

Top-m identification for linear bandits

Motivated by drug repurposing, we propose a generic family of algorithms to tackle the identification of the $m \geq 1$ arms with largest means in a linear bandit model.

Fixed-confidence linear Top- m identification

The goal is to identify the $m < K$ arms with the largest expected rewards, using fixed time-independent arm contexts. In a linear bandit model, the expected reward μ_a of each arm a is a linear function of context $x_a \in \mathbb{R}^N$

for any arm a , $\mu_a = \theta^T x_a$ for some vector $\theta \in \mathbb{R}^N$

A (ϵ, m, δ) -PAC (Probably Approximately Correct) algorithm for fixed-confidence linear Top- m returns a subset of m arms \hat{S}_m^t at round t s.t. if $\mu_{(m)}$ is the m^{th} largest mean reward:

$\text{Prob}(\hat{S}_m^t \subseteq S^{*\epsilon}) \geq 1 - \delta$

where $S^{*\epsilon} = \{a : \mu_a \geq \mu_{(m)} - \epsilon\}$

Contributions

GIFA (Gap-Index Focused Algorithm) family for Top- m

We propose a generic family of algorithms aimed at Top- m identification, which encompasses known algorithms for Top- m and best arm identification (LUCB¹, UGapE², LinGapE³, ...). These algorithms rely on good gap indices $B_{ij}(t)$ for any arm pair (i, j) s.t.

$\text{Prob} \{ \forall t > 0, \forall j \in (S^{*\epsilon})^c, \forall k \in S^{*0}, B_{kj}(t) \geq \mu_k - \mu_j \} \geq 1 - \delta$

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1:  $t \leftarrow 1$ 
2: while  $\text{stopping}(t) > \epsilon$  do
3:   // estimated m best arms at t
4:    $J(t) \leftarrow \text{compute\_Jt}(t)$ 
5:   // estimated m-best arm at t
6:    $b_t \leftarrow \text{compute\_bt}(t)$ 
7:   // challenger to  $b_t$  at t
8:    $c_t \leftarrow \arg \max_{a \notin J(t)} B_{a,b_t}(t)$ 
9:   // sampling arms
10:   $l_t \leftarrow \text{selection}(b_t, c_t)$ 
11:   $r_t \leftarrow \text{sample}(l_t)$ 
12:  Update gap indices  $B_{i,j}(t+1)$ 
13:   $t \leftarrow t + 1$ 
14: end while
15:  $\hat{S}_m^t \leftarrow J(t)$ 
16: return  $\hat{S}_m^t$ 
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Alg 1: General structure of a GIFA algorithm

* Correctness for partially specified GIFA algorithms:

Theorem 1. If using good gap indices, (i) any GIFA algorithm with $b_t = \arg \max_{j \in J(t)} \max_{i \in J(t)} B_{ij}(t)$ and the LUCB stopping rule, or (ii) any GIFA algorithm using $b_t \in J(t)$ with the UGapE stopping rule, is (ϵ, m, δ) -PAC.

* Existence of good gap indices:

Lemma 3. Gap indices using confidence bounds for linear bandits in ⁴ are good gap indices.

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Proposals for linear Top- m identification: m -LinGapE and LinGIFA

$\mu_i(t)$ is the empirical mean for arm i at time t .

Algorithm	compute_Jt	compute_bt	selection	stopping at t
m -LinGapE	$\underset{j \in [K]}{\text{argmax}} \mu_j(t)$	$\underset{j \in J(t)}{\text{argmax}} \max_{i \in (J(t))^c} B_{ij}(t)$	greedy, optimized	$B_{ct, b_t}(t) \leq \epsilon$ (LUCB rule)
LinGIFA	$\underset{j \in [K]}{\text{argmin}} \max_{i \neq j} B_{ij}(t)$	$\underset{j \in J(t)}{\text{argmax}} \max_{i \neq j} B_{ij}(t)$	largest variance, greedy	$\max_{j \in J(t)} \max_{i \neq j} B_{ij}(t) \leq \epsilon$ (UGapE rule)

Tab. 1: New proposals for linear Top- m identification

where

Greedy³ $\underset{a \in [K]}{\text{argmin}} (x_{ct} - x_{bt})^T (V_{t-1} + x_a x_a^T)^{-1} (x_{ct} - x_{bt})$
Optimized³ $\underset{a \in [K]}{\text{argmax}} n_a(t) \|w^*(b_t, c_t)\|_1 / |w^*(b_t, c_t)|$
and $w^*(b_t, c_t) = \underset{w \in \mathbb{R}^K}{\text{argmin}} \{ \|w\|_1 \mid x_{bt} - x_{ct} = \sum_{a \in [K]} w_a x_a \}$
Largest variance² $\underset{a \in \{ct, bt\}}{\text{argmax}} x_a^T V_t x_a$
and $V_t = \lambda I_N + \sum_{a \in [K]} n_a(t) x_a x_a^T$, $n_a(t)$: # of times a pulled until round t

Unified sample complexity analysis for LUCB-*like* GIFA algorithms

This subclass of algorithms comprises LUCB, LinGapE and m -LinGapE (same rules for estimating $J(t)$, b_t and stopping). Analysis leads to upper bounds on stopping time of the form

$C(H) = \inf_{u>0} \{ u > 1 + H \cdot D(\delta, u) + O(K) \}$

* this term corresponds to an initialization phase where all arms are sampled once, $D(\delta, u)$ controls the width of the gap indices

Algorithm	Complexity constant $H^\epsilon(\mu)$
LUCB	$2 \sum_{a \in [K]} \max(\epsilon/2, \Delta_a)^{-2}$ $\Delta_a = \mu_a - \mu_{(m+1)}$ if $a \in S^{*0}$, $\mu_{(m)} - \mu_a$ otherwise
m -LinGapE (largest variance rule)	$4\sigma^2 \sum_{a \in [K]} \max(\epsilon, (\epsilon + \Delta_a)/3)^{-2}$ σ is the variance on the noise model
m -LinGapE (optimized rule)	$\sigma^2 \sum_{a \in [K]} \max_{i,j} w_a^*(i,j) (\epsilon + \max(\Delta_i, \Delta_j)/3)^{-2}$

Tab. 2: Comparaison between upper bounds on stopping time for LUCB-*like* algorithms



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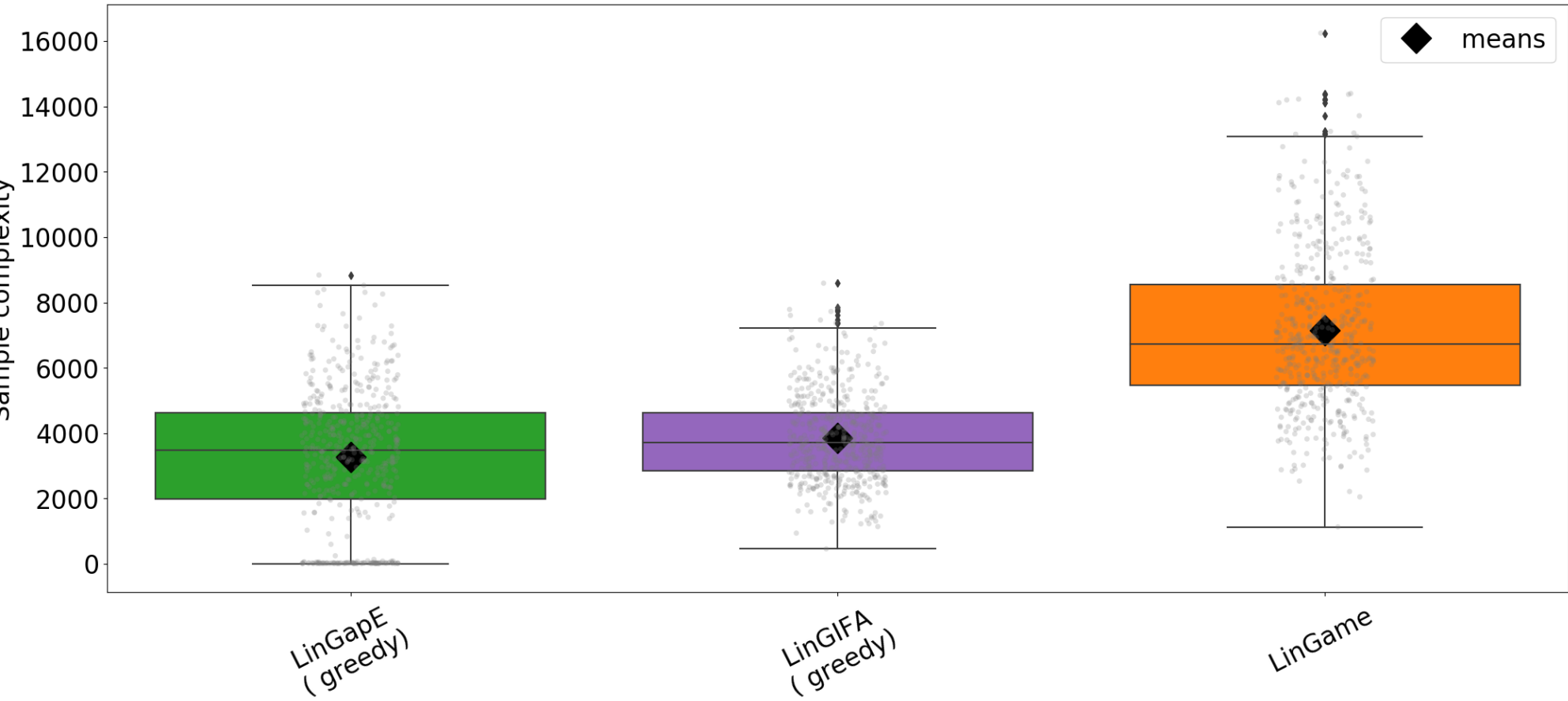
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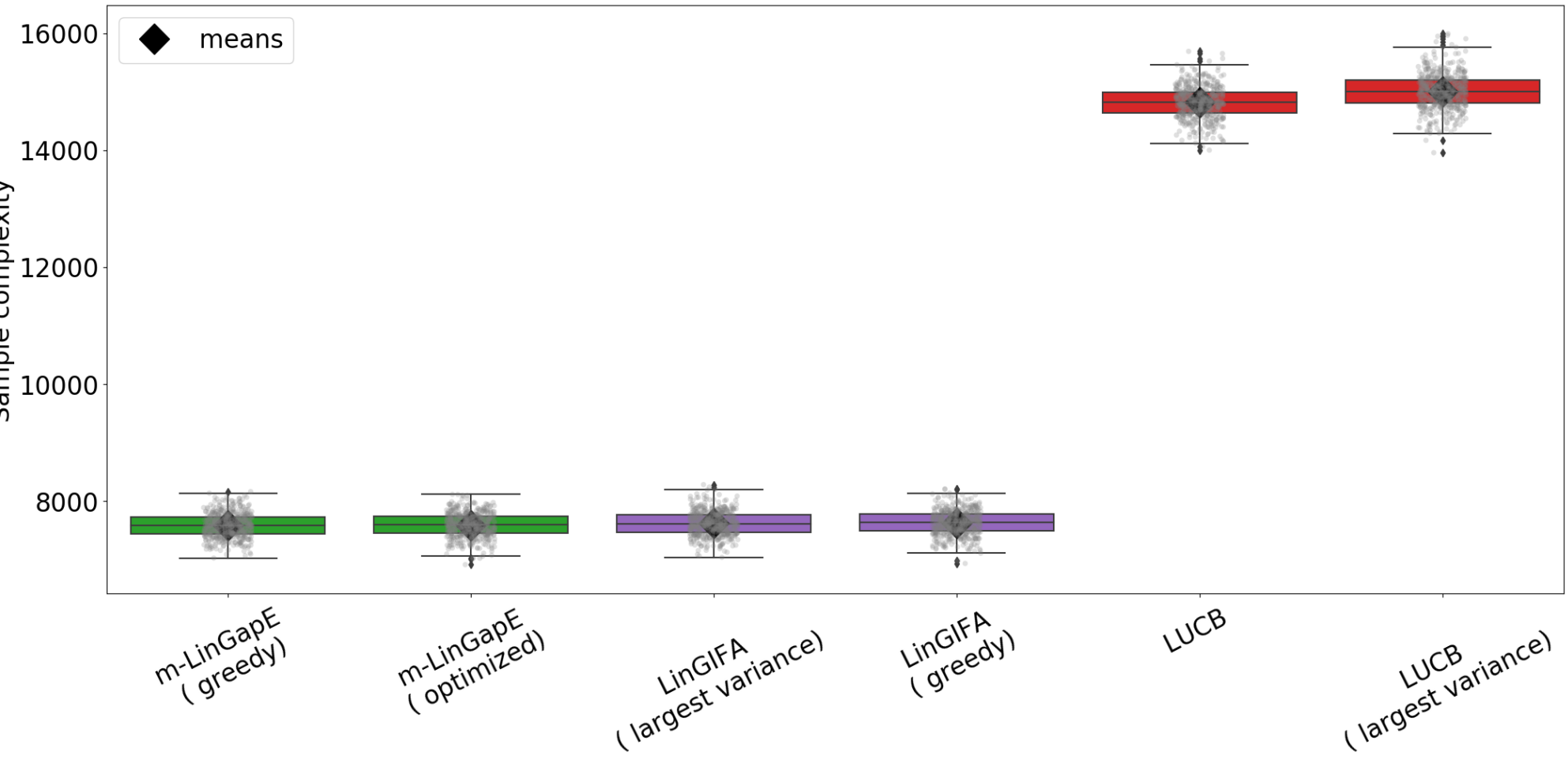
Experimental study (across 500 simulations)

Simulated data: hard best arm (Top-1) identification instance⁵



m -LinGapE and LinGIFA remain competitive even against asymptotically optimal (i.e., when $\delta \rightarrow 0$) such as LinGame. ($K=3$, $\mu_{(1)} - \mu_{(2)} = \cos(0.1)$, $N=3$)

Biological data: application to drug repurposing with $m > 1$



In a drug repurposing instance for epilepsy ($m = 5$), sample complexity is improved by a factor $\frac{1}{2}$ compared to LUCB. ($K=10$, $\mu_{(m)} - \mu_{(m+1)} \sim 0.066$, $N=71$)

References

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