

# Constrained Binary Identification Problem

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## Abstract

We consider the problem of building a binary decision tree, to locate an object within a set by way of the least number of membership queries. This problem is equivalent to the “20 questions game” of information theory and is closely related to lossless source compression. If any query is admissible, Huffman coding is optimal with close to  $H[P]$  questions on average, the entropy of the prior distribution  $P$  over objects. However, in many realistic scenarios, there are constraints on which queries can be asked, and solving the problem optimally is NP-hard.

We provide novel polynomial time approximation algorithms where constraints are defined in terms of “graph”, general “cost”, and “submodular” functions. In particular, we show that under graph constraints, there exists a constant approximation algorithm for locating the target in the set. We then extend our approach for scenarios where the constraints are defined in terms of general cost functions that depend only on the size of the query and provide an approximation algorithm that can find the target within  $O(\log(\log n))$  gap from the cost of the optimum algorithm. Submodular functions come as a natural generalization of cost functions with decreasing marginals. Under submodular set constraints, we devise an approximation algorithm that can find the target within  $O(\log n)$  gap from the cost of the optimum algorithm. The proposed algorithms are greedy in a sense that at each step they select a query that most evenly splits the set without violating the underlying constraints. These results can be applied to network tomography, active learning and interactive content search.

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## 1 Introduction

Constructing a binary search tree is one of the fundamental problems in discrete mathematics. Formally, we are given a set  $\mathcal{N} = \{1, \dots, n\}$  of objects, identified with integers from 1 to  $n$ , as well as a probability distribution  $P = (p_1, \dots, p_n)$  over  $\mathcal{N}$ ,  $p_i \geq 0$ ,  $i = 1, \dots, n$ , and  $\sum_{i=1}^n p_i = 1$ . The goal is to locate an object within a set  $\mathcal{N}$  by way of the least number of membership queries. Each binary split is a partition of  $\mathcal{N}$  into a subset  $Q \subset \mathcal{N}$  and its complement  $\mathcal{N} \setminus Q$ . Given that a membership oracle provides answers without noise, what is the best we can do in a computationally feasible manner? This problem is equivalent to the “twenty questions game” of information theory [7], where a player has to determine the identity of an object from a set (say, a famous person) by asking a minimum number of yes/no questions. If any split is an admissible query, Huffman coding is optimal with close to  $H[P]$  questions on average [7], the entropy of the distribution  $P$  from which the target object is drawn at random.



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However, in realistic scenarios, there are *constraints* on which queries can be asked, and solving the problem optimally is NP-hard [15]. Instances of these constraints arise frequently in practice such as network tomography and interactive content search.

### Network Tomography

A frequent situation that arises in sensor network tomography can be briefly explained as follows [4]. Suppose we are to remotely maintain a sensor network, whose nodes are severely restricted. First, being connected in a graph  $G$ , they can communicate with their neighbours only. Second, although they can receive broadcast messages from a base station, communication from any node back to the base must be kept to an absolute minimum due to power constraints. As nodes fail regularly, it is necessary to periodically search for faults. For simplicity, we assume that at most one node is faulty. To minimize the communication back to base, we should use the least number of broadcasts of the form “To nodes in  $Q$ : Is one of you faulty?”. If  $Q$  is a connected subgraph of  $G$ , the question can be answered by a simple local message passing protocol, where a designated root in  $Q$  reports positively to the base only if it receives messages from all neighbours in  $Q$ , etc. Given that this does not happen, the root can poll the branch in question, whereby the faulty node can be identified recursively and passed back to the root, which reports it back to base. Finally, if the base does not get any reply after a reasonable time, it concludes that the chosen root must be faulty.

We address the constraints of this type under the general and realistic notion of *graph constraints* where we assume that the set  $\mathcal{N}$  is furnished with an undirected graph structure  $G$ , and  $Q \subset \mathcal{N}$  is an admissible query if and only if the subgraph  $G|_Q$  induced by  $Q$  (containing all edges of  $G$  between vertices in  $Q$ ) is connected. We then provide an efficient constant factor approximation to the “graph-constrained” twenty questions problem, which finds the target in  $\leq 4H[P] + 2$  queries on average.

### Interactive Content Search

In content search applications with humans involved, it is of obvious value to keep the number of queries as small as possible, as human interventions are disproportionately costly. Examples include visual recognition [2] and pattern classification [11], where questions are restricted to simple visual attributes. Whenever people are in the loop, queries are limited by their ability to meaningfully disambiguate in the presence of many attributes. As these examples show, it is usually not practical to allow for *any* query  $Q \subset \mathcal{N}$ . In these scenarios, it is more natural to define the constraints in terms of a cost rather than through a graph.

More formally, we are given a *non-decreasing cost function*  $C : \{1, 2, \dots, n\} \rightarrow \mathbb{R}$  where the cost of making an oracle query on set  $S$  is  $C(|S|)$ . Notice that the cost only depends on the size of the set and not the elements it contains. When a human plays the role of the membership oracle, it is more difficult to answer a query with a bigger set than one with a smaller set [2]. This is why in many interactive search strategies such a cost function arises naturally. In this case, we provide an approximation algorithm that can find the target within  $O(\log(\log n))$  gap from the cost of the optimum algorithm.

To generalize our analysis beyond non-decreasing cost functions, we consider the case where constraints are defined in terms of a *submodular function*  $Cost : 2^{\mathcal{N}} \rightarrow \mathbb{R}$ . In this setting, for all subsets  $S_1 \subseteq S_2 \subseteq \mathcal{N}$  and an item  $i \in \mathcal{N}$  we have  $Cost(S_1 \cup \{i\}) - Cost(S_1) \geq Cost(S_2 \cup \{i\}) - Cost(S_2)$ . In words, adding a new item  $i$  to a larger set should produce an incremental cost no more than adding  $i$  to a smaller set. Under submodular constraints,

we provide an approximation algorithm that can find the target within  $O(\log(n))$  gap from the cost of the optimum algorithm. Further applications of submodular functions in active learning can be found in [12].

## 2 Related Work

The problem studied here can be seen as a special case of the binary identification problem (BIP) [10]. Suppose that we are given a set of objects  $\mathcal{N}$  and a set of tests  $\{t_1, \dots, t_k\}$  where one of the objects is marked. Each test determines whether the marked object is in the test set or not. The goal is to define a strategy that minimizes the number of tests to find the marked object. It is known that both the average case minimization and worse case minimization are NP-complete [15]. Moreover, it is even NP-Hard to have an  $o(\log n)$ -approximation algorithm for the average case [5]. In both cases, there exist heuristic algorithms that admit  $O(\log n)$ -approximation [20, 6]. The closest work to ours is [14] where the authors study the same problem in a setting that a cost is associated with each test and the goal is to minimize the average total cost. They propose an  $O(\log n)$ -approximation algorithm. Unfortunately, there is no straight way to follow their approach and obtain similar performance guarantees for the submodular cost functions (for the other two problems that we consider in this paper, our approximation factors are better). Note that in this case, there are  $2^n$  number of tests and a naive reduction admits an exponential running time. To overcome this barrier we propose a novel algorithm with a similar approximation guarantee.

Adding some structure to the set of tests leads to interesting special cases. For instance, if we let the set of tests be the power set of objects, then the optimal average case strategy is attained by Huffman coding [7]. Another basic variant of BIP is finding a marked element in a totally ordered set, a problem that is very well studied in the literature [19]. This can be generalized to searching in structures where the input has a partial order between its elements instead of a total order [1, 3]. It is known that searching in posets for the worst case minimization is NP-hard [3]. However, in [20], the authors showed a greedy algorithm with  $O(\log n)$  approximation which was further improved to a constant factor approximation by [21]. It has been recently shown in [17] that the average case minimization is also NP-hard for the class of trees with diameter at most 4.

Constrained BIP is strongly related to *active learning* [8, 22, 13, 9, 18] in which a *hypothesis space*  $\mathcal{H}$  is defined as a set of binary valued functions defined over a finite set  $\mathcal{Q}$ , called the *query space*. Each hypothesis  $h \in \mathcal{H}$  generates a label from  $\{-1, +1\}$  for every query  $q \in \mathcal{Q}$ . A target hypothesis  $h^*$  is sampled from  $\mathcal{H}$  according to some prior  $P$ ; asking a query  $q$  amounts to revealing the value of  $h^*(q)$ , thereby restricting the possible candidate hypotheses. The goal is to determine  $h^*$  by asking as few queries as possible. In our setting, the hypothesis space  $\mathcal{H}$  is the set  $\mathcal{N}$ , and the query space  $\mathcal{Q}$  is the set of all admissible subsets. The target hypothesis sampled from  $P$  is the target  $t_*$ . A well-known algorithm for determining the true hypothesis in the active-learning setting is the *generalized binary search* (GBS) or *splitting* algorithm [8, 22, 9]. Define the *version space*  $V \subseteq \mathcal{H}$  to be the set of possible hypotheses that are consistent with the query answers observed so far. At each step, GBS selects the query  $q \in \mathcal{Q}$  that minimizes  $|\sum_{h \in V} P(h)h(q)|$ . Recently, Golovin and Krause [12] showed that GBS makes at most  $OPT \cdot (H_{\max}(\mu) + 1)$  queries in expectation to identify hypothesis  $h^* \in \mathcal{N}$ , where  $OPT$  is the minimum expected number of queries made by any adaptive policy and  $H_{\max}(P) = \max_{x \in \text{supp}(P)} \log \frac{1}{P(x)}$ . In our setting, the version space  $V$  comprises all possible objects in  $z \in \mathcal{N}$  that are consistent with answers given so far. Under graph, cost, and submodular constraints, we replace  $H_{\max}$  (which in practice can be

**Algorithm 1** The role of size

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**Input:** Connected constraint graph  $G = (\mathcal{N}, \mathcal{E})$ , root node  $v_R \in \mathcal{N}$ .  
 Grow a connected subtree  $T \subset G$ , starting from  $v_R$ , by a breadth-first search (BFS), until  $T$  spans  $\lceil n/2 \rceil$  vertices.  
 Query the vertex set of  $T$  (size  $\lceil n/2 \rceil$ ).  
**if**  $I_{\{t_* \in T\}} = 1$  **then**  
   Recurse on  $T$ , root  $v_R$ .  
**else**  
   Collapse all nodes in  $T$  into a single node  $v_T$ , which is connected to all  $v \in \mathcal{N}$  adjacent to nodes in  $T$  in the original  $G$ . Create  $G'$  by connecting all neighbours of  $v_T$  to one another and removing  $v_T$ . Recurse on  $G'$ .  
**end if**

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quite large) by a constant,  $O(\log(\log(n)))$ , and  $O(\log(n))$ , respectively.

The use of interactive methods for searching in a dataset has a long history in literature. Relevance feedback [24] is a method for interactive image retrieval, in which users mark the relevance of image search results, which then used to create a refined search query. We use the membership oracle to model the role of a user for identifying a target in a database. In practice, the cost of a query depends on the characteristics of the query, e.g., its size [11, 2]. However, due to the lack of analytical results, in many such applications only heuristic methods were proposed. To close this gap, we introduce general constraints, in terms of graph, cost and submodular functions, on the set of queries and establish analytical guarantees associated with them.

### 3 Graph Constraints

Let us assume that the set  $\mathcal{N}$  is endowed with undirected graph structure,  $G = (\mathcal{N}, \mathcal{E})$ , where  $\mathcal{E}$  contains the edges. For any subset  $A \subset \mathcal{N}$ ,  $G - A$  denotes the graph obtained from  $G$  by removing all vertices  $A$  and all edges adjacent to  $A$ , while  $G|_A$  is the graph induced by  $A$  (vertex set  $A$ , edge set those  $e \in \mathcal{E}$  between vertices in  $A$ ).

Given  $G$ , a query  $Q \subset \mathcal{N}$  is *admissible* if and only if the graph  $G|_Q$  is connected. Our detection algorithm can submit any admissible query  $Q$  to a membership oracle, which initially sampled the target  $t_*$  at random from  $P$ : it will answer by one bit of information,  $I_{\{t_* \in Q\}} = 1$  if  $t_* \in Q$ , 0 otherwise. Our goal is to detect  $t_*$  with as few queries to the oracle as possible. Here, we restrict ourselves to deterministic policies which terminate only once the version space for  $t_*$  consistent with all previous queries  $Q_1, \dots, Q_k$  (formally,  $(\cap_{k|t_* \in Q_k} Q_k) \cap (\cap_{k|t_* \notin Q_k} (\mathcal{N} \setminus Q_k))$ ) reaches size one. Throughout this section, we assume that the algorithm knows the distribution  $P$  the target is drawn from.

We begin with Algorithm 1, a simple divide-and-conquer scheme which detects the target with no more than  $\lceil \log n \rceil$  queries, if seeded with an arbitrary  $v_R \in \mathcal{N}$  as root node. Here and elsewhere, we use binary logarithms (base 2). Essentially, this algorithm mirrors the principle of binary search by way of intermediate sets produced during a breadth-first-search (BFS). Given adequate backpointer structures, its running cost is that of a single BFS.

► **Lemma 1.** *Presented with a connected constraint graph  $G = (\mathcal{N}, \mathcal{E})$ , where  $|\mathcal{N}| = n$ , Algorithm 1 detects the target  $t_*$  with no more than  $\lceil \log n \rceil$  queries to the membership oracle.*

While Algorithm 1 is simple and efficient, it has the obvious drawback of not exploiting knowledge about the distribution  $P$  at all. However, it is useful as subroutine for our main

algorithm developed next.

Suppose w.l.o.g. that  $P = (p_1, \dots, p_n)$  is such that  $p_1 \geq p_2 \geq \dots \geq p_n$ . Define the nested subsets  $A_1 \subset A_2 \subset \dots \subset \mathcal{N}$  by

$$A_i = \{1, \dots, j_i\}, \quad j_i = \min \left\{ j \mid \sum_{k>j} p_k \leq 2^{-i} \right\}.$$

Put differently,  $A_i$  is the smallest subset of  $\mathcal{N}$  so that  $\sum_{k \in A_i} p_k \geq 1 - 2^{-i}$ . Also, define  $a_i = \log(1/p_{j_i})$ , so that  $p_k \geq 2^{-a_i}$  for all  $k \in A_i$ , moreover  $a_0 = 0$ . The intuition behind this choice is that  $A_1$  contains about half of the probability mass,  $A_2 \setminus A_1$  half of the remaining mass, and so on.

Our algorithm processes the  $A_i$  in order,  $i = 1, 2, \dots$ . For  $A_i$ , it calls Algorithm 1, which will detect  $t_*$  if it is in  $A_i$ , otherwise we move to the next set. However, we cannot simply pass the induced subgraph  $G|_{A_i}$  to Algorithm 1, as it may well not be connected. In this case, we generate  $G_i = (A_i, V_i)$  passed to the algorithm as follows. We initialize  $G_i \leftarrow G|_{A_i}$  and cluster the nodes into connected components. We then join each pair of disjoint components by a shortest path  $\pi$ , a central part of which necessarily features nodes  $V(\pi)$  with  $V(\pi) \cap A_i = \emptyset$ . We collapse this part of  $\pi$  into a new *virtual* edge between nodes in  $A_i$ , which is labelled by the vertices  $V(\pi)$  and added to  $V_i$ . This process stops when  $G_i$  is connected. Finally, when running Algorithm 1 on  $G_i$ , we need to translate queries back to connected subgraphs of the original  $G$ . For a query  $Q \subset A_i$  in question, this is done by adding nodes with which virtual edges of  $G_i|_Q$  are labelled. Our construction of  $G_i$  from  $A_i$  and labeling of virtual edges ensures that any query processed in this way corresponds to a connected subgraph of  $G$ , therefore is admissible.

Note that due to the presence of virtual edges, Algorithm 1 may receive positive answers from the oracle, even though  $t_* \notin A_i$ . After all,  $t_*$  might be among the vertices on virtual edges. However, in this case, Algorithm 1 simply descends to a single vertex  $v_i \in A_i$ , so that one more query  $\{v_i\}$  settles the question “ $t_* \in A_i$ ”. Our main result is as follows.

► **Theorem 2.** *Given a connected constraint graph  $G$  with  $n$  vertices and any distribution  $P$  over  $\{1, \dots, n\}$ , our algorithm finds a target  $t_*$  sampled at random from  $P$  with no more than  $2 + 4\mathbb{H}[P]$  admissible queries, on average over draws of  $t_*$ .*

**Proof.** We prepare our proof with a lemma whose proof can be found in the full version.

► **Lemma 3.** *For the numbers  $a_k$  defined above, we have that  $\sum_{k=1}^{\infty} a_k/2^{k+1} \leq \mathbb{H}[P]$ .*

To establish Theorem 2, recall that we visit sets  $A_i$  in order,  $i = 1, 2, \dots$ , calling Algorithm 1 on  $G_i = (A_i, V_i)$  constructed as detailed above. Since  $p_j \geq 2^{-a_i}$  for all  $j \in A_i$ , the size of  $A_i$  is bounded by  $2^{a_i}$ , and Algorithm 1 returns after  $\leq a_i$  queries. As noted above, due to “virtual edge” complications, we may have to query one more single node, yet after  $\leq a_i + 1$  questions we know whether  $t_* \in A_i$  or not, and in the former case will have detected  $t_*$ . Now, the probability of not finding  $t_*$  in  $A_i$  or earlier is bounded by  $\sum_{j|p_j < 2^{-a_i}} p_j \leq 2^{-i}$ . This means that the expected number of queries in our algorithm is at most  $\sum_{i \geq 1} (a_i + 1)/2^{i-1} \leq 2 + 4\mathbb{H}[P]$ , where this inequality is due to Lemma 3. ◀

## 4 Cost Function Constraints

In this section we analyze another variant of the binary identification problem where the constraints are defined in terms of a cost function rather than a graph. More formally, we are given a non-decreasing cost function  $C : \{1, 2, \dots, n\} \rightarrow \mathbb{R}$  where the cost of making an

**Algorithm 2 Binary Identification Algorithm with Cost Function Constraints**

**Input:**  $n$  objects with a probability distribution on them, and a cost function  $C : \{1, 2, \dots, n\} \rightarrow \mathbb{R}$ , fixed constant  $\epsilon > 0$ .

Create clusters  $S_1, S_2, \dots, S_l$  for  $l = \log(n^2/\epsilon)$ .

(Phase one) Use Procedure 3 to determine which cluster contains the target  $t_*$ .

(Phase two) If cluster  $S_i$  contains the target, find it by using the dynamic program 2.

(Phase three) If the target is not found in any of the above clusters, query each of the non-clustered objects.

oracle query on set  $S$  is  $C(|S|)^1$ . Notice that the cost only depends on the size of the set not the elements it contains. When a human plays the role of the membership oracle, it is more difficult to answer a query with a bigger set than one with a smaller set. This is why in many interactive search strategies such a cost function arises naturally. To avoid confusion in using term "cost" for queries and algorithms, we formally define the notion of cost for sets and algorithms as follows.

► **Definition 4.** We refer to the cost of making a query on a set  $S$  by the "cost of set  $S$ ". On the other hand, we use the term "expected search cost" of an algorithm to represent the expected value of total cost of queries the algorithm makes. Formally, An algorithm  $A$  consists of a family of possible queries  $\mathcal{F}_A$ . Algorithm  $A$  asks each query  $S \in \mathcal{F}_A$  with probability  $\Pr(A, S)$  which is a function of both the algorithm  $A$ , and the probability distribution of objects,  $P$ . We define the expected search cost of algorithm  $A$  to be  $\sum_{S \in \mathcal{F}_A} \Pr(A, S)C(|S|)$ .

By the above definition, it makes sense to talk about the expected search cost of Algorithm 2 or any other algorithm such as the optimum algorithm. We can also refer to the expected cost associated with a part of Algorithm 2 (it has three phases) and we can similarly define its expected cost in each phase. Note that the expected search cost of an algorithm is exactly the expected value of its total cost according to distribution  $P$ .

► **Definition 5.** Let  $\epsilon > 0$  to be a small and fixed constant. We place objects in  $l = \log(n^2/\epsilon)$  clusters based on their probabilities as follows. Let  $S_i$  be the cluster of objects with probability in range  $(1/2^i, 1/2^{i-1}]$  for  $1 \leq i \leq l$ . For any  $1 \leq i \leq l$ , we define  $\Pr(S_i)$  to be the sum of probabilities of objects in cluster  $S_i$ , and we define  $\Pr[i, j] \doteq \sum_{x=i}^j \Pr(S_x)$  for  $1 \leq i \leq j \leq l$ . We let  $\Pr[i, j] = 0$  for  $i > j$ .

The choice of  $\epsilon$  in the above definition is for ensuring that non-clustered objects have negligible probabilities, namely, at most  $\epsilon/n^2$ .

Our algorithm for finding the target is shown in Algorithm 2. In phase one, we run a recursive algorithm that uses procedure *ClusterFinder* (shown in Algorithm 3) as a subroutine. The procedure *ClusterFinder* gets two numbers  $i$  and  $j$ , and finds the cluster containing the target only if the target is in one of the clusters  $\{S_i, S_{i+1}, \dots, S_j\}$ . More specifically, Algorithm 3 finds a number  $k$  and a collection of clusters  $Q$  to query as follows:

$$Q = \bigcup_{i \leq m \leq k} S_m, \quad k = \max \left\{ k' \mid \Pr[i, k'] < \frac{\Pr[i, j]}{2} \right\} \quad (1)$$

<sup>1</sup> Note that there are two ways to determine whether or not the target lies in some set  $S$ . We can query set  $S$  or we can query the complement set  $\mathcal{N} \setminus S$ . Here, we assume that the cost is always a function of the size of the queried set and it is the job of the algorithm to determine which set needs to be queried.

**Algorithm 3** ClusterFinder( $i, j$ )**Input:** Clusters  $S_i, S_{i+1}, \dots, S_j$ 

- 1: Find the number  $k$  and the set  $Q$  according to Equation 1. Query set  $Q$ .
- 2: If  $t_* \in Q$  call  $ClusterFinder(i, k)$ . Otherwise, query  $S_{k+1}$ .
- 3: If  $t_* \in S_{k+1}$  return  $S_{k+1}$ . Otherwise, call  $ClusterFinder(k+2, j)$ .

If the target is in  $Q$ , the algorithm calls procedure  $ClusterFinder(i, k)$ . Otherwise, it queries set  $S_{k+1}$ , and if the target is not there either, it calls  $ClusterFinder(k+2, l)$ . Note that in Eq. 1, we might have  $\Pr(S_i) = \Pr[i, i] \geq \Pr[i, j]/2$  which causes  $k$  to be  $i-1$ . In this case, the set  $Q$  will be  $\emptyset$ .

► **Definition 6.** The procedure calls of Algorithm 3 can be represented by a binary tree  $T$  as follows. The root node is the procedure  $ClusterFinder(1, l)$ . The left child of the root is  $ClusterFinder(1, k)$ , and its right child is  $ClusterFinder(k+2, l)$ . In the same fashion, we can define the rest of the tree nodes based on the recursive calls.

In phase two of Algorithm 2, we are given a cluster  $S_i$  that contains the target, and we want to find the target object. Let  $x$  be the number of objects in  $S_i$ . Our algorithm assumes that the probabilities of all objects in  $S_i$  are identical. Since they are in the same cluster, their probabilities are close to each other (we prove later that this assumption does not incur much cost for us). By this assumption we have symmetry among objects in cluster  $S_i$ . Therefore, the only relevant characterization of a subset of  $S_i$  is its size. We define a dynamic programming array  $A[j]$ , for  $1 \leq j \leq x$ , which is the expected search cost of optimal strategy to find the target given that the target is in a subset of  $S_i$  with size  $j$  under the assumption that these objects have the same probability, and therefore are identical. We can fill up the entries of array  $A$  as follows. It is clear that  $A[1]$  is zero. We can update each  $A[j]$  using the lower entries  $A[1], A[2], \dots, A[j-1]$  as follows:

$$A[j] = \min_{1 \leq j' < j} \{C(j') + A[j'](j'/j) + A[j-j']((j-j')/j)\}. \quad (2)$$

The optimal strategy chooses some  $1 \leq j' < j$ , and a subset of size  $j'$  among the remaining objects (it does not matter which subset it chooses, the only important factor is the size of the subset because of the identical probabilities assumption). Making a query on the selected subset of size  $j'$  has cost  $C(j')$ . With probability  $j'/j$ , the target is in the selected set, and we have to pay  $A[j']$  in this case. With probability  $(j-j')/j$ , the target is not in the selected set, and we have to pay  $A[j-j']$  in this case. Since we assume that  $S_i$  contains  $x$  elements, the optimal cost for finding the target in cluster  $S_i$  is therefore  $A[x]$ .

Finally, phase three of Algorithm 2 makes sure that if we have not found the target in the previous phases, we query each nonclustered object until we find the target.

► **Theorem 7.** Let  $C_{Greedy}$  and  $C_{OPT}$  denote the expected search costs of Algorithm 2 and the optimum algorithm, respectively. Then, we have  $C_{Greedy} = O(\log(\log(n))C_{OPT})$ .

Before we proceed to the proof of this theorem, let us consider the following definitions for *general* cost functions. Note that in general, the cost of a query is a function of the query set and not necessarily its size.

► **Definition 8.** Let  $I(X)$  denote an instance of the search problem on a subset  $X \subset \mathcal{N}$  where the probabilities of objects in  $X$  are normalized to make sure their sum is one, and we know that the target is in  $X$ . We denote by  $Cost(X')$  the cost of making a query on

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$X' \subset X$ . Let also  $Opt(I(X))$  represent the expected search cost of the optimum algorithm for instance  $I(X)$ . We define

$$Cost_X^i = \min \{ Cost(X') | X' \subseteq X, \Pr(X') \geq 1 - 1/2^i \}.$$

Let  $S(X, i)$  denote the subset for which  $Cost(S(X, i)) = Cost_X^i$  and  $\Pr(S(X, i)) \geq 1 - 1/2^i$ .

Note that the particular cost function we consider in this section is  $Cost(X') = C(|X'|)$ . The following lemma will provide us with a general lower bound on the expected search cost of the optimum solution.

► **Lemma 9.**  $Opt(I(X)) \geq \sum_{i=1}^{\infty} Cost_X^i / 2^{i+1}$ .

Lemma 9 help us bound the expected search cost of the optimum algorithm from below.

**Proof of Theorem 7.** We first show that the expected search cost caused by nonclustered objects is negligible.

► **Lemma 10.** *The expected search cost incurred by the nonclustered objects in phase three of Algorithm 2 is at most  $\epsilon \cdot C_{OPT}$ .*

**Proof.** The probability of each nonclustered objects is at most  $1/2^l = \epsilon/n^2$ . There are at most  $n$  non-clustered objects, so the probability of the target is one of these non-clustered objects is at most  $\epsilon/n$ . With probability at most  $\epsilon/n$ , we make a query for each non-clustered object. Hence, its expected search cost is at most  $\epsilon n \times nC(1) = \epsilon C(1)$ . On the other hand, note that  $C(1)$  is a lower bound for the optimum solution because no matter where the target is we always have to make at least a query of size at least one to find the target. ◀

Lemma 10 entails that the expected search cost incurred by nonclustered objects is much less than the optimal expected search cost. Thus for simplicity we can assume that there exists no nonclustered object. In order to bound the expected search cost of the first phase of Algorithm 2 we need a few intermediate results. Let us start with the following definition.

► **Definition 11.** A set of nodes  $S$  of a tree  $T'$  is sparse, if and only if there do not exist three nodes  $v_1, v_2$  and  $v_3$  in  $S$  such that  $v_1$  is one of the ancestors of  $v_2$  and  $v_3$ , and neither of  $v_2$  and  $v_3$  is an ancestor of one another.

Hence, in a sparse set  $S$  there cannot be two nodes such that they don't have an ancestor/-descendant relationship and simultaneously have a common ancestor in  $S$ . The following lemma bounds the number of possible sparse partitions of any tree

► **Lemma 12.** *The nodes of any tree  $T'$  can be partitioned into  $\lceil \log(|T'|) \rceil$  sparse sets where  $|T'|$  is the number of nodes in  $T'$ .*

To link the expected search cost of the first phase of Algorithm 2 to nodes of the tree  $T$ , we need the following definition.

► **Definition 13.** Let a node  $v \in T$  represent procedure  $ClusterFinder(i, j)$  for some  $1 \leq i \leq j \leq l$ . With probability  $\Pr[i, j]$ , we go to this node, and we make a query on the set  $Q$  as defined in Eq. 1. If the target is not in  $Q$  (i.e., it is in one of the clusters  $S_{k+1}, S_{k+2}, \dots, S_j$ ), we make a second query on the set  $S_{k+1}$ . Hence, the expected search cost of our algorithm at node  $v$  is  $\Pr[i, j]C(|Q|) + \Pr[k+1, j]C(|S_{k+1}|)$ . We define this quantity as the price of node  $v$ .

It is clear that the expected search cost of our algorithm in phase 1 is the sum of the prices of all nodes in tree  $T$ . In Lemma 14, we bound the sum of prices of nodes of a sparse set which provide us with the key result to bound the expected search cost of the first phase. The proof of Lemma 14 heavily relies on the observation we made in Lemma 9.

► **Lemma 14.** *The sum of prices of nodes of a sparse set is  $O(C_{OPT})$ .*

Due to the above lemma, we know that the total prices of the nodes of every sparse set is  $O(C_{OPT})$ . By recalling Lemma 12 we also know that we can partition the nodes of  $T$  into  $O(\log(\log(n)))$  sparse sets. Hence, Lemma 14 together with Lemma 12 readily bounds the expected search cost of the first phase which is the sum of prices of all nodes in tree  $T$ .

► **Lemma 15.** *The expected search cost for identifying which cluster contains the target  $t_*$  (first phase of Algorithm 2) is  $O(\log(\log(n)) \cdot C_{OPT})$ .*

We are ready to analyze the second phase of algorithm 2. In this part, we know the cluster, say  $S_i$ , that contains the target  $t_*$  and we just need to find it. Let us define  $I_i$  to be this instance: we are given the information that the target is in  $S_i$ , and we have to find it. Following Definition 8, we denote the optimum expected search cost to identify the target in  $I_i$  by  $OPT(I_i)$ . The following lemma bounds the expected search cost of Algorithm 2 (phase two) in terms of  $OPT(I_i)$ .

► **Lemma 16.** *Assume that the target  $t_*$  is in cluster  $S_i$ . Then, the expected search cost of phase two of Algorithm 2 to find the target is at most  $2\Pr(S_i)OPT(I_i)$ .*

Lemma 16 helps us relate the expected search cost of phase two to  $C_{OPT}$ .

► **Lemma 17.** *The expected search cost of phase two of Algorithm 2 is at most  $2C_{OPT}$ .*

By combining Lemmas 10, 15, and 17 we can conclude that the expected search cost of Algorithm 2 is  $O(\log(\log(n)) \cdot C_{OPT})$ . ◀

## 5 Submodular Constraints

In this section we analyze another variant of the binary identification problem where the constraints are defined in terms of a submodular function. More specifically, we are given a set of objects  $\mathcal{N} = \{1, 2, \dots, n\}$  and a non-decreasing submodular function  $C_{sub} : \mathcal{F}_n \rightarrow \mathbb{R}$  where  $\mathcal{F}_n = 2^{\mathcal{N}}$  is the family of all subsets of  $\mathcal{N}$ . We denote the cost of making a query on the set  $S \subseteq \mathcal{N}$  by  $C_{sub}(S)$  where we assume that the cost function is submodular:

$$\forall S_1 \subseteq S_2 \subseteq U, i \in U : C_{sub}(S_1 \cup \{i\}) - C_{sub}(S_1) \geq C_{sub}(S_2 \cup \{i\}) - C_{sub}(S_2).$$

Intuitively, this property insures that adding an object  $i$  to a set  $S_1$  increases its cost at least as much as adding  $i$  to a superset  $S_2$ . Another equivalent definition that we later use reads as follows:  $\forall S_1, S_2 \subseteq \mathcal{N}, C_{sub}(S_1) + C_{sub}(S_2) \geq C_{sub}(S_1 \cup S_2) + C_{sub}(S_1 \cap S_2)$ . In view of Definition 8 the cost function of a subset  $S \in \mathcal{N}$  is  $Cost(S) = C_{sub}(S)$ .

Before elaborating on our new algorithm, we should note that it is possible to mimic the first phase of Algorithm 2 and obtain an approximation factor of  $O(\log(\log(n)))$ . However, it is no longer possible to run the second phase with the same approximation factor for submodular functions. Recall that in the second phase, we relied on a dynamic program that only depends on the size of the queried sets. Due to the fact that in our new setup, the cost of a query depends on the set (an not only on its size), we need to reformulate the dynamic program: it matters which subset is chosen. To do so, we ought to construct an

**Algorithm 4 Bicriteria Approx. Alg.****Input:**  $\epsilon > 0$ .

- 1: Find all  $\alpha_k$ 's in (3) and their corresponding sets  $S_{\alpha_k}$ .
- 2: Among all  $S_{\alpha_k}$ , keep only those that have  $\Pr(S_{\alpha_k}) \geq 1/6$ .
- 3: Among the remaining sets, choose the one with minimum  $C_{sub}(S_{\alpha_k})$ . Call this set  $S'$ .
- 4: Keep removing objects from  $S'$  while its probability remains at least  $1/6$ .

**Algorithm 5 BIP with Submod. Cnst.****Input:** Set of objects  $\mathcal{N}$ .

- 1: Find set  $S' \subset \mathcal{N}$  (and  $\bar{S}' = \mathcal{N} \setminus S'$ ) by using Algorithm 4. Query  $S'$ .
- 2: **if** set  $S'$  contains the target  $t_*$  **then**
- 3:   Query one of the objects  $i \in S'$ . Either  $i$  is the target, or otherwise recurs on  $S' \setminus \{i\}$ .
- 4: **else**
- 5:   Recurs on  $\bar{S}'$ .
- 6: **end if**

exponential size array (one for each subset) and as a result, the algorithm will no longer run in polynomial time. To avoid this drawback, we can devise a  $\log(n)$ -approximation algorithm for the second phase of Algorithm 2 which can incorporate submodular constraints. Instead, in this section we present a much simpler algorithm with the same approximation guarantee. To do so, we first need an intermediate step.

**Submodular Functions Under Linear Constraints**

The problem of “minimizing submodular functions under linear constraints” is to find a subset  $S^* \in \mathcal{N}$  such that  $\Pr(S^*) \geq 1/2$  and its cost, namely  $C_{sub}(S^*)$ , is minimum. Here, we present a bicriteria approximation algorithm that finds a set  $S'$  with  $C_{sub}(S') \leq 3C_{sub}(S^*)$  and  $\Pr(S') \geq 1/6$ . To do so, we first need to define another submodular function  $f^\alpha : 2^{\mathcal{N}} \rightarrow \mathbb{R}$  for which  $f^\alpha(S) = C_{sub}(S) + \alpha \cdot \Pr(\bar{S})$ , where  $\bar{S} = \mathcal{N} \setminus S$ , and  $\alpha > 0$  is a real number. The reason that  $f^\alpha$  is a submodular function simply follows from the fact that we only added a linear term to the submodular function  $C_{sub}$  (check the equivalent definition we provided earlier). It is a folklore result that submodular function minimization problem has strongly polynomial time exact algorithms due to [23] and [16]. Hence, one can find a subset  $S_\alpha \subseteq \mathcal{N}$  in polynomial time that admits the minimum value of  $f^\alpha$ . Let  $c_{\min} = \min_{i \in \mathcal{N}} C_{sub}(\{i\})$ . For a fixed  $\epsilon > 0$  we also define

$$\alpha_k = 3c_{\min}(1 + \epsilon)^k, \quad \text{for all } k = 0, 1, \dots, \lceil \log_{1+\epsilon}(C_{sub}(\mathcal{N})/c_{\min}) \rceil. \quad (3)$$

Note that there exists a  $k = \kappa$  for which  $3C_{sub}(S^*) \leq \alpha_\kappa \leq 4C_{sub}(S^*)$ .

► **Lemma 18.** *Let  $S_{\alpha_\kappa}$  denote the set that admits the minimum of the function  $f^\alpha$  for  $\alpha = \alpha_\kappa$ . Then  $C_{sub}(S_{\alpha_\kappa}) \leq 3C_{sub}(S^*)$  and  $\Pr(S_{\alpha_\kappa}) \geq 1/6$ .*

Since the value of  $C_{sub}(S^*)$  is not known a priori, we cannot apply Lemma 18 without modification. To this end, we propose Algorithm 4. It first calculates all values of  $\alpha_k$ . For each  $\alpha_k$  it finds the corresponding set  $S_{\alpha_k}$  that minimizes  $f^{\alpha_k}$ . It then finds among those  $S_{\alpha_k}$  with  $\Pr(S_{\alpha_k}) \geq 1/6$ , the one that has minimum cost  $C_{min}(S_{\alpha_k})$ . Once such a set  $S'$  is found, it starts removing objects from  $S'$  while keeping  $\Pr(S') \geq 1/6$ .

► **Lemma 19.** *Algorithm 4 outputs a set  $S' \subset \mathcal{N}$  with  $C_{sub}(S') \leq 3C_{sub}(S^*)$  and  $\Pr(S') \geq 1/6$ . Moreover, for any  $i \in S'$ , we have  $\Pr(S' \setminus \{i\}) < 1/6$ .*

**Approximation Algorithm with Submodular Constraints**

Algorithm 5 starts by finding a set  $S' \subset \mathcal{N}$ . To this end, it calls on Algorithm 4. According to Lemma 19, for the set  $S'$  we have  $\Pr(S') \geq 1/6$  and  $C_{sub}(S') \leq 3C_{sub}(S^*)$ . Remember  $S^*$

is the set with minimum cost and the probability at least a half. Ideally, we would like to query the set  $S^*$ . Unfortunately, it is not easy to find such a set. Instead, Algorithm 5 queries the set  $S'$  that approximate our ideal candidate. If the target  $t_*$  is in  $S'$ , then Algorithm 5 chooses an arbitrary object  $i \in S'$  and see whether  $i$  is the target or not. In case it is not the target, the whole procedure repeats now from the set  $S' \setminus \{i\}$ , namely, Algorithm 4 will be called for the set  $S' \setminus \{i\}$ . If the target is not in  $S'$  from the beginning, the algorithm recurs on set  $\bar{S}' = \mathcal{N} \setminus S'$ . Note that by making a singleton query before recursion, Algorithm 5 makes sure that the probability of the remaining set will shrink by a factor of  $1/6$  (Lemma 19) if the target is in set  $S' \setminus \{i\}$ . Otherwise, the probability shrinks by a factor of  $5/6$  because we have that  $\Pr(\bar{S}') = 1 - \Pr(S') \leq 1 - 1/6$ .

► **Theorem 20.** *Let  $C_{Greedy}$  and  $C_{OPT}$  denote the expected search costs of Algorithm 5 and the optimum algorithm, respectively. Then, we have  $C_{Greedy} = O(\log(n) \cdot C_{OPT})$ .*

**Proof.** The proof of this theorem is similar, *mutatis mutandis*, to the proof of phase one of Theorem 7. Let us first modify Definition 8 as follows. Let  $Cost_X^i = \min_{X' \subseteq X | \Pr(X') \geq 1 - (5/6)^i} Cost(X')$ . Then, we can show

► **Lemma 21.**  $Opt(I(X)) \leq \sum_{i=1}^{\infty} \frac{Cost_X^i}{(5/6)^{i+1}/5}$ .

The search mechanism portrayed in Algorithm 5 defines a binary tree  $T$ , similar to the one we saw in the previous section. In brief, the tree starts from the root  $\mathcal{N}$ . In each internal node  $S$  with probability at least  $1/6$  we go to the left child (representing set  $S' \setminus \{i\}$ ) and with the remaining probability we go to the right child (representing  $S \setminus S'$ ). Note that there are at most  $2n$  nodes in this tree where the probability of each child is at most  $5/6$  of the probability of its father (instead of  $1/2$  in the previous section). Because the tree  $T$  has at most  $2n$  nodes, instead of  $O(\log \log(n))$  sparse sets, we need  $\log(2n)$  sparse sets to cover the expected cost of all nodes. As a result one can provide a similar proof showing that the expected cost of Algorithm 5 with submodular constraints is at most  $O(\log(n))$  times the cost of the optimum solution. ◀

## 6 Conclusion

In this work we considered the problem of binary identification problem (BIP) under the general notion of graph, cost and submodular constraints. We also provided novel polynomial time approximation algorithms with provable analytical guarantees. Even though we believe that the three variants of BIP we considered are all NP-hard, we did not provide any rigorous proof. This is an interesting future direction we would like to pursue.

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