

Exact ensemble properties in combinatorial dynamic programming schemes

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Optimization problem =

- Search space S
- Objective function f

Problem: Find element $e \in S$ which min(max)-imizes $f(s)$?

Dynamic programming scheme relates the minimal value of f to its minimal value(s) on some *smaller* search space(s) (Substructure property).

DP scheme = Efficient factorization (traversal) of S .

Alt.: DP scheme is **generating** search space.

In order to say something general, let us be specific. . .

Definition (Combinatorial DP)

A **combinatorial DP** scheme computes functions that are **locally additive** and relies on a decomposition that is:

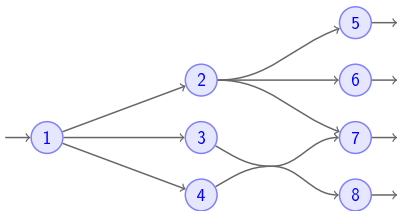
- **Unambiguous**: Each solution generated **at most** once!
- **Complete**: Each solution generated **at least** once!

Based on this property, DP schemes for optimization readily translate into *DP* schemes for counting, generating. . .

Remark: None of this above is strictly required by DP!

What is a decomposition?

Hypergraphs as decompositions



Hypergraphs generalize directed graphs to arcs of arbitrary in/out degrees.

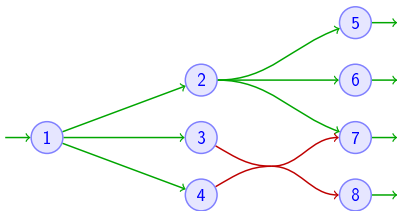
Definition (Hypergraph)

A directed hypergraph \mathcal{H} is a couple (V, E) such that:

- V is a set of vertices
- E is a set of hyperarcs $e = (t(e) \rightarrow h(e))$ such that $t(e), h(e) \subset V$

Forward hypergraphs, or F-graphs, are hypergraphs whose arcs have incoming degree exactly 1.

Hypergraphs as decompositions



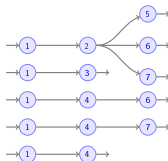
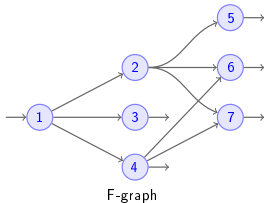
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All F-paths starting from vertex 1

Definition (F-path)

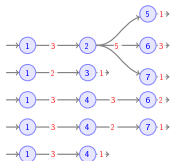
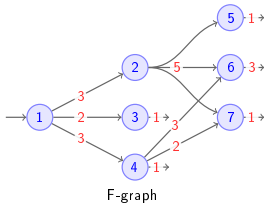
A **F-path** is a tree having root $s \in V$, whose children are F-paths built from the **outgoing vertices** of some arc $e = (s \rightarrow \mathbf{t}) \in E$.

Remark: Vertices of out degree 0 ($\mathbf{t} = \emptyset$) provide an elegant terminal case to the above recursive definition.

F-graph is **independent** iff each F-path sees at most once each arc.

A numerical **valued-function** $\pi : E \rightarrow \mathbb{R}$ can be assigned to each arc:

- **Weight** of a path is the **product** of its arcs' values
- **Score** of a path is the **sum** of its arcs' values



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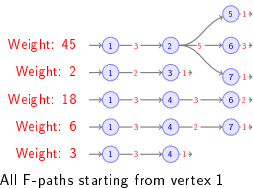
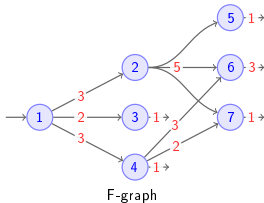
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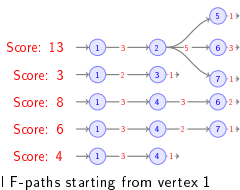
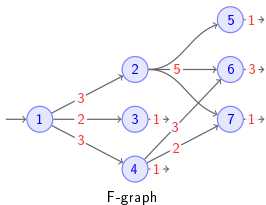
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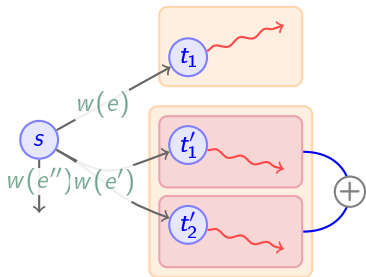
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Basic algorithms

$\mathcal{H} = (s_0, V, E, \pi)$: acyclic F-graph s_0 : Init. node π : value fun.

$$m_s = \min_{e=(s \rightarrow t)} \left(w(e) + \sum_{u \in t} m_u \right)$$

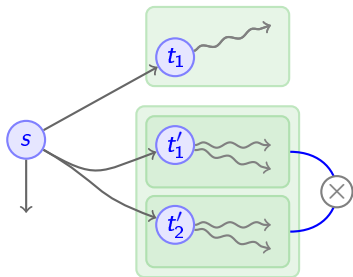


Problem	Recurrence	Complexities time/space
Min score	$m_s = \min_{e=(s \rightarrow t)} \left(w(e) + \sum_{u \in t} m_u \right)$	$\Theta(E + V) / \Theta(V)$
Num. paths	$n_s = \sum_{(s \rightarrow t)} \prod_{u \in t} n_u$	$\Theta(E + V) / \Theta(V)$
Total weight	$w_s = \sum_{e=(s \rightarrow t)} w(e) \cdot \prod_{s' \in t} w_{s'}$	$\Theta(E + V) / \Theta(V)$

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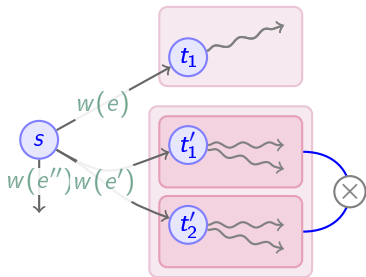


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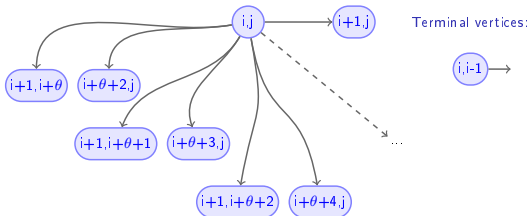
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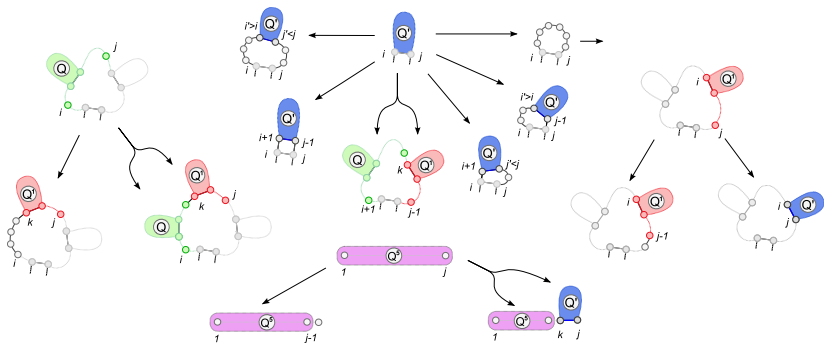
Nussinov's algorithm revisited

$$\begin{array}{c} \bullet \text{-----} \bullet \\ i \qquad \qquad \qquad j \end{array} = \begin{array}{c} \bullet \text{---} \bullet \text{-----} \bullet \\ i \quad i+1 \qquad \qquad \qquad j \end{array} + \begin{array}{c} \text{---} \text{---} \text{---} \text{---} \text{---} \text{---} \text{---} \text{---} \text{---} \text{---} \\ i \quad i+1 \quad \quad \quad k-1 \quad k \quad k+1 \quad j \\ \text{---} \text{---} \text{---} \text{---} \text{---} \text{---} \text{---} \text{---} \text{---} \text{---} \\ \geq \theta \end{array}$$

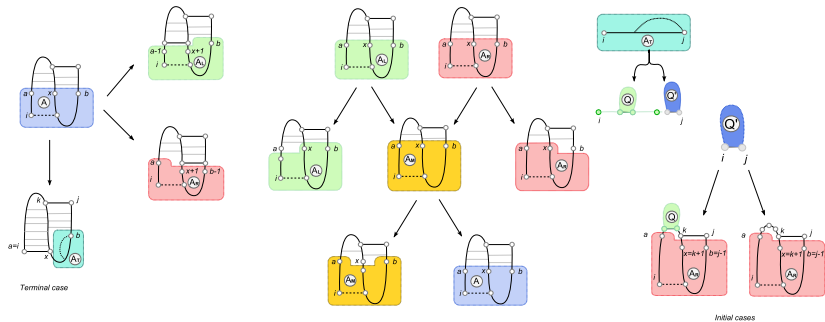


This decomposition is unambiguous!
(Proof may use length generating functions).

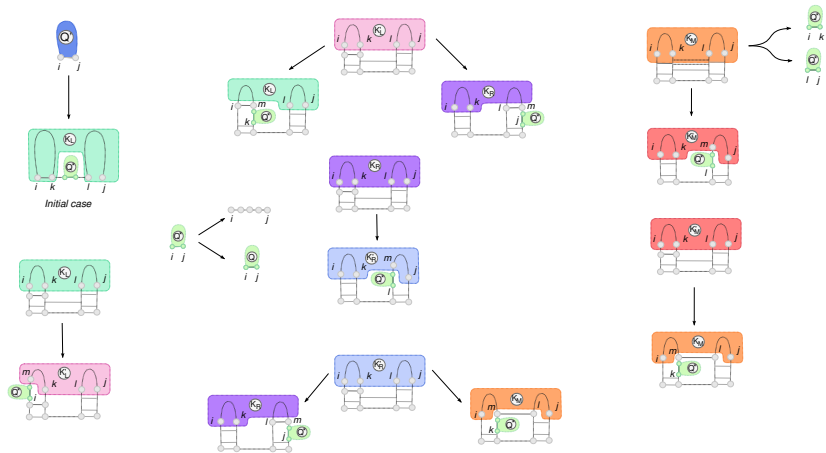
Mfold/Unafold decomposition



Akutsu's simple pseudoknots



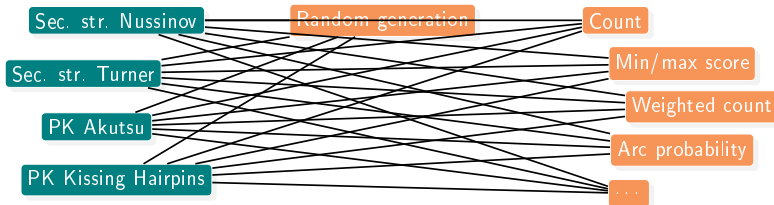
Kissing hairpins



Half time summary

Message #1

Applications of DP could (and should) be detached from the equation, and be expressed at an abstract – combinatorial – level.



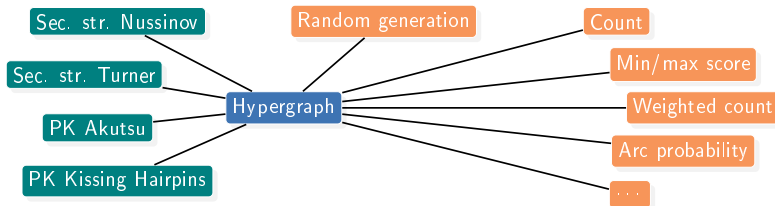
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Let us extend applications of DP...

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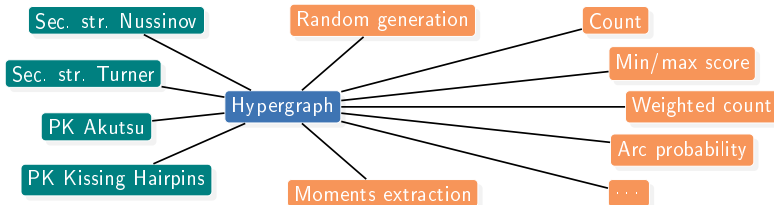


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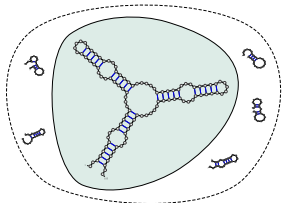
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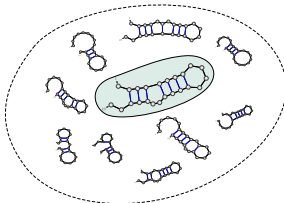
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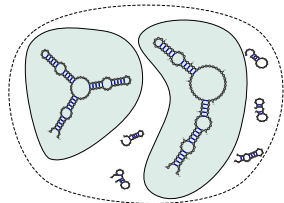
Distribution of solutions



Functional folding?



Ill-defined folding:
mRNA?



Bistable RNA
Kinetics?

- An acyclic F-graph $\mathcal{H} = (s_0, V, E, \pi)$, defining a *search space* \mathcal{T} the set of F-paths (trees) in \mathcal{H} .
- Feature functions $\alpha_1, \dots, \alpha_k : E \rightarrow \mathbb{R}$ extended **additively** on \mathcal{T} .

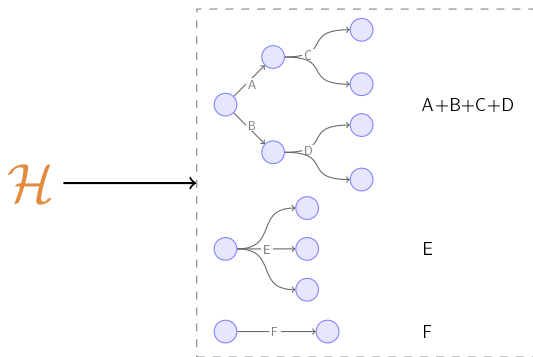
What can we say about the distribution(s) of $\alpha_i(s)$?

Naive approach: Compute the distribution exactly, accessible values being choices of $\mathcal{O}(n)$ values among $|E|$, $n = \#$ arcs in largest F-path.

\Rightarrow DP count the $\#$ ways of **accessing** each value in exponential time...

Example: Mean value of a feature function

What is the average values of α assuming a uniform distribution on \mathcal{T} ?

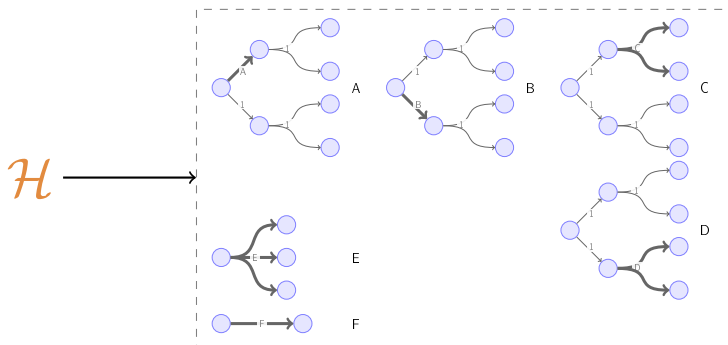


$$\text{Mean value} = \text{Weighted sum} = \frac{\sum_{t \in \mathcal{T}} \alpha(t)}{|\mathcal{T}|}$$

Remark: Dropping the terminal edges (Alt. $\alpha(e) = 0$)...

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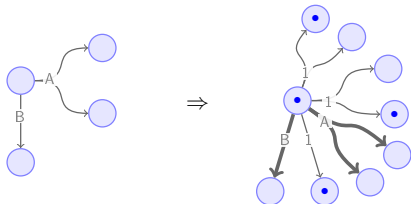


$$\text{Mean value} = \text{Weighted sum} = \frac{\sum_{t \in \mathcal{T}} \alpha(t)}{|\mathcal{T}|}$$

Principle and main equation

By introducing a **controlled ambiguity**, one can extract the feature expectation from a combinatorial DP scheme.

Idea: Passing a dot \bullet along or drop it ($\equiv \alpha(e)$) on some arc e .



$$\Rightarrow c_s^{\bullet\alpha} = \sum_{o \in \mathcal{T}} \alpha(o) = \sum_{e=(s \rightarrow t)} \left(\alpha(e) \cdot \prod_{t_i \in t} c_{t_i} + \sum_{t_i \in t} c_{t_i}^{\bullet\alpha} \prod_{t_j \neq t_i \in t} c_{t_j} \right)$$

$\Rightarrow \mathbb{E}(\alpha) := c_{s_0}^{\bullet\alpha} / c_{s_0}$ obtained in $\Theta(|V| + |E| \cdot D^2) / \Theta(|V|)$ time/space, with $D := \max_{e \in E} (|h(e)|)$

Remark: Similar to the *folklore* pointing operator in enumerative combinatorics, and to the formal derivative.

$D: \max_{e \in E} (|h(e)|)$

- Boltzmann-like distribution

Unconvinced by the uniform distribution? Got an energy function $\Delta : E \rightarrow \mathbb{R}$ extended additively, and a good reason to expect convergence toward a Boltzmann distribution $e^{-\frac{\Delta}{kT}}$? (We do...)

\Rightarrow Expectation in the Boltzmann distribution is just a weight away.

$$\begin{aligned} c_s^{\bullet\alpha} &= \sum_{o \in \mathcal{T}} \alpha(o) \cdot e^{-\Delta(o)/RT} \\ &= \sum_{e=(s \rightarrow t)} e^{-\frac{\Delta(e)}{kT}} \left(\alpha(e) \cdot \prod_{t_i \in \mathbf{t}} c_{t_i} + \sum_{t_i \in \mathbf{t}} c_{t_i}^{\bullet\alpha} \prod_{t_j \neq t_i \in \mathbf{t}} c_{t_j} \right) \end{aligned}$$

$\Rightarrow \Theta(|V| + |E| \cdot D^2) / \Theta(|V|)$

- Higher-order moments
- Exact correlations

Nb: Specializes into Miklos *et al* 2005 for RNA secondary structures.

Low-hanging fruits

D : $\max_{e \in E} (|h(e)|)$

- Boltzmann-like distribution $\Rightarrow \Theta(|V| + |E| \cdot D^2) / \Theta(|V|)$
- Higher-order moments

Want to go beyond the expected value of α ? No problem (almost), compute higher-order moments.

$$\begin{aligned} c_s^{\bullet \alpha^m} &= \sum_{o \in \mathcal{T}} \alpha(o)^m \cdot e^{-\Delta(o)/RT} \\ &= \sum_{e=(s \rightarrow t)} e^{-\frac{\Delta(e)}{kT}} \left(\sum_{\substack{(m'_1, \dots, m'_{|t|}) \\ \text{s.t. } \sum m'_i = m' \leq m}} \alpha(e)^{m-m'} \prod_{t_i \in t} c_{t_i}^{\bullet \alpha^{m'_i}} \right) \end{aligned}$$

$$\Rightarrow \Theta(|V| + |E| \cdot m^{D+2}) / \Theta(m \cdot |V|)$$

- Exact correlations

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Low-hanging fruits

$$D: \max_{e \in E} (|h(e)|)$$

- Boltzmann-like distribution $\Rightarrow \Theta(|V| + |E| \cdot D^2) / \Theta(|V|)$
- Higher-order moments $\Rightarrow \Theta(|V| + |E| \cdot m^{D+2}) / \Theta(m \cdot |V|)$
- Exact correlations Given two α and α' , correlation is given by

$$c_s^{\alpha \cdot \alpha'} = \sum_{o \in \mathcal{T}} \alpha(o) \cdot \alpha'(o) \cdot e^{-\Delta(o)/RT}$$

$$= \sum_{e=(s \rightarrow t)} e^{-\frac{\Delta(e)}{kT}} \left(\sum_{\substack{((m_1 \cdot m_{|t|}), \\ (m'_1 \cdot m'_{|t|})) \\ m := \sum m_i \leq 1, \\ m' := \sum m'_i \leq 1}} \alpha(e)^{1-m} \alpha'(e)^{1-m'} \prod_{t_i \in t} c_{t_i}^{\alpha \cdot \alpha'} \right)$$

$$\Rightarrow \Theta(|V| + |E| \cdot D^2) / \Theta(|V|)$$

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Conclusion

- Many algorithms in Bioinformatics rely on dynamic programming
- Adopting a combinatorial vision over the search space greatly, and hypergraph representations, helped:
 - Prove correctness.
 - Instant transposal of new applications.
- Implementation tricky (avoid explicit representations).
- Use in combination with machine-learning as classifier/scanner for ncRNAs.
- How to transpose usual DP tricks (four-russians, cutting corners, sparsification)?
- Are there even more compact ways to describe more specific search spaces (e.g. CFGs, split-types??)