FAKULTÄT FÜR MATHEMATIK, INFORMATIK UND STATISTIK INSTITUT FÜR INFORMATIK

LEHRSTUHL FÜR DATENBANKSYSTEME UND DATA MINING

#### Lecture Notes to Big Data Management and Analytics Winter Term 2018/2019 Stream Analytics

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LUDWIG-

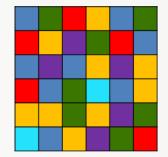
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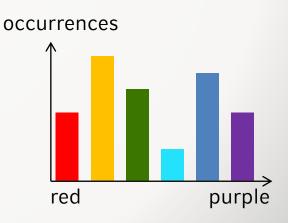
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#### Outlook

- Maintaining Histograms
- Change Detection
- Frequent Itemset Mining
- Clustering
- Classification

- Histograms are graphical representations of the distribution of numerical data
- Histograms estimate the probability distribution of a random variable
- Used for approximate query processing with error guarantees





- Histograms are defined by non-overlapping intervals or buckets
- A bucket is defined by its boundaries and its frequency count
- In case of streams:
   One never observes all values of a random variable
- → k-bucket histogram defined as  $] - \infty, b_1], ]b_1, b_2], ..., ]b_{k-1}, \infty[$  buckets with frequency counts  $f_1, f_2, ..., f_k$

In general: two types of histogram maintenance techniques

- 1. Equal-width histograms: The range of observed values is divided into equi-sized intervals  $(\forall i, j: (b_i, b_{i+1}) = (b_j, b_{j+1}))$
- 2. Equal-frequency histograms: The range of observed values is divided into k intervals such that the counts in each interval are equal  $(\forall i, j: (f_i = f_j))$

*K*-buckets Histograms (Gibbons et al., 1997)

- Incremental maintenance of histograms applicable for Insert-Delete Models
- Setting: Pre-defined number of intervals k and continuously occurring inserts and deletes as given in a sliding window approach
- Histogram maintenance based on two operations
  - Split & Merge Operation
  - Merge & Split Operation

K-buckets Histograms (Gibbons et al., 1997)

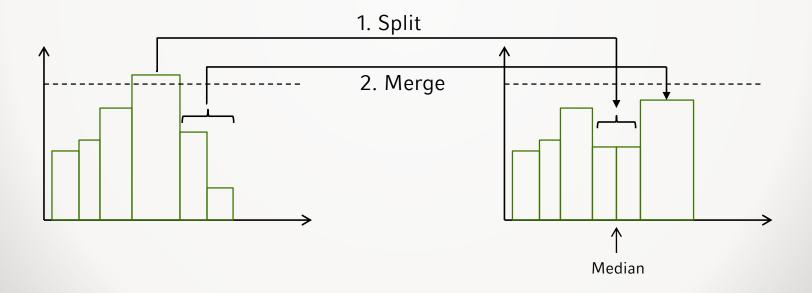
- 1. Split & Merge Operation:
  - Occurs with inserts
  - Triggered whenever the count in a bucket is greater than a given threshold
  - Split overflowed bucket into two and merge two consecutive buckets

#### **Stream Applications and Algorithms**

### **Maintaining Histograms**

K-buckets Histograms (Gibbons et al., 1997)

1. Split & Merge Operation:



K-buckets Histograms (Gibbons et al., 1997)

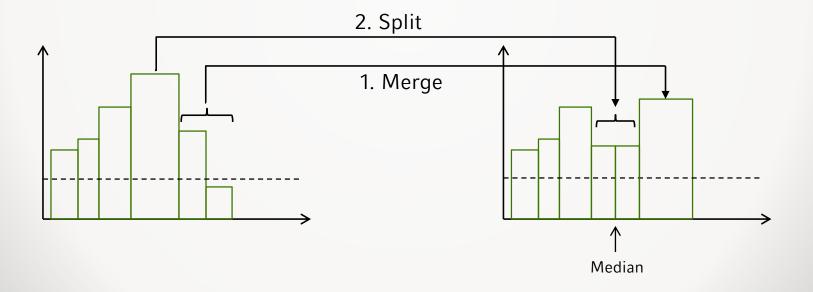
- 1. Merge & Split Operation:
  - occurs with deletes
  - triggered whenever the count in a bucket is below a given threshold
  - merge "underflowed" bucket with a neighbor bucket and split the bucket with the highest count

#### **Stream Applications and Algorithms**

### **Maintaining Histograms**

*K*-buckets Histograms (Gibbons et al., 1997)

1. Merge & Split Operation:



Exponential Histograms (Datar et al., 2002)

- task: count the number of 1s among the last N readings of a bit stream
- trivial solution: maintain a window of N Bits as a ring buffer
   => remove last bit and decrement count if it was a 1
   => insert new bit and increment count if it is a 1

Can we reduce the memory requirement to log(N)?

- exponential histograms estimate the number of 1s
- memory consumption is log(N)
- compresses older readings stronger than fresh ones

Exponential Histograms (Datar et al., 2002)

- varying bucket sizes and interval sizes
- each bucket consists of *size* and *timestamp*
- uses two additional variables, i.e. *LAST* and *TOTAL*, to estimate the number of elements in the sliding window

#### Exponential Histograms (Datar et al., 2002)

```
Algorithm Exponential Histogram Maintenance
Input: data stream S, window size N, error param. \epsilon
begin
                                                     Algorithm Exponential Histogram Count Estimation
 TOTAL \coloneqq 0
                                                     Input: current Exponential Histogram EH
 LAST \coloneqq 0
                                                     Output: estimate number of 1's within EH.N
 while S do
                                                     begin
   x_i \coloneqq S.next
                                                      return EH. TOTAL – EH. LAST/2
  if x_i == 1 do
                                                     end
    create new bucket b_i with timestamp t_i
    TOTAL += 1
    while t_i \leq t_i - N do
      TOTAL = b_1. size
      drop the oldest bucket b_l
      b_l \coloneqq b_{l-1}
      LAST \coloneqq b_1. size
    while exist |1/\epsilon|/2 + 2 buckets of the same size do
      merge the two oldest buckets of the same size with the largest timestamp of both buckets
      if last bucket was merged do
       LAST := size of the new created last bucket
end
```

S = (1,0,1,0,1,1,1,0,0,0,1,1) N = 8 (2<sup>3</sup>=> 3 Buckets),  $\epsilon = \frac{1}{2}$ 

Timest. <i>t</i> <sub>i</sub>	Buckets <i>b<sub>i</sub></i>	Element <i>x<sub>i</sub></i>	TOTAL	LAST	# buckets of same size = τ ?
1	11	1	1	0	no
2	11	0	1	0	no
3	1 <sub>1</sub> , 1 <sub>3</sub>	1	2	0	no
4	1 <sub>1</sub> , 1 <sub>3</sub>	0	2	0	no
5	$1_1, 1_3, 1_5 \\ 2_3, 1_5$	1	3	2	yes
6	2 <sub>3</sub> , 1 <sub>5</sub> , 1 <sub>6</sub>	1	4	2	no
7	$2_3, 1_5, 1_6, 1_7$ $\rightarrow 2_3, 2_6, 1_7$	1	5	2	yes
8				•••	

**General Assumptions:** 

- for static datasets:
  - data generated by a fixed process
  - data is a sample of a fixed distribution
- for data streams:
  - additional temporal dimension
  - underlying process can change over time
  - $\rightarrow$  challenge: detection and quantification of changes

Impact of changes on data processing algorithms:

• Data Mining:

data that arrived before a change can bias the model due to characteristics that no longer hold after the change

• Query Processing:

query answers for time intervals with stable underlying data distributions might be more meaningful

The nature of changes

 Concept Drifts: gradual change in target concept

 Concept Shifts: abrupt change in target concept



mmmmmm

Two general approaches

- Monitoring the evolution of performance indicators (Klinkenberg et al., 1998), e.g.
  - accuracy of the current classifier
  - attribute value distribution
  - monitoring *top* attributes (according to any ranking)
- monitoring distribution on two different time-windows

CUSUM Algorithm (Page, 1954)

- monitors the cumulative sum of instances of a random variable
   Algorithm CUSUM
- detects a change if the cumulative difference between observation and likelihood gets larger than threshold α

```
Algorithm CUSUM

Input: data stream S, threshold param. \alpha

begin

G_0 \coloneqq 0

while S do

x_t \coloneqq next instance of S

compute likelihood \omega_t

G_t \coloneqq \max(0, G_{t-1} - \omega_t + x_t)

if G_t > \alpha then

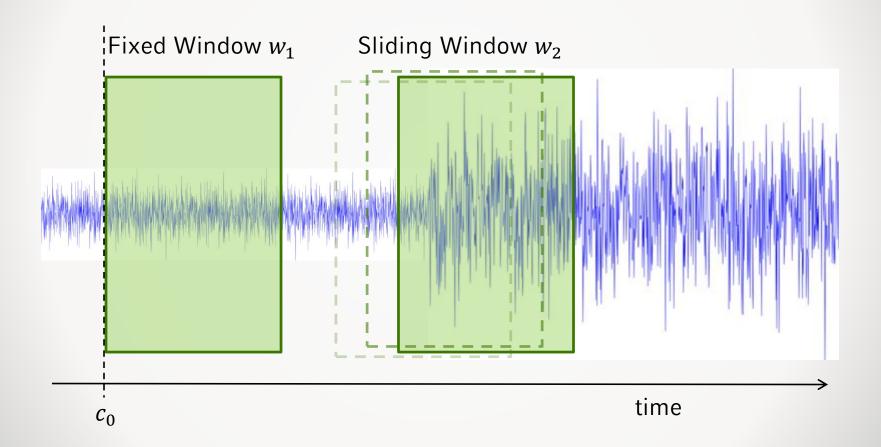
report change at time t

G_t \coloneqq 0

end
```

•  $\omega_t$  commonly represents the likelihood function

#### Two Windows Approach (Kifer et al., 2004)



#### Two Windows Approach (Kifer et al., 2004)

```
Algorithm Two Windows Approach

Input: data stream S, window sizes m_1 and m_2, distance func. d: D \times D \to R, threshold param. \alpha

begin

c_0 \coloneqq 0

W_1 \coloneqq \text{first } m_1 \text{ points from time } c_0

W_2 \coloneqq \text{most recent } m_2 \text{ points from } S

while S do

slide W_2 by 1 point

if d(W_1, W_2) > \alpha then

c_0 \coloneqq \text{current time}

report change at time c_0

W_1 \coloneqq \text{first } m_1 \text{ points from time } c_0

W_2 \coloneqq \text{most recent } m_2 \text{ points from S}

end
```

#### d measures the distance between two probability distributions

- Let  $A = \{a_1, a_2, \dots, a_n\}$  be a set of *items* (e.g. products)
- Any subset  $I \subseteq A$  is called an *itemset*
- Let  $T = (t_1, t_2, ..., t_m)$  be a set of *transactions* with  $t_i$  being a pair  $\langle TID_i, I_i \rangle$  where  $I_i \subseteq A$  is a set of items (e.g. the set of products bought by a customer within a certain period in time)
- The support  $\sigma_{min}$  of an itemset  $I \subseteq A$  is the number/fraction of transactions  $t_i \in T$  that contain I

#### Example:

Given the set of items  $A = \{a, b, c, d, e\}$ , the set of transactions T, and a relative support  $\sigma_{min} = 0.3$ , determine the set of frequent item sets that is  $\{I \subseteq A | \sigma_T(I) \ge \sigma_{min}\}$ .

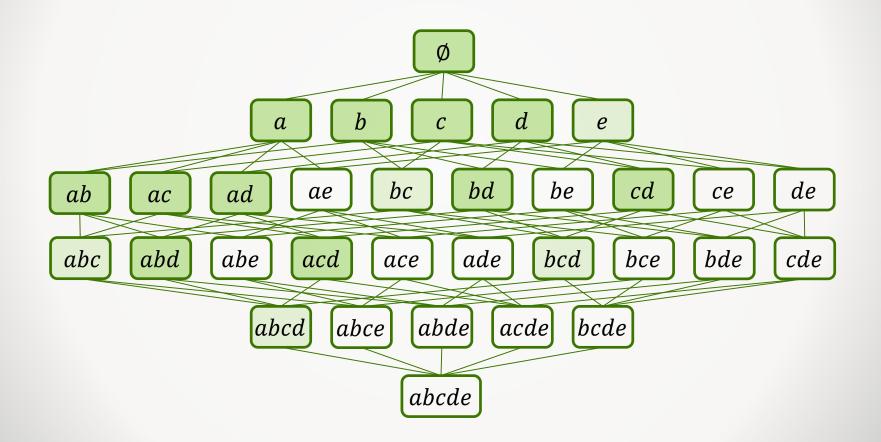
T:	TID <sub>i</sub>	I <sub>i</sub>			
	1	$\{a, b, c, d\}$	0 items	1 item	2 items
	2	{ <i>b</i> , <i>d</i> , <i>e</i> }	Ø: 7	{ <i>a</i> }:5	{a, b}: 3
	3	$\{a, b, d\}$		{ <i>b</i> }:5	{ <i>a</i> , <i>c</i> }: 4
	4	{ <i>a</i> , <i>b</i> , <i>c</i> , <i>d</i> , <i>e</i> }		{ <i>c</i> }:5	${a,d}:4$
	5	{ <i>a</i> , <i>c</i> }		$\{d\}: 6$	$\{b, d\}: 4$
	6	{ <i>c</i> , <i>d</i> }			${c, d}: 4$
	7	{ <i>a</i> , <i>c</i> , <i>d</i> }			

3 items

 ${a, c, d}: 3$ 

 ${a, b, d}: 3$ 

search space



LossyCounting Algorithm (Manku et al., 2002)

- One-pass algorithm for computing frequency counts that exceed a user-specified threshold
- Approximate error but guaranteed to be below a userspecified boundary
- $\rightarrow$  Two parameters:
  - Support threshold  $s \in [0,1]$
  - Error threshold  $\epsilon \in [0,1]$
  - $-\epsilon \ll s$

LossyCounting Algorithm (Manku et al., 2002)

- Setup:
  - Stream S is divided into buckets of width  $\omega = \left|\frac{1}{\epsilon}\right|$
  - The current bucket id  $b_{curr} = \left[\frac{N}{\omega}\right]$
  - For element e, the true frequency seen so far is  $f_e$
  - The data structure D is a set of entries of the form  $(e, f, \Delta)$ 
    - e is the element
    - *f* is the frequency seen since *e* is in *D*
    - $\Delta$  is the maximum possible error, resp. the estimated frequency of *e* in buckets b = 1 to  $b_{curr}$ -1

#### LossyCounting Algorithm (Manku et al., 2002)

```
Algorithm LossyCounting
Input: data stream S, error threshold \epsilon
begin
 D = \emptyset, N = 0, \omega = \left|\frac{1}{\epsilon}\right|
 while S do
   e_i \coloneqq next object from S
    N += 1
    b_{curr} = \left[\frac{N}{\omega}\right]
    if e_i \in D then
      increment e_i's frequency by 1
    else
     D.add((e_i, 1, b_{curr} - 1))
    whenever N \equiv 0 \mod \omega do
      foreach entry (e, f, \Delta) in D do
        if f + \Delta \leq b_{curr} then
          delete (e, f, \Delta)
end
```

```
Algorithm LossyCounting – User request
Input: lookup table D, support threshold s
begin
```

 $S = \emptyset$ foreach entry  $(e, f, \Delta)$  in D do if  $f \ge (s - \epsilon)N$  then add  $(e, f, \Delta)$  to Sreturn Send

f is the exact frequency count of e since the entry was inserted into D

 $\Delta$  is the maximum number of times *e* could have occurred in the first  $b_{curr} - 1$  buckets

**Clustering** is the process of grouping objects into different groups, such that the similarity of data in each subset is high, and between different subsets is low.

**Clustering from data streams** aims at maintaining a continuously consistent good clustering of the sequence observed so far, using a small amount of memory and time.

General approaches to clustering

- *Partitioning*: Fixed number of clusters, new object is assigned to closest cluster center (k-means/k-medoid)
- Density-based: Take connectivity and density functions into account (DBSCAN)
- *Hierarchical*: Find a tree-like structure representing the hierarchy of the cluster model (Single Link/Complete Link)
- *Grid-based*: Partition the space into grid cells (STING)
- Model-based: Take a model and find the best fit clustering (COBWEB)

Requirements for stream clustering algorithms

- Compactness of representation
- Fast, incremental processing (one-pass)
- Tracking cluster changes

   (as clusters might (dis-)appear over time)
- Clear and fast identification of outliers

LEADER algorithm (Spath, 1980)

- Simplest form of partitioning based clustering applicable to data streams
- Depends on the order of incoming objects
- Depends on a good choice of the threshold parameter  $\delta$

```
Algorithm LEADER

Input: data stream S, threshold param. \delta

begin

while S do

x_i \coloneqq next object from S

find closest cluster c_{clos} to x_i

if d(c_{clos}, x_i) < \delta then

assign x_i to c_{clos}

else

create new cluster with x_i

end
```

Stream K-means (O'Callaghan et al., 2002)

- Partition data stream S into chunks X<sub>1</sub>, ..., X<sub>n</sub>, ... so that each chunk fits in memory
- Apply k-means for each chunk X<sub>i</sub> and retrieve k cluster centers each weighted with the number of points it compresses
- Apply k-means on the cluster centers to get an overall kmeans clustering when demanded

Microcluster-based Clustering

- Common approach to capture temporal information for being able to deal with cluster evolution
- A *microcluster* (or *cluster feature CF*) is a triple (*N*,*LS*,*SS*) that stores the sufficient information of a set of points
  - *N* is the number of points
  - LS is the linear sum of the N points, i.e.  $\sum_{i=1}^{N} \vec{x_i}$
  - SS is the square sum of the N points, i.e.  $\sum_{i=1}^{N} \overrightarrow{x_i}^2$

Microcluster-based Clustering

- The properties of cluster features are:
  - Incrementality:

$$N_i = N_i + 1$$
,  $LS_i = LS_i + \vec{x}$ ,  $SS_i = SS_i + \vec{x}^2$ 

- Additivity:  

$$N_k = N_i + N_j$$
,  $LS_k = LS_i + LS_j$ ,  $SS_k = SS_i + SS_j$ 

- Centroid: 
$$\overrightarrow{X_c} = \frac{LS_i}{N}$$

- Radius: 
$$r = \sqrt{\frac{SS_i}{N_i} - \left(\frac{LS_i}{N_i}\right)^2}$$

BIRCH (Zhang et al., 1996)

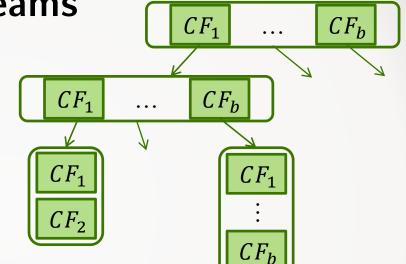
- Usage of Microclusters within CF-Tree
  - B<sup>+</sup>-Tree like structure
  - Two user specified parameters:
    - Branching factor B
    - Maximum diameter (or radius) T of a CF
  - Each non-leaf node contains at most B entries of the form [CF<sub>i</sub>, child<sub>i</sub>] where
    - *CF<sub>i</sub>* is the CF representing the subcluster that child forms
    - *child<sub>i</sub>* is a pointer to the i-th child node
  - Each leaf node contains entries of the form [CF<sub>i</sub>, prev, next]

# **Stream Applications and Algorithms**

# **Clustering from Data Streams**

BIRCH (Zhang et al., 1996)

- Inserts into CF-Tree
  - At each non-leaf node, the new object follows the *closest-CF* path



- At leaf node level, the closest-CF tries to absorb the object (which depends on diameter threshold T and the page size)
  - If possible: update *closest-CF*
  - If not possible: make a new CF entry in the leaf node (split the parent node if there is no space)

# **Clustering from Data Streams**

BIRCH (Zhang et al., 1996)

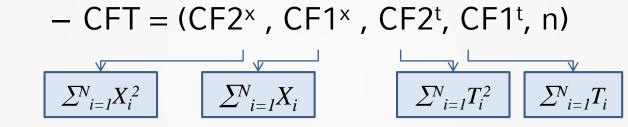
Two step algorithm:

- 1. Online component:
  - Microclusters are kept locally
  - Maintenance of the hierarchical structure
  - Optional: Condense by building smaller CF-Tree (requires scan over leaf entries)
- 2. Offline component:
  - Apply global clustering to all leaf entries
  - Optional: Cluster refinement to the cost of additional passes (use centroids retrieved by global clustering and re-assign data points)

### CluStream [AggEtAl03]

Assume that the data stream consists of a set of multi-dimensional records  $X_{1}, ..., X_{n_i}, ...,$  arriving at  $T_{1}, ..., T_{n_i}, ...; X_i = (x_i^{1}, ..., x_i^{d})$ 

• The micro-cluster summary for a set of d-dimensional points ( $X_1$ ,  $X_2$ , ...,  $X_n$ ) arriving at time points  $T_1$ ,  $T_2$ , ...,  $T_n$  is defined as:



• Easy calculation of basic measures to characterize a cluster:

• Center: 
$$\frac{CF1^x}{n}$$
 • Radius:  $\sqrt{\frac{CF2^x}{n} - \left(\frac{CF1^x}{n}\right)^2}$ 

- Important properties of micro-clusters:
  - Incrementality:  $CFT(C_1 \cup p) = CFT(C_1) + p$
  - Additivity:  $CFT(C_1 \cup C_2) = CFT(C_1) + CFT(C_2)$
  - Subtractivity:  $CFT(C_1 C_2) = CFT(C_1) CFT(C_2)$ ,  $C_1 \supseteq C_2$

### CluStream: overview

- A fixed number of *q* micro-clusters is maintained over time
- Initialize: apply q-Means over initPoints, built a summary for each cluster
- Online micro-cluster maintenance as a new point p arrives from the stream
  - Find the closest micro-cluster *clu* for the new point *p* 
    - If *p* is within the max-boundary of *clu*, *p* is <u>absorbed</u> by *clu*
    - o.w., a <u>new cluster</u> is created with *p*
  - The number of micro-clusters should not exceed q
    - <u>Delete</u> most obsolete micro-cluster or <u>merge</u> the two closest ones
- Periodic storage of micro-clusters snapshots into disk
  - At different levels of granularity depending upon their recency
- Offline macro-clustering
  - Input: A user defined time horizon h and number of macro-clusters k to be detected
  - Locate the valid micro-clusters during h
  - Apply k-Means upon these micro-clusters  $\rightarrow$  k macro-clusters

### **CluStream:** Initialization step

- Initialization
  - Done using an offline process in the beginning
  - Wait for the first InitNumber points to arrive
  - Apply a standard k-Means algorithm to create q clusters
    - For each discovered cluster, assign it a unique ID and create its microcluster summary.
- Comments on the choice of *q* 
  - much larger than the natural number of clusters
  - much smaller than the total number of points arrived

### CluStream: Online step

- A fixed number of *q* micro-clusters is maintained over time
- Whenever a new point *p* arrives from the stream
  - Compute distance between p and each of the q maintained micro-cluster centroids
  - $clu \leftarrow$  the closest micro-cluster to p
  - Find the max boundary of *clu*
    - It is defined as a factor of *t* of *clu* radius
  - If p falls within the maximum boundary of *clu* 
    - *p* is absorbed by *clu*
    - Update clu statistics (incremental property)
  - Else, create a new micro-cluster with p, assign it a new ID, initialize its statistics
    - To keep the total number of micro-clusters fixed (i.e., *q*):
      - Delete the most obsolete micro-cluster or
        - If its safe based on its time statistics
      - Merge the two closest ones (Additivity property)
        - When two micro-clusters are merged, a list of ids is created. This way, we can identify the component micro-clusters that comprise a micro-cluster.

### CluStream: storing micro-cluster storage

- all micro-clusters are saved to disc for a given time frame (snapshot)
- not all snapshots are kept. goal: provide a high granularity for recent snapshots and maintain a rough picture of older snapshots
- ⇒ Pyramidal Storage

**idea**: Maintain the last *a<sup>i</sup>+1* snapshots for multiple levels of storage granularity. => larger level cover a longer history with less granularity

- For each level i, we store a snapshot if the current timestamp t mod a<sup>i</sup> = 0, except t mod a<sup>i+1</sup> = 0 holds as well (no redundancy)
- At most a<sup>b</sup>+1 snapshots are stored at each order; if a new snapshot arrives the oldest one is deleted.
- #levels: log<sub>a</sub>(t)
- #stored snapshots: (a<sup>b</sup>+1)log<sub>a</sub>(t)

level	clock times
0	<del>60</del> 59 <del>58</del> 57 <del>56</del>
1	<del>60</del> 58 <del>56</del> 54 <del>52</del>
2	60 56 52 4 <del>8</del> 44
3	48 40 <del>32</del> 24 <del>16</del>
4	48 <del>32</del> 16
5	32

Snapshots stored at t = 60, a=2, b=2

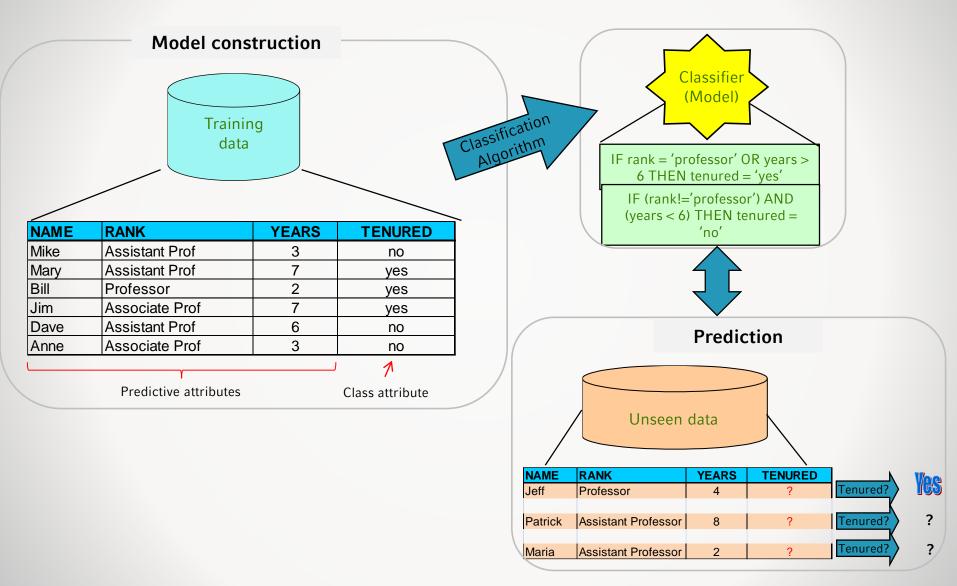
### CluStream: Offline step

- The offline step is applied on demand
- User input: time horizon h, # macro-clusters k to be detected
- Step 1: Find the active micro-clusters during *h*:
  - We exploit the subtractivity property to find the active micro-clusters during *h*:
    - Suppose current time is  $t_c$ . Let  $S(t_c)$  be the set of micro-clusters at  $t_c$ .
    - Find the stored snapshot which occurs just before time  $t_c$ -h. We can always find such a snapshot h'. Let  $S(t_c-h')$  be the set of micro-clusters.
    - For each micro-cluster in the current set  $S(t_c)$ , we find the list of its component micro-cluster ids. For each of the list of ids, find the corresponding micro-clusters in  $S(t_c-h')$ .
    - Subtract the CF vectors for the corresponding micro-clusters in  $S(t_c-h')$
    - This ensures that the micro-clusters created before the user-specified horizon do not dominate the result of clustering process
- Step 2: Apply k-Means over the active micro-clusters in h to derive the k macroclusters
  - Initialization: centers are not picked up randomly, rather sampled with probability proportional to the number of points in a given micro-cluster
  - Distance is the centroid distance
  - New centers are defined as the weighted centroids of the micro-clusters in that partition

### **CluStream:** discussion

- + CluStream clusters large evolving data streams
- + Views the stream as a changing process over time, rather than clustering the whole stream at a time
- + Can characterize clusters over different time horizons in changing environment
- + Provides flexibility to an analyst in a real-time and changing environment
- Fixed number of micro-clusters maintained over time
- Sensitive to outliers/ noise

### The (batch) classification process



### **Stream vs batch classification 1/2**

- So far, classification as a batch/ static task
  - The whole training set is given as input to the algorithm for the generation of the classification model.
  - The classification model is static (does not change)
  - When the performance of the model drops, a new model is generated from scratch over a new training set.
- But, in a dynamic environment data change continuously
   Batch model re-generation is not appropriate/sufficient anymore

# Stream vs batch classification 2/2

Need for new classification algorithms that

- have the ability to incorporate new data
- deal with non-stationary data generation processes (concept drift/shift)

 $\circ\,$  ability to remove obsolete data

# – subject to:

resource constraints (processing time, memory)

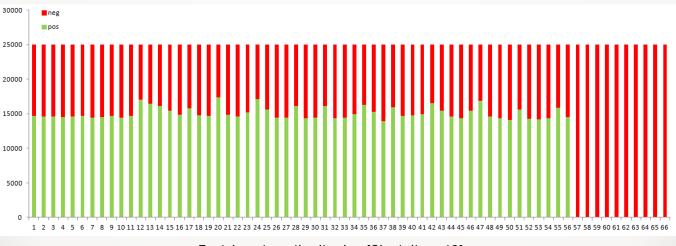
 $\circ$  single scan of the data (one look, no random access)

### Non-stationary data distribution → Concept drift

- In dynamically changing and non-stationary environments, the data distribution might change over time yielding the phenomenon of concept drift
- Different forms of change:
  - The input data characteristics might change over time
  - The relation between the input data and the target variable might change over time
- Concept drift between  $t_0$  and  $t_1$  can be defined as  $\exists X : p_{t_0}(X,y) \neq p_{t_1}(X,y)$ 
  - P(X,y): the joint distribution between X and y
- According to the Bayesian Decision Theory:  $p(y|X) = \frac{p(y)p(X|y)}{p(X)}$
- So, changes in data can be characterized as changes in:
  - The prior probabilities of the classes p(y)
  - The class conditional probabilities p(Xly).
  - The posterior p(y|X) might change

### Example: Evolving class priors

- E.g., evolving class distribution
  - The class distribution might change over time
  - Example: Twitter sentiment dataset
    - o 1.600.000 instances split in 67 chunks of 25.000 tweets per chunk
    - Balanced dataset (800.000 positive, 800.000 negative tweets)
    - The distribution of the classes changes over time
    - Dataset online at: https://sites.google.com/site/twittersentimenthelp/forresearchers



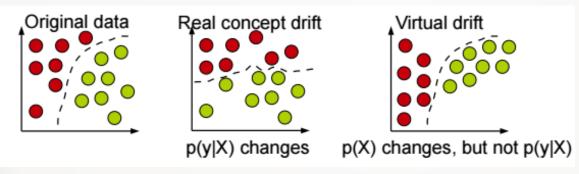
Evolving class distribution [Sinelnikova12]

# Real vs virtual drift

- Real concept drift
  - Refers to changes in p(y|X). Such changes can happen with or without change in p(X).

Source: [GamaETAI13]

- E.g., "I am not interested in tech posts anymore"
- Virtual concept drift
  - If the p(X) changes without affecting p(y|X)



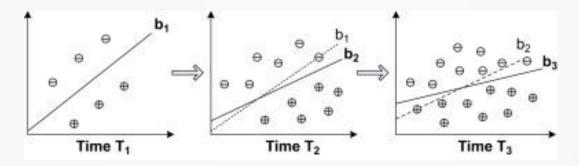
Drifts (and shifts)

0

- Drift more associated to gradual changes
- Shift refers to abrupt changes

# Model adaptation

- As data evolve over time, the classifier should be updated to "reflect" the evolving data
  - Update by incorporating new data
  - Update by forgetting obsolete data



The classification boundary gradually drifts from  $b_1$  (at  $T_1$ ) to  $b_2$  (at  $T_2$ ) and finally to  $b_3$  (at  $T_3$ ). (Source: A framework for application-driven classification of data streams, Zhang et al, Journal Neurocomputing 2012)

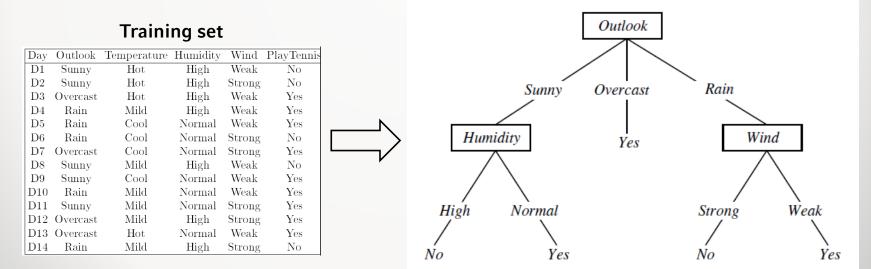
# Data stream classifiers

- The batch classification problem:
  - Given a finite training set  $D=\{(x,y)\}$ , where  $y=\{y_1, y_2, ..., y_k\}$ , |D|=n, find a function y=f(x) that can predict the y value for an unseen instance x
- The data stream classification problem:
  - Given an infinite sequence of pairs of the form (x,y) where  $y=\{y_1, y_2, ..., y_k\}$ , find a function y=f(x) that can predict the y value for an unseen instance x
    - the label y of x is not available during the prediction time
    - but it is available shortly after for model update Supervised scenario

- Example applications:
  - Fraud detection in credit card transactions
  - Churn prediction in a telecommunication company
  - Sentiment classification in the Twitter stream \_
  - Topic classification in a news aggregation site, e.g. Google news

### (Batch) Decision Trees (DTs)

- Training set: D = {(x,y)}
  - predictive attributes: x=<x<sub>1</sub>, x<sub>2</sub>, ..., x<sub>d</sub>>
  - class attribute:  $y=\{y_1, y_2, ..., y_k\}$
- Goal: find y=f(x)
- Decision tree model
  - nodes contain tests on the predictive attributes
  - leaves contain predictions on the class attribute



#### Big Data Management and Analytics

### (Batch) DTs: Selecting the splitting attribute

- Basic algorithm (ID3, Quinlan 1986)
  - Tree is constructed in a top-down recursive divide-and-conquer manner
  - At start, all the training examples are at the root node

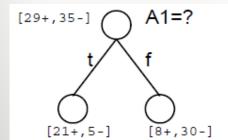
# 1. $A \leftarrow$ the "best" decision attribute for next node2. Assign A as decision attribute for node3. For each value of A, create new descendant of

- 4. Sort training examples to leaf nodes
- 5. If training examples perfectly classified, Then STOP, Else iterate over new leaf nodes

#### – But, which attribute is the best?

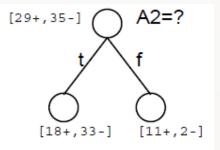
#### Attribute selection measures:

- Information gain
- Gain ratio
- Gini index



Main loop:

node



Goal: select the most "useful" attribute

 i.e., the one resulting in the purest partitioning

### (Batch) DTs: Information gain

- Used in ID3
- It uses entropy, a measure of pureness of the data
- The information gain Gain(S,A) of an attribute A relative to a collection of examples S measures the gain reduction in S due to splitting on A:

$$Gain(S, A) = Entropy(S) - \sum_{v \in Values(A)} \frac{|S_v|}{|S|} Entropy(S_v)$$
  
Before splitting After splitting on A

Gain measures the expected reduction in entropy due to splitting on A

• The attribute with the higher entropy reduction is chosen

### (Batch) DTs: Entropy

- Let S be a collection of positive and negative examples for a binary classification problem, C={+, -}.
- p<sub>+</sub>: the percentage of positive examples in S
- p\_: the percentage of negative examples in S
- Entropy measures the impurity of S:

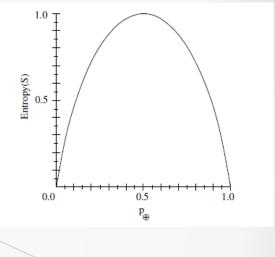
$$Entropy(S) = -p_{+} \log_{2}(p_{+}) - p_{-} \log_{2}(p_{-})$$

• Examples :

- Let S: [9+,5-] 
$$Entropy(S) = -\frac{9}{14}\log_2(\frac{9}{14}) - \frac{5}{14}\log_2(\frac{5}{14}) = 0.940$$

- Let S: [7+,7-] 
$$Entropy(S) = -\frac{7}{14}\log_2(\frac{7}{14}) - \frac{7}{14}\log_2(\frac{7}{14}) = 1$$

- Let S: [14+,0-]  $Entropy(S) = -\frac{14}{14}\log_2(\frac{14}{14}) - \frac{0}{14}\log_2(\frac{0}{14}) = 0$ 

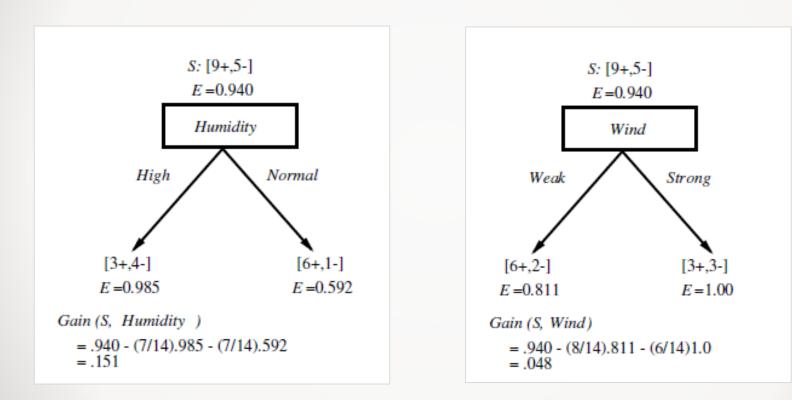


in the general case (k-classification problem) Entropy(S) =  $\sum_{i=1}^{k} - p_i \log_2(p_i)$ 

- Entropy = 0, when all members belong to the same class
- Entropy = 1, when there is an equal number of positive and negative examples

### (Batch) DTs: Information gain example

• Which attribute to choose next?



### From batch to stream DT induction

- Thus far, in order to decide on which attribute to use for splitting in a node (essential operation for building a DT), we need to have all the training set instances resulting in this node.
- But, in a data stream environment
  - The stream is infinite
  - We can't wait for ever in a node
- Can we make a valid decision based on some data?
  - Hoeffding Tree or Very Fast Decision Tree (VFDT) [DomingosHulten00]

### Hoeffding Tree [DomingosHulten00]

- Idea: In order to pick the best split attribute for a node, it may be sufficient to consider only a small subset of the training examples that pass through that node.
  - No need to look at the whole dataset
  - (which is infinite in case of streams)
- **Problem**: How many instances are necessary?
  - Use the Hoeffding **bound**!

### The Hoeffding bound

- Consider a real-valued random variable *r* whose range is *R* 
  - e.g., for a probability the range is 1
  - for information gain the range is  $log_2(c)$ , where c is the number of classes
- Suppose we have *n* independent observations of *r* and we compute its mean  $\bar{r}$
- The Hoeffding bound states that with confidence  $1-\delta$  the true mean of the variable,  $\mu_r$ , is at least  $\bar{r}$ - $\epsilon$ , i.e.,  $P(\mu_r \ge \bar{r}-\epsilon) = 1-\delta$

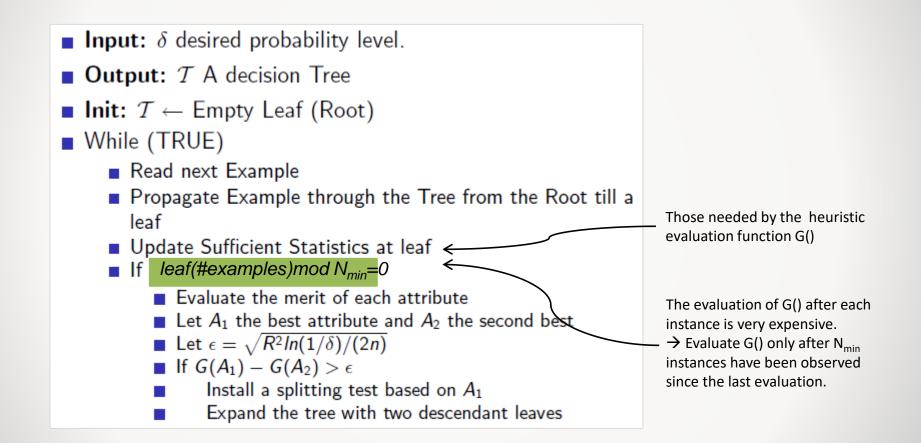
• The 
$$\varepsilon$$
 is given by:  $\varepsilon = \sqrt{\frac{R^2 \ln(1/\delta)}{2n}}$ 

- This bound holds true regardless of the distribution generating the values, and depends only on the range of values, number of observations and desired confidence.
  - A disadvantage of being so general is that it is more conservative than a distributiondependent bound

Using the Hoeffding bound to select the best split at a node

- Let *G()* be the heuristic measure for choosing the split attribute at a node
- After seeing *n* instances at this node, let
  - $X_a$ : be the attribute with the highest observed G()
  - $X_b$ : be the attribute with the second-highest observed G()
- $\Delta \bar{G} = \bar{G}(X_a) \bar{G}(X_b) \ge 0$  the difference between the 2 best attributes
- $\Delta \bar{G}$  is the random variable being estimated by the Hoeffding bound
- Given a desired  $\delta$ , if  $\Delta \overline{G} > \varepsilon$  after seeing n instances at the node
  - the Hoeffding bound guarantees that with probability 1- $\delta$ ,  $\Delta \overline{G} \ge \Delta \overline{G} \cdot \varepsilon > 0$ .
  - Therefore we can confidently choose  $X_a$  for splitting at this node
- Otherwise, i.e., if  $\Delta \overline{G} < \varepsilon$ , the sample size is not enough for a stable decision.
  - With R and  $\delta$  fixed, the only variable left to change  $\epsilon$  is n
  - We need to extend the sample by seeing more instances, until  $\epsilon$  becomes smaller than  $\Delta \bar{G}$

### Hoeffding Tree algorithm

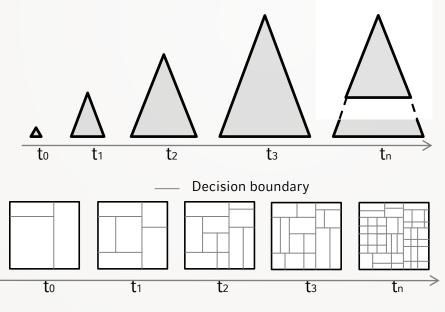


Hoeffding tree algorithm more details

- Breaking ties
  - When ≥2 attributes have very similar G's, potentially many examples will be required to decide between them with high confidence.
  - This is presumably wasteful, as it makes little difference which is chosen.
  - Break it by splitting on current best if  $\Delta G < \epsilon < \tau$ ,  $\tau$  a user-specified threshold
- Grace period (MOA's term)
  - Recomputing G() after each instance is too expensive.
  - A user can specify # instances in a node that must be observed before attempting a new split

### Hoeffding Tree overview

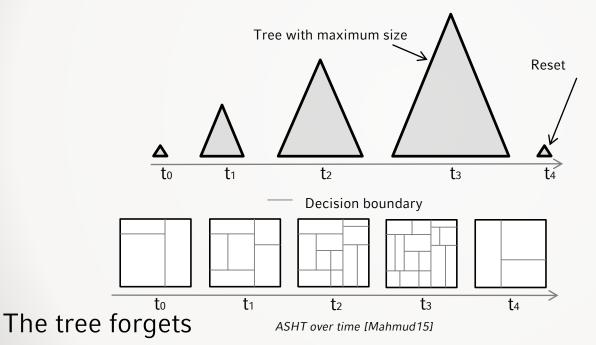
- The HT accommodates new instances from the stream
- But, doesn't delete anything (doesn't forget!)
- With time
  - The tree becomes more complex (overfitting is possible)
  - The historical data dominate its decisions (difficult to adapt to changes)



HT over time [Mahmud15]

### Adaptive Size Hoeffding Tree (ASHT) [BifetEtAl09]

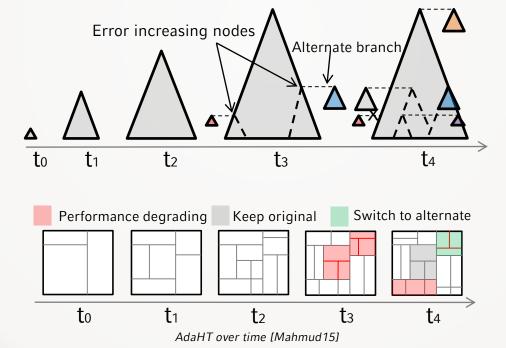
- Introduces a maximum size (#splitting nodes) bound
- When the limit is reached, the tree is reset
  - Test for the limit, after node's split



- but, due to the reset, it looses all information learned thus far

## Concept-Adapting Hoeffding Tree [HultenEtAl01]

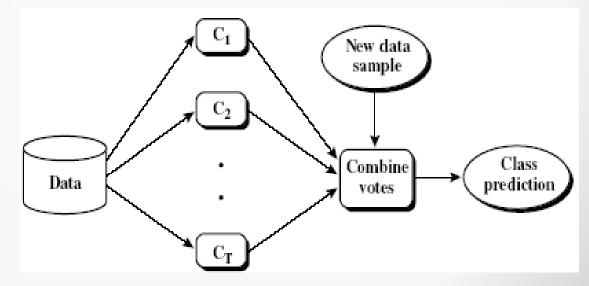
- Starts maintaining an alternate sub-tree when the performance of a node decays
- When the new sub-tree starts performing better, it replaces the original one
- If original sub-tree keeps performing better, the alternate sub-tree is deleted and the original one is kept



### Ensemble of classifiers

Idea:

- Instead of a single model, use a combination of models to increase accuracy
- Combine a series of T learned models, M<sub>1</sub>, M<sub>2</sub>, ..., M<sub>T</sub>, with the aim of creating an improved model M\*
- To predict the class of previously unseen records, aggregate the predictions of the ensemble

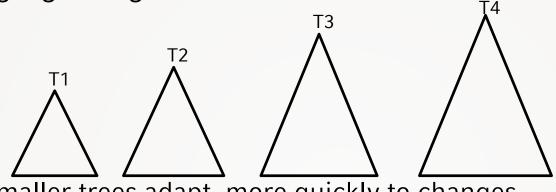


### Many methods

- Bagging
  - Generate training samples by sampling with replacement (bootstrap)
  - Learn one model at each sample
- Boosting
  - At each round, increase the weights of misclassified examples
- Stacking
  - Apply multiple base learners
  - Meta learner input = base learner predictions

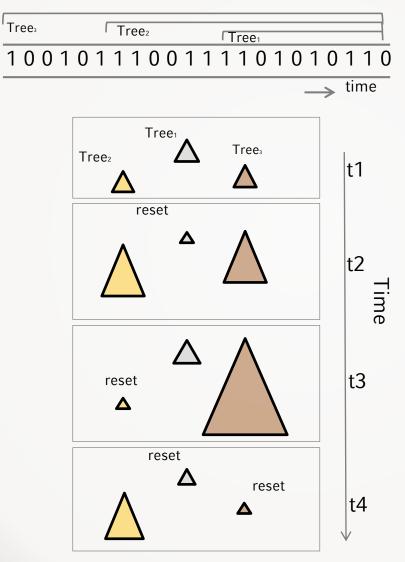
Ensemble of Adaptive Size Hoeffding Trees (ASHT) [BifetEtAl09] 1/2

Bagging using ASHTs of different sizes.



- Smaller trees adapt more quickly to changes
- Larger trees perform better during periods with no or little change
- The max allowed size for the n<sup>th</sup> ASHT tree is twice the max allowed size for the (n-1)<sup>th</sup> tree.
- Each tree has a weight proportional to the inverse of the square of its error
- The goal is to increase bagging performance by tree diversity

### Ensemble of Adaptive Size Hoeffding Trees (ASHT) [BifetEtAI09] 1/2



### Hoeffding Tree family overview

- All HT, AdaHT, ASHT accommodate new instances from the stream
- HT does not forget
- ASHT forgets by resetting the tree once its size reaches its limit
- AdaHT forgets my replacing sub-trees with new ones
- Bagging ASHT uses varying size trees that respond differently to change

# Summary: Stream Classification

- Extending traditional classification methods for data streams implies that
  - They should accommodate new instances
  - They should forget obsolete instances
- Typically, all methods incorporate new instances from the model
- They differ mainly on how do they forget
  - No forgetting, sliding window forgetting, damped window forgetting,...
- and which part of the model is affected
  - Complete model reset, partial reset, ...
- So far, we focused on fully-supervised learning and we assumed availability of class labels for all stream instances
  - Semi-supervised learning
  - Active learning
- Dealing with class imbalances, rare-classes
- Dealing with dynamic feature spaces

# further reading

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