

Exercise: Repeat Resolution

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1 Algorithm of Tammi

Given the following multiple alignment and corresponding error rates find all pairs of DNPs with $p^{corr} \leq p_{max}^{corr} = 0.25$. Why is this threshold not appropriate?

Note: Since there are many calculation steps, it is recommended to implement an algorithm or use a spreadsheet e.g. Excel.

A	T	G	C	T	.01	.01	.001	.01	.1
C	T	A	A	A	.01	.01	.01	.01	.01
A	T	G	C	A	.001	.001	.001	.001	.01
A	T	G	C	A	.001	.001	.001	.001	.01
C	T	T	A	A	.001	.001	.001	.001	.001
C	T	C	A	A	.001	.01	.01	.01	.001
A	A	T	C	A	.01	.1	.1	.01	.01

2 Algorithm of Kececioglu

Given the following multiple alignment: The columns 2 and 5 are identified as DNPs and

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C A T C A
C A T C C
C T G C T
C A T C A
C T T G T
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$k=2$, that means the DNPs can be split into 2 groups.

Build a K_n graph and formulate an ILP for the problem. The ILP is going to have a lot of constraints. Solve it with a LP solver (e.g. soplex).