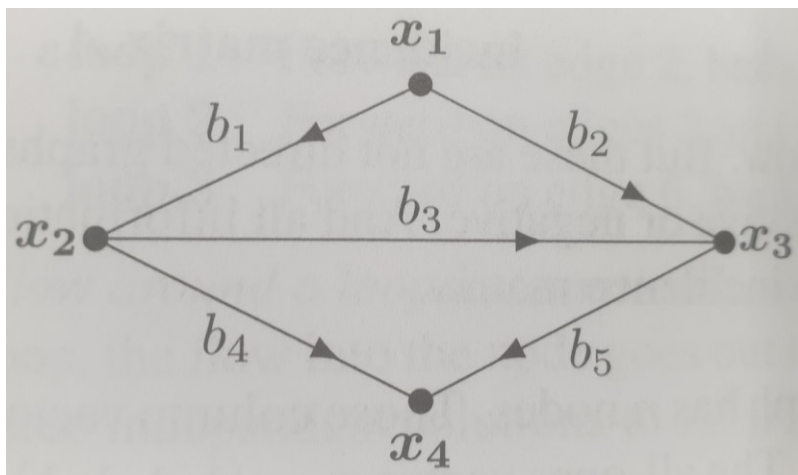


Graph (1)

- Consist of a set of nodes and a set of edges between those nodes
- Incidence matrix A ($m \times n$)
 - m edges and n nodes
 - Dimensions of the four subspaces
- Graph Laplacian matrix $A^T A$
 - Symmetric, positive semidefinite
 - Degree matrix D , adjacency matrix B

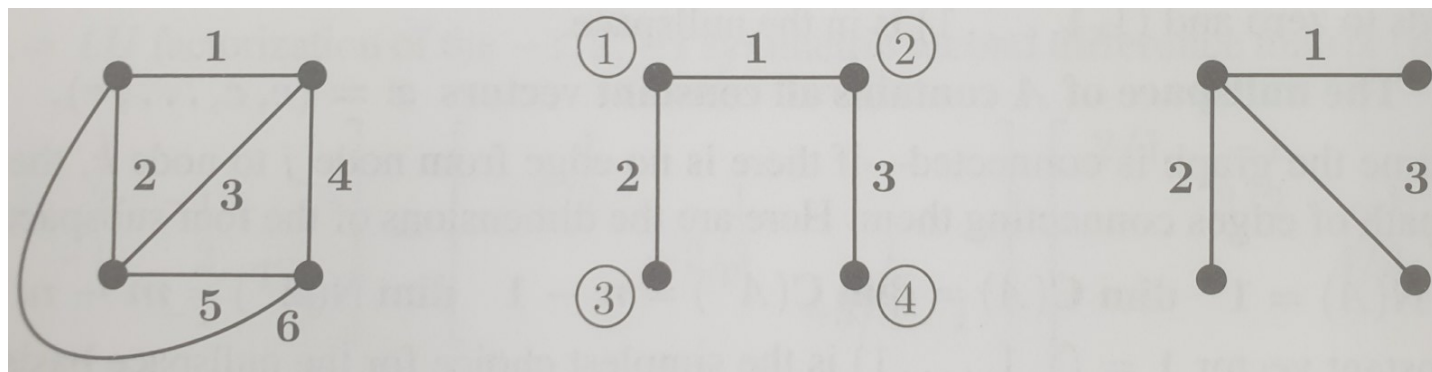
continuous	discrete
function	vector
derivative	difference
integral	sum
calculus	Linear algebra



$$\left\{ \begin{array}{l} N(\mathbf{A}): \text{constant vector } \mathbf{1} \\ C(\mathbf{A}^T): (n-1) \text{ rows of } \mathbf{A} \text{ that produce a tree in the graph (a tree has no loop)} \\ C(\mathbf{A}): (n-1) \text{ columns of } \mathbf{A} \\ N(\mathbf{A}^T): \text{flows around the } (m-n+1) \text{ small loops in the graph} \end{array} \right.$$

Graph (2)

- Complete graph: every pair of nodes is connected by an edge
- Tree: there are no loops in the connected graph



$$\mathbf{A}_1 = \begin{bmatrix} -1 & 1 & 0 & 0 \\ -1 & 0 & 1 & 0 \\ 0 & -1 & 1 & 0 \\ 0 & -1 & 0 & 1 \\ 0 & 0 & -1 & 1 \\ -1 & 0 & 0 & 1 \end{bmatrix}$$

complete graph: $m = \frac{1}{2}n(n-1)$

$$\mathbf{A}_2 = \begin{bmatrix} -1 & 1 & 0 & 0 \\ -1 & 0 & 1 & 0 \\ 0 & -1 & 0 & 1 \end{bmatrix}, \quad \mathbf{A}_3 = \begin{bmatrix} -1 & 1 & 0 & 0 \\ -1 & 0 & 1 & 0 \\ -1 & 0 & 0 & 1 \end{bmatrix}$$

tree: $(n-1)$

→ any graph: $(n-1) \leq m \leq \frac{1}{2}n(n-1)$

Kirchhoff's Current Law

- KCL = balance of currents (forces, money)
 - flow into each node equals flow out from that node
 - Key to solving $A^T y = 0$ is to look at the small loops in the graph
 - $(m-n+1)$ independent solutions
 - $(\text{number of nodes}) - (\text{number of edges}) + (\text{number of loops}) = 1$

$$\mathbf{A}_1 = \begin{bmatrix} -1 & 1 & 0 & 0 \\ -1 & 0 & 1 & 0 \\ 0 & -1 & 1 & 0 \\ 0 & -1 & 0 & 1 \\ 0 & 0 & -1 & 1 \\ -1 & 0 & 0 & 1 \end{bmatrix} \rightarrow \mathbf{A}_1 \mathbf{x} = 0, \mathbf{A}_1^T \mathbf{y} = \begin{bmatrix} -1 & -1 & 0 & 0 & 0 & -1 \\ 1 & 0 & -1 & -1 & 0 & 0 \\ 0 & 1 & 1 & 0 & -1 & 0 \\ 0 & 0 & 0 & 1 & 1 & 1 \end{bmatrix} \begin{bmatrix} y_1 \\ y_2 \\ y_3 \\ y_4 \\ y_5 \\ y_6 \end{bmatrix} = 0 \rightarrow \mathbf{y}_1 = \begin{bmatrix} -1 \\ 1 \\ -1 \\ 0 \\ 0 \\ 0 \end{bmatrix}, \mathbf{y}_2 = \begin{bmatrix} 0 \\ 0 \\ 1 \\ -1 \\ 1 \\ 0 \end{bmatrix}, \mathbf{y}_3 = \begin{bmatrix} 0 \\ -1 \\ 0 \\ 0 \\ -1 \\ 1 \end{bmatrix}$$

$$\mathbf{A}_2^T \mathbf{y} = \begin{bmatrix} -1 & -1 & 0 \\ 1 & 0 & -1 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} y_1 \\ y_2 \\ y_3 \end{bmatrix} = 0$$

A^TCA Framework in Applied Mathematics

- Graphs are perfect examples for three equations in engineering, science, economics
- Weighted graph Laplacian
- Describe a system in steady state equilibrium

{ voltages $\mathbf{x} = (x_1, x_2, x_3, x_4)$ at the four nodes

{ currents $\mathbf{y} = (y_1, y_2, y_3, y_4, y_5, y_6)$ at along the six edges

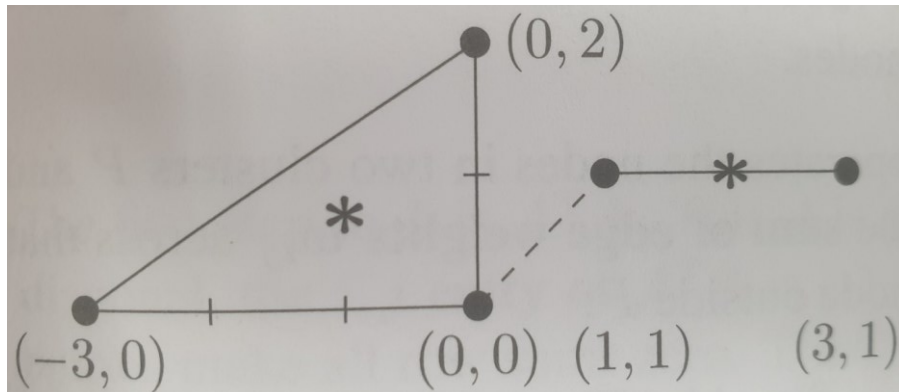
{ Voltage differences across edges $\mathbf{e} = \mathbf{A}\mathbf{x}$ $e_1 = (\text{voltage at end node 2}) - (\text{voltage at end node 1})$
 Ohm's law on each edge $\mathbf{y} = \mathbf{C}\mathbf{e}$ current $y_1 = (\text{conductance})(\text{voltage})$
 Kirchhoff's Law with current sources $\mathbf{f} = \mathbf{A}^T\mathbf{y}$ current sources \mathbf{f} into nodes balance the internal currents \mathbf{y}

$$\rightarrow \mathbf{A}^T\mathbf{C}\mathbf{A}\mathbf{x} = \mathbf{f} \rightarrow \mathbf{K}\mathbf{x} = \mathbf{f}$$

\mathbf{K} : symmetric, positive **semi**definite $\xrightarrow[x_4=0]{\text{boundary condition}}$ reduced \mathbf{K} : symmetric, positive definite

Example with Two Clusters

- How to understand a graph with many nodes?
 - Separate nodes into two or more clusters
 - Human Genome project: cluster genes that show highly correlated
- Break a graph in two pieces
 - For load balancing in high computing, assign equal work to two processors
 - For social networks, identify two distinct groups
 - Segment an image
 - Reorder rows and columns of a matrix to make off-diagonal blocks sparse



$$\mathbf{A} = \begin{bmatrix} 0 & 1 & 3 & 0 & -3 \\ 0 & 1 & 1 & 2 & 0 \end{bmatrix} \approx \begin{bmatrix} -1 & 2 & 2 & -1 & -1 \\ 2/3 & 1 & 1 & 2/3 & 2/3 \end{bmatrix}$$

Approximate an $m \times n$ matrix of \mathbf{A} by $\mathbf{CR} = (m \times k)(k \times n)$

$$\mathbf{A} \approx \mathbf{CR} = \begin{bmatrix} -1 & 2 \\ 2/3 & 1 \end{bmatrix} \begin{bmatrix} 1 & 0 & 0 & 1 & 1 \\ 0 & 1 & 1 & 0 & 0 \end{bmatrix}$$

Four Methods for Clustering

- Find the Fiedler vector z that solves $A^T C A z = \lambda D z$. D normalizes the Laplacian. Positive and negative components of eigenvector of λ_2 indicate two clusters of nodes.
- Replace the graph Laplacian by the modularity matrix $M = (\text{adjacency matrix}) - dd^T/2m$. Choose the eigenvector that comes with the largest eigenvalue of M . vector d gives the degrees of the n nodes.
- Find the minimum normalized cut that separates the nodes in two clusters P and Q . The unnormalized measure of a cut is the sum of edge weights w_{ij} across that cut. Those edges connect a node in P to a node outside P .
- K-means

Spectral Clustering

$$\mathbf{A}^T \mathbf{C} \mathbf{A} \xrightarrow{\text{normalized}} \mathbf{L} = \mathbf{D}^{-1/2} \mathbf{A}^T \mathbf{C} \mathbf{A} \mathbf{D}^{-1/2} = \mathbf{I} - \mathbf{N} \quad \text{where } n_{ij} = \frac{w_{ij}}{\sqrt{d_i d_j}} \text{ (normalized weights)}$$

$\mathbf{L} = \mathbf{I} - \mathbf{N}$ is like a correlation matrix in statistics

\mathbf{L} is symmetric positive semidefinite

The eigenvectors for $\lambda = 0$ is $\mathbf{u} = (\sqrt{d_1}, \dots, \sqrt{d_n})$. Then $\mathbf{L}\mathbf{u} = \mathbf{D}^{-1/2} \mathbf{A}^T \mathbf{C} \mathbf{A} \mathbf{1} = \mathbf{0}$.

The second eigenvector \mathbf{v} of \mathbf{L} minimizes the Rayleigh quotient on a subspace.

$$\left(\lambda_2 = \text{smallest nonzero eigenvalue of } \mathbf{L} \rightarrow \min_{\substack{\text{subject to} \\ \mathbf{x}^T \mathbf{u} = 0}} \frac{\mathbf{x}^T \mathbf{L} \mathbf{x}}{\mathbf{x}^T \mathbf{x}} = \frac{\mathbf{v}^T \mathbf{L} \mathbf{v}}{\mathbf{v}^T \mathbf{v}} = \lambda_2 \text{ at } \mathbf{x} = \mathbf{v} \right)$$

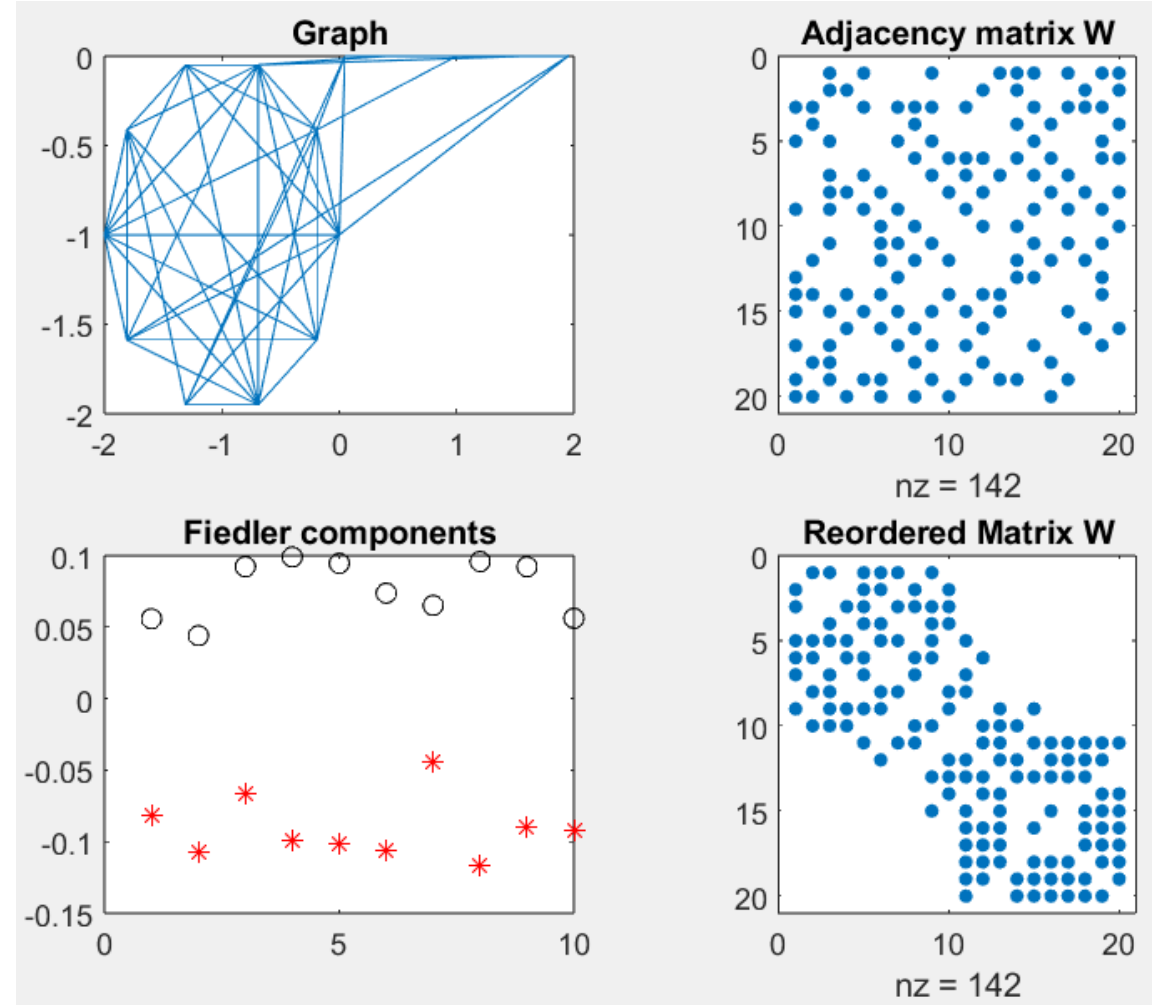
$$\mathbf{L}\mathbf{v} = \mathbf{D}^{-1/2} \mathbf{A}^T \mathbf{C} \mathbf{A} \mathbf{D}^{-1/2} \mathbf{v} = \lambda \mathbf{v} \xrightarrow[\text{normalized Fiedler vector}]{\mathbf{z} = \mathbf{D}^{-1/2} \mathbf{v}} \mathbf{A}^T \mathbf{C} \mathbf{A} \mathbf{z} = \lambda \mathbf{D} \mathbf{z} \quad \text{with } \mathbf{1}^T \mathbf{D} \mathbf{z} = 0$$

$$\min_{\substack{\text{subject to} \\ \mathbf{x}^T \mathbf{u} = 0}} \frac{\mathbf{x}^T \mathbf{L} \mathbf{x}}{\mathbf{x}^T \mathbf{x}} \xrightarrow{\mathbf{x} = \mathbf{D}^{1/2} \mathbf{y}} \min_{\substack{\text{subject to} \\ \mathbf{1}^T \mathbf{D} \mathbf{y} = 0}} \frac{\mathbf{y}^T \mathbf{A}^T \mathbf{C} \mathbf{A} \mathbf{y}}{\mathbf{y}^T \mathbf{D} \mathbf{y}} = \frac{\sum \sum w_{ij} (y_i - y_j)^2}{\sum d_i y_i^2} = \lambda_2 \text{ at } \mathbf{y} = \mathbf{z}$$

Code: MATLAB

```
N=10; W=zeros(2*N,2*N); % Generate 2N nodes in two clusters
rand('state',100) % rand repeats to give the same graph
for i=1:2*N-1
for j=i+1:2*N
p=0.7-0.6*mod(j-i,2); % p=0.1 when j-i is odd, 0.7 else
W(i,j)=rand<p; % Insert edges with probability p
end % The weights are wi,j=1 (or 0)
end % So far W is strictly upper triangular
W=W+W'; D=diag(sum(W)); % Adjacency matrix W, degrees in D
G=D-W; [V,E]=eig(G,D); % Eigenvalues of Gx=(lambda)Dx in E
[a,b]=sort(diag(E)); z=V(:,b(2)); % Fiedler eigenvector z for (lambda)2
plot(sort(z),'-'); % Show +- groups of Fiedler components
```

```
theta=[1:N]*2*pi/N; x=zeros(2*N,1); y=x; % Angles to plot graph
x(1:2:2*N-1)=cos(theta)-1; x(2:2:2*N)=cos(theta)+1;
y(1:2:2*N-1)=sin(theta)-1; x(2:2:2*N)=sin(theta)+1;
print theta,x,y
subplot(2,2,1), gplot(W,[x,y]), title('Graph')
subplot(2,2,2), spy(W), title('Adjacency matrix W')
subplot(2,2,3), plot(z(1:2:2*N-1),'ko'), hold on
plot(z(2:2:2*N),'r*'), hold off, title('Fiedler components')
[c,d]=sort(z); subplot(2,2,4), spy(W(d,d)), title('Reordered Matrix W')
```



Minimum Cut

(edge) weight across cut: $links(P) = \sum w_{ij}$ for i in P and j not in P

size of cluster: $size(P) = \sum w_{ij}$ for i in P

normalized cut weight: $Ncut(P, Q) = \frac{links(P)}{size(P)} + \frac{links(Q)}{size(Q)}$

normalized K -cut: $Ncut(P_1, \dots, P_k) = \sum_{i=1}^K \frac{links(P_i)}{size(P_i)}$

[cuts connected to eigenvectors]

perfect indicator of a cut: vector \mathbf{y} with all components equal to p or $-q$ (two values only) \rightarrow node i goes $\begin{cases} \text{in } P \text{ if } y_i = p \\ \text{in } Q \text{ if } y_i = -q \end{cases}$

$\mathbf{1}^T \mathbf{D} \mathbf{y}$ will multiply one group of d_i by p and the other group by $-q$.
 The first d_i add to $size(P) = \text{sum of } d_i$ (i in P).
 The second group of d_i add to $size(Q)$

$$\left. \begin{array}{l} \mathbf{1}^T \mathbf{D} \mathbf{y} = 0 \rightarrow psize(P) = qsize(Q) \end{array} \right\}$$

$$\frac{\mathbf{y}^T \mathbf{A}^T \mathbf{C} \mathbf{A} \mathbf{y}}{\mathbf{y}^T \mathbf{D} \mathbf{y}} = \frac{\sum \sum w_{ij} (y_i - y_j)^2}{\sum d_i y_i^2} = \frac{(p+q)^2 links(P, Q)}{p^2 size(P) + q^2 size(Q)} = \frac{(p+q) links(P, Q)}{psize(P)} = \frac{links(P, Q)}{size(P)} + \frac{links(P, Q)}{size(Q)} = Ncut(P, Q)$$

Clustering by k-means

n points $\mathbf{a}_1, \dots, \mathbf{a}_n$ in d -dimensional space \rightarrow partition those points into k clusters

clusters P_1, \dots, P_k have centroids $\mathbf{c}_1, \dots, \mathbf{c}_k$

$\mathbf{c}_j = \frac{\text{sum of } \mathbf{a}'\text{s}}{\text{number of } \mathbf{a}'\text{s}} \rightarrow \text{minimize } \sum \|\mathbf{c} - \mathbf{a}\|^2 \text{ for all } \mathbf{a}'\text{s in cluster } P_j$

clustering: minimize $D = \sum_{j=1}^k D_j = \sum_{j=1}^k \|\mathbf{c}_j - \mathbf{a}_i\|^2$ for \mathbf{a}_i in cluster P_j

step 1: find the **centroids** \mathbf{c}_j of the (old) clustering P_1, \dots, P_k .

step 2: find the **(new) clustering** that puts \mathbf{a} in P_j if \mathbf{c}_j is the closest centroid.

Weights and Kernel Method

weights in the distance: $d(\mathbf{x}, \mathbf{a}_i) = w_i \|\mathbf{x} - \mathbf{a}_i\|^2$, $\mathbf{c}_j = \frac{\sum w_i \mathbf{a}_i}{\sum w_i}$ (\mathbf{a}_i in P_j)

$$\|\mathbf{c}_j - \mathbf{a}_i\|^2 = \mathbf{c}_j \cdot \mathbf{c}_j - 2\mathbf{c}_j \cdot \mathbf{a}_i + \mathbf{a}_i \cdot \mathbf{a}_i$$

Kernel method: weighted kernel matrix \mathbf{K} has entries $\mathbf{a}_i \cdot \mathbf{a}_l$

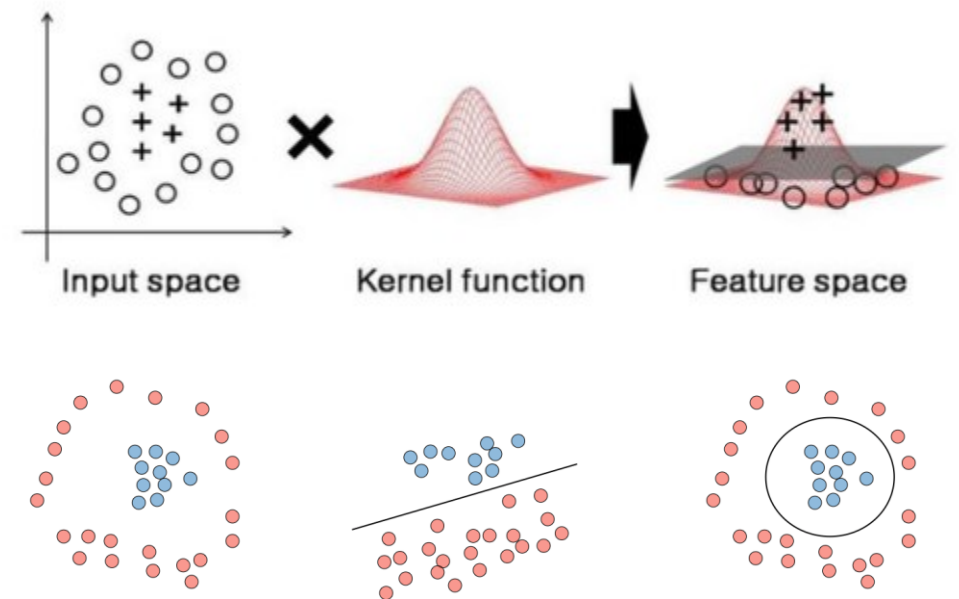
nodes are point \mathbf{x}_i in input space $\rightarrow \mathbf{a}_i = \phi(\mathbf{x}_i)$ points in a high-dimensional **feature space**

$$\sum \|\mathbf{c}_j - \mathbf{a}_i\|^2 = \frac{\sum w_i w_l \mathbf{K}_{il}}{(\sum w_i)^2} - 2 \frac{\sum w_i \mathbf{K}_{il}}{\sum w_i} + \sum \mathbf{K}_{ii}$$

(vision) polynomial $\mathbf{K}_{il} = (\mathbf{x}_i \cdot \mathbf{x}_l + c)^d$

(statistics) Gaussian $\mathbf{K}_{il} = \exp\left(-\frac{\|\mathbf{x}_i - \mathbf{x}_l\|^2}{2\sigma^2}\right)$

(neural networks) Sigmoid $\mathbf{K}_{il} = \tanh(c\mathbf{x}_i \cdot \mathbf{x}_l + \theta)$



Non-linear separability \rightarrow Use of a kernel mapping ϕ \rightarrow Decision boundary in the original space

Applications of Clustering

- Learning theory, training sets, neural networks, Hidden Markov Models
- Classification, regression, pattern recognition, Support Vector Machines
- Statistical learning, maximum likelihood, Bayesian statistics, spatial statistics, kriging, time series, ARMA models, stationary processes
- Social networks, organization theory
- Data mining, document indexing, image retrieval, kernel-based learning
- Bioinformatics, microarray data, systems biology
- Cheminformatics, drug design, decision trees
- Information theory, vector quantization, rate distortion theory
- Image segmentation, computer vision, texture, min cut
- Predictive control, feedback samples, robotics, adaptive control