Graph Kernels and Support Vector Machines for Pattern Recognition

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Summary

- Introduction
- 2 Methodology
- 3 Experiments
- 4 Conclusion
 - References

Motivation

- A lot of data can be represented as graphs such as proteins or social networks
- Being able to compare them would be useful (classification, clustering)



Figure 1: A fragment of a protein transformed into a graph[1]

Current Methods

Support Vector Machine

SVMs are models used in classification introduced about 25 years ago. They have several advantages

- Great accuracy
- Great capacity of generalization
- Allows the use of kernels in its dual form

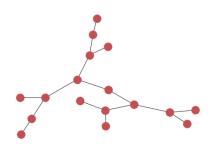
Kernels

The kernel trick can replace the dot product while implicitly projecting data to a feature space and combine very well with the SVMs

- Computes data projection faster implicitly (ex. RBF kernel)
- Improve the accuracy of SVM by making linear separation easier

Objective

- These methods are adapted to vector data
- Graphs and their adjacency matrices aren't and vectorizing implies a loss of information
- New types of kernels were discovered
- However, the complexity is a big problem, these kernels have to be accelerated



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Background: graphs

Definition

A graph[2] is a type of mathematical structure that represents connections between objects. It is more precisely an ordered pair G=(V,E) of two sets: vertices V (or nodes) and edges E that connect two vertices together.

$$E \subseteq \{(u,v) : (u,v) \in V^2\}$$

Properties

- Undirected
- Labeled or not
- Degree
- Path and Cycle

- Connected
- Tree
- Subgraph
- Dual Graph

Background: support vector machines

SVM

Background: kernels

Definition

In its dual form, the SVM problem only requires a dot product between the observations' vectors.

$$\max \sum_{i=1}^{n} \alpha_i - \frac{1}{2}$$

This means the vectors can be mapped to higher dimensions with a function ϕ . Moreover, even the dot product itself can be replaced by a function κ without explicitly specifying the map ϕ as long as the function is positive semi definite.

$$\kappa(\mathbf{x_i}, \mathbf{x_j}) = e^{-\frac{\|\mathbf{x_i} - \mathbf{x_j}\|^2}{2\sigma^2}}$$

An example of kernel: the RBF kernel

Graph Kernels

Random Walks

Random Walks

A random walk is a path obtained from a chosen vertex, randomly picking an edge to follow iteratively, until the trail reaches a certain length. Comparing common random walks between graphs is an acceptable metric.

Product graph

It has been shown[3] that computing random walks on two separate graphs is equivalent to computing it on the product graph of the two. The product graph is computed using the Kronecker (or tensor) product \otimes





Figure 3: A graph G_1 , a graph G_2 and the product graph $G_1 \otimes G_2$

Random Walk

Definition

kernel def

Acceleration methods

Cas particulier

Inverse Kernel

inv ker

On va vouloir accelerer machin ce kernel va être l'objectif de plusieurs méthodes

Sylvester Equation

Conjugate Gradient

Spectral Decomposition

Nearest Kronecker Product Approximation

- The idea is to approximate two matrices S and T such that $\|W_{\times} A \otimes B\|_F$ is minimized.
- Labeled-graph kernel computation can be turned into into an unlabeled one with some loss in accuracy, but gain in computation time.
- Computed in $O(dn^2)$ time
- All methods such as Spectral Decomposition can then be applied

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Frame

Conclusion

conclu

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