Finding Eulerian paths/tours

Problem solved?

• Number of possible Eulerian/Chinese Postman tours in a graph

Generally an exponential number of compatible sequences

- Value computed by application of the BEST theorem (Hutchinson, ¹⁹⁷ $\mathcal{W}(G,t) = (\det L) \left\{ \prod_{u \in V} (r_u - 1)! \right\} \left\{ \prod_{(u,v) \in E} a_{uv}! \right\}^{-1}$

L = n x n matrix with r_u - a_{uu} along the diagonal and $-a_{uv}$ in entry uv

 $r_u = d^+(u) + 1$ if u=t, or $d^+(u)$ otherwise

 a_{uv} = multiplicity of edge from u to v

Introduction to genome assembly

There are no shortcuts in assembly



<u>Theorem:</u> Must try all possible assemblies before finding the correct one Peltola et al. 1970s Myers et al. 1990s Medvedev et al. 2000s Nagarajan et al. 2000s

Two approaches

- Greedy
 - Keep visiting non-visited edges and fix mistakes later
- Using a guide-tree
 - Build spanning tree
 - Traverse graph using spanning tree as escape route



Spanning tree

