

# Algorithm graph theory: a practical approach

## Master de Bioinformática (UV)

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Bioinformatic at the Computational Genomics Institute (CIPF)

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- Library for complex network research
- R & python implementations
- Core implementation: C
- <http://igraph.sourceforge.net>
- Python API:  
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- Debian:
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- Open Suse
  - `sudo yum install python-igraph`
- easy\_install or pip
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  - Edge
  - Weight
  - DAG: Directed Acyclic Graph

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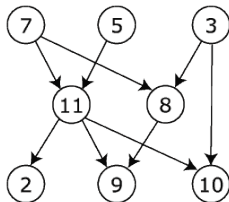


Figura: DAG

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The algorithm uses a queue data structure to store intermediate results as it traverses the graph, as follows:

- 1 Create empty array “path” and a queue
- 2 Enqueue the root node
- 3 Dequeue a node and examine it
  - Add node to path
  - Enqueue any successors (the direct child nodes) that have not yet been discovered.
- 4 If the queue is empty, every node on the graph has been examined.
- 5 If the queue is not empty, repeat from Step 3.

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## Example

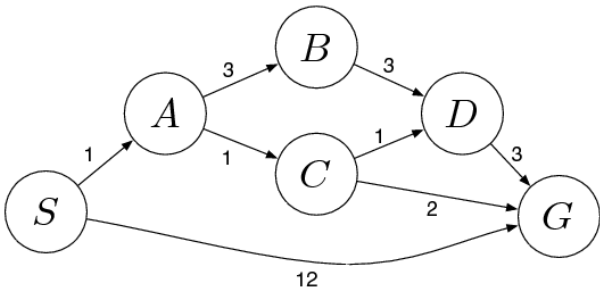


Figura: BFS example

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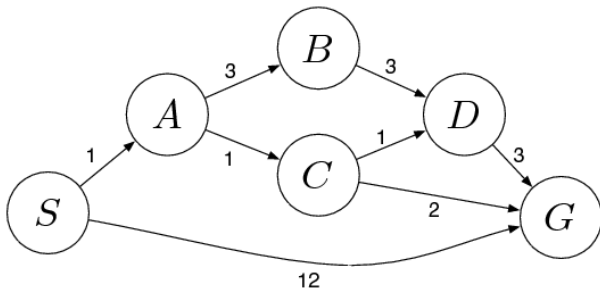


Figura: DFS example

## Is there any path?

A path in a graph is a sequence of vertices such that from each of its vertices there is an edge to the next vertex in the sequence

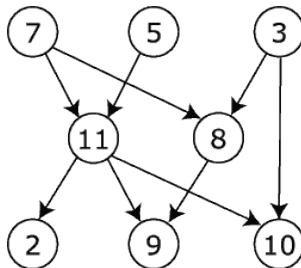


Figura: Is there any path? Example

# Dijkstra

Dijkstra algorithm is a graph search algorithm that solves the single-source shortest path problem for a graph with non-negative edge path costs, producing a shortest path tree.

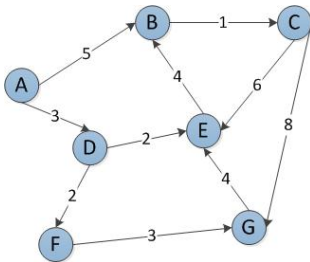


Figura: Dijkstra



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# Thank you