Random generation of capacities

Michel GRABISCH^{*a,b*}, Christophe LABREUCHE^{*c*} Peiqi SUN^{*a*}

^aUniversité Paris I Panthéon-Sorbonne, Centre d'Economie de la Sorbonne ^bParis School of Economics, Paris, France ^cThales Research and Technology, Palaiseau, France • Capacities = monotone set functions vanishing on the empty set

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- Another problem: How to measure the performance of a capacity generator?

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Outline

1. The problem of uniform random generation

- 2. The 2-layer approximation method
- 3. Measure of performance

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Order polytopes

 (P, ≼): (finite) poset, with P a finite set and ≼ a partial order (reflexive, antisymmetric, transitive)

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- The order polytope (Stanley, 1986) associated to (P, \preccurlyeq) , denoted by $\mathcal{O}(P)$, is the set

$$\mathcal{O}(P) = \{ f : P \longrightarrow [0,1] \mid f(x) \leqslant f(y) \text{ if } x \preccurlyeq y \}.$$

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- (P, ≼): (finite) poset, with P a finite set and ≼ a partial order (reflexive, antisymmetric, transitive)
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• $\mathcal{O}(P)$ is a polytope of dimension p := |P|.

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- Notation: To each linear extension ≤ on P = {x₁,..., x_p} corresponds a permutation σ on {1,..., p} such that x_{σ(1)} < ··· < x_{σ(p)}.

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- Each linear extension σ defines a region in $\mathcal{O}(P)$:

 $R_{\sigma} := \{ f \in \mathcal{O}(P) \mid 0 \leqslant f(x_{\sigma(1)}) \leqslant f(x_{\sigma(2)}) \leqslant \cdots \leqslant f(x_{\sigma(p)}) \leqslant 1 \}$

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- All regions R_{σ} are identical (up to a change of coordinates) *p*-dimensional simplices with volume $\frac{1}{p!}$
- Vertices of R_σ are the p+1 functions given by

$$0 = f(x_{\sigma(1)}) = \cdots = f(x_{\sigma(k)}), \ f(x_{\sigma(k+1)}) = \cdots = f(x_{\sigma(p)}) = 1,$$

 $k = 1, \dots, p - 1$, and the two constant functions 0 and 1.

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Uniform random generation of a point in O(P) = uniform random selection of a linear extension σ of P + uniform random selection of a point in R_{σ}

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 - 3 Put $f(x_{\sigma(1)}) = z_1, \ldots, f(x_{\sigma(p)}) = z_p$.

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 - **1** $\mu(\emptyset) = 0, \ \mu(N) = 1 \ (normalization)$

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- Consequence: the problem of the uniform generation of capacities is solved!

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The number of linear extensions on $(2^N \setminus \{\emptyset, N\})$ (equal to the number of linear extensions on $(2^N, \subseteq)$ is given in the table below:

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This is sequence A046873 in the Online Encyclopedia of Integer Sequences. $e(2^N)$ is not known beyond n = 7. Some bounds are known (Brightwell and Winkler, 1991).
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 - supermodular capacities (Beliakov, 2022)

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 - supermodular capacities (Beliakov, 2022)
 - 2-additive capacities (Miranda and Garcia-Segador, 2020a)

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The Random Node Generator

Algorithm:

- $\ \textbf{Pick} \ \ S \in 2^N \setminus \mathcal{L}$
- Compute $\mu_{\min}(S) = \max_{T \in \mathcal{L}, T \subseteq S} \mu(T)$, $\mu_{\max}(S) = \min_{T \in \mathcal{L}, T \supseteq S} \mu(T)$
- Draw uniformly a number t in $[\mu_{\min}(S), \mu_{\max}(S)]; \mu(S) \leftarrow t$
- Goto step 2 while $\mathcal{L} \neq 2^N$

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Algorithm:

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$$\mathcal{L} \leftarrow \{ \varnothing, N \}; \ \mu(N) = 1; \ \mu(\varnothing) = 0$$

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Advantages: very simple and fast Drawbacks: yields very biased distribution

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- Two linear extensions σ, τ are *neighbors* if they differ only by a single transposition of neighbor elements:

$$\sigma: \quad x_{\sigma(1)}, \dots, x_{\sigma(k)}, x_{\sigma(k+1)}, \dots, x_{\sigma(p)}$$

$$\tau: \quad x_{\sigma(1)}, \dots, x_{\sigma(k+1)}, x_{\sigma(k)}, \dots, x_{\sigma(p)}$$

for some $k \in [1, p-1]$. Denote by $n(\sigma)$ the number of neighbors of σ (at most p-1).

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for some $k \in [1, p-1]$. Denote by $n(\sigma)$ the number of neighbors of σ (at most p-1).

• The order Markov chain M is defined on the set of states E(P) with transition probabilities:

$$egin{aligned} \mathcal{P}(\sigma, au) &= egin{cases} 1/(2p-2) & ext{if } \sigma, au ext{ are neighbors} \ 1-n(\sigma)/(2p-2) & ext{if } \sigma= au \ 0 & ext{otherwise.} \end{aligned}$$

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• The order Markov chain M is ergodic time-reversible and converges to the uniform distribution on E(P).

Algorithm

- **●** Input: a poset (P, \preccurlyeq) with $P = \{x_1, \ldots, x_p\}$, an integer T
- 2 Find a linear extension σ on P
- For i = 1 to T do:
 - choose at random an integer $k \in [1, 2p 2]$
 - if $k \leqslant p-1$ and $\operatorname{not}[x_{\sigma(k)} \prec x_{\sigma(k+1)}]$ then swap $x_{\sigma(k)}$ and $x_{\sigma(k+1)}$ in σ

• Output σ

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Estimation of T to get almost uniformity: $T = O(p^5 \log(e(P)))$.

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- Similarly, $x_{\sigma(2)} \in Min(P \setminus \{x_{\sigma(1)}\})$, and $x_{\sigma(p-1)} \in Max(P \setminus \{x_{\sigma(p)}\})$.
- Based on this observation, the probability that a linear extension of P starts with $m \in Min(P)$ (resp., ends with $M \in Max(P)$) is

$$\Pr(m \mid P) = \frac{e(P \setminus \{m\})}{e(P)}; \quad \Pr(M \mid P) = \frac{e(P \setminus \{M\})}{e(P)}$$

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 the correct probability given above, a minimal or a maximal element
 of a poset which is diminished by one element at each step.

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 of a poset which is diminished by one element at each step.
- As this probability directly depends on e(P), the computation can be exact only when P becomes small enough.
- Idea: take the lower part of the poset for choosing minimal elements, and the upper part for choosing maximal elements, thus neglecting minimal and maximal elements which are outside these two subparts.

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- Layer of H: all nodes (subsets) of same cardinality

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• For y in the 2nd layer, pred(y) is the set of its predecessors in the 1st layer (similarly with succ(x), x in the 1st layer)

In a dual way, we introduce B_H (denoted also B_H[h', k', |I'|]), the poset of the two bottom layers of H.

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- In a dual way, we introduce B_H (denoted also B_H[h', k', |I'|]), the poset of the two bottom layers of H.
- Consider a maximal element M of H belonging to T_H, and a minimal element m of H belonging to B_H. We put

$$\Pr(M \mid H) \approx \frac{e(T_H \setminus \{M\})}{e(T_H)} = \Pr(M \mid T_H)$$
$$\Pr(m \mid H) \approx \frac{e(B_H \setminus \{m\})}{e(B_H)} = \Pr(m \mid B_H)$$

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Rationale: If in average a node has l predecessors in the layer just above, a node in layer k has therefore O(l^{k-1}) predecessors in H. Hence, a node in the 3d layer has very little probability to become maximal, since all its predecessors must be eliminated first, without eliminating all nodes of the 1st layer

generate-linext(P, I)**Input:** a poset P subset of $2^N \setminus \{\emptyset, N\}$ **Output:** a linear extension I of P generated with a uniform distribution $H \leftarrow P$: Imin $\leftarrow \varnothing$: Imax $\leftarrow \varnothing$ While height of H > 2 do Compute the basic parameters of T_{H} : k, h, |I|Select $M \in T_H[h, k, |I|]$ with probability $\Pr(M \mid T_H[h, k, |I|])$ Add M at the beginning of ImaxCompute the basic parameters of B_H : k', h', |I'|Select $m \in B_H[h', k', |I'|]$ with probability $\Pr(m \mid B_H[h', k', |I'|])$ Add *m* at the end of *lmin* $H \leftarrow H \setminus \{M, m\}$

end while

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% Now *H* is reduced to two layers: B_H and T_H coincide While height of H = 2 do

If number of nodes in the upper layer \leqslant number of nodes in the lower layer then

Select $M \in T_H[h, k, |I|]$ with probability $\Pr(M \mid T_H[h, k, |I|])$ Add M at the beginning of *Imax* $H \leftarrow H \setminus \{M\}$

otherwise

Select $m \in B_H[h', k', |I'|]$ with probability $\Pr(m \mid B_H[h', k', |I'|])$ Add m at the end of *lmin*

$$H \leftarrow H \setminus \{m\}$$

end if

end while

% Now H is reduced to one layer, which is an antichain whose elements have % the same probability

While $H \neq \emptyset$ do

```
Select uniformly at random an element x \in H
```

```
Add x at the end of lmin
```

$$H \leftarrow H \setminus \{x\}$$

end while

```
I \leftarrow \mathit{Imin} ; concatenate \mathit{Imax} to the end of I
```



Random generation of capacities


3

Computation of the probabilities

It remains to compute $Pr(M | T_H)$ and $Pr(m | B_H)$. This is made possible through a simplifying assumption.

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Definition

Let x be a node of the upper layer of T_H .

The function f_x assigns to every node y of the lower layer an integer as follows:

$$f_x(y) = egin{cases} |\mathsf{pred}(y)|, & y \in \mathsf{succ}(x) \ 0, & ext{otherwise}. \end{cases}$$

The function n_x : N → N is defined from f_x as follows: n_x(r) is the number of occurences of f_x(y) = r, i.e., n_x(r) = |f_x⁻¹(r)|. When r > 0, it is the number of successors of x having r predecessors, otherwise when r = 0 it is the number of nodes in the lower layer which are not successors of x.

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Computation of the probabilities

Definition

We say that T_H is *regular* if n_x is invariant with x, i.e., $n_x(r) = n_{x'}(r)$ for every r and every two nodes x, x' in the upper layer.

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Definition

We say that T_H is *regular* if n_x is invariant with x, i.e., $n_x(r) = n_{x'}(r)$ for every r and every two nodes x, x' in the upper layer.



 $n_{124}(0) = 3 = n_{234}(0), n_{124}(1) = 2 = n_{234}(1), n_{124}(2) = 1 = n_{234}(2),$ hence the above T_H is regular. Every T_H closed under intersection and balanced is regular. Dual definitions exist for B_H .

Proposition

Consider the poset $T_H[h, k, |I|]$ and suppose that it is regular. Then the probabilities $\mathbb{P}_u(T_H[h, k, |I|])$ that node x of the upper layer terminates a linear extension, and $\mathbb{P}_l(T_H[h, k, |I|])$ that isolated node y of the lower layer terminates a linear extension are given by

$$\begin{split} \mathbb{P}_{u}(T_{H}[h, k, |l|]) &= \frac{1}{h} \frac{\prod_{i=1}^{|l'|} (h - 1 + k - |l'| + i)}{\prod_{i=1}^{|l'|} (h - 1 + k - |l'| + i) + |l| \times \prod_{i=1}^{|l''|} (h - 1 + k - |l'| + i) \prod_{i=1}^{|l|-1} (h + k - |l| + i)} \\ \mathbb{P}_{l}(T_{H}[h, k, |l|]) &= \frac{\prod_{i=1}^{|l''|} (h - 1 + k - |l'| + i) \prod_{i=1}^{|l|-1} (h + k - |l| + i)}{\prod_{i=1}^{|l'|} (h - 1 + k - |l'| + i) + |l| \times \prod_{i=1}^{|l''|} (h - 1 + k - |l'| + i) \prod_{i=1}^{|l|-1} (h + k - |l| + i)}, \end{split}$$

where I' is the set of isolated nodes in the poset $T_{H\setminus\{x\}}$, and $I \cup I'' = I'$.

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Outline

- 1. The problem of uniform random generation
- 2. The 2-layer approximation method
- **3. Measure of performance**

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Uniform distribution of μ in C(N) does not mean that the distribution of μ(S) for a given S is uniform!

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- Denote by μ the r.v. with uniform distribution on C(N). Take a linear extension σ and consider the associated region R_σ ⊆ C(N).

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- Uniform distribution of μ in C(N) does not mean that the distribution of μ(S) for a given S is uniform!
- Exact distribution of $\mu(S)$ seems to be very hard to obtain.
- Denote by μ the r.v. with uniform distribution on C(N). Take a linear extension σ and consider the associated region R_σ ⊆ C(N).
- Given that µ ∈ R_σ, we know that µ(S_{σ(k)}) follows the distribution of the kth order statistics on [0, 1]. It is known that the probability density function f_(k) of the kth order statistics on [0, 1] when the underlying 2ⁿ − 2 random variables are i.i.d. and uniform is a Beta distribution:

$$f_{(k)}(u) = (2^n - 2) \binom{2^n - 3}{k - 1} (1 - u)^{2^n - 2 - k} u^{k - 1} = \text{Beta}(k, 2^n + k - 1)$$

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Denoting by OS_k the corresponding cumulative distribution function, it follows that for any $S \in 2^N \setminus \{\emptyset, N\}$, the distribution $F_{\mu(S)}(\alpha)$ is given by

$$F_{\mu(S)}(\alpha) = \Pr(\mu(S) \leq \alpha) = \sum_{\sigma \in E(2^N \setminus \{\emptyset, N\})} \Pr(\mu(S) \leq \alpha \mid \mu \in R_{\sigma}) \Pr(\mu \in R_{\sigma})$$
$$= \frac{1}{e(2^N)} \sum_{\sigma \in E(2^N \setminus \{\emptyset, N\})} \Pr(\mu(S) \leq \alpha \mid \mu \in R_{\sigma})$$
$$= \frac{1}{e(2^N)} \sum_{\sigma \in E(2^N \setminus \{\emptyset, N\})} \operatorname{OS}_{k(S, \sigma)}(\alpha), \tag{1}$$

where $E(2^N \setminus \{\emptyset, N\})$ is the set of permutations corresponding to linear extensions, and $k(S, \sigma)$ is such that $S = S_{\sigma(k)}$.

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Lemma

Assume μ is uniformly distributed and take $\varnothing \neq S, S' \subset N$. Then

- **(**) $\mu(S)$ and $\mu(S')$ for |S| = |S'| are identically distributed.
- **2** $\mu(S)$ and $1 \mu(N \setminus S)$ are identically distributed.

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Figure: Histograms of $\mu(S)$ for n = 4 and the exact method

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Figure: Histograms of $\mu(S)$ for n = 4 and the 2-layer approximation method

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Figure: Histograms of $\mu(S)$ for n = 4 and the Markov chain method

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Figure: Histograms of $\mu(S)$ for n = 4 and the Random Node Generator

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$$c = \sum_{\sigma \in E(2^N)} b_{\sigma}$$

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 - $c(N \setminus S) = 1 c(S)$ and c(S) depends on |S| only.

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- centroid for n = 3:

c = (0.298, 0.298, 0.298, 0.702, 0.702, 0.702).

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• centroid for n = 4:

c = (0.1810, 0.1810, 0.1810, 0.1810, 0.5000, 0.5000, 0.5000, 0.5000, 0.5000, 0.5000, 0.8190, 0.8190, 0.8190, 0.8190).

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 Idea: compare with the Markov chain method (asymptotically exact when T tends to infinity), by choosing T so that the performance of the 2-layer approximation method is approximately the same as the one of the Markov chain method

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- When n ≤ 4, we take as performance the L₁ distance between the theoretical centroid and the obtained centroid. We obtain T = 1170.
- When n > 4, we use the symmetry properties of the centroid (c(S) depends only on |S|). The performance is measured by the standard deviation of c(S) when |S| is constant. We obtain T = 9000 for n = 5.

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- Given two discrete probability distributions *p*, *q* on the same universe *X*, the *Kullback-Leibler divergence* is defined as

$$\mathbb{D}_{\mathcal{K}L}(p||q) = \sum_{x \in X} p(x) \log \frac{p(x)}{q(x)}$$

The smaller the value, the closer are the two distributions.

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The smaller the value, the closer are the two distributions.

• Distributions of $\mu(S)$ are discretized with $\delta = 0.01$ on [0, 1]. We call $\mu_{MC}(S), \mu_{2L}(S)$ the discrete distributions obtained by the Markov chain method and the 2-layer approximation, respectively.

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 With n ≤ 4, q is the exact distribution and p is the distribution to be tested. We compute

$$S^{4}_{\mathcal{KL}}(\mu_{MC}) = \sum_{S \in 2^{N}} \mathbb{D}_{\mathcal{KL}}(\mu_{MC}(S) || \boldsymbol{\mu}(S))$$
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• With *n* > 4, we use symmetry properties of the distributions. We compute

$$S_{KL}^{N}(\mu_{MC}) = \sum_{S,S' \in 2^{N} \text{ s.t. } |S| = |S'|} \mathbb{D}_{KL}(\mu_{MC}(S) || \mu_{MC}(S'))$$

$$S_{KL}^{N}(\mu_{2L}) = \sum_{S,S' \in 2^{N} \text{ s.t. } |S| = |S'|} \mathbb{D}_{KL}(\mu_{2L}(S) || \mu_{2L}(S'))$$

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Results

$S_{KL}^4(\mu_{MC})$	$S^4_{KL}(\mu_{2L})$	$S_{KL}^5(\mu_{MC})$	$S_{KL}^5(\mu_{2L})$
0.061	0.059	2.41	2.24

Comparison of CPU time (s) for generating 10,000 capacities ($3.2~\mbox{GHz}$ PC with 16 GB of RAM)

Method		<i>n</i> = 4	<i>n</i> = 5	<i>n</i> = 6	<i>n</i> = 7
2 layer approximation		2.58	11.51	60.06	330.17
Markov Chain	CPU time	20.46	161.33	pprox 1500	pprox 9000
	Т	1170	9000	80,000	500,000

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- Naive methods yield poor results
- Good methods try to generate a representative sample of linear extensions: Markov chain method, 2-layer approximation
- The Markov chain method and the 2-layer approximation method yield similar results, with high accuracy.
- The 2-layer approximation method is much faster.

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Thank you for your attention!

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