Subsequences with Generalised Gap Constraints Upper and Lower Complexity Bounds

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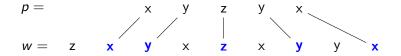
June 27, 2024

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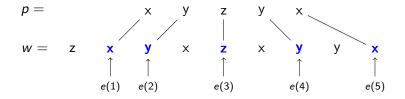
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$$p =$$
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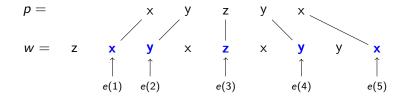


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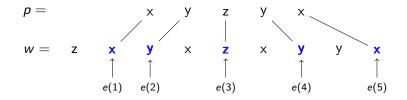
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Notation: $p \leq_e w$ with $e : [m] \rightarrow [n]$ (or just $p \leq w$)

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Properties of e:

e(1) < e(2) < · · · < e(m)
 w[e(i)] = p[i]

Definition

Given positions $i, j \in [m], i < j$, define

$$gap_{w,e}[i,j] := w[e(i) + 1..e(j) - 1]$$

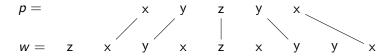
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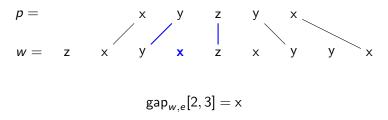


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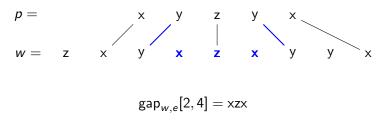


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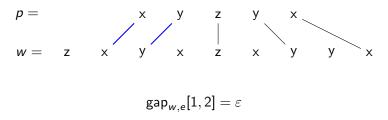


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A gap constraint is defined as a triple C = (i, j, L) with $i, j \in [m], i < j$ and $L \subseteq \Sigma^*$. An embedding *e* is said to satisfy the constraint if and only if

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Semilinear sets

Definition

A subset $L \subseteq \mathbb{N}$ is called *linear*, if it is of the form

$$L = L(x_0; x_1, \dots, x_m) = \{x_0 + \sum_{i=1}^m k_i x_i \mid k_1, \dots, k_m \in \mathbb{N}_0\}$$

with $x_0 \in \mathbb{N}_0, x_1, \dots, x_k \in \mathbb{N}$. A *semilinear* set is a finite union of linear sets.

Size of a (semi-)linear set:

- A linear set $L = L(x_0; x_1, ..., x_m)$ has size size(L) = m + 1.
- A semilinear set $L = \bigcup_{i=1}^{k} L_i$ with L_i linear sets has size size $(L) = \sum_{i=1}^{k} \text{size}(L_i)$.

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- $p \leq_e w$ and
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We use $\rm MATCH_{REG}$ and $\rm MATCH_{SLS}$ to denote the variants of the matching problem with regular and semilinear length constraints respectively.

Polynomial solutions for MATCH with constant $|\mathcal{C}|$

 $\mathrm{MATCH}_{\mathsf{REG}}:$ Construct an NFA to solve the problem:

Theorem

MATCH_{REG} can be solved in polynomial time for constant |C| and is fixed parameter tractable for the combined parameter (|p|, gapsize(C)).

 $\rm Match_{SLS}$: Enumerate all possible partial embeddings of the positions that have a constraint, then "fill the gaps":

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MATCH_{SLS} can be solved in polynomial time for constant |C|.

Hardness of $\mathrm{MATCH}_{\mathsf{SLS}}$

Definition (k-CLIQUE)

Given a graph G = (V, E) with *n* vertices, decide whether there is a subset $K \subseteq V$ of *k* vertices that are pairwise adjacent.

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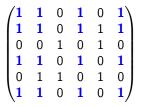
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- Constrain the gap between the anchor and each element on the diagonal to be a multiple of n+1.
- Constrain the gap between elements on subsequent rows and in the same column to be one smaller than a multiple of *n*.

/1	1	0	1	0	1
1 1 0	1	0	1	1	1 1
1 0 1 0	0		0	1	0
1	1	0	1	0	1
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- Constrain the gap between the first and last element of each row to be smaller than n-1.

Theorem

MATCH_{SLS} parameterised by |p| is W[1]-hard, even for constant gapsize(C) and binary alphabet Σ .

Hardness of $\mathrm{Match}_{\mathsf{REG}}$

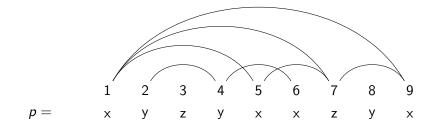
Definition (1-in-3-3SAT)

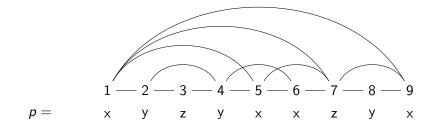
Given a set of variables $A = \{x_1, \ldots, x_n\}$ and clauses $c_1, \ldots, c_m \subseteq A$ with $|c_j| = 3$, find a subset $B \subseteq A$, such that $|c_i \cap B| = 1$ for every $j \in [m]$.

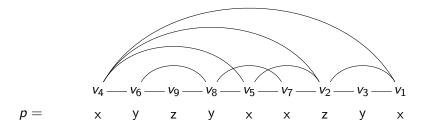
- Pattern $p = (b\#)^n (b\#)^m$
- Text $w = (bb\#)^n (bbb\#)^m$
- First part: decide $x_i \in B$ for $i \in [n]$.
- Second part: decide which of the 3 variables from c_j is in B for $j \in [m]$ (need some ordering on the variables).
- Use constraints to ensure that assignments are consistent.

Theorem

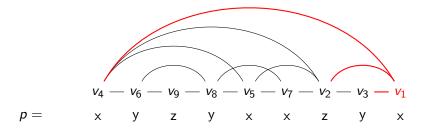
 $Match_{REG}$ is NP-complete, even for binary alphabet Σ and with gap-constraints that can be represented by DFAs with at most 8 states.



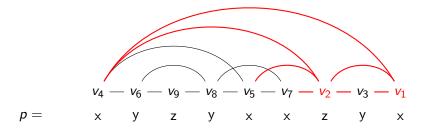




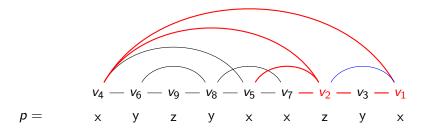
- Construct partial embeddings by adding the positions one-by-one in order σ = (v₁,..., v_m).
- At each time t, we compute all possible partial embeddings, but we only care about the the values e(v_i), where i ≤ t and (v_i, v_j) ∈ E for some j > t.
- When adding $e(v_t)$ to the embedding, combine all previously possible embeddings with all possible values of $e(v_t)$, then remove all embeddings that violate any constraint involving v_t .



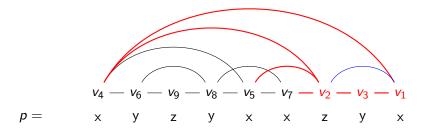
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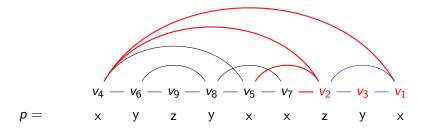
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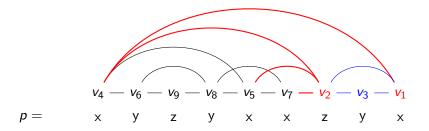
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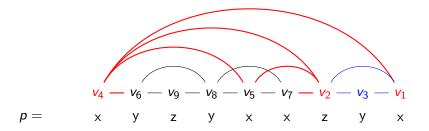
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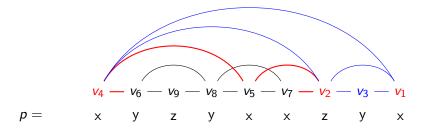
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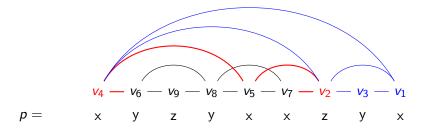
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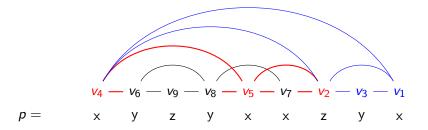
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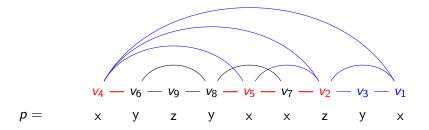
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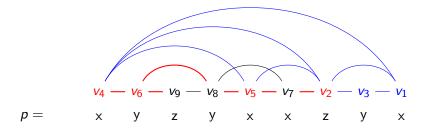
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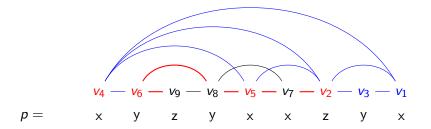
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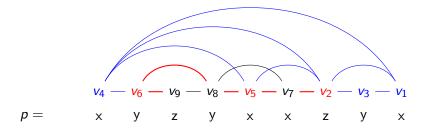
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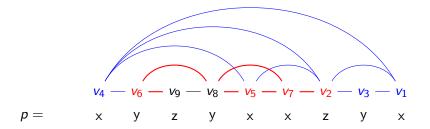
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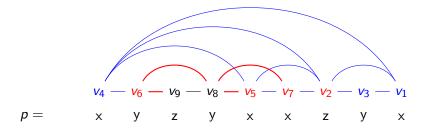
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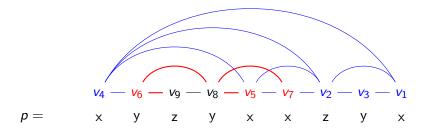
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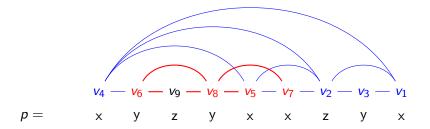
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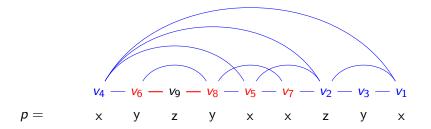
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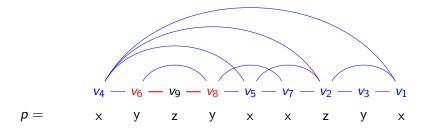
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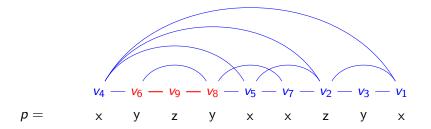
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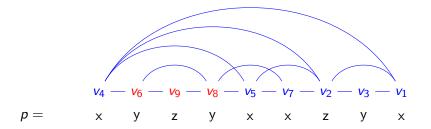
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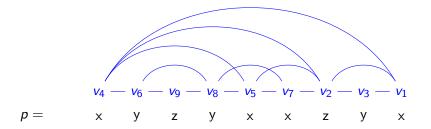
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Definition (Vertex separation number)

Given a linear ordering $\sigma = (v_1, \ldots, v_n)$ of vertices in a graph, the vertex separation number of σ is the smallest number s such that, for each vertex v_i at most s vertices of v_1, \ldots, v_{i-1} have some v_j with $j \ge i$ as neighbour. The vertex separation number of a graph is the minimum vertex separation number over all linear orderings of the graph.

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Theorem

If the vertex separation number of the constraint graph is bound by k, MATCH_{REG} and MATCH_{SLS} can be solved in $\mathcal{O}(m^2n^{k+1} + m^2n^2\log\log n)$ and $\mathcal{O}(m^2n^{k+1})$ time respectively.

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The second part is witnessed by the k-CLIQUE reduction from earlier.

The Interval Structure of Constraints

For C = (i, j, L) define *interval*(C) = [i, j - 1].

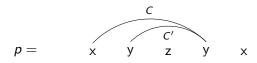
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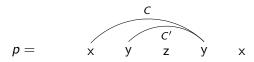
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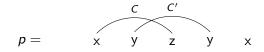
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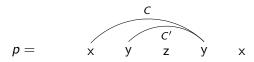
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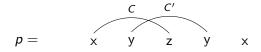
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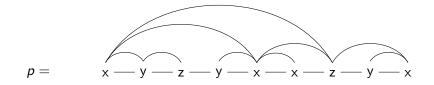
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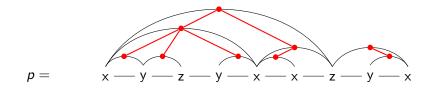
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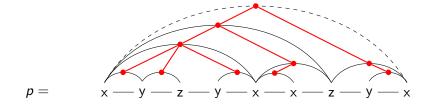


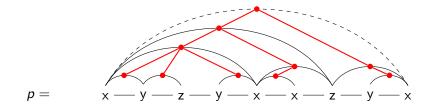
What about MATCH with pairwise non-intersecting constraints?



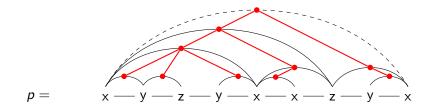








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Theorem

If constraints are pairwise non-intersecting, $MATCH_{REG}$ and $MATCH_{SLS}$ can be solved in $\mathcal{O}(n^{\omega}k + n^2k \log \log n)$ and $\mathcal{O}(n^{\omega}k)$ time respectively, where $\mathcal{O}(n^{\omega})$ is the time needed to multiply two boolean matrices of size $n \times n$.

Definition (3-OV)

Given three sets $A = \{\vec{a}_1, \dots, \vec{a}_n\}, B = \{\vec{b}_1, \dots, \vec{b}_n\}, C = \{\vec{c}_1, \dots, \vec{c}_n\}$ of *d*-dimensional boolean vectors, are there indices $i, j, k \in [n]$, such that $\vec{a}_i \cdot \vec{b}_j \cdot \vec{c}_k = \vec{0}$?

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$$p = \overline{\mathsf{C}}_{p}(\vec{a}_{n}) \dots \overline{\mathsf{C}}_{p}(\vec{a}_{i}) \dots \overline{\mathsf{C}}_{p}(\vec{a}_{1}) \quad \S \quad \mathsf{C}_{p}(\vec{a}_{1}) \dots \mathsf{C}_{p}(\vec{a}_{i}) \dots \mathsf{C}_{p}(\vec{a}_{n})$$
$$w = w_{0} \ \overline{\mathsf{C}}_{w}(\vec{b}_{n}) \dots \overline{\mathsf{C}}_{w}(\vec{b}_{1}) \ w_{0} \qquad \S \qquad w_{0} \ \mathsf{C}_{w}(\vec{c}_{1}) \dots \mathsf{C}_{w}(\vec{c}_{n}) \ w_{0}$$

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Theorem

Both variants of MATCH with pairwise non-intersecting constraints cannot be solved in $O(n^g k^h)$ time with g + h < 3, unless the Strong Exponential Time Hypothesis fails.

Thank you for your attention!

Summary:

- Problem: Matching subsequences with gap constraints
- Two types of constraints: regular and semilinear length constraints
- Polynomial solutions for constant amount of constraints
- Hardness of the problem parameterized by the length of the pattern, witnessed by reductions from *k*-CLIQUE and 1-in-3-3-SAT.
- Graph structure of Constraints: Relation between complexity of the problem and vertex separation number of constraint graph
- Interval structure of Constraints: Efficient solution for the case of non-intersecting constraints, lower bound via fine-grained reduction from 3-OV.

Do you have any questions?