Eulerian and Hamiltonian Cycles

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Toy Genome Assembly Problem

Find a string whose all substrings of length 3 are

AGC, ATC, CAG, CAT, CCA, GCA, TCA, TCC.

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How is it related to cycles in graphs?..

Outline

Eulerian Cycles

Hamiltonian Cycles

Genome Assembly

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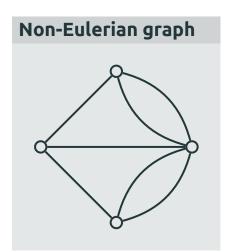
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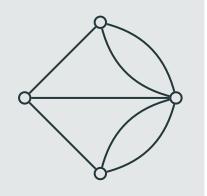
- The definition works for both directed and undirected graphs
- A cycle must have the same starting and ending nodes
- While in a path the starting and ending node should not necessarily be equal

Example

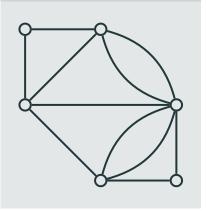


Example

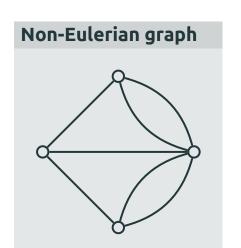
Non-Eulerian graph

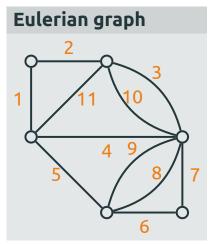


Eulerian graph



Example





Criteria

Theorem

A connected *undirected* graph contains an Eulerian cycle, if and only if the degree of every node is even.

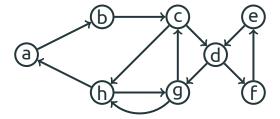
Criteria

Theorem

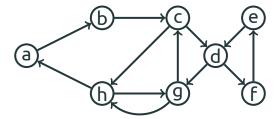
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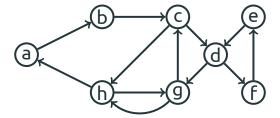
A strongly connected *directed* graph contains an Eulerian cycle, if and only if, for every node, its in-degree is equal to its out-degree.



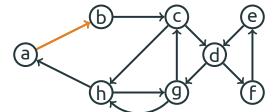
if some node is imbalanced, there is clearly no Eulerian cycle

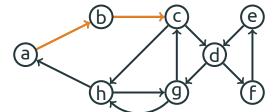


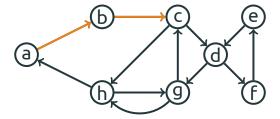
thus, assume that the graph is balanced



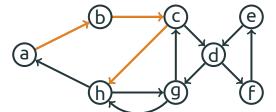
start walking from some node

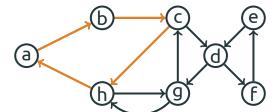


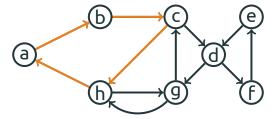




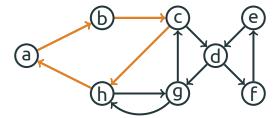
since the graph is balanced, at some point we'll return back to the starting node



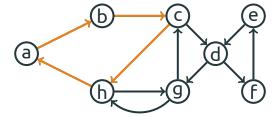




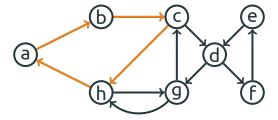
OK, what's next? we haven't traversed all the edges and now we are stuck



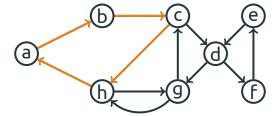
since the graph is strongly connected, there must be untraversed edges going out of a node from the current cycle



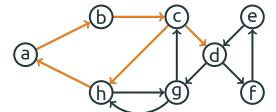
the current cycle has neither beginning nor end

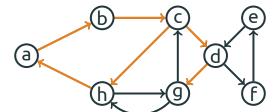


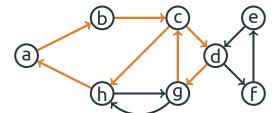
so, we may assume that its starting and ending point is the node c

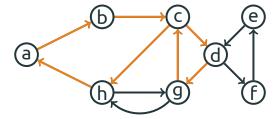


we then continue exploring the graph from the node c

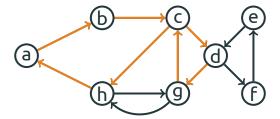




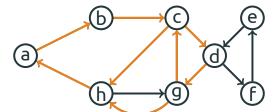


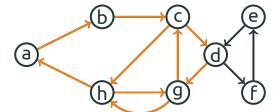


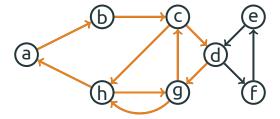
we got larger cycle



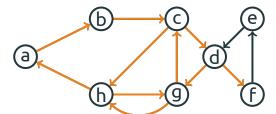
continue from g

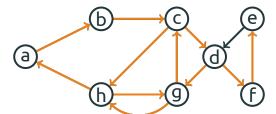




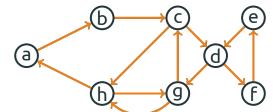


continue from d



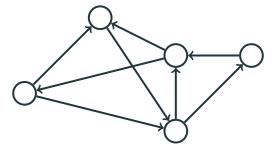


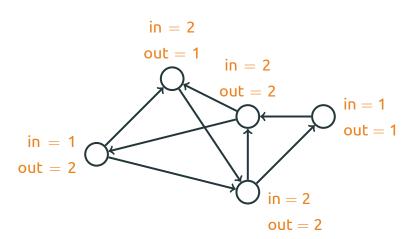
Proof (Directed Case)

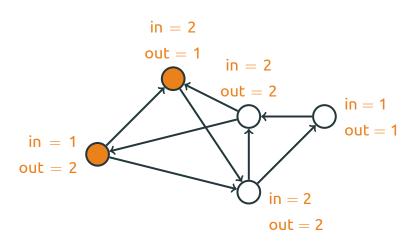


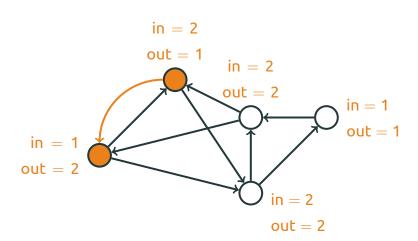
Path Instead of Cycle

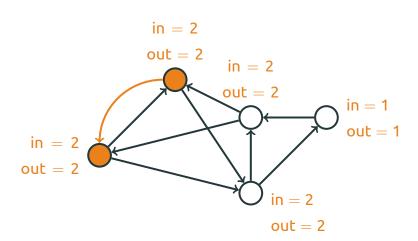
- The criteria for a path is similar
- A graph is allowed to contain two imbalanced nodes: one for the starting point and one for the ending point of a path
- By adding an edge between these two nodes, one gets a graph with an Eulerian cycle











Efficient Algorithms

The proof of existence of an Eulerian cycle can be transformed into an efficient algorithm for constructing an Eulerian cycle

Outline

Eulerian Cycles

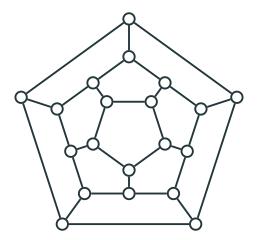
Hamiltonian Cycles

Genome Assembly

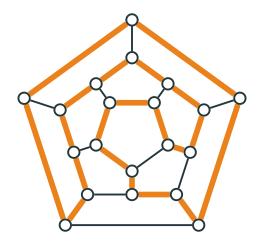
Hamiltonian Cycle

A Hamiltonian cycle visits every node of a graph exactly once.

Example



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Simple Criteria?

 No simple criteria is known for the Hamiltonian cycle problem

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- No polynomial time algorithm known

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- No simple criteria is known for the Hamiltonian cycle problem
- · No polynomial time algorithm known
- The question whether there is a polynomial time algorithm for the Hamiltonian cycle problem is the P versus NP problem, the most important open problem in Computer Science, with a prize of \$1M from the Clay Mathematics Institute (http://www.claymath.org/ millennium-problems)

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All Substrings of Length 3

```
DISCRETE
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ISC
SCR
CRE
RET
ETE
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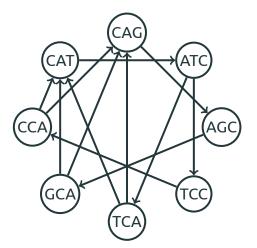
Every two neighbor 3-substrings have a common part, called an overlap, of length 2

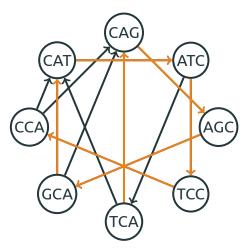
Finding a Permutation

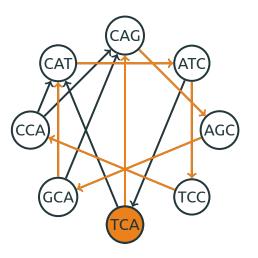
 Goal: Find a string whose all substrings of length 3 are AGC, ATC, CAG, CAT, CCA, GCA, TCA, TCC

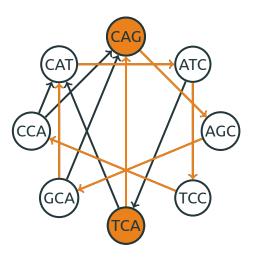
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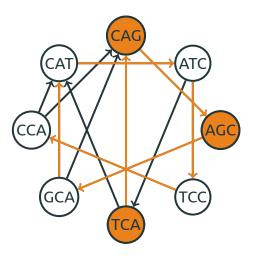
- Goal: Find a string whose all substrings of length 3 are AGC, ATC, CAG, CAT, CCA, GCA, TCA, TCC
- Hence, we need to find an order of these
 3-strings such that the overlap between
 any two consecutive strings is equal to 2



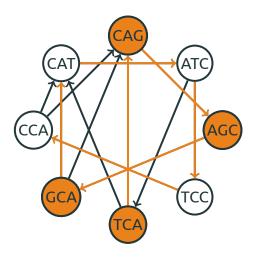




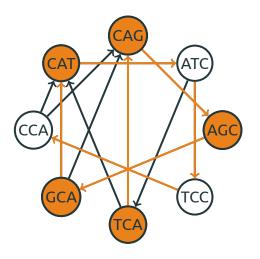




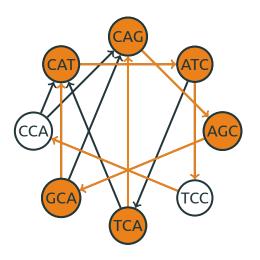
TCAGC



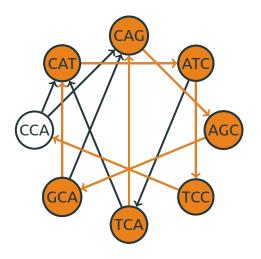
TCAGCA



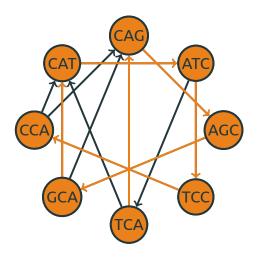
TCAGCAT



TCAGCATC



TCAGCATCC



TCAGCATCCA

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- But we don't have efficient algorithms for the Hamiltonian cycle problem!
- The approach is useless for the case when there are thousands or millions of input strings

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- Let's instead try to represent each string by an edge (De Bruijn; Pevzner, Tang, Waterman)

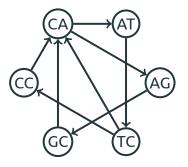
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- E.g., represent the string CAT as an edge $CA \rightarrow AT$

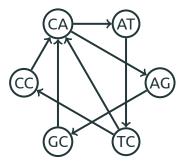
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- Used in state-of-the-art genome assemblers

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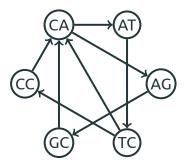


AGC, ATC, CAG, CAT, CCA, GCA, TCA, TCC



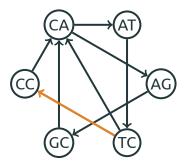
now, we need to find an order of edges

AGC, ATC, CAG, CAT, CCA, GCA, TCA, TCC



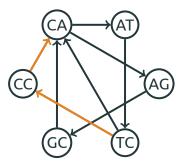
that is, an Eulerian cycle

AGC, ATC, CAG, CAT, CCA, GCA, TCA, TCC



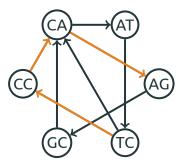
TCC

AGC, ATC, CAG, CAT, CCA, GCA, TCA, TCC



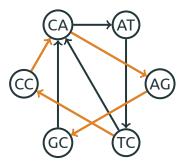
TCCA

AGC, ATC, CAG, CAT, CCA, GCA, TCA, TCC



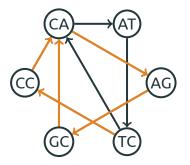
TCCAG

AGC, ATC, CAG, CAT, CCA, GCA, TCA, TCC



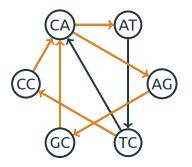
TCCAGC

AGC, ATC, CAG, CAT, CCA, GCA, TCA, TCC



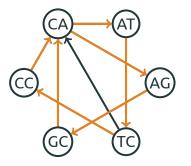
TCCAGCA

AGC, ATC, CAG, CAT, CCA, GCA, TCA, TCC



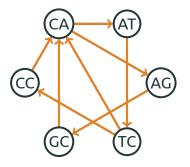
TCCAGCAT

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TCCAGCATC

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TCCAGCATCA

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- Look similar to each other, but differ drastically from the computational point of view

- Eulerian cycle visits every edge exactly once
- Hamiltonian cycle visits every node exactly once
- Look similar to each other, but differ drastically from the computational point of view
- Genome assembly: the right problem formulation makes all the difference!