# Trading Time and Space in Catalytic Branching Programs 

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Joint work with James Cook

## Space-bounded computation

$B P(w, \ell)$ : layered branching programs of width $w$ and length $\ell$


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$B P$ is non-uniform: no restrictions on the description

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|  |  | $(1,0)$ |
| :---: | :---: | :---: |
| 1 |  | $(1,1)$ |
| 2 |  | $(2,0)$ |
| $i$ | $f(x)=0$ | $(i, 0)$ |
|  | $f(x)=1$ | $(i, 1)$ |
| $m$ |  |  |

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## Catalytic computation

$\operatorname{CSPACETIME}(s, t, c)$ : space-bounded Turing Machines with an extra worktape ( $c$ bits) of full memory

work tape


## catalytic tape

| 0 | 1 | 1 | $\cdots$ | 0 | 1 |
| :--- | :--- | :--- | :--- | :--- | :--- |

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- $m \cdot w$ nodes in a layer $\leftrightarrow \log m+\log w$ bits in memory
- $m$ sources plus source-sink pairing requirement $\leftrightarrow$ resetting $\log m$ catalytic memory)


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Two interpretations of reducing $w$ and $m$ (non-uniform):

1) amortized space: reducing the amortized space ( $w=(w \cdot m) / m$ ) needed to compute $f$, or the number of copies $(m)$ needed for amortization to help
2) catalytic space: reducing the amount of space ( $\log w$ ) and catalytic space $(\log m)$ needed to compute $f$

## Known results

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In contrast, [Potechin'17] gives (asymptotically) optimal amortized width $w=O(1)$ and length $\ell=O(n)$ simultaneously
...but we need $\mathbf{m}=\mathbf{2}^{\mathbf{2}^{\mathbf{n}}-\mathbf{1}}$ to get it!

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## Our results (permutation branching programs)

[Potechin'17]': every function $f$ can be computed by a read-4 permutation branching program of width $2^{2^{n}+1}$.

Main result 1': for any $\epsilon>0$, every function $f$ can be computed by a read $-O_{\epsilon}(1)$ permutation branching program of width $2^{2^{\epsilon n}}$.

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Setup: catalytic space $\log m=2^{n}-1$ in some initial state $\tau_{1} \ldots \tau_{2^{n}-1}$, plus $\log 4=2$ bits of free work space

$(11 \ldots 1) \bigcirc$
$\bigcirc((11 \ldots 1), 1)$
[Potechin'17] in two slides
0) First free bit: $\overrightarrow{0}$ entry of $g$


## [Potechin'17] in two slides

1) $g\left(\alpha_{1} \ldots \alpha_{i} \ldots \alpha_{n}\right) \rightarrow g\left(\alpha_{1} \ldots \alpha_{i}^{x_{i}} \ldots \alpha_{n}\right)$


## [Potechin'17] in two slides

2) $g(y) \rightarrow g(y)+f(y)$


## [Potechin'17] in two slides

3) $g\left(\alpha_{1} \ldots \alpha_{i}^{x_{i}} \ldots \alpha_{n}\right) \rightarrow g\left(\alpha_{1} \ldots \alpha_{i} \ldots \alpha_{n}\right)$


## [Potechin'17] in two slides

4) Second free bit (output): copy the answer from first free bit


## [Potechin'17] in two slides

5) Undo steps 1-3 (do steps 3-1)


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Catalytic algorithms give us a way to compute $f$ over the monomial basis only using catalytic memory.

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- compute each monomial into separate memory in parallel
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2) Cook-Mertz algorithm (branching program edition)

- compute each monomial directly into the output register in series
- exponential time, linear space


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Main result 1: for any $\epsilon>2 / n$, every function $f$ can be computed by an $m$-catalytic branching program of width $2 m$ and length $2^{1 / \epsilon} \cdot 2 \epsilon n$, where $m=2^{n+\frac{1}{\epsilon} \cdot 2^{\epsilon n}}$.

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Proof idea (new): monomial version of Potechin algorithm again, but now only compute monomials which actually appear in $f\left(\binom{n}{\leq d}\right.$ by assumption).

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Main result 2: for any $\epsilon>2 / d$, if $f$ is a degree $d$ polynomial over $\mathbb{F}_{2}$, then $f$ can be computed by an $m$-catalytic branching program of of width $2 m$ and length $2^{1 / \epsilon} \cdot 2 n$, where $m=2^{n+\frac{1}{\epsilon} \cdot\left({ }_{\leq \epsilon d}^{n}\right)}$.

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Proof idea: same* time-space tradeoff as before, now with $\epsilon d$ instead of $\epsilon n$.

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Main result 4: any permutation* branching program calculating the AND function which reads any variable less than three times requires length at least $4 n-4$.

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Optimal permutation branching program length for any function

- somewhere between $3 n^{*}$ and $4 n-4$
- can get a read-3 program for $\operatorname{AND}\left(x_{1}, x_{2}, x_{3}\right)$

