

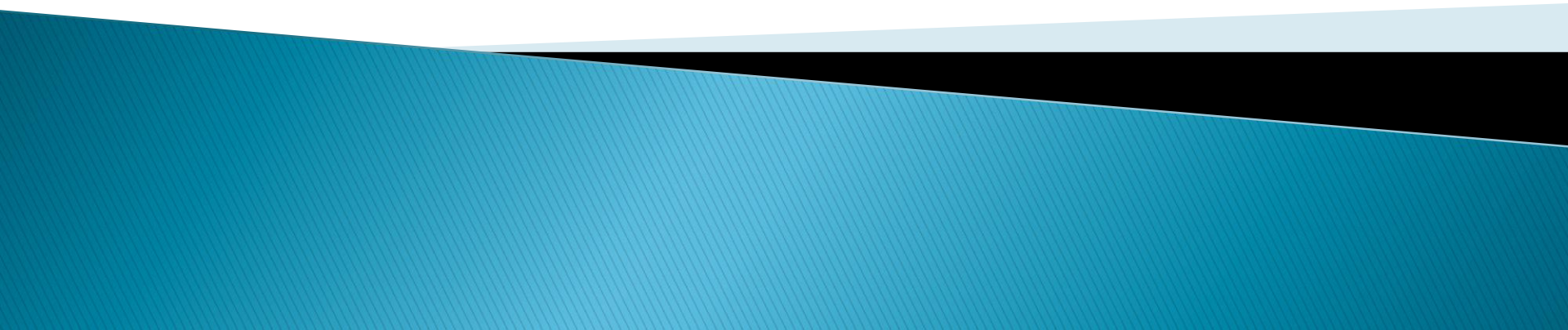
Relationship Between Edit Distance and Q-Gram Distance

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Motivation: Read Mapping to Reference Genomes

- ▶ DNA sequencing

...AGCTGCCATTGCT...

adenine guanine thymine cytosine



Motivation: Read Mapping to Reference Genomes

- ▶ Read mapping (aligning) to reference genome: matching shorter sequences (reads) to a longer reference sequence



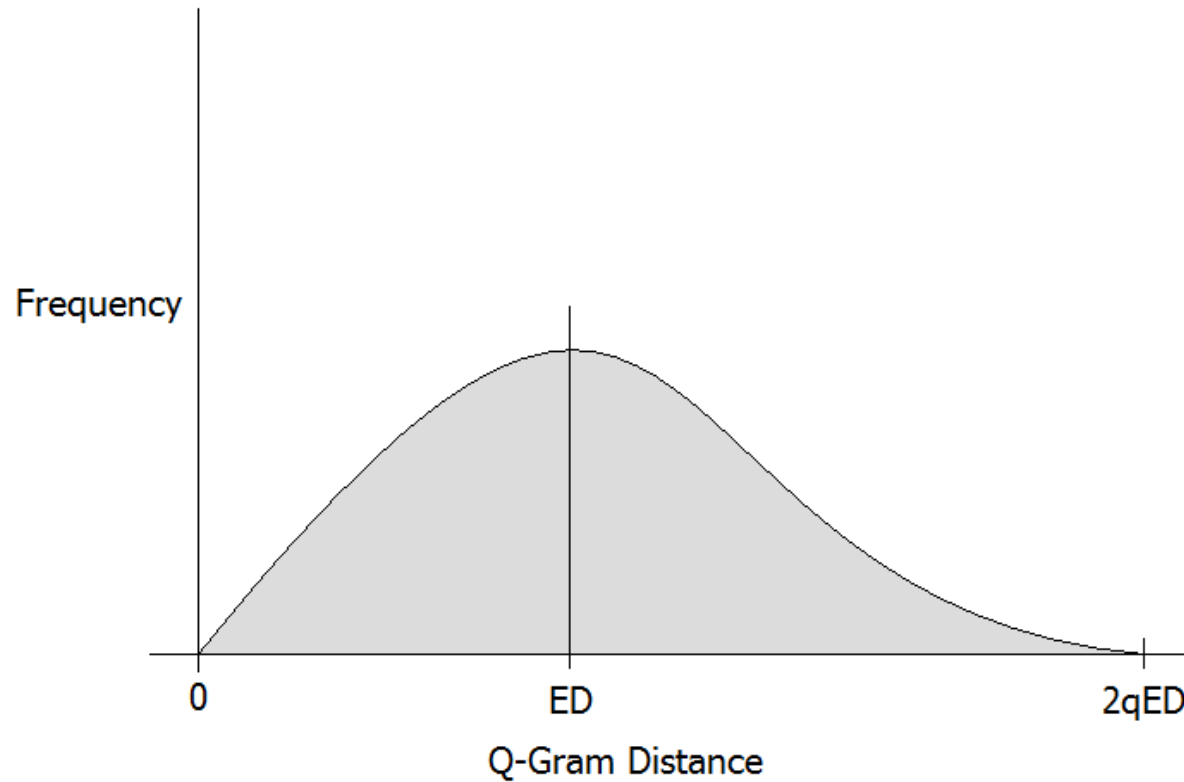
TCAGTCGTAATATACG — TAAGCCATGCGCTCGTACGTTAC
ATGTGCTAA — AT

Edit Distance and Q-Gram Distance

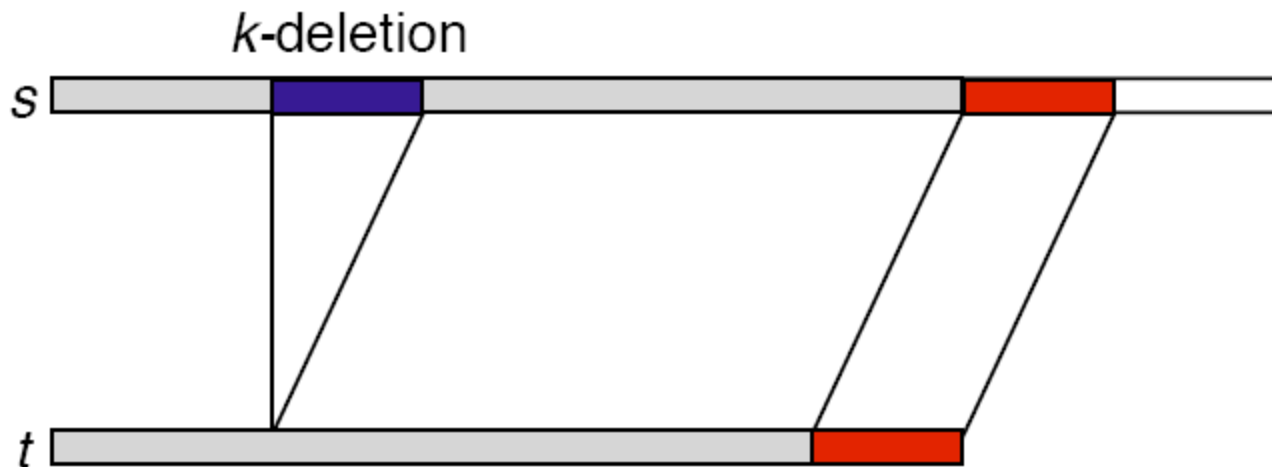
- ▶ **Edit Distance:** Minimal number of edits (insertions, deletions, substitutions) required to transform one character string into another.
- ▶ **Q-Gram Distance:** L1 norm of the difference of the q-gram profiles
 - Example: AAA ($f_{AA}:2$) and AAG ($f_{AA}:1, f_{AG}:1$), q-gram distance of 2

Edit Distance and Q-Gram Distance

- ▶ $QD \leq 2qED$



K-Indel Problem



- ▶ $L1(s,t)$ = q-gram distance between s and t
- ▶ Probability of $L1(s,t) = 0$? $L1(s,t) = 2x$?

1-Gram Case

$$\begin{aligned} &P[L_1(s, t) = 0 | ED(s, t) = k, \text{one deletion}] \\ &\leq \binom{k}{c_A, c_C, c_G, c_T} P_1^* \leq \frac{k!}{(\frac{k}{4}!)^4} (P_2^*)^k \end{aligned}$$

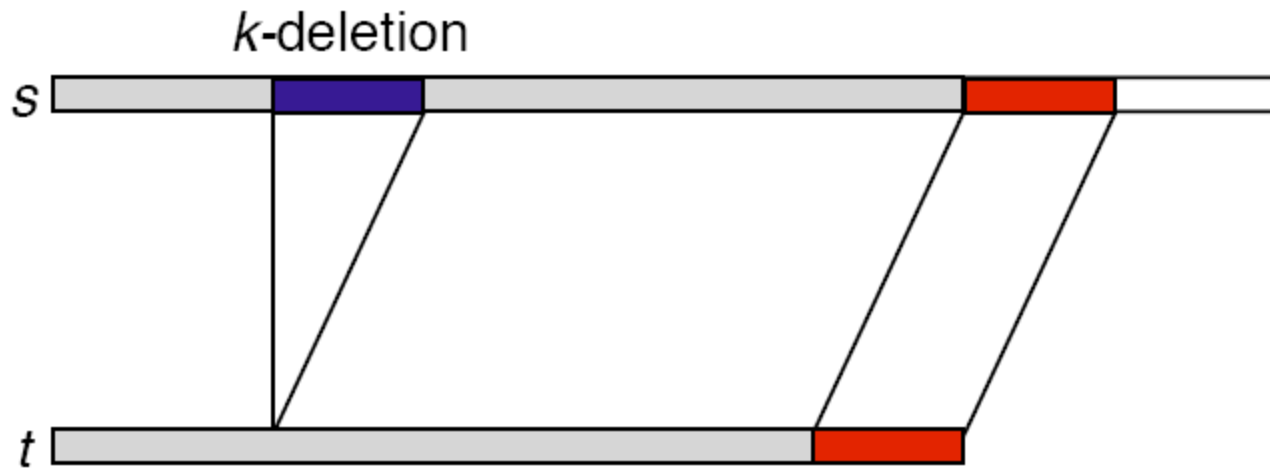
1-Gram Case

$$\begin{aligned} & P[L_1(s, t) = 2x | ED(s, t) = k, \text{onedeletion}] \\ & \leq \left[\binom{4}{1} \binom{x+2}{2} + \binom{4}{2} \binom{x-1}{1} \binom{x+1}{1} + \binom{4}{3} \binom{x-1}{2} \right] \frac{k!}{(\frac{k}{4}!)^4} (P_2^*)^k \end{aligned}$$

► Simplifies to...

$$[L_1(s, t) = 2x | ED(s, t) = k, \text{one deletion}] \leq (10x^2 + 2) \frac{k!}{(\frac{k}{4}!)^4} (P_2^*)^k$$

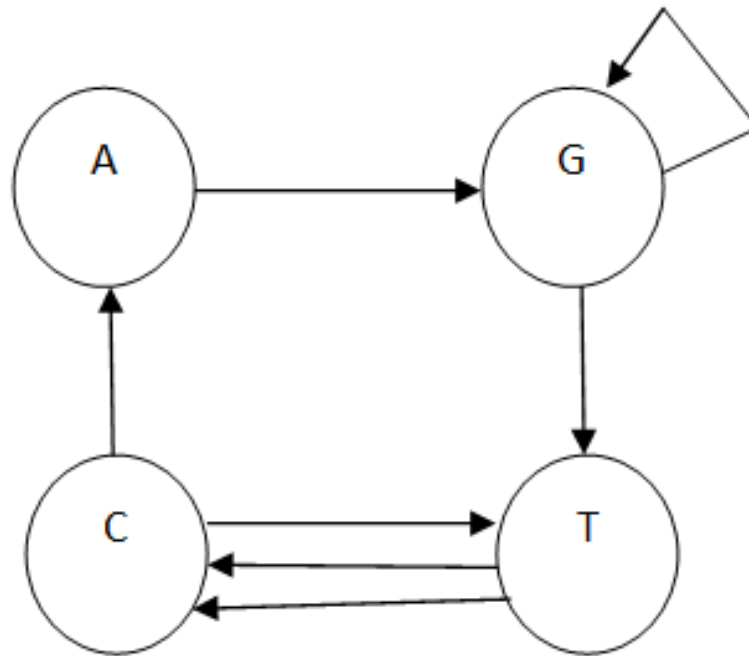
2-Gram Case: Previous Attempts



- ▶ 1-gram case as an upper bound
- ▶ 2-gram as independent
- ▶ 2-gram frequency vector as a state for a Markov Chain

2-Gram Case: Eulerian Graphs

► AGGTCTCA



2-Gram Case: Eulerian Graphs

- ▶ BEST Theorem

$$ec(G) = t_w(G) \prod_{v \in V} (deg(v) - 1)!$$

- ▶ Requires some modification

$$ec^*(G) = \frac{f^* t_w(G) \prod_{v \in V} (deg(v) - 1)!}{\prod_{a,b \in \Sigma} f_{ab}!}$$

2-Gram Case: Eulerian Graphs

- ▶ Example: AGGTCTCA
- ▶ Using $ec^*(G)$, # of sequences with the same 2-gram composition = 2
- ▶ Using 1-gram upper bound, # of sequences with the same 1-gram composition = 2520

2-Gram Case: Eulerian Graphs

$$\begin{aligned} &P[L_1(s, t) = 0 | ED(s, t) = k, \text{one deletion}] \\ &\leq \frac{f^* t_w(G) \prod_{v \in V} (\deg(v) - 1)!}{\prod_{a, b \in \Sigma} f_{ab}!} \left(\prod_{u \in \Sigma^2} P_u^{f_u} \right) \end{aligned}$$

Further Investigation

- ▶ Refine the Eulerian graph approach
- ▶ Generalize to $L1(s,t) \neq 0$
- ▶ Generalize to $q > 2$
- ▶ Generalize to other cases of modified strings (>1 indel, mismatches)

References

- ▶ *Circuits and Trees in Oriented Linear Graphs*, T. van Aardenne-Ehrenfest, N.G. de Bruijn
- ▶ *On Unicursal Paths in a Network of Degree 4*, W.T. Tutte, C.A.B. Smith
- ▶ *Approximate string-matching with q -grams and maximal matches*, Esko Ukkonen
- ▶ *Indel-tolerant Read Mapping with Trinucleotide Frequencies using Cache-Oblivious kd -Trees*, Md Pavel Mahmud, John Wiedenhoeft, Alexander Schliep