:
    E = \{e_1, e_2, .... e_k\}
    k(number of clusters)
    MaxIters(limit of iterations)
Output:
    C = \{c_1, c_2, ...c_k\}
    L = \{l(e)|e = 1, 2, ...n\} (set of cluster labels of E)
 1: for each c_i \in C do
        c_i \leftarrow e_j \in E(\text{e.g random selection})
 2:
 3: end for
 4: for each e_i \in E do
        l(e_i) \leftarrow argminDistance(e_i, c_j)j \in \{1..k\}
 6: end for
 7:\ changed \leftarrow false
 8: iter \leftarrow 0
    while changed = true and iter \le = MaxIters do
        for c_i \in C do
10:
            UpdateCluster(c_i)
11:
        end for
12:
13:
        for c_i \in E do
            minDist \leftarrow argminDistance(e_i, e_j)j \in \{1..k\}
14:
            if minDist \neq l(e_i) then
15:
                l(e_i) \leftarrow minDist
16:
17:
                changed \leftarrow true
            end if
18:
        end for
19:
        iter ++
20:
21: end while
```

```
Algorithm 24 DBSCAN link:42 [Ram et al., 2010]
Input:
    D: Data
    \epsilon:Threshold distance
    MinPts: Minimum number of points required to form a cluster
Output: Clustered Data
 1: function DBSCAN(D, \epsilon, minPts)
       C = 0
       for each point Pin datasetD do
 3:
 4:
           if P is visited then
              continue next point
 5:
           end if
 6:
           mark P as visited
 7:
           NeighborPts = regionQuery(P, \epsilon)
 8:
           if sizeof(NeighborPts) < MinPts then
 9:
              mark P as NOISE
10:
           else
11:
              C = \text{next cluster}
12:
              expandCluster(P, NeighborPts, C, \epsilon, MinPts)
13:
14:
       end if
15: end for
and function
function EXPANDCLUSTER(P, NeighborPts, C, \epsilon, MinPts)
18: add P to Cluster C
    {\bf for} \ {\bf each} \ {\bf point} \ P'inNeighborPts \ {\bf do}
20:
       if P' is not visited then
           mark P' as visited
21:
           NeighborPts' = regionQuery(P', eps)
22:
           if sizeof(NeighborPts) >= MinPts then
23:
              NeighborPts = NeighborPts joined with NeighborPts'
24:
           end if
25:
26:
       end if
       if P' is not yet member of any cluster then
27:
           add P' to cluster C
28:
       end if
29.
30: end for
end function
function REGIONQUERY(P, \epsilon)
```

```
Algorithm 25 Principle Component Analysis [?] Link:41,69,72

Input:

x_1, .....x_nd length vector k

Output: Transform matrix R

1: X \leftarrow n \times d data matrix with x_i in each row
2: x \leftarrow \frac{1}{n} \sum_{i=1}^{n} x_i
3: X \leftarrow subtract x from each row x_i i n X
4: COV \leftarrow \frac{1}{n-1} X^T \times XCompute eigenvalue e_1, ...., e_d of COV, and sort them
5: Compute matrix V which satisfy V^{-1} \times COV \times V = D, D is the diagonal matrix of eigenvalue of COV
```

33: return all points within $P's \epsilon neighborhood$

6: $R \leftarrow$ the first k column of V

end function

Algorithm 26 Logistic Regression

Input:

Training data of the form $\{(x_1, 1), (x_2, 0), ...\}$

x: unknown sample

Output: The output is a probability that the given input point belongs to a certain class

- 1: $0 \leftarrow \beta$
- 2: Compute y by setting its elements to

$$y = \begin{cases} 1, & \text{if } g_i = 1\\ 0, & \text{if } g_i = 2 \end{cases}$$
 (7)

i = 1.2...N

3: Compute p by setting its elements to

$$p(x_i, \beta) = \frac{e^{\beta^T x_i}}{1 + e^{\beta^T x_i}}$$

i = 1,2,...N

- 4: Compute the diagonal matrix W. The ith diagonal element is $p(x_i, \beta)(1 p(x_i; \beta))$
- 5: $z \leftarrow X\beta + W^{-1}(y-p)$ 6: $\beta \leftarrow (X^TWX)^{-1}X^TWz$
- 7: If the stopping criteria, stop; otherwise go back to step 3

Algorithm 27 Gaussian Process [., b]

Input:

$$X = \begin{bmatrix} x_1^T \\ \dots \\ x_n^T \end{bmatrix} \in \mathbb{R}^{n \times D}, \text{m training inputs}$$

$$y = \begin{bmatrix} y_1^T \\ \dots \\ y_n^T \end{bmatrix} \in \mathbb{R}^n$$

$$k(.,.) : \mathbb{R}^{D \times D}$$

 x_* test input

 σ^2 noise level on the observations

$$[y(x) = f(x) + \epsilon, \epsilon \sim N(0, \sigma^2)]$$

Output:

$$f_*$$
 $cov(f_*)$

 $cov(f_*)$ 1: $K \in \mathbb{R}^{n \times n}$ Gram matrix. $K_{ij} = k(x_i, x_j)$

$$k(x_*) = k_* = k(X, x_*) = \begin{bmatrix} k(x_1, x_*) \\ \dots \\ k(x_n, x_*) \end{bmatrix} \in \mathbb{R}$$

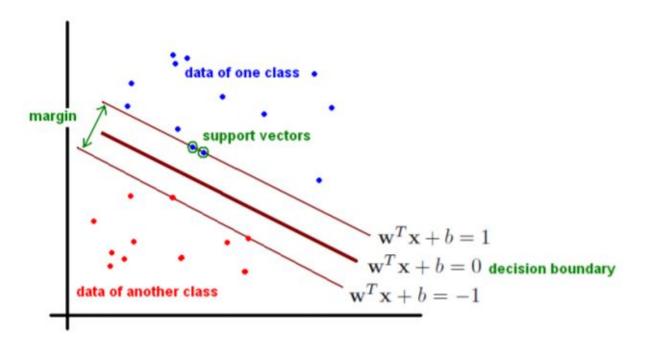
2: $\alpha = (K + \sigma_2 \mathbb{I}_n)^{-1} y$ 3: $f_* = k_*^T \alpha \in \mathbb{R}$

4: $cov(f_*) = k(x_*, x_*) - k_*^T [K + \sigma_2 \mathbb{I} n]^{-1} k_*$

Algorithm 28 Support Vector Machines [Gavrilov,

Input:

Set of N input-output pairs $\{x,y\}^{N_1}$ x: input vectors of the same dimension and y : set of output target labels $y_i = \{0,1\}$



1 Simple case: linearly-separable data, binary classification

Goal:we want to find the hyperplane (i.e. decision boundary) linearly separating our classes. Our boundary will have the equation: $\mathbf{w}^{\mathbf{T}}\mathbf{x} + \mathbf{b} = 0$

Anything above the decision boundary should have label 1 i.e., $\mathbf{w}^{\mathsf{T}}\mathbf{x}_i + b > 0$ will have corresponding $y_i = 1$

Similarly, anything below the decision boundary should have label -1 i.e. $\mathbf{w^T}\mathbf{x}_i + b < 0$ will have corresponding $y_i = -1$

The reason for this labeling scheme is that it lets us condense for the decision function to

$$f(x) = sign(\mathbf{w^T} + b)$$

since f(x) = +1 for all x above the boundary, and f(x) = -1 for all x below the boundary.

Thus, we can figure out if an instance has been classified properly by checking that $y(\mathbf{w^T} + b) \ge 1$ (which will be the case as long as either both $y, \mathbf{w^T} + b > 0$ or else $y, \mathbf{w^T} + b < 0$)

You'll notice that we will now have some space between our decision boundary and the nearest data points of either class. Thus,let's rescale the data such that anything on or above the boundary $\mathbf{w}^{\mathbf{T}}\mathbf{x} + b = 1$ is of one class(with label 1), and anything on or below the boundary $\mathbf{w}^{\mathbf{T}}\mathbf{x} + b = -1$ is of another class (with label -1)

What is the distance between these newly added boundaries?

First note that the two lines are parallel, and thus share their parameters w, b.Pick an arbitary point x_1 to lie on line $\mathbf{w^Tx} + b = -1$. Then the closest point on line $\mathbf{w^Tx} + b = 1$ is the point $\mathbf{x}_2 = \mathbf{x}_1 + \lambda \mathbf{w}$ (since the closest point will always lie on the perpendicular; recall that the vector \mathbf{w} is perpendicular to both lines). Using this formulation, $\lambda \mathbf{w}$ will be the line segment segment connecting \mathbf{x}_1 and \mathbf{x}_2 , and thus, $\lambda \|\mathbf{w}\|$, the distance between \mathbf{x}_1 and \mathbf{x}_2 is the shortest distance between the two lines/boundaries.

References

[., a] . (.a). Bagging.

[., b] . (.b). Gaussian process.

[., c] . (.c). Logistic regression.

[., d] . (.d). Naive bayes.

[., e] . (.e). Page rank.

[., 2015a] . (2015a). Decision trees.

[., 2015b] . (2015b). Gradient descent.

[Bernstein, 2016] Bernstein, M. (2016). Random forests.

[Brownlee, 2015a] Brownlee, J. (2015a). Back-propagation.

[Brownlee, 2015b] Brownlee, J. (2015b). Clever algorithms: Nature-inspired programming recipes.

[Brownlee, 2015c] Brownlee, J. (2015c). Learning vector quantization.

[Brownlee, 2015d] Brownlee, J. (2015d). Perceptron.

[Dai and Ji, 2014] Dai, W. and Ji, W. (2014). A mapreduce implementation of c4.5 decision tree algorithm.

[Gavrilov,] Gavrilov, Z. (.). Support vector machines.

[Hertzmann et al., 2015] Hertzmann, A., Fleet, D., and Brubaker, M. (2015). Adaboost.

[Mnih et al., 2013] Mnih, V., Kavukcuoglu, K., Silver, D., Graves, A., Antonoglou, I., Wierstra, D., and Riedmiller, M. A. (2013). Playing atari with deep reinforcement learning. *CoRR*, abs/1312.5602.

[Ng,] Ng, A. Coursera machine learning.

[Ram et al., 2010] Ram, A., Jalal, S., Jalal, A. S., and Kumar, M. (2010). A density based algorithm for discovering density varied clusters in large spatial databases. *International Journal of Computer Applications*, 3(6):1–4.

[Schapire, 2014] Schapire, R. (2014). Machine learning algorithms for classification.

[Stein, 2016a] Stein, B. (2016a). Unit hierarchial cluster analysis.

[Stein, 2016b] Stein, B. (2016b). Unit hierarchial cluster analysis.

[Tay et al., 2014] Tay, B., Hyun, J., and Sejong, O. (2014). A machine learning approach for specification of spinal cord injuries using fractional anisotropy values obtained from diffusion tensor images.

Algorithm 28 Support Vector Machines (continued)

```
Solving for \lambda: \mathbf{w}^{\mathbf{T}}\mathbf{x}_{2} + b = 1 where \mathbf{x}_{2} = \mathbf{x}_{1} + \lambda \mathbf{w} \mathbf{w}^{\mathbf{T}}(\mathbf{x}_{1}\lambda\mathbf{w}) + b = 1 \mathbf{w}^{\mathbf{T}}(\mathbf{x}_{1}\lambda\mathbf{w}) + b = 1 \mathbf{w}^{\mathbf{T}}\mathbf{x}_{1} + b + \lambda \mathbf{w}^{\mathbf{T}}\mathbf{w} = 1 where \mathbf{w}^{\mathbf{T}}\mathbf{x}_{1} + b = -1
   -1 + \lambda \mathbf{w}^{\mathbf{T}} \mathbf{w} = 1
\lambda \mathbf{w}^{\mathbf{T}} \mathbf{w} = 2
\lambda = \frac{2}{\mathbf{w}^{\mathbf{T}} \mathbf{w}} = \frac{2}{\|\mathbf{w}\|^2}
And, so the distance \lambda \|w\| is \frac{2}{\|\mathbf{w}\|^2} \|w\| = \frac{2}{\|w\|} = \frac{2}{\sqrt{\mathbf{w}^T \mathbf{w}}}
```

It's intuitive that we would want to maximize the distance between the two boundaries demarcating the classes(Why?We want to be as sure that we are not making classification mistakes and thus we want our data points from the two classes to lie as far away from each other as possible). This distance is called the margin, so went to obtain the maximal margin.

Thus, we want to maximize $\frac{2}{\sqrt{\mathbf{w}^T\mathbf{w}}}$, which is equivalent to minimizing $\frac{\sqrt{\mathbf{w}^T\mathbf{w}}}{2}$ which is in turn equivalent to minimizing $\frac{\mathbf{w}^T\mathbf{w}}{2}$ (since square root is a monotonic function)

```
This quadratic programming problem is expressed as:
subject to: y_i(\mathbf{w}^T\mathbf{x} + b) \ge 1(\forall \text{ data points } \mathbf{x}_i)
```

$\mathbf{2}$ Soft-margin extension

Consider the case that your data isn't linearly separable. For instance, maybe you aren't guaranteed that all your data points are correctly labelled, so you want to allow some data points of one class to appear on the other side of the boundary.

```
We can introduce slackvariables an -\epsilon_i \geq 0. Our quadratic programming problem becomes: min_{\mathbf{w},b,\epsilon} \frac{\mathbf{w^T}\mathbf{w}}{2} + C\sum_i \epsilon_i subject to : y_i(\mathbf{w^T}\mathbf{x}_i + b) \geq 1 - \epsilon
```

3 Nonlinear decision boundary

Mapping your data vectors, x_i , into a higher-dimension (even infinite) feature space may make them linearly separable in that space (whereas they may not be linearly separable in the original space). The formation of the quadratic programmatic problem is as above, but with all \mathbf{x}_i replaced with $\phi(\mathbf{x}_i)$, where ϕ provides the higher-dimensional mapping. So we have the standard SVM formulation:

```
min_{\mathbf{w},b,\epsilon} \frac{\mathbf{w}^T \mathbf{w}}{2}
subject to: y_i(\mathbf{w}^T \phi(\mathbf{x}_i) + b) \ge 1 - \epsilon and \epsilon_i \ge 0 (\forall \text{ data points } \mathbf{x}_i)
```

Reformulating as a Lagrangian 4

We can introduce Lagrange multipliers to represent the condition:

 $y_i(\mathbf{w}^T\phi(\mathbf{x}_i+b))$ must be as close to 1 as possible. This condition is captured by: $\max_{ai\geq 0}\alpha_i[1-y_i(\mathbf{w}^T\phi(\mathbf{x}_i+b))]$ b))] This ensures that when $y_i(\mathbf{w}^T\phi(\mathbf{x}_i+b)) \geq 1$, the expression above is maximal when $\alpha_i = 0$ (since [1 $y_i(\mathbf{w}^T\phi(\mathbf{x}_i+b))$ ends up being negative). Otherwise, $y_i(\mathbf{w}^T\phi(\mathbf{x}_i+b)) < 1$, so $[1-y_i(\mathbf{w}^T\phi(\mathbf{x}_i+b))]$ is a positive value, and the expression is maximal when $a_i \to \infty$. This has the effect of penalizing any misclassified data points, while assigning 0 penalty to properly classified instances.

We thus have the following formulation:

$$min_{w,b}\left[\frac{\mathbf{w}^T\mathbf{w}}{2} + \sum_{i} max_{\alpha \geq 0} \alpha_i \left[1 - \mathbf{w}^T \phi(\mathbf{x}_i + b)\right]\right]$$

 $min_{w,b}[\frac{\mathbf{w}^T\mathbf{w}}{2} + \sum_i max_{\alpha \geq 0}\alpha_i[1 - \mathbf{w}^T\phi(\mathbf{x}_i + b)]]$ To allow for slack(soft-margin), preventing the α variables from going to ∞ , we can impose constraints on the Lagrange multipliers to lie within: $0 \le \alpha_i \le C$. We can define the dual problem by interchanging the max and min as follows (i.e minimize after fixing alpha):

$$max_{alpha \geq zero}[min_{w,b}J(\mathbf{w},b;\alpha)]$$
 where $=\frac{\mathbf{w}^T\mathbf{w}}{2} + \sum_i \alpha_i[1 - y_i(\mathbf{w}^T\phi(x_i) + b)]$

Algorithm 28 Support Vector Machines (continued)

Since, we're solving an optimization problem, we set $\frac{\partial J}{\partial \mathbf{w}} = 0$ and discover that the optimal setting of \mathbf{w} is $\sum_i \alpha_i y_i \phi(x_i)$, while seeting $\frac{\partial J}{\partial \mathbf{b}} = 0$ yields the constraint $\sum_i \alpha_i y_i = 0$

```
Thus, after substituting and simplifying,
we get:  \min_{w,b} J(\mathbf{w},b,\alpha) = \sum_i \alpha_i \alpha_j y_i y_j \phi(\mathbf{x}_i)^T \phi(\mathbf{x}_j) \text{ And thus our dual is: } \\  \max_{\alpha \geq 0} [\sum_i \alpha_i - \frac{1}{2} \sum_{i,j} \alpha_j y_i y_j \phi(\mathbf{x}_i)^T \phi(\mathbf{x}_j)] \\  \text{Subject to: } \sum_i \alpha_i y_i = 0 \text{ and } 0 \leq \alpha_i \leq C
```

5 Kernel trick

Because we're working in a higher-dimension space(and potentially even an infinite-dimensional space), calculating $\phi(\mathbf{x}_i)^T\phi(\mathbf{x}_j)$ may be intractable. However, it turns out there are special kernel functions that operate on the lower dimension vectors \mathbf{x}_i and \mathbf{x}_j to produce a value equivalent to the dot-product of the higher dimensional vectors. For instance, consider the function $\phi \colon \mathbb{R}^3 \longmapsto \mathbb{R}^{10}$, where : $\phi(x) = (1, \sqrt{2}x^{(1)}, \sqrt{2}x^{(2)}, \sqrt{2}x^{(3)}, [\mathbf{x}^{(1)}]^2, [\mathbf{x}^{(2)}]^2, [\mathbf{x}^{(3)}]^2, \sqrt{2}x^{(1)(2)}, \sqrt{2}x^{(1)(3)}, \sqrt{2}x^{(2)(3)})$ Instead, we have the the kernel trick, which tells us that $K(\mathbf{x}_i, \mathbf{x}_j) = (1 + \mathbf{x}_i^T\mathbf{x}_j)^2 = \phi(\mathbf{x}_i)^T\phi(\mathbf{x}_j)$ for the given ϕ . Thus, we can simplify our calculations. Re-writing the dual in terms of the kernel yields: $\max_{\alpha \geq 0} [\sum_i \alpha_i - \frac{1}{2} \sum_{i,j} \alpha_i \alpha_j y_i y_j]$

6 Decision function

To classify a novel instance \mathbf{x} once you've learned the optimal α_i parameters, all you have to do is calculate $f(\mathbf{x}) = sign(\mathbf{w}^T\mathbf{x} + b) = \sum_i \alpha_i y_i \phi(\mathbf{x}_i)$ and using the kernel trick). Note that α_i is only non-zero for instances $\phi(\mathbf{x}_i)$ on or near the boundary-those are called the *supportvector* since they alone specify the decision boundary. We can toss out the other data points once training is complete. Thus, we only sum over the x_i which constitute the support vectors.