Minimizing State Space Partitionings Using Decision Trees

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***Abstract*— Lorem ipsum dolor sit amet, consectetuer adipiscing elit. Ut purus elit, vestibulum ut, placerat ac, adipiscing vitae, felis. Curabitur dictum gravida mauris. Nam arcu libero, nonummy eget, consectetuer id, vulputate a, magna. Donec vehicula augue eu neque. Pellentesque habitant morbi tristique senectus et netus et malesuada fames ac turpis egestas. Mauris ut leo. Cras viverra metus rhoncus sem. Nulla et lectus vestibulum urna fringilla ultrices. Phasellus eu tellus sit amet tortor gravida placerat. Integer sapien est, iaculis in, pretium quis, viverra ac, nunc. Praesent eget sem vel leo ultrices bibendum. Aenean faucibus. Morbi dolor nulla, malesuada eu, pulvinar at, mollis ac, nulla. Curabitur auctor semper nulla. Donec varius orci eget risus. Duis nibh mi, congue eu, accumsan eleifend, sagittis quis, diam. Duis eget orci sit amet orci dignissim rutrum.**

I. Introduction

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II. Preliminaries

We start by defining some relevant concepts.

Definition 1 (Partitionings): A partitioning $\mathcal{A}$ of the state space $\mathcal{S} \in \mathbb{R}^K$ is a set of regions ν that divides $\mathcal{S}$ such that $\bigcup_{\nu \in \mathcal{A}} \nu = \mathcal{S}$ and for any two regions $\nu, \nu' \in \mathcal{A}$ where $\nu \neq \nu'$ it holds that $\nu \cap \nu' = \emptyset$.

For an *axis aligned* partitioning, each region ν can be expressed in terms of two corner points, $s^{\min}, s^{\max} \in \mathcal{S}$, so that for each $s = (s_1, \ldots, s_K) \in \nu$ it holds that $s_i^{\min} < s_i \leq s_i^{\max}$ for $i = 1, \ldots, K$. In this work we exclusively consider axis aligned partitionings and we define all regions as a tuples $\nu = (s^{\min}, s^{\max})$. Note that the entire state space $\mathcal{S} \in \mathbb{R}^K$ can be described as a region: if $\mathcal{S}$ is unbounded in all dimensions (meaning its limits are positive and negative infinity) then $s_i^{\min}$ and $s_i^{\max}$ for the entire state space is $-\infty$ and ∞ respectively for $i = 1, \ldots, K$.

Definition 2 (Decision tree): A binary decision tree over the domain $\mathcal{S} \in \mathbb{R}^K$ is a tuple $\mathcal{T} = (\eta_0, \mathcal{N}, \mathcal{L}, \mathcal{D})$ where $\eta_0 \in \mathcal{N}$ is the root node of the tree, $\mathcal{N}$ is a set of branching nodes and $\mathcal{L}$ is a set of leaf nodes, each of which is assigned a decision δ from the set of decisions $\mathcal{D}$. Each branch node $\eta \in \mathcal{N}$ consists of two child nodes and a predicate function of the form $\rho_\eta(s) = s_i \leq c$ with $s \in \mathcal{S}$ and c being a constant.

Given a state $s \in \mathcal{S}$ and a decision tree $\mathcal{T}$, we can evaluate $\mathcal{T}(s)$ to obtain a decision δ by following the *path* from the root node to a leaf node given by the repeated evaluation of the predicate function $\rho_\eta(s)$ at each node η, starting with the root node and continuing with the left child if $\rho_\eta(s)$ evaluates to true and with the right child if it evaluates to false. When we encounter a leaf node ℓ, we return the decision assigned to ℓ. Further, we also allow evaluating a region of $\mathcal{S}$. Given a region $\nu = (s^{\min}, s^{\max})$, $[\delta]_\nu = \mathcal{T}(\nu)$ is the set of all decisions that can be obtained evaluating configurations of ν, ie. $\mathcal{T}(\nu) = \{\mathcal{T}(s) \mid s \in \nu\}$.

We denote the path $\lambda(\ell)$ and define it as an ordered list of tuples of the form (η_j, b) where η_j is the jth node on the path (η_0 will always be the root node) and b is a binary value indicating wether the path continues with the left child ($b = 1$) or right child ($b = 0$).

Not sure if it makes sense or is necessary to state which value of b means what. The reasoning for my choice is that $b = 1$ indicates 'true' and it is when $\rho_{\eta_j}(s)$ is true, that we choose the left path.

The path can then be said to define a region where the corner points $s^{\min}$ and $s^{\max}$ are given by compiling the bounds on each dimension $i = 1, \ldots, K$ given by the

predicate function $\rho_{\eta_j} = i_{\eta_j} \leq c_{\eta_j}$ for each $\eta_j \in \lambda(\ell)$ into points. The coordinates for each point is given by

$$s_i^{\min} = \max(\{\, c_\eta \mid (\eta, b) \in \lambda(\ell),\ i_\eta = i,\ b = 1 \,\})$$
$$s_i^{\max} = \min(\{\, c_\eta \mid (\eta, b) \in \lambda(\ell),\ i_\eta = i,\ b = 0 \,\})$$

I struggled a lot with coming up with a good way of writing this definition. Do let me know if it works (and if it is even necessary to describe how a region is constructed from a leaf node).

for all $i = 1, \ldots, K$. We write ν_ℓ to denote the region associated with the leaf node ℓ.

The set of regions obtained from all the leaf nodes of a decision tree constitutes a complete partitioning of a state space $\mathcal{S}$ in accordance with Definition1. We thus say that $\mathcal{T}$ induces a partitioning $\mathcal{A}_\mathcal{T} = \{\nu_\ell \mid \ell \in \mathcal{L}\}$. For any region ν and a decision tree $\mathcal{T}$ we say, that ν has *singular mapping* in $\mathcal{T}$ if for all $p \in \nu$, $\mathcal{T}(p) = \delta$ for some $\delta \in \mathcal{D}$. Naturally, all regions in $\mathcal{A}_\mathcal{T}$ has singular mapping in $\mathcal{T}$. For any partitioning $\mathcal{B}$ of the same state space, we say $\mathcal{B}$ *respects* $\mathcal{T}$ if and only if every region $\nu \in \mathcal{B}$ has singular mapping in $\mathcal{T}$.

III. MaxPartitions algorithm

Since state space discretization for Reinforcement Learning is usually done *before* any learning takes place, it tends to be conservative. For this reason, discretization is likely to create adjacent discrete states that are mapped to the same optimal action. The question we would then like to answer is this: if $\mathcal{T}$ is a decision tree representing a trained strategy and $\mathcal{A}_\mathcal{T}$ is its induced partitioning, can we find another partitioning $\mathcal{B}$ which is smaller than $\mathcal{A}_\mathcal{T}$ but still respects $\mathcal{T}$?

As an example, consider a case where we have a state space $\mathcal{S} \in \mathbb{R}^2$ over variables x and y, both of which are defined in the interval $[0, 9]$, and a set of actions $Act = \{action1, action2\}$. Before learning, we might decide to discretize both x and y into 3 distinct bins, giving us a partitioning of $\mathcal{S}$ with $3 \times 3 = 9$ regions. After training, we end up with a Q-table that maps states to action in a way that is shown in Figure 1a. This same mapping can also be represented by a decision tree, as shown in Figure 1b.

However, we can (in this small toy example) easily see that we do not need 9 regions to represent this exact state-action mapping. For example, the two regions given by $((3,0),(6,3))$ and $((6,0),(9,3))$, respectively, both assign $action2$ as the optimal action, but this mapping would still be preserved if we replaced those two regions with a larger one $((3,0),(9,3))$. After a little bit of inspection, we can actually see here that we could represent the same state-action mapping with a partitioning consisting of only 5 regions.

This example showcases how discretization techniques can easily end up with redundancy in the partitioning. This can be very difficult to anticipate before learning, especially since a very fine-grained discretization is typically needed for the learning to capture essential information and details for the strategy. Furthermore, for other learning techniques, such as the online partitioning refinement scheme of UPPAAL Stratego [1], regions can be created on-the-fly, not be arranged in a straight grid and/or vary in size, which can enhance the problem of redundancy.

We propose MaxPartitions, an algorithm that post-processes a decision tree inducing a partitioning of a state space in order to minimize the partitioning by *maximizing* the size of the individual regions (or partitions). The output of MaxPartitions is a new partitioning, ie. a list of regions with associated actions, which can then be arranged into a new decision tree.

A. Details of the algorithm

We write $\mathcal{T}_i$ for the (ascendingly) sorted list of bounds on dimension i in the policy given by the tree $\mathcal{T}$. The first bound in the list is defined to be negative ∞ and the last is positive ∞. By $\mathcal{T}_{i,j}$ we write the jth smallest bound on dimension i for each $j = 1, 2, \ldots, |\mathcal{T}_i|$. This can be precomputed as a matrix in log-linear time by collecting and sorting the bounds on all branch nodes in $\mathcal{T}$ and allows accessing $\mathcal{T}_{i,j}$ in constant time. Further, in a slight abuse of notation, we define $\mathcal{T}_{i,|\mathcal{T}_i|+1}$ to be some *sentinel* value representing that we are outside the boundaries of dimension i. Correspondingly, we define a sentinel action α, and we say that $\mathcal{T}(s^p_\mathcal{T}) = \alpha$ if and only if $\exists p_i \in p,\ p_i = |\mathcal{T}_i| + 1$.

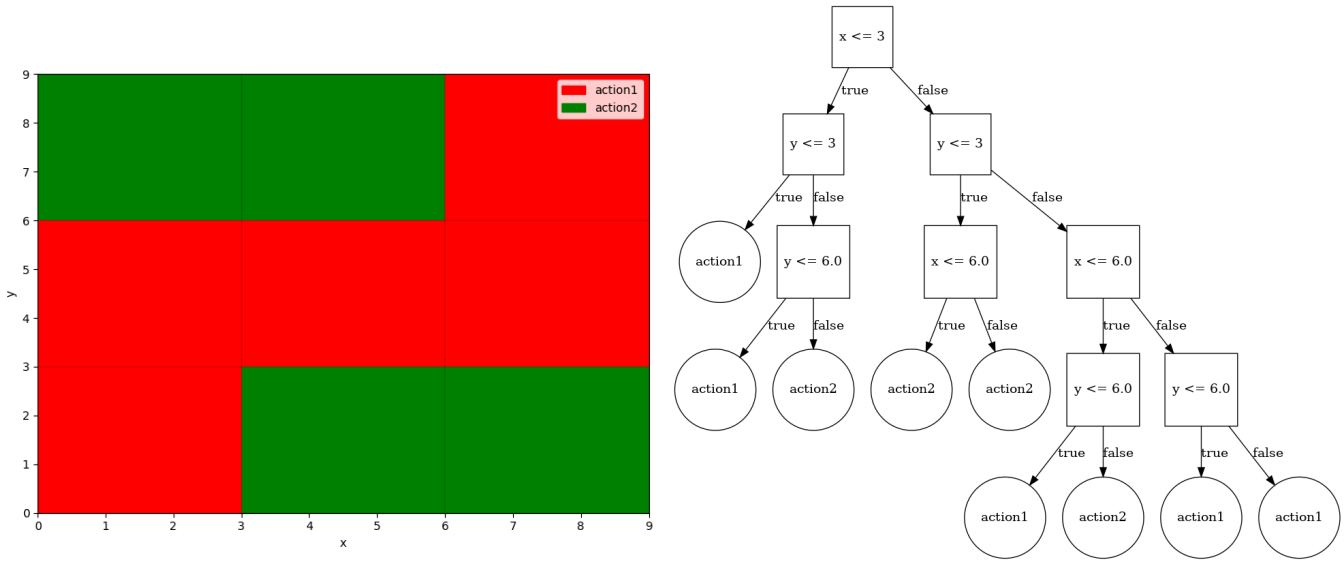
Exploiting this notation, let p be a K-dimensional vector of integers, such that $p_i \leq |\mathcal{T}_i| + 1$ for all $i = 1, \ldots, K$, then we can define a point at an intersection of bounds in $\mathcal{T}$ as $s^p_\mathcal{T} = (\mathcal{T}_{1,p_1}, \mathcal{T}_{2,p_2}, \ldots, \mathcal{T}_{K,p_K})$. To avoid cluttering the notation, will for the most part omit the subscript $\mathcal{T}$ on $s^p_\mathcal{T}$. To make things clear, we will write s to refer to an actual point in the state space of $\mathcal{T}$, ie. $s \in \mathcal{S}$, and we will write p to refer to a vector of integers representing indicies of bounds in $\mathcal{T}$.

The algorithm (given in pseudo-code in Algorithm 1) works by maintaining a pair $(p^{\min}, p^{\max})$, and iteratively incrementing $p^{\max}$ in one dimension at a time until a region $\nu = (s^{p^{\min}}, s^{p^{\max}})$ cannot be expanded further. When this happens, the region is added to a tree $\mathcal{T}_{track}$, which is used to track which areas of the state space have been covered and to provide new a starting point for each iteration of the algorithm. Expansion in dimension i is disallowed if one of the following three *expansion rules* are violated by the expanded region ν':

Definition 3 (Expansion rules): Let ν' be a candidate region for a new partitioning derived from $\mathcal{A}_T$. Then ν' is valid if it adheres to the following rules:

1) ν' does not have singular mapping in $\mathcal{T}$
2) ν' intersects with one or more regions already in $\mathcal{T}_{track}$
3) ν' intersects with a region ν_o in the original partitioning, such that the disjunction $\nu' \cap \nu_o$ cannot be described by a single region of the form $(s^{\min}, s^{\min})$

The first two cases are directly related to the definition of the problem, ie. the produced partitioning should respect $\mathcal{T}$ and



(a) (b)

Fig. 1: Two representations of a strategy mapping states to actions. In a the strategy is represented as a partitioning with colors representing the assigned action. In b the strategy is given as a decision tree, which also induces the partitioning. For this small example, it is obvious to see that an equivalent state-action mapping could be achieved with fever regions/leaves.

only have non-overlapping regions. The third case is required in order to guarantee that in each iteration the algorithm on average will add *at least* one region from the original partitioning to the new partitioning. To see this, consider ...

Make an example to show

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How do we determine this expansion? Let $(p^{\min}, p^{\max})$ define region in the original partitioning (or potentially the remains of a region previously cut in two). We then want to find a $\Delta_p \in \mathbb{Z}^K$ such that $(p^{\min}, p^{\min} + \Delta_p)$ defines a region that follows the three expansion rules and such that incrementing in any one dimension would result in a violation. By definition, a valid value for Δ_p is when $\Delta_p = p^{\max} - p^{\min}$, since this would just produce the original region. We are therefore guaranteed to at least find this region. This gives rise to the following definition.

Definition 4 (The expansion vector Δ_p): Given $p^{\min} \in \mathbb{Z}^K$, a decision tree $\mathcal{T}$ over a K-dimensional space and a decision tree $\mathcal{T}_{track}$ of already found regions, $\Delta_p \in \mathbb{Z}^K$ is a vector such that for $p^{\max} = p^{\min} + \Delta_p$ the region $\nu = (s_{\mathcal{T}}^{p^{\min}}, s_{\mathcal{T}}^{p^{\max}})$ does not violate any of the expansion rules in Definition 3 and where for any other $\Delta'_p = (\Delta_{p_1}, \ldots, \Delta_{p_i} + 1, \ldots, \Delta_{p_K})$ this would not hold.

A greedy approach to finding Δ_p starts with $\Delta_p = p^{\max} - p^{\min}$, for some (remainder of a) region $\nu = (s_{\mathcal{T}}^{p^{\min}}, s_{\mathcal{T}}^{p^{\max}})$. We then iteratively increment a single dimension chosen non-deterministically untill the invariants are violated. Let $\hat{\mathbf{e}}_i$ denote the unit vector parallel to axis i, such that $\Delta_p + \hat{\mathbf{e}}_i = (\Delta_{p_1}, \ldots, \Delta_{p_i} + 1, \ldots, \Delta_{p_K})$. At each increment, we define a candidate region ν' from $p^{\min}$ and $p^{\max} = p^{\min} + \Delta_p$ and check for singular mapping (Rule 1) and no overlap with regions in $\mathcal{R}$ (Rule 2). If any of these two do not hold, we mark dimension i as exhausted, roll back the increment and continue with a new dimension not marked as exhausted.

If Rule 3 is violated, the algorithm wil initiate an attempt at *healing* the candidate expansion, by continuing the expansion to the largest bound in the expansion dimension of any of the broken regions. This way we try to see if the violation can be overcome by simply expanding more aggressively. However, care is required to ensure, that we can roll back this extra expansion if it did not work (or if we inadverdently broke any of the other rules in the process).

When all dimensions have been exhausted, Δ_p adheres to Defintion III-A.

B. Analyzing the algorithm

In the following we provide an upper bound of the running time of MAXPARTITIONS and a proof of correctness.

1) Running time:

2) Proof of correctness:

C. From regions to decision tree

The output of the MAXPARTITIONS algorithm is a list of regions with associated actions. For this to be of any use, we need to construct a new decision tree to represent these state-action pairs. To this goal, we face the issue that it is not given (and in fact, very unlikely) that the suggested partitioning can be perfectly represented by a decision tree, as this would require the existence of enough 'clean splits' (ie. predicates on some variable that perfectly divides the regions into two sets with an empty intersection) to arrange the entire set of regions.

Therefore, we suggest a brute-force algorithm that tries to separate the regions as cleanly as possible. Let $\mathbf{R}$ be a list of regions and let a_ν be the action associated with the

Algorithm 1 MaxPartitions

Require: $\mathcal{T}$: A binary decision tree over the domain $\mathbb{R}^K$ inducing the partitioning $\mathcal{A}_\mathcal{T}$

$\mathcal{T}_{track} \leftarrow$ empty tree
$\mathcal{R} \leftarrow \{\}$
while $\mathcal{T}_{track}$ has unexplored regions **do**
  $(p^{\min}, p^{\max}) \leftarrow$ select region bounds of unexplored region
  $\Delta_p \leftarrow p^{\max} - p^{\min}$
  while not all dimensions have been exhausted **do**
    $d \leftarrow$ randomly select unexhausted dimension
    $\Delta_{p_d} \leftarrow \Delta_{p_d} + 1$
    $\nu' \leftarrow (s_\mathcal{T}^{p^{\min}}, s_\mathcal{T}^{p^{\min}+\Delta_p})$
    if ν' violates Rule 1 or 2 (Definition 3) **then**
      $\Delta_{p_d} \leftarrow \Delta_{p_d} - 1$
      mark d as exhausted
    else if ν' violates Rule 3 (Definition 3) **then**
      attempt $healing$ in dimension d
  $\nu' \leftarrow (s_\mathcal{T}^{p^{\min}}, s_\mathcal{T}^{p^{\min}+\Delta_p})$
  PUT($\mathcal{T}_{track}$, ν')
  $\mathcal{R} \leftarrow \mathcal{R} \cup \{\nu'\}$
return $\mathcal{R}$

region $\nu = (s^{\min}, s^{\min})$. In the following, we refer to $s^{\min}$ and $s^{\max}$ of a region ν by $\nu_{\min}$ and $\nu_{\max}$ respectively, and to the value of a specific dimension i in one such boundary point as $\nu_{\min,i}$ or $\nu_{\max,i}$.

We iteratively create a branch node that splits $\mathbf{R}$ into two, $\mathbf{R}_{low}$ and $\mathbf{R}_{high}$, based on a predicate function $\rho(x) = x_i \leq c$ with $c \in \mathbb{R}$ so that $\mathbf{R}_{low} = \{\nu \in \mathbf{R} \mid \rho(\nu_{\min}) \text{ is True}\}$ and $\mathbf{R}_{high} = \{\nu \in \mathbf{R} \mid \rho(\nu_{\max}) \text{ is False}\}$. When the list only contains a single element ν, we create a leaf node with action a_ν and return.

The question is how to determine $\rho(x)$, more specifically which dimension i to predicate on and at which value c. Ideally, we want to split $\mathbf{R}$ in two equally sized subsets and in a way that no single region would have to occur in both, ie. we would like $\mathbf{R}_{low} \cap \mathbf{R}_{high} = \emptyset$. For this we define an impurity measure $I(\mathbf{R}_{low}, \mathbf{R}_{high})$ that penalises the difference in size between $\mathbf{R}_{low}$ and $\mathbf{R}_{high}$ and the size of the intersection between the two. Let $abs(a)$ be the absolute value of a and let $|b|$ denote the size of a set b, then

$$I(\mathbf{R}_{low}, \mathbf{R}_{high}) = abs(|\mathbf{R}_{low}| - |\mathbf{R}_{high}|) + |\mathbf{R}_{low} \cap \mathbf{R}_{high}|$$

Our brute-force way of finding the predicate that minimizes I is to iterate over the dimensions in $\mathcal{S}$ and for each dimension i we sort the regions according to their upper bound. Let $\mathbf{R}_i = \{\nu^1, \nu^2, \ldots, \nu^n\}$ be the list sorted according to the i th dimension so that for all $\nu^j, \nu^{j+1} \in \mathbf{R}_i$ it holds that $\nu^j_{\max,i} \leq \nu^{j+1}_{\max,i}$. If we then let $\rho(x) = x_i \leq c$ with $c = \nu^j_{\max,i}$ we have $|\mathbf{R}_{low}| = j$ and —$\mathbf{R}_{high}| = n - j$. For determining the size of $\mathbf{R}_{low} \cap \mathbf{R}_{high}$ we simply need to count the number of regions ν^{j+m} for $m = 1, 2, \ldots, n - j$ whose lower bound is less than our predicate bound c, since these regions will appear both in $\mathbf{R}_{low}$ (because then, by definition, $\rho(x) = x_i \leq c$ will be true for $x_i = \nu^{j+m}_{\min,i}$ and $c = \nu^j_{\max,i}$) and in $\mathbf{R}_{high}$ (because our sorting ensures that for all ν^j, ν^{j+m} it holds that $\nu^j_{\max,i} \leq \nu^{j+m}_{\max,i}$).

Now we can write our impurity measure in terms of these quantities:

$$I(\mathbf{R}_{low}, \mathbf{R}_{high}) = abs(j-(n-j)) + \sum_{m=1}^{n} \mathbb{1}(\rho(\nu^{j+m}_{\min})), \quad \text{for all } \nu^j \in \mathbf{R}_i$$

where $\mathbb{1}$ is the indicator function, $\mathbf{R}_i$ is the list of regions sorted according to upper bounds in dimension i and $\mathbf{R}_{low}$ and $\mathbf{R}_{high}$ are the subsets resulting from splitting on the predicate function $\rho(x) = x_i \leq c$ with $c = \nu^j_{\max,i}$ so that $\mathbf{R}_{low} \subsetneq \mathbf{R}$, $\mathbf{R}_{high} \subsetneq \mathbf{R}$ and $\mathbf{R} \supseteq \mathbf{R}_{low} \cup \mathbf{R}_{high}$.

Finding the best split, ie. the one that minimizes the impurity, is a $O(Kn^2)$ operation, as it requires a nested loop through all the regions for each of the Kdimensions (the nested loop being the final summation term over $m = 1, 2, \ldots, n - j$ for all $j = 1, 2, \ldots, n - 1$). In this work, we have not attempted to find a faster implementation as we found that the size of $\mathbf{R}$ obtained by using our MAXPARTITIONS algorithm did not cause performance issues.

IV. EXPERIMENTS

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TABLE I: Comparing MAXPARTITIONS and VIPER for pure shields and shielded oracles.

Model	Pure shield			Shielded oracle		
	Input	MAXPARTITIONS	VIPER	Input	MAXPARTITIONS	VIPER
Bouncing ball	xxx	xxx	xxx	xxx	xxx	xxx
Random walk	xxx	xxx	xxx	xxx	xxx	xxx
Cartpole	xxx	xxx	xxx	xxx	xxx	xxx
Cruise	xxx	xxx	xxx	xxx	xxx	xxx
DCDC boost control	xxx	xxx	xxx	xxx	xxx	xxx

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APPENDIX