

# Machine Learning and Data Mining Clustering

**UCSC**extension  
Silicon Valley



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# Clustering

## Main Sources

- K-Means (Andrew Ng)
  - <http://cs229.stanford.edu/notes/cs229-notes7a.pdf>
- Expectation – Maximization (Andrew Ng)
  - <http://www.youtube.com/watch?v=ey2PE5xi9-A&feature=related>
  - <http://cs229.stanford.edu/notes/cs229-notes2.pdf>
- EM Algorithm (Andrew Ng)
  - <http://cs229.stanford.edu/notes/cs229-notes8.pdf>
  - <http://cs229.stanford.edu/notes/cs229-notes7b.pdf>
- Discriminant Analysis (Andrew Ng)
  - <http://cs229.stanford.edu/notes/cs229-notes2.pdf>
- R and Expectation Maximization
  - [http://en.wikibooks.org/wiki/Data\\_Mining\\_Algorithms\\_In\\_R/Clustering/Expectation\\_Maximization\\_%28EM%29](http://en.wikibooks.org/wiki/Data_Mining_Algorithms_In_R/Clustering/Expectation_Maximization_%28EM%29)
- Paper Describing mclust package
  - <http://www.stat.washington.edu/fraley/mclust/tr504.pdf>

# R Packages

- CRAN list of Cluster Packages
  - <http://cran.r-project.org/web/views/Cluster.html>
- CRAN K-Means `kmeans()`
  - <http://127.0.0.1:57279/rhelp/browse/user-local--1vj6d1s7n4kx5/library/stats/html/kmeans.html>
- CRAN Agglomerative Hierarchical Clustering
  - `stat` package `hclust()` Installed with R
- CRAN Divisive Hierarchical Clustering `diana()`
  - <http://cran.r-project.org/web/packages/cluster/cluster.pdf>
- CRAN Mixture Model Package `mclust()`
  - <http://cran.r-project.org/web/packages/mclust/mclust.pdf>
  - <http://www.stat.washington.edu/research/reports/2006/tr504.pdf>
- CRAN Density Based Clustering `dbscan()`
  - <http://cran.r-project.org/web/packages/fpc/index.html>

# Cluster Analysis

- Huge variety in R

<http://cran.r-project.org/web/views/Cluster.html>

- Methods

- Partitioning (e.g., kmeans, pam)
- Hierarchical Agglomerative (e.g., average, ward, single, complete) (hclust() and diana())
- Model Based (e.g., ML estimation, Bayesian estimation) (mclust())
- Density Based (e.g., ML estimation, Bayesian estimation) (dbscan()) page 35 of fpc package

# Example Code

The Iris Data is globular with ellipsoidal covariance

- `kMeans.R` `kmeans()` Prototype Based Clustering
  - Randomly Generated Gaussian Data
  - Iris Data Set 89% correct
- `hclustExample.R` Hierarchical Clustering
  - USArrests Data
  - Iris Data Set
    - `hclust()` Agglomerative Hierarchical 91% correct
    - `diana()` Divisive Hierarchical 85% correct
- `mclustExample` `mclust()` Expectation - Maximization
  - Iris Data 97% Correct

# Lessons

- Measures.pdf
- Chap8\_basic\_cluster\_analysis.pdf (kmeans)
- Video of kmeans
  - <http://www.youtube.com/watch?v=74rv4snLI70&feature=endscreen&NR=1>
- Example Code Kmeans.r
- Chap8\_basic\_cluster\_analysis.pdf (Hierarchical Clustering)
- Expectation – Maximization
- Video of Expectation – Maximization
  - <http://www.youtube.com/watch?v=v-pg8VCQk4M&feature=related>
- Example Code (mclustExample.r)

# Classes vs. Clusters

- **Supervised:**  $X = \{ \mathbf{x}^j, y^j \}_t$
- Classes  $C_i, i=1, \dots, K$

$$p(\mathbf{x}) = \sum_{i=1}^K p(\mathbf{x} | C_i) P(C_i)$$

where  $p(\mathbf{x} | C_i) \sim N(\boldsymbol{\mu}_i, \boldsymbol{\Sigma}_i)$

- $\Phi = \{P(C_i), \boldsymbol{\mu}_i, \boldsymbol{\Sigma}_i\}_{i=1}^K$

$$\hat{P}(C_i) = \frac{\sum_j y_i^j}{N} \quad \mathbf{m}_i = \frac{\sum_j y_i^j \mathbf{x}^j}{\sum_j y_i^j}$$

$$\mathbf{S}_i = \frac{\sum_j y_i^j (\mathbf{x}^j - \mathbf{m}_i)(\mathbf{x}^j - \mathbf{m}_i)^T}{\sum_j y_i^j}$$

**Unsupervised:**  $X = \{ \mathbf{x}^j \}_j$

- Clusters  $G_i, i=1, \dots, k$

$$p(\mathbf{x}) = \sum_{i=1}^k p(\mathbf{x} | G_i) P(G_i)$$

where  $p(\mathbf{x} | G_i) \sim N(\boldsymbol{\mu}_i, \boldsymbol{\Sigma}_i)$

- $\Phi = \{P(G_i), \boldsymbol{\mu}_i, \boldsymbol{\Sigma}_i\}_{i=1}^k$

**No target Labels,  $y^j$**

(where the covariance matrix =  $\boldsymbol{\Sigma}_i$  )

# EM - Clusters

**Unsupervised:**  $X = \{ \mathbf{x}^j \}_j$

- Clusters  $G_i$   $i=1, \dots, k$

$$p(\mathbf{x}) = \sum_{i=1}^k p(\mathbf{x} | G_i) P(G_i)$$

where  $p(\mathbf{x} | G_i) \sim N(\boldsymbol{\mu}_i, \boldsymbol{\Sigma}_i)$

- $\Phi = \{P(G_i), \boldsymbol{\mu}_i, \boldsymbol{\Sigma}_i\}_{i=1}^k$

**No target Labels,  $y^j$**

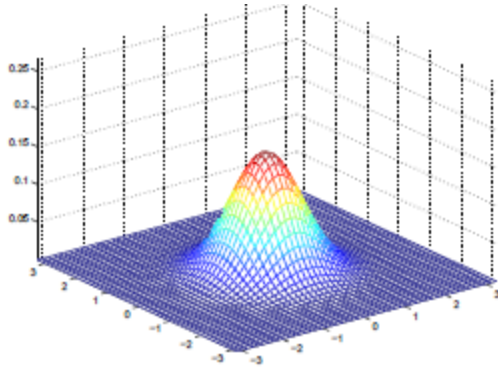
(where the covariance matrix =  $\boldsymbol{\Sigma}_i$  )

- $G_i$  are the mixture components – group or clusters
- $P(G_i)$  are the mixture proportions
- $k$  is the number of components
  - specified beforehand
- $p(\mathbf{x} | G_i)$  and  $\Phi$  are the parameters that should be estimated from the iid sample  $X = \{ \mathbf{x}^j \}_j$

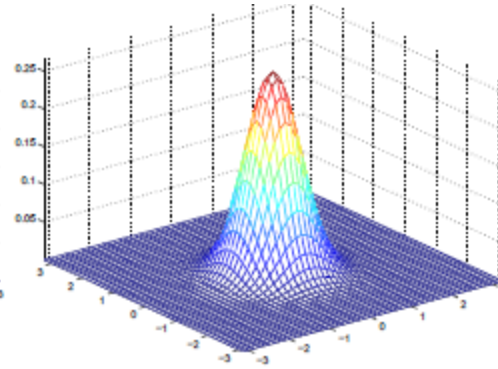
Goal: Find the component density parameters that maximize the likelihood of the sample. That is, find the parameter vector  $\Phi$  that maximizes the likelihood of the observed values of  $X = \{ \mathbf{x}^j \}_j$



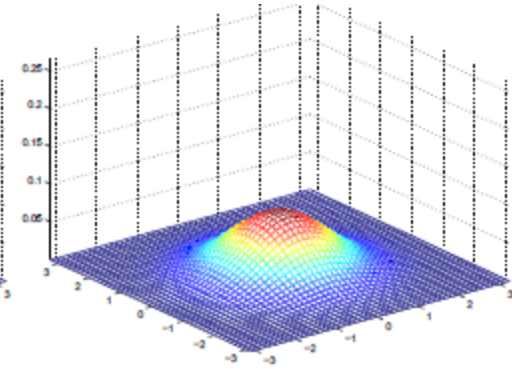
# Gaussian Distributions



$$\mu = 0; \Sigma = I$$



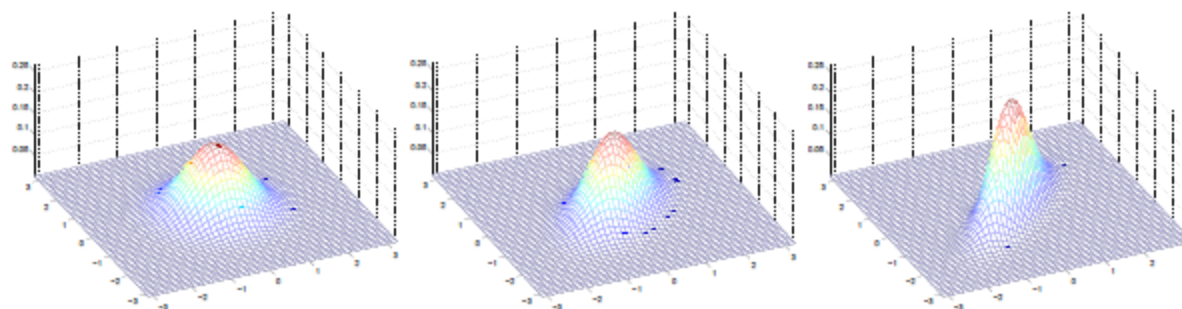
$$\mu = 0; \Sigma = (0.6)I$$



$$\mu = 0; \Sigma = 2I$$

( $I = 2 \times 2$  identity matrix)

# Gaussians with mean 0



$$\Sigma = \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix}; \quad \Sigma = \begin{bmatrix} 1 & 0.5 \\ 0.5 & 1 \end{bmatrix}; \quad \Sigma = \begin{bmatrix} 1 & 0.8 \\ 0.8 & 1 \end{bmatrix}$$

Covariance Matrices as indicated

# Covariance Matrix $\Sigma$

- Let  $X$  be a set of  $n$  random variables
  - $X = (X_1, X_2, \dots, X_n)^T$
- Covariance between  
 $X_i$  and  $X_j = E[(X_i - \mu_i)(X_j - \mu_j)]$
- The  $ij$ th entry of the covariance matrix  
 $\Sigma_{ij} = \text{cov}(X_i, X_j) = E[(X_i - \mu_i)(X_j - \mu_j)]$
- Note that  $\Sigma_{ii}$  is the variance of  $X_i$
- $\Sigma = E[(X - E[X])(X - E[X])^T]$

# Gaussian Mixture Model

- Each  $x_j$  is generated randomly
  - choose  $l \in \{1 \text{ to } k\}$  to select  $G_l$ , one of the Gaussians
  - Use the parameters of  $G_l$  to generate  $x_j$
- So, assuming your data was generated as a Gaussian Mixture Model, the Expectation – Maximization Algorithm can be used with Gaussians
- Note: other distributions can be used
  - Usually  $z$  is used to denote the hidden target (unsupervised)
    - $z_i = G_l$
  - For supervised methods, target was denoted by  $y$

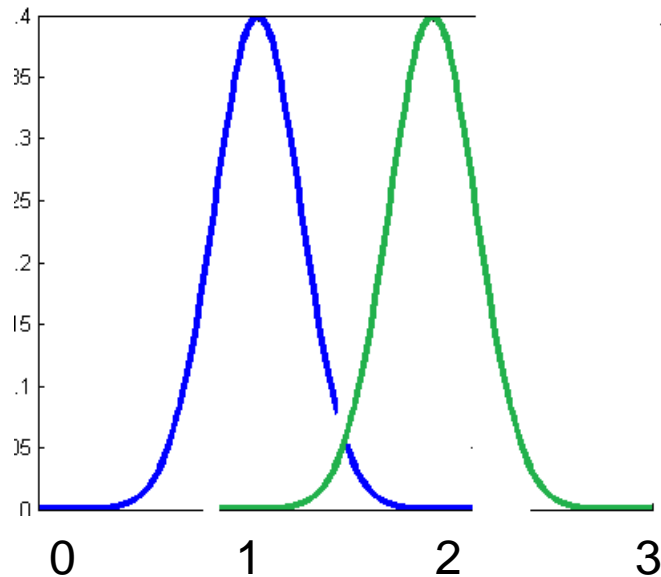
# Mixture of Mixtures

- In classification, the input comes from a mixture of classes (supervised).
- If each class is also a mixture, e.g., of Gaussians, (unsupervised), we have a mixture of mixtures:

$$p(\mathbf{x} | C_i) = \sum_{j=1}^{k_i} p(\mathbf{x} | G_{ij}) P(G_{ij})$$

$$p(\mathbf{x}) = \sum_{i=1}^K p(\mathbf{x} | C_i) P(C_i)$$

# Example – 2 Gaussian Distributions



- Classify Data less than 1.5 as Blue
- Classify Data greater than 1.5 as Green

$P(x = 0.5 \in \Phi_{\text{blue}}) > P(x = 0.5 \in \Phi_{\text{green}})$  so classify  $x = 0.5$  as blue

Assume there are two Classes, each with a Gaussian Distribution with known mean and variance. Classify a given observation by choosing the Gaussian distribution which maximizes the probability .

# Intuition for Gaussian Distribution

## (Where does Maximization Step Come From?)

**Expectation:** Given  $\mu$  and  $\sigma$  the probability distribution function for a point  $x$ :

$$\text{Probability Distribution Function} = \frac{1}{\sigma\sqrt{2\pi}} \exp\left[-\frac{(x-\mu)^2}{2\sigma^2}\right]$$

**Maximization:** Given a set of points  $x_j$ ,

$$\text{mean } \mu = \sum_{j=1}^k x_j$$

$$\text{standard deviation } \sigma = \left(\sum_{j=1}^k (x_j - \mu)^2\right)^{1/2}$$

# Maximum Likelihood Derivation of mean and std for Gaussian Distribution

- Maximize the Likelihood function:

$$l(\phi) = p(\mathbf{X}|\phi) = \prod_{j=1}^N p(x_j|\phi)$$

- Easier to Maximize the Log of the Likelihood function:

$$L(\phi|\mathbf{X}) = \log l(\phi|\mathbf{X}) = \sum_{j=1}^N \log p(x_j|\phi)$$

- For Gaussian Distributions:

$$L(\mu, \sigma|\mathbf{X}) = -\frac{N}{2} \log(2\pi) - N \log \sigma - \frac{\sum_j (x_j - \mu)^2}{2\sigma^2}$$



# Maximum Likelihood Estimation

- Suppose  $X = \{x_j\}_{j=1}^N$  where  $x_j$  are independent identically distributed (iid) samples from some known probability density family,  $p(x | \Phi)$  defined up to parameters,  $\Phi : x_j \sim p(x | \Phi)$
- Find  $\Phi$  that makes sampling  $x_j$  from  $p(x | \Phi)$  as likely as possible.
- The “likelihood” of the sample  $X$  given the parameter  $\Phi$  is the product of the likelihoods of the individual points:

$$l(\phi) = p(\mathbf{X}|\phi) = \prod_{j=1}^N p(x_j|\phi)$$

(note throughout  $\Phi$  is the same as  $\phi$ )

# Maximum Likelihood Derivation of mean for Gaussian Distribution

- Holding  $\mathbf{X}$  fixed, maximize the likelihood =  
$$\max_{\mu, \sigma} \left( L(\mu, \sigma | \mathbf{X}) = -\frac{N}{2} \log(2\pi) - N \log \sigma - \frac{\sum_j (x_j - \mu)^2}{2\sigma^2} \right)$$
- Set the Derivative with respect to  $\mu$  equal to 0 and solve for  $\mu$ :

$$\frac{\partial L}{\partial \mu} = 0 \Rightarrow \sum_j x_j = \sum_{j=1}^N \mu \Rightarrow \sum_j x_j = N\mu \Rightarrow$$
$$\mu = \frac{\sum_j x_j}{N}$$

Similar derivation for Standard Deviation

# Clustering Using Mixture Models

- Assume the data belongs to a set of specific distributions  $\{G_i\}$  for  $i=1$  to  $k$ 
  - Example: Gaussians
- Each distribution corresponds to a cluster
- The parameters of each distribution provide a description of the corresponding cluster
- Estimate the Parameters  $\Phi_i$  for each Distribution
  - Gaussian – mean & standard deviation ie  $\Phi_i = \{\mu_i, \sigma_i\}$
- Classify an object by putting it in the cluster for which it has the maximum probability

# Expectation Maximization Algorithm

**Initialize** the set of model parameters

For Gaussian Class  $G_i$ ,  $\Phi_i = \{\mu_i, \sigma_i\}$  for  $i = 1$  to  $K$

**Repeat:**

**Expectation Step:** For each  $x_j$  in the data set, calculate the probability that  $x_j$  belongs to Class  $G_i$ , i.e., For each  $i$  calculate:  $P(x_j \in \text{Class } G_i | \Phi_i)$

**Maximization Step:** Given the probabilities from the expectation step, find the new estimates of the parameters  $\Phi_i$  that maximize the expected likelihood for the data set.

**Until:** The change in parameters,  $\Phi_i$ , is below a set threshold.

# Expectation Maximization Details

**Expectation Step:**  $w_{ij} = p(x_j \in \text{Class } G_i | \Phi_i)$

$w_{ij}$  is the probability that observation  $x_j$  is in class  $G_i$

**Maximization Step:**

$$w_i := \frac{1}{m} \sum_{j=1}^N w_{ij}$$

$$\mu_i := \frac{\sum_{j=1}^N w_{ij} x_j}{\sum_{j=1}^N w_{ij}}$$

$$\text{the covariance matrix}(\sum_i) := \frac{\sum_{j=1}^N w_{ij} (x_j - \mu_i)(x_j - \mu_i)^T}{\sum_{j=1}^N w_{ij}}$$

# Bayesian Information Criterion (BIC)

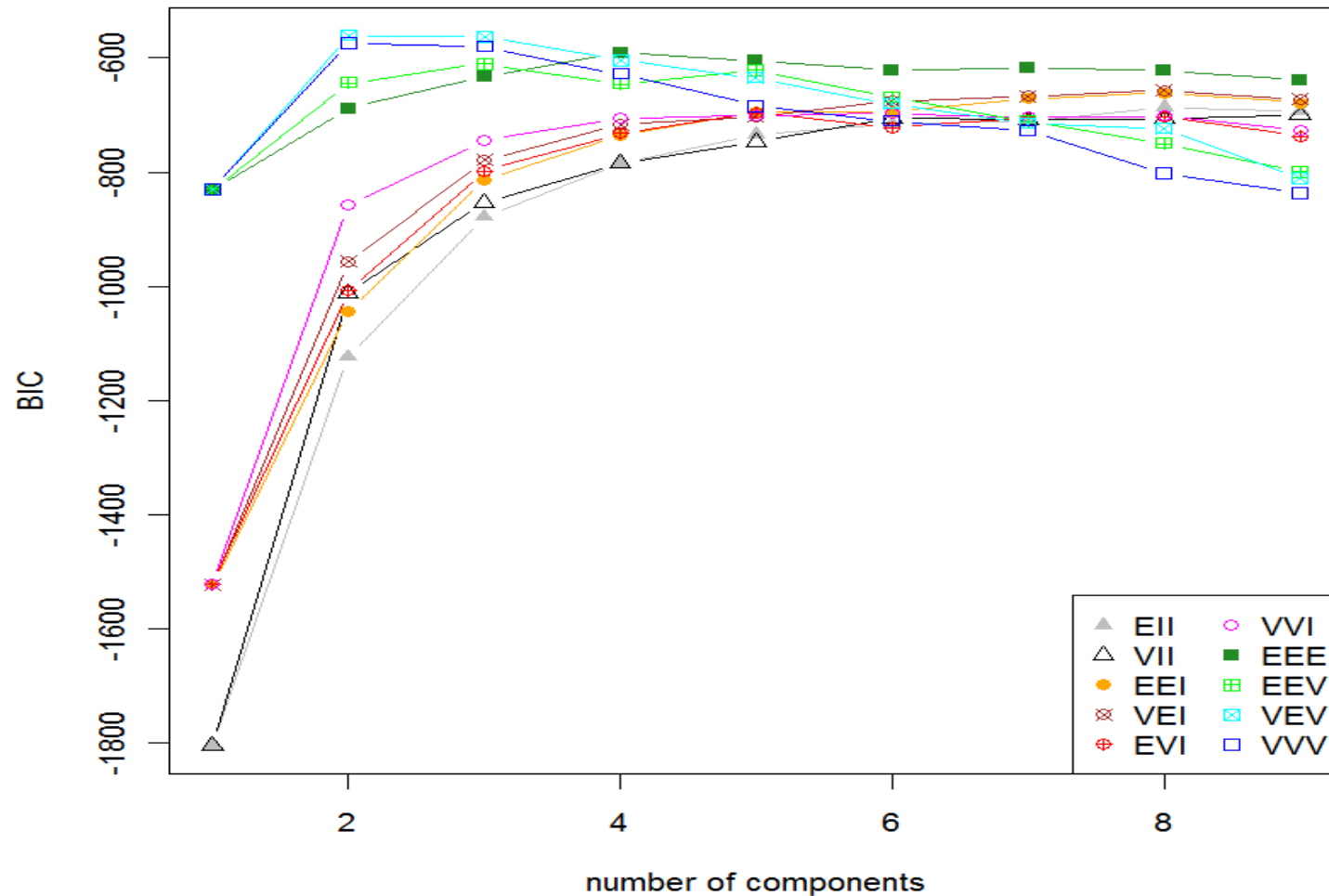
- BIC = maximized log of likelihood with penalty for number of parameters in model.
- BIC allows comparison of models with differing parameterizations and/or differing number of clusters.
- $BIC = -2\ln L + m\ln(N)$  where
  - $L$  = Maximized value of the likelihood
  - $m$  = number of free parameters to estimate
  - $N$  = the number of observations
- Lower BIC is better – requires fewer explanatory variables

# Iris Data – BIC chooses VEV

Parameterizations of the covariance matrix  $\Sigma$  currently available in MCLUST for hierarchical clustering (HC) and/or EM for multidimensional data.  
( • indicates availability).

identifier	Model	HC	EM	Distribution	Volume	Shape	Orientation
E		•	•	(univariate)	equal		
V		•	•	(univariate)	variable		
EII	$\lambda I$	•	•	Spherical	equal	equal	NA
VII	$\lambda_k I$	•	•	Spherical	variable	equal	NA
EEI	$\lambda A$		•	Diagonal	equal	equal	coordinate axes
VEI	$\lambda_k A$		•	Diagonal	variable	equal	coordinate axes
EVI	$\lambda A_k$		•	Diagonal	equal	variable	coordinate axes
VVI	$\lambda_k A_k$		•	Diagonal	variable	variable	coordinate axes
EEE	$\lambda D A D^T$	•	•	Ellipsoidal	equal	equal	equal
EEV	$\lambda D_k A D_k^T$		•	Ellipsoidal	equal	equal	variable
VEV	$\lambda_k D_k A D_k^T$		•	Ellipsoidal	variable	equal	variable
VVV	$\lambda_k D_k A_k D_k^T$	•	•	Ellipsoidal	variable	variable	variable

# BIC Plot for Iris Data (VEV)



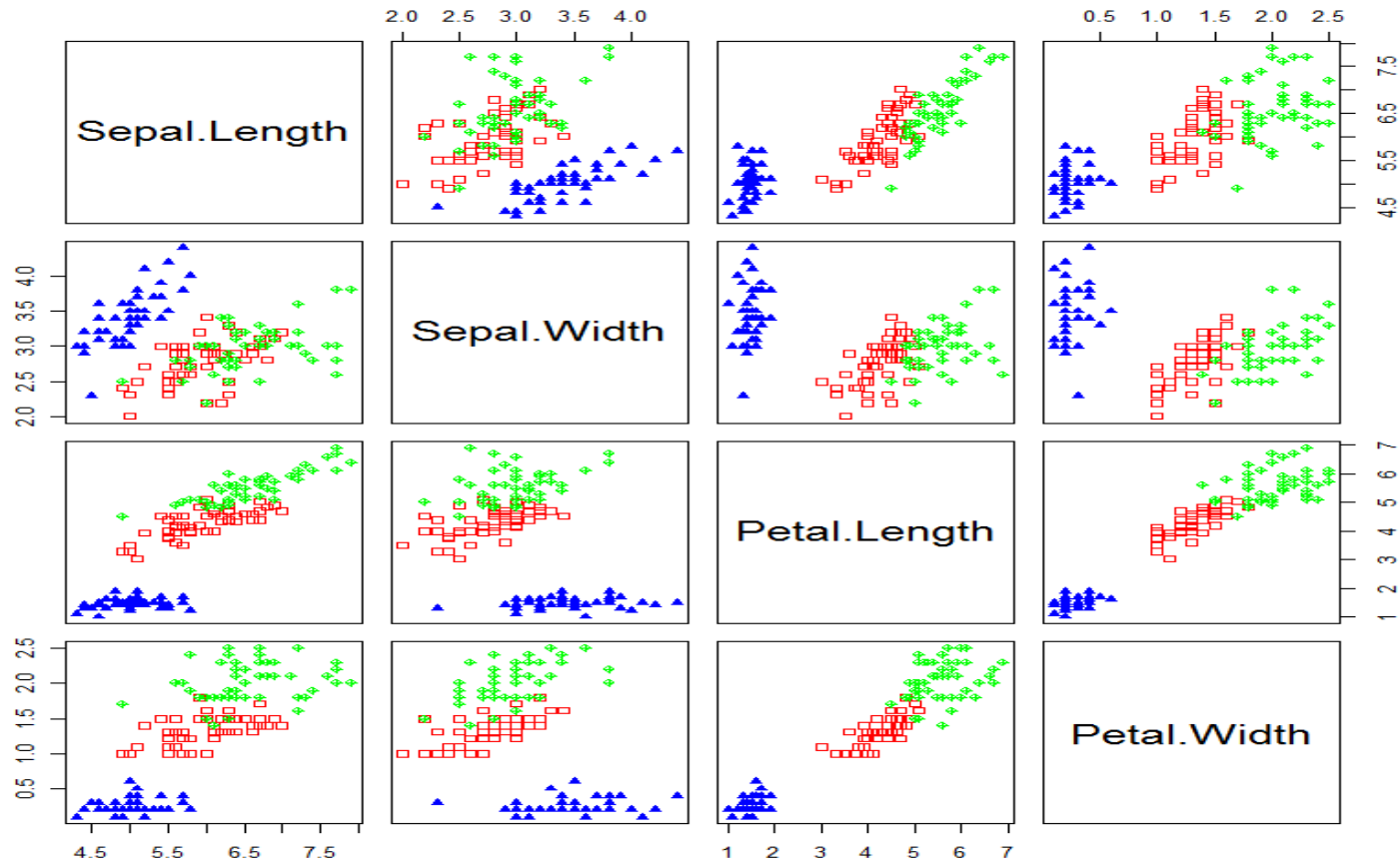


# Iris Data and BIC

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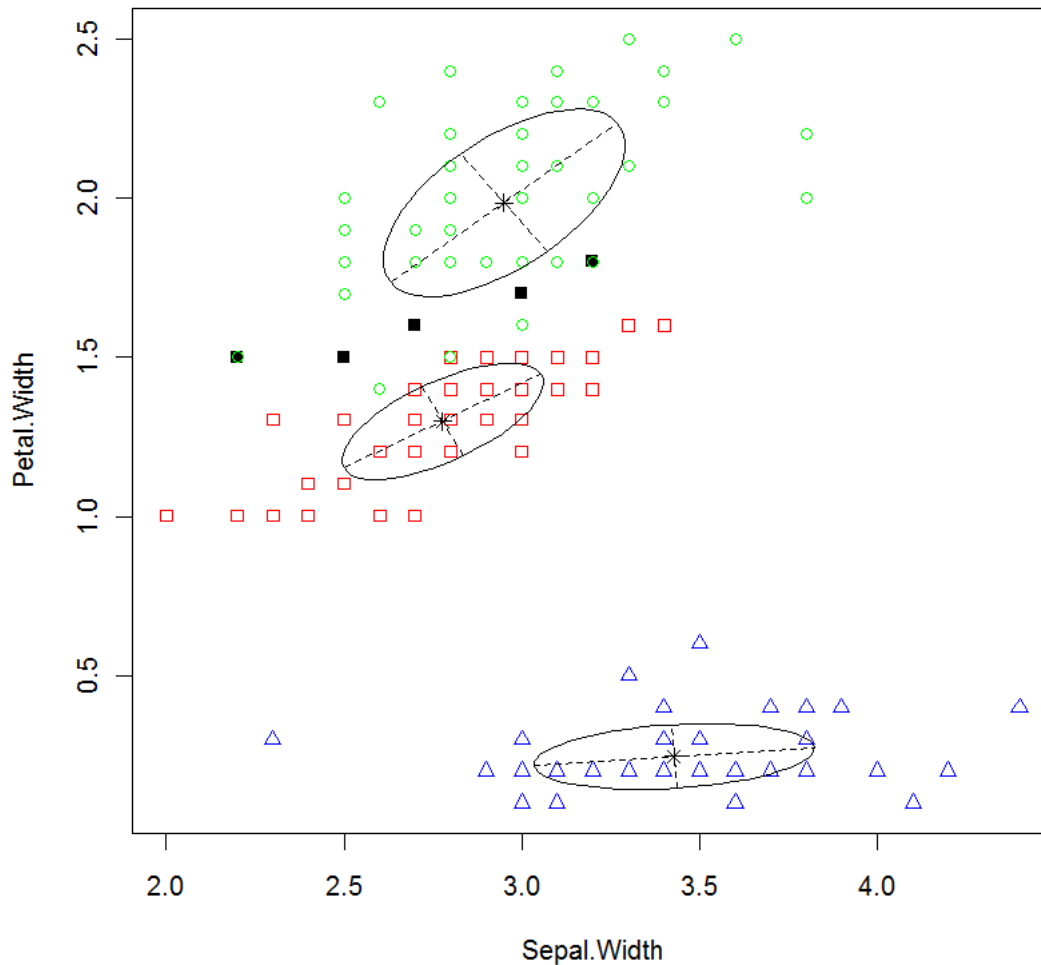
- Best Model for Iris Data is VEV using 2 clusters
  - `mclustBIC(iris[, -5])`
- However Iris Data has 3 classes
  - Must force 3 clusters
    - `mclustBIC(iris[, -5], G = 3)`

# Pairs Plot for Iris Data



# Iris Data

Notice that covariance are ellipsoid not oriented with axis



VEV – volume and orientation  
are variable

classification table:

1	2	3
50	45	55

# EM characteristics

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- Objects being classified must be elements of a finite dimensional vector-space over the real numbers
- Wide variety of shapes (long skinny clusters, clusters with holes punched in the middle, etc.)
- Less sensitive to large density variations than k-means
- Generative model
  - maximum likelihood parameters for a probabilistic model

# K- Means Characteristics

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- Works for any data for which you can define a distance or a similarity measure
- Roughly spherical (convex polyhedral) and roughly same volume

# After Clustering

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- Dimensionality reduction methods find correlations between features and group features
- Clustering methods find similarities between instances and group instances
- Allows knowledge extraction through
  - number of clusters,
  - prior probabilities,
  - cluster parameters, i.e., center, range of features.

Example: CRM, customer segmentation