Al Project Paper

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General Definitions

The following are definitions that are shared between our two search paradigms.

$$G = Games = \{g_0, ..., g_n\}$$

$$P = Practices = \{p_0, ..., p_n\}$$

$$GP = Games \ and \ Practices = G \cup P$$

$$Sl = Slots$$

$$Constr(f) = \begin{cases} Fact \ meets \ all \ hard \ constraints \land Complete(f) & True \\ Otherwise & False \end{cases}$$

$$PartConstr(f) = \begin{cases} Fact \ meets \ all \ hard \ constraints^* & True \\ Otherwise & False \end{cases}$$

$$Complete(f) = \begin{cases} \forall \ sl \in f, sl \neq \$ & True \\ Otherwise & False \end{cases}$$

PartEval(f), $f \in F'$: Returns the sum of penalties from all violated soft constraints.

AConstr(a): Evaluates whether an individual assignment is valid. This is false if a game is in a practice-only slot, or vice versa.

 $Assign = \{(g, sl) | g \in GP, sl \in Sl, AConstr(a)\}$: Ordered pairs representing possible assignments from games or practices to their corresponding slots.

 $assign = GP \rightarrow Sl$ such that $\forall g (g \in GP \rightarrow (g, assign(g)) \in Assign)$: Surjection mapping a game or practice to a slot. Must be in Assign, to meet constraints. $partassign: GP \times (Sl \cup \{\$\})$: Initial assignment provided by input

 $passign = assign \cup GP \rightarrow \$$: Modified version of assign which allows null assignments

Set-Based Search

Informal Description

Our set-based search model is an evolutionary algorithm. We use an Or-Tree-based search to find some initial valid solutions (to create our initial population), then run our evolutionary algorithm of both crossovers and mutations on the set of valid solutions provided to us from the or tree search, to produce new facts for our population. We continue this process until we reach a point where the number of facts we have in the

population exceeds some cap, at which point we delete some facts in the population. Between the selective pressure applied from delete, the carry-on of "good" traits from crossover, and the genetic diversity applied from mutation, we can predict that after several generations (iterations of our extension functions) we will likely reach a population containing a fact that is close to the optimal solution to our problem. When the search is complete, we choose the fact in our population with the best evaluation to be considered our "solution".

Search Model

$$A_{set} = (S_{set}, T_{set})$$

$$S_{set} = \{s | s \in 2^F\}$$

$$T_{set} = \{(s, s') | \exists A \to B \in Ext \text{ with } A \subseteq s \text{ and } s' = (s - A) \cup B\}$$

Let F be the set of all facts. A fact f is a complete, valid schedule. We will first define F' which contains f': both complete and incomplete facts, where there may be some games or practices not assigned to slots, making F' the set of all incomplete and complete schedules. This will be used in our or-tree.

$$F' \equiv \{f'|f' = \{(gp, passign(gp))|gp \in GP\}, PartConstr(f)\}$$

From F' we will construct our set of facts, which must also satisfy Constr(f), meaning they must be fully assigned. These are fully assigned schedules – complete facts.

$$F \equiv \{f | f \in F', Constr(f), f \in Assign\}$$

Extension Rules

Our extension rules are the core of the genetic algorithm. They take our set of solutions (our population) and add a new fact. We have crossover, which takes two different facts from the population, and considers their assigned slots to games or practices to build up a new child fact, which is then added to the population. Mutation, which takes a fact and changes a single assignment in it, and adds this new mutated fact to the population, and finally, we have delete, which runs when our population gets too large. When this happens, the worst of the population gets removed.

$$Ext = \left\{ A \to B \middle| A, B \in S \land \left(\left(Crossover(A, B) \lor Mutation(A, B) \right) \lor Delete(A, B) \right) \right\}$$

We will define a useful variable a_n as the assignment

 $a_n = (gp_n, sl_n | sl_n = assign(gp_n))$: the individual tuples of valid games/practices and their assigned slot.

Crossover(A,B)

Crossover is an extension function which uses 2 facts (parents) from our current population, to create a new child fact, which is then added in to our population s. We use crossover to generate child facts that are made up of 2 fit parent facts, which will hopefully allow us to carry on the "good" traits of 2 parents into a new generation.

Reference our search control section for formal definitions on the contents of A, a set of 2 facts provided to crossover(A,B).

Let k = |GP|. In our search control, f_{select} will provide crossover with 2 victims f_a and f_b , making

$$A = \{f_a, f_b\}$$

$$A = \{\{a_{a1}, \dots, a_{ak}\}, \{a_{b1}, \dots, a_{bk}\}\}$$

$$A = \{\{(gp_{a1}, sl_{a1}), \dots, (gp_{ak}, sl_{ak})\}, \{(gp_{b1}, sl_{b1}), \dots, (gp_{bk}, sl_{bk})\}\}$$

We will be using an OR tree search to complete this crossover and create a child fact. The OR tree search will select first a random slot from either parent a or parent b. If the selection of the slot from the random parent violates our hard constraints, the OR tree will go on to select the slot from the other parent. If this too violates our hard constraints, the OR tree will select slots at random, until we reach a valid slot. For more information on this process, please reference section SECTION on the or tree. Additionally, note that OR tree will always return a valid slot, as both parents f_a and f_b are valid facts, and OR tree will simply return one of the parents as a child fact if there are no other valid possibilities to create a new fact under.

OrTree(A, Crossover): This is our inputs into our or tree. See ORTREE for more details This or tree will return:

 $f_c = \{a_{c1}, ..., a_{ck}\} = \{(gp_{c1}, sl_{c1}), ..., (gp_{ck}, sl_{ck})\}$. The child fact (see 1.N OR TREE for more specific definitions on what f_c is composed of, and how we make f_c).

As a result, we have

$$B = A \cup \{f_c\}$$

Mutation(A,B)

Mutation is an extension function which uses a single fact from our current population, to preform a single point mutation on one of its current slots, creating a new modified fact, which is then added in to our population.

We receive an A from f_{select} . Let k = |GP|

$$A = \{a_1, \dots, a_k\}$$

$$A = \{(gp_1, sl_1), \dots, (gp_k, sl_k)\}$$

Let r be some random integer such that there exists some slot $sl_r \in a_r$ with $1 \le r \le k$. We take this slot sl_r and alter it to $sl_r' = \$$, to update A to

 $A' = \{(gp_1, sl_1'), \dots, (gp_k, sl_k') | \forall sl_i' \in A', sl_i' = sl_i, \text{ except for a single } sl_r' \in A' \text{ where } 1 \leq r \leq k, \text{ such that } sl_r' \neq sl_r, \text{ and } sl_r' = \$\}.$

Essentially, we convert the assignment of one slot in A to be null, as this will be the point of mutation.

We will be using an OR tree search to complete the mutation. The OR tree will take the empty slot $sl_r{'}$ and randomly search until we find a slot $sl \in Sl$ such that the slot can be filled into A at a_r , and will fit all hard constraints. We prioritize filling the slot $sl_r{'}$ with some slot $sl_i \in Sl$ such that $sl_i \neq sl_r$. However, if we find no possible slot that fits the constraints and is not sl_r , then we will be forced to return A from the or tree as our mutated fact. Otherwise, we will receive a new fact $f_{new} \in F$.

$$Otree({A, A'}, mutation)$$

This tree will return a new complete fact, giving us the following for B:

 $B = A \cup f_{new} = \{(gp_1, sl_1), \dots, (gp_k, sl_k)\} \cup \{(gp_1, sl_1''), \dots, (gp_k, sl_k'')\}$ where $sl_i = sl''_i$ except for the $sl_r \in A$, where we have performed a single point mutation, such that $sl''_r \in Sl$, but not nessecairly $sl_r = sl_r''$.

Essentially, this extension function returns Our original A from f_{select} , unioned with the output of the Or tree.

Note that if the output of the or tree is the same as the original A, we are simply just returning the input we received from f_{select} . Also, note that this process can be expanded to

fit multi-point mutations – although it would require some further considerations into the relationships between slots (MWF, TH, F, for example).

Delete(A,B)

Delete is an extension function which will remove a set of facts from our current population.

We receive A, a set of facts, from f_{select} (see search control for more specific details on the selected facts) such that

$$A = \{f_1, ... f_n \}$$

We simply delete these facts, creating a set B such that

$$B = \{\}$$

This effectively removes these facts from our population.

Search Process

Our search process defines how a search can take place. It takes our model, an environment, and a search control.

$$P_{set} = (A_{set}, Env, K_{set})$$

Our environment specifies some static properties, listed below, which are used by our search control.

Our search control is defined as follows:

$$K_{set}: S_{set} \times Env \rightarrow S_{set}$$

$$K_{set}(s,e) = (s-A) \cup B$$

$$Env = \{Cap, P_{mut}, c, g, lim\}$$

 ${\it Cap}$: A pre-determined integer value for the capacity of complete facts f in our current population

 P_{mut} : The probability of choosing mutation where $0 \le P_{mut} \le 1$

c: A decimal representing the percentage $0 < c \le 1$ of the facts with the lowest partEval(f) value in the current population that should be considered for crossover.

g: A decimal representing the percentage 0 < c < 1 of the facts with the highest partEval(f) values, in our current population, that should be considered for delete. Note that a g value that is too small is unfavorable, as the smaller g is, the more often we will be running the delete function.

lim: Some integer $0 \le lim < \infty$, representing how many iterations we run of our algorithm.

Our f_{wert} sets some value to each extension function of crossover, mutation or delete. Note that $s \in S$ is our current population.

$$f_{wert}(Crossover) = f_{wert}(Mutation) = \begin{cases} |s| \le Cap: & 0\\ Otherwise: & 1 \end{cases}$$
$$f_{wert}(Delete) = 1 - f_{wert}(Crossover)$$

 f_{select} will select the extension rule containing the lowest value from f_{wert}

In the event of the tie between crossover and mutation, we break this using f_{select} .

Let *RNG* be a random number generator.

 f_{select} uses a RNG to generate a random number r where $0 \le r \le 1$.

 f_{select} chooses mutation(A, B) when $0 \le r \le P_{mut}$

 f_{select} chooses crossover(A,B) when $P_{mut} < r \le 1$

Note that there will never be a tie in values when cap is surpassed, so we do not require an f_{select} tiebreaker when delete(A, B) is the chosen extension rule.

Selecting Victims

Let r be some random integer, generated by RNG, such that $1 \le r \le |s|$. When selecting facts to be victims of crossover or mutation, f_{select} chooses $f_r \in s$, where s is the current state (our current population), making f_r a random fact from our current population.

For Mutation(A,B), our victim is simply any random $f_r \in s$.

For Crossover(A,B), however, our victims are f_{r1} and f_{r2} , where we first ensure that $f_{r1} \neq f_{r2}$ to reduce redundant crossovers, are selected according to their fitness evaluation. We consider only the best k facts, such that $k = \lceil c * |s| \rceil$. in the current population s, giving us

 $f_{r1}, f_{r2} \in \{f_1, ..., f_k\} \subset s$ where for any $f_j \notin \{f_1, ..., f_k\}$, and $\forall i$ s. t. $1 \le i \le k$, $partEval(f_j) \ge partEval(f_i)$. Saying, any fact not in the set of facts $\{f_1, ..., f_k\}$ has a higher partEval value that any fact in the current set of facts. Any fact not in the set of the facts $\{f_1, ..., f_k\}$ will not be considered as victims for our crossover -- we want to crossover only the best p% of our current population.

However, if there are less than 2 distinct facts in the set from the best p% of the population, we multiply p*2 to create a new value for p and repeat the process. There is a risk our given p value could be small enough that there does not exist 2 facts within this percentage, and we need to account for that.

If for any reason our population is down to 1 fact, we will run $orTree(\{partAssign, partAssign\}, initiate)$ to generate a new fact, before proceeding with selecting victims for our crossover. Note that, due to the fact that delete operates on $\lfloor g* \vert s \vert \rfloor$ facts, and g<1, we will never be in a position where our population has no facts following an extension rule.

For Delete(A,B) we will do the opposite. We consider only the worst h facts, such that $h = \lfloor g^* | s \rfloor \rfloor$. Here, we send in the crossover function a set of facts $\{f_h, \dots, f_{|s|}\}$ such that for any $f_i \notin \{f_h, \dots f_{|s|}\}$, and $\forall h \leq i \leq |s|$, $EVAL(f_i) \leq EVAL(f_i)$.

Saying, any fact not in the set of facts $\{f_h, ..., f_{|s|}\}$ has a lower *partEval* value that any fact in the selected set of facts. Any fact not in the set of the facts $\{f_h, ..., f_{|s|}\}$ will not be considered as victims for our delete -- we want to delete only the worst g% of our current population.

Search Instance

To define our search instance, we will use s_0 for the starting state and G_{set} for the goal conditions, then the search instance is defined as:

$$Ins_{set} = (s_0, G_{set})$$

The starting state $s_0 \in S_{set}$ is an assignment with the hard constraints satisfied. We will use an Or-Tree based search to get these initial states, which is defined below.

The goal $s_{goal} \in S_{set}$ is any assignment of games and practices that meet all hard constraints and maximizes the soft constraints. The goal condition is defined by:

$$G_{set}: S \rightarrow \{yes, no\}$$

Where $G_{set}(s_i) = yes$ if and only if $s_{goal} \subseteq s_i$ where s_{goal} has an Eval of zero, or the amount of times we run our algorithm reaches a set limit $lim \in Env$.

Otherwise, $G_{set}(s_i) = no$.

Or-tree Search Model

We use an or tree to return a single valid solution. This will be used to develop the initial population for our set based search, to generate a child for crossover from 2 parents, and to create a new mutated fact from a fact with a single empty slot assignment. Our or tree search model will use specific search methods to create each of these 3 outputs, and will return the leaf containing the complete fact fitting all hard constraints, or will run until it has visited every possible node, and determined that there does not fit any valid facts within our constraints.

Model Description

We first need to define our model $A_{\rm V}=(S_{\rm V},T_{\rm V})$. Having the given definitions for $S_{\rm V}$ and $T_{\rm V}$, we only need to define Prob to complete our definition of A. We define Prob as F'.

Now, with our definition of *Prob*, we can defined S_{V} as

 $S_{\vee} \subseteq \textit{Otree}$ where Otree is defined recursively by

$$(pr,sol) \in \textbf{Otree}$$
 for $pr \in Prob, sol \in \{yes,?,no\}$ $(pr,sol,b_1,...,b_n) \in \textbf{Otree}$ for $pr \in Prob, sol \in \{yes,?,no\}, b_i \in \textbf{Otree}$

And we can define T_{V} as

$$T_{\vee} \subseteq S_{\vee} \times S_{\vee}$$

$$T_{\vee} = \{(s_1, s_2) | s_1, s_2 \in S_{\vee} \text{ and } Erw_{\vee}(s_1, s_2)\}$$

Where Erw_{V} is a relation on Otree defined by

$$Erw_{\vee}((pr,?),(pr,yes))$$

 $Erw_{\vee}((pr,?),(pr,no))$

$$Erw_{\vee}\left((pr,?),(pr,?,(pr_1,?)\ldots,(pr_n,?))\right) \text{ if Altern}(pr,pr_1,\ldots,pr_n) \text{ holds}$$

$$Erw_{\vee}\left((pr,?,b_1,\ldots,b_n),(pr,?,b_1',\ldots,b_n')\right) \text{ if for an } i : Erw_{\vee}(b_i,b_i') \text{ and } b_j=b_j' \text{ for } i\neq j$$

Defining Altern Relationship

By the definition of Prob, we know that

$$Altern \subseteq Prob^+$$

Altern is a function that takes a partial schedule prob and generates a new prob with one more assignment. It does this by selecting some arbitrary game or practice that is not assigned, removes that ordered pair, and replaces it with that same game or practice assigned to a slot.

$$Altern(prob) = (prob - \{(gp, \$)\}) \cup \{(gp, assign(gp)\}\}$$
 where $gp \in GP$

The environment

We define the environment as

$$Env = (w_{set}, type)$$

Where w_{set} is a set of 2 facts such that

$$W_{set} = \{f_1, f_2 | f_1, f_2 \in F'\}$$

The specific contents of f_1 and f_2 depend on the *type* variable. Let k = |GP|

1. If type = crossover then:

We receive 2 facts from w_{set} —parent a and parent b, which will be used to help us develop our child fact

$$f_1 = f_a = \{a_{a1}, \dots, a_{ak}\} = \{(g_{a1}, sl_{a1}), \dots (gl_{ak}, sl_{ak}) | \ \forall sl_i \in a_i, sl_i = assign(g_{ai})\}$$

$$f_2 = f_b = \{a_{b1}, \dots, a_{bk}\} = \{(g_{b1}, sl_{b1}), \dots (gl_{bk}, sl_{bk}) | \forall sl_i \in b_i, sl_i = assign(g_{bi})\}$$

2. and by our victim definitions in f_{select} ,

$$f_a \neq f_b$$

If type = mutation then:

If we have a mutation, then f_1 is going to be a complete fact f_{og} , wherein $f_{stripped}$ is simply f_{og} , with a random one of the slots in f_{og} associated to a game set to null.

$$\begin{split} f_1 &= f_{og} = \{a_1, \dots, a_k\} = \{(g_1, sl_1), \dots (gl_k, sl_k) | \ \forall sl_i \in a_i, sl_i = assign(g_i)\} \\ f_2 &= f_{stripped} = \{a_1, \dots, a_k\} = \{(g_1, sl_1'), \dots (gl_k, sl_k') | \ \forall sl_i' \in f_2, \forall sl_i \in f_1, sl_i' = sl_i, \\ & \text{except for one random } sl_r \in a_r, \text{ where } sl_r = \$\} \end{split}$$

3. If type = initiate

In *initiate*, we do not require any other input facts than *partAssign*, meaning the fact f_2 is redundant.

$$f_1 = partAssign$$

 $f_2 = partAssign$

And type is the type of or tree we will be implementing

type = one of crossover, mutation or initialize

Defining the solved state

We now need to define when a *pr* in our tree is solved to a *yes* (we have found a valid complete fact), a *no* (we have violated some hard constraints), or a ? (when we have yet to violate hard constraints, and we have not reached a complete fact). These are defined as the following:

$$sol = yes$$
 when $partialConstr(pr) \land Complete(pr)$

Our node is labeled as "yes" when we reach a complete solution that passes all of the hard constraints

$$sol = no$$
 when $\neg partialConstr(pr)$

Our node is labeled as "no" when we violate any hard constraints, regardless of if our branch is complete or not

$$sol = ?$$
 when $partialConstr(pr) \land \neg Complete(pr)$

Our node is labeled as "?" when we have not yet violated any partial constraints, however we have not reached a full solution

Search Control

Now that we have defined the search model A_{V} , we can define the search process,

$$P_{\vee} = (A_{\vee}, Env, k_{v}).$$

The search control K_V chooses which leaf to transition to using the following evaluation functions:

Mutation f_{eval}

When type=mutation, we receive a set of facts f_{og} and $f_{stripped}$ (reference the environment in or tree). Let k=|GP|, and let sl_n be some slot associated with a game or practice gp_n such that $sl_n=assign(gp_n)$. To restate what was already defined in our environment

$$f_{og} = \{a_1, \dots, a_k\}$$

$$f_{og} = \{(gp_1, sl_1), \dots, (gp_k, sl_k)\}$$

 $f_{stripped} = \{(gp_1, sl_1'), ..., (gp_k, sl_k') | \exists \text{ a single random } sl_r' \in f_{stripped} \text{ where } 1 \leq r \leq k, s. t \ sl_r = \$ \}.$

We use these definitions to represent $f_{mutEval}$

$$f_{mutEval}(pr) = \begin{cases} 0 & if \ for \ sl'_r \in pr, sl_r \in f_{og}, sl_r' \neq sl_r \\ 1 & otherwise \end{cases}$$

Essentially, if our empty slot $sl'_r \in f_{stripped}$ is replaced with the slot $sl_r \in f_{og}$, then we give the associated pr a partEval value of 1. We would rather explore a pr with a slot that is different that the one associated with original fact that mutation is provided with from f_{select} , to help us increase genetic diversity.

Crossover f_{eval}

When type=crossover, we receive a set of facts $\{f_a, f_b\}$ —the two parent facts. Let k=|GP|, and let sl_n be the empty slot in pr of which we are currently assigning a slot $sl \in Sl$ to, and let $sl_{an} \in f_a$ be the corresponding slot in parent a, and let $sl_{bn} \in f_b$ be that of parent b.

$$f_{crossEval}(pr) = \begin{cases} 0 \text{ if } for \, sl_n \in pr, sl_n = sl_{an} \text{ or } sl_n = sl_{bn} \\ 1 \text{ otherwise} \end{cases}$$

Essentially, we are favoring nodes that contain a slot from either parent 1 of parent 2, above slots with random entries.

$f_{leafExt}(s)$ Crossover and mutation

When type = crossover or type = mutation we proceed with this $f_{leafExt}$.

When type = crossover, we use $f_{crossEval}$ to calculate our leafEval value, such that $f_{crossEval}(pr) = leafEval$.

When type = mutation, we use $f_{mutEval}$ to calculate our leafEval value, such that $f_{mutEval}(pr) = leafEval$.

 $f_{leafExt}$ will return us the node in our state s with the lowest leafEval value. In the case of a tie, it will choose the deepest of the tied nodes. In the case of another tie between these, it will select one of the tied nodes randomly.

$f_{leafInit}(s)$ initialize

When type = initialize, we proceed with this $f_{leafInit}$.

This will simply choose the current deepest node in our state s to explore. In the case of a tie between nodes, it will select one of the tied nodes randomly.

 $f_{trans}\left((pr,?)\right)$ selects the only Altern of pr that holds.

Or-Tree Search Instance

The s_0 of our or tree is the starting state. This will take in a fact we receive from w_{set} to help us initialize our search.

Mutation s_0

$$s_0 = (f_{stripped},?)$$

In our mutation search instance, our starting state s_0 is the fact $f_{stripped}$ that the or tree received from the mutation extension function. $f_{stripped}$ is simply a fact with a single null slot, of which our or tree will assign a slot $sl \in Sl$ to.

Crossover s_0

In our crossover search instance, our starting state s_0 is an empty fact f_{empty} , as we will build up the empty fact one by one to create our child fact to return to *crossover*.

Let k = |GP| Let $f_{empty} \in F'$ be a fact such that

$$f_{empty} = \{a_1, \dots a_k\} = \{(gp_1, sl_1), \dots, (gp_k, sl_k) | \forall s_i \in f_{empty}, s_i = \$\} \,.$$

It is a fact containing games and practices, all of which are all mapped to a null slot.

$$s_0 = (f_{empty},?)$$

Initialize s_0

In our initialize search instance, our starting state s_0 is the fact PartAssign, the provided partial assignment of games and practices to slots, with the remaining unassigned games or practices being mapped to null.

$$s_0 = (PartAssign,?)$$

Goal

Note that for all s_0 we have defined above, $s_0 = (pr,?)$

 $G_{\wedge}(s) = y$ es, if and only if

- 1. s = (pr', yes) or
- 2. $s = (pr', ?, b_1, ..., b_n), G_{\wedge}(b_1) = ... = G_{\wedge}(b_n) = yes \text{ or }$
- 3. All leaves of our tree have either the sol-entry *no* or cannot be processed using *Altern*

We stop our or tree search when we reach either of these conditions.

Now that we have defined P_V , A_V and Ins_V , we have defined everything needed for our or tree model.

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Visualization and walkthrough

For this example, lets first define all initial variables of our GA. In our environment

$$Env = \{Cap, P_{mut}, c, g, lim\}$$

Suppose the following:

$$Cap = 3$$

$$P_{mut} = 0.20$$

c: 0.6

g: 0.40

lim: 3

Suppose, in our example, $GP = \{gp_1, gp_2, gp_3\}$, and $Sl = \{sl_1, sl_2, sl_3\}$.

Suppose we receive some partassign such that

$$partassign = \{(gp_1, sl_1), (gp_2, \$), (gp_3, \$)\}.$$

1. Creating initial population of size CAP

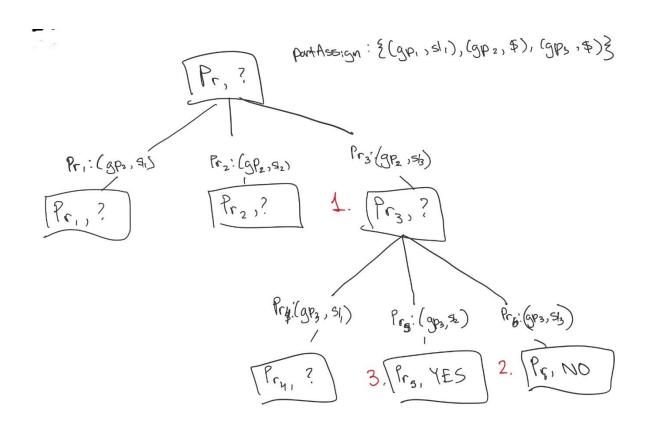
Given partassign is simply some initial configuration of games to slots provided to us, we can now begin to create our $population_0$ using partassign as a guide for our or tree. To do so, we call

We insert 2 partassigns, knowing the second one will be redundant. Refer to (ORTREE) for further details on the parameters accepted.

Having fed all relevant information to the Or Tree, we begin the or tree process.

Please note: although our or tree search selects a random $gp_i \in Pr$ to assign a slot to, for the sake of spacing and ease, our diagrams assign a slot to an empty gp_i in numerical order. It is important to know that this may not always be the case, as the decision as far as which game or practice to assign to a slot is made randomly in our tree.

Also note: at each definition in the diagram, where pr_i : (gp_n, sl_j) , we omit the redundant associated a_n components in the set in pr for spacing purposes.



Step 1:

We start with a single leaf node (pr,?), which gets expanded into its alternatives pr_1, pr_2 and pr_3 , where gp_1 get assigned to sl_1, sl_2 and sl_3 respectively.

Step 2:

We select one of the possible three nodes randomly. Our selection, in this example, is pr_3 , which assigns, sl_3 to gp_2 .

Step 3:

 pr_4 , pr_5 and pr_6 are the deepest leaves that we can now explore. We select one randomly. Our selection, in this example, is pr_6 , which assigns slot sl_3 to gp_3 . However, in this example, the fact $f_i = \{(gp_1, sl_1), (gp_2, sl_3), (gp_3, sl_3)\}$ violates a hard constraint. As a result, we set the variable sol to no.

Step 4:

 pr_4 , pr_5 are now the deepest unnassigned leaves that we can now explore. Our next random selection is pr_5 , which assigns slot sl_2 to gp_3 . This assignment creates the fact $f_i = \{(gp_1, sl_1), (gp_2, sl_3), (gp_3, sl_2)\}$ which, in our example, passes all hard constraints, and is complete. As a result, we can set sol = yes, and we are done our search.

We repeat the initialize function *Cap* times, to create a population of size *Cap* (in this example, to create a population of size 3).

Let this population be:

$$population_0 = \{f_1, f_2, f_3\}$$

$$population_0 = \{$$

$$\{(gp_1, s_1), (gp_2, s_2), (gp_3, s_3)\},$$

$$\{(gp_1, s_2), (gp_2, s_1), (gp_3, s_3)\}$$

$$\{(gp_1, s_1), (gp_2, sl_3), (gp_3, sl_2)\}\}$$

2. Choosing an extension function

Immediately after our initialization of our initial population, we have cap (in this example, 3) chromosomes (facts) in our population. As a result, $f_{wert}(crossover) = f_{wert}(mutation) = 0$, and $f_{wert}(delete) = 1 - f_{wert}(crossover) = 1$.

Given $f_{wert}(crossover) = f_{wert}(mutate)$, f_{select} is tasked with breaking the tie between the two extension rules. To do so, we generate some random number $0 \le r \le 1$. Suppose r = 0.6

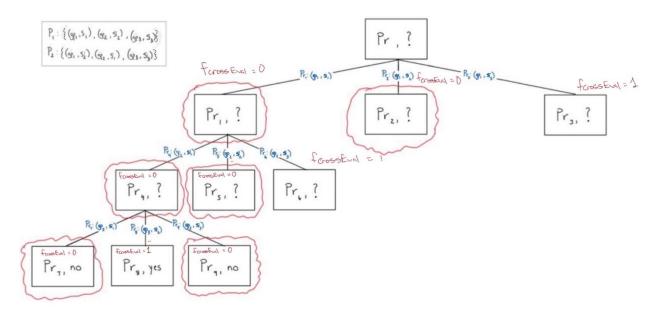
According to our definition of f_{select} , and its relationship to $P_{mut} \in Env$, we select mutate(A, B) when $0 \le r \le 0.2$, and we select crossover(A, B) when $0.2 < r \le 1$. Given r = 0.6, we are selecting crossover(A, B) for **iteration 1**.

The victims crossover receives is also part of our definition of f_{select} , and its relationship to $c \in Env$. In this example, c = 0.60. This means that we will be selecting our victims (parents) from the fittest 60% of the population. The number of victims we take is

[0.6*3] = 2. Suppose in our population, $eval(f_1)$, $eval(f_2) < eval(f_3)$. This means that the set $\{f_1, f_2\}$ contains the two parents to be considered from crossover. As a result, we are able to proceed with $Crossover(\{f_1, f_2\}, B)$.

3. Crossover or tree

In the crossover function, we call $OrTree(\{f_1, f_2\}, crossover)$ to complete the crossover.



Step by Step Example

Step 1: We start with a single leaf node (pr,?), which gets expanded into its alternatives pr_1, pr_2 , and pr_3 , where gp_1 gets assigned to sl_1, sl_2 , and sl_3 , respectively

Step 2: $(pr_1,?)$ and $(pr_2,?)$ are the two deepest leaves that last chose an assignment, $f_{crossEval}=0$, that is consistent with one of their parents. We randomly chose pr1 between them to expand into its alternatives pr_4 , pr_5 , and pr_6 , which assigns gp_2 to s_1 , s_2 , and s_3 respectively.

Step 3: $(pr_4,?)$ and $(pr_5,?)$ are the two deepest leaves that last chose an assignment, $f_{crossEval}=0$, that is consistent with one of their parents. We randomly chose pr_4 between them to expand into its alternatives pr_7 , pr_8 , and pr_9 , which assigns gp_3 to s_1 , s_2 , and s_3 respectively.

Step 4: We notice that pr_7 is unsolvable, so we change that leaf to (pr_7, no) .

Step 5: We notice that pr_9 is unsolvable, so we change that leaf to (pr_9, no) .

Step 6: Since pr_8 is the deepest leaf, we choose it to consider next, even though its last assignment is not consistent with either of its parents. Since pr8 can be solved, we solve this leaf and turn into (pr_8, yes)

4. Choose next extension function

Immediately after the crossover, we have 4 chromosomes (facts) in our $population_1$. Given we have reached 4 > Cap facts, we set $f_{wert}(crossover) = f_{wert}(mutation) = 1$, and $f_{wert}(delete) = 1 - f_{wert}(crossover) = 0$.

As a result, f_{select} will select delete(A, B) as the extension function for iteration 2.

The victims delete(A, B) receives is part of our definition of f_{select} , and its relationship to $g \in Env$. In this example, g = 0.40. This means that we will be selecting our victims (parents) from the least fit 40% of the population. The number of victims we take is

[0.4 * 4] = 1. Suppose in our population, $eval(f_1)$, $eval(f_2)$, $eval(f_3) < eval(f_4)$. This means that the set $\{f_4\}$ contains the fact to be considered for delete. As a result, we are able to proceed with $delete(\{f_4\}, B)$.

5. Delete

Delete will simply remove the fact f_4 from our population, creating

$$population_2 = \{f_1, f_2, f_3\}$$

Immediately after its completion.

6. Choose next extension function

Following delete, according to the same logic in step 2, we choose a new extension function. Suppose, this time, the function is mutate(A,B) and we randomly select a fact from our population such that

$$A = f_1 = \{(gp_1, s_1), (gp_2, s_2), (gp_3, s_3)\}$$

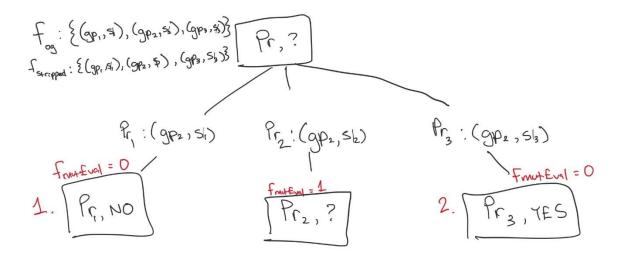
7. Mutation

Mutation takes in a fact, in this case f_1 , and strips a random single slot $sl \in Sl$ from the fact f_1 to create $f_{stripped} = \{(gp_1, s_1), (gp_2, \$), (gp_3, s_3)\}$. Let $f_1 = f_{og}$

Mutation then sends these two facts into our or tree as

$$orTree(\{f_{og}, f_{stripped}\}, mutation)$$

From here, we can begin or or tree search, with the initial state s_0 being $f_{stripped}$



Step 1:

We start with a single leaf node (pr, ?), which gets expanded into its alternatives pr_1, pr_2 and pr_3 , where the unnasigned gp_2 get assigned to sl_1, sl_2 and sl_3 respectively.

Step 2:

 $f_{mutEval}(pr_1) = f_{mutEval}(pr_3) = 0$, as the slot being assigned in each of these branches is different than the slot associated with gp_2 in our original fact f_{og} . As a result, $f_{mutEval}(pr_2) = 1$.

 $f_{leafExt}$ will need to select and tie break between pr_1 and pr_3 . It does so by selecting one of the possible two nodes randomly. Our selection, in this example, is pr_1 , which assigns, sl_1 to gp_2 .

Step 3:

In our example, the fact generated from the leaf $f_i = \{(gp_1, sl_1), (gp_2, sl_1), (gp_3, sl_3)\}$ violates some hard constraint. As a result, we set the variable sol in pr_1 to no.

Step 4:

 pr_2 , pr_3 are now the deepest unassigned leaves that we can now explore. Our next random selection is pr_3 , which assigns slot sl_3 to gp_2 . This assignment creates the fact $f_{mut} = \{(gp_1, sl_1), (gp_2, sl_3), (gp_3, sl_3)\}$ which, in our example, passes all hard constraints, and is complete. As a result, we can set sol = yes, and we are done our search.

We now have a new population of

$$population_3 = \{f_1, f_2, f_3, f_{mut}\}$$

8. Final population

Considering lim=3, and we have completed 3 iterations of our extension rules, we have now reached the state s_{goal} , and have completed our set based search.

And-Tree-Based Search

Re-iteration of Definition Legend:

$$G = Games = \{g_0, ..., g_n\}$$

$$P = Practices = \{p_0, ..., p_n\}$$

$$GP = Games \ and \ Practices = G \cup P$$

$$Sl = Slots$$

$$Constr(f) = \begin{cases} Fact \ meets \ all \ hard \ constraints \land Complete(f) & True \\ Otherwise & False \end{cases}$$

$$PartConstr(f) = \begin{cases} Fact \ meets \ all \ hard \ constraints^* & True \\ Otherwise & False \end{cases}$$

$$Complete(f) = \begin{cases} \forall \ sl \in f, sl \neq \$ & True \\ Otherwise & False \end{cases}$$

PartEval(f), $f \in F'$: Returns the sum of penalties from all violated soft constraints.

AConstr(a): Evaluates whether an individual assignment is valid. This is false if a game is in a practice-only slot, or vice versa.

 $Assign = \{(g, sl) | g \in GP, sl \in Sl, AConstr(a)\}$: Ordered pairs representing possible assignments from games or practices to their corresponding slots.

 $assign = GP \rightarrow Sl$ such that $\forall g (g \in GP \rightarrow (g, assign(g)) \in Assign)$: Surjection mapping a game or practice to a slot. Must be in Assign, to meet constraints. $partassign: GP \times (Sl \cup \{\$\})$: Initial assignment provided by input

 $passign = assign \cup GP \rightarrow \$$: Modified version of assign which allows null assignments

And-Tree-Based Search

Informal Description

Our other search paradigm is an And-Tree search, which utilizes a branch and bound search algorithm to find the best assignment of games and practices to slots assuming a valid input.

We start the search with whatever partial solution we are given. From there, we build from this parent solution by dividing the problem into n different sub-solutions, where each sub-

solution assigns the next game/practice to one of the n slots. Each sub-solution is then checked if it violates PartialConstr, which checks if the sub-solution violates one of the hard constraints. If this partial solution does violate PartialConstr, then we say that the sub-problem is solved, since we cannot build a valid full solution from it.

After this, we look at all the leaves we have in our tree that have not been solved yet and calculate a lower bound for each leaf based off what the best Eval evaluation you can theoretically get from turning that partial solution into a full solution; We call this function PartialEval. We continue the search on the various leaves until we reach a leaf that is a valid full solution to our problem. At this point, if it is a better solution than whatever we have in our Environment (Env) we replace it with the better one.

While we are searching through our tree, if we ever come across a partial solution that has a larger PartialEval than the Eval we get from the solution we have stored in our Env, we can prune, and instantly say that leaf is solved, since it is impossible for any full solutions built off it to be better than whatever solution we have stored in our Env.

At the end of the search, whatever solution is stored in our Env is the optimal solution to our problem. If no solution is stored in our environment, we can conclude that there are no solutions to the problem given.

Formal Description

We start our definition of the And-Tree search process by defining our model $A_{\wedge}=(S_{\wedge},T_{\wedge})$. Using the definitions for S_{\wedge} and T_{\wedge} given in lecture, we only need to define Prob for the definition A_{\wedge} of to be completed.

We define the problem space *Prob* as the following:

$$Prob := \{f'|f': GP \rightarrow Sl \cup \{\$\}, PartialConstr(f)\}$$

- Where f' is an assignment of Games & Practices to slots or null assignments "\$" and f' meets all hard constraints and Prob is a set of assignments.

To manage hard constraints on an assignment f we define PartialConstr(f) as follows:

$$PartialConstr(f') := \begin{cases} true & \textit{No hard constraints are currently violated by } f' \\ false & \textit{Otherwise} \end{cases}$$

Now that we have defined Prob, we want to define when Prob is solved (sol-entry "yes") or unsolved (sol-entry "?"). Before we can do that, however, we must define the function PartialEval, as well as our Env.

PartialEval(f')

 $:= \frac{\text{Counts how many soft constraints are currently violated by } f' \text{ and sum's their}}{\text{penalties together, not including the penalties for gamemin and practicemin}}$

- PartialEval(f') will be used as a way for us to measure a problem in a leaf.

$$Env = \left\{ (Bestsol, assign) | assign \in Prob, Bestsol = \left\{ \begin{matrix} \infty & assign = \$ \\ PartialEval(assign) & Otherwise \end{matrix} \right\} \right\}$$

Bestsol is set to infinity at the start to signify that no assignments have been made.

Our initial environment is $e_0 = (\infty, \$)$.

Now that we have defined everything in our search model, we need to determine when a problem is marked as sol-entry "yes" or sol-entry "?". Let $pr \in Prob$, and

$$e = (Bestsol, assign) \in Env.$$

We say pr is marked as sol-entry "yes" if and only if:

$$\neg PartialConstr(pr) \lor pr : GP \rightarrow Sl \lor (PartialEval(pr) > Bestsol)$$

- pr doesn't meet the hard constraints or
- pr is a complete schedule or
- pr has a worse (higher) evaluation score than our current best schedule.

And that $pr \in Prob$ is never marked as sol-entry "?" if it can't be expanded.

Now, to define how we can divide a leaf into its sub-problems we have the following definitions and their descriptions:

- 1. Let $prob \in Prob \ s.t. \exists i \in \mathbb{N}, 1 \le i \le n$ with $prob(a_i) = \$$ (Note: If this isn't true then prob is solved.)
 - Define a subproblem as an element of the larger problem such that it does not have an assignment at an index "i" in the range of possible assignments.
- 2. Let $h \in \mathbb{N}$, $1 \le h \le n$ s. t. $\forall j \in \mathbb{N}$, $1 \le j \le n$, $prob(a_j) \ne \$ \land prob(a_h) = \$$.
 - Then, define an "h" and a "j" such that the subproblem has a partial assignment at index "j" and no assignment at index "h".
- 3. $Divide(prob) := \{f : GP \to Sl \cup \{\$\} | f(a_i) = prob(a_i), \forall i \in \mathbb{N}, 1 \le i \le n, i \ne h, f(a_h) \ne \$\}$
 - Then, we define a "Divide" function that maps a partial assignment of Games and Practices to an available set of Slots such that this assignment corresponds to a subproblem for every index of that subproblem "prob" that can be expanded upon.

- 4. $Div(prob, pr_1, ..., pr_n)$ holds if and only if $Divide(prob) = \{pr_1, ..., pr_n\}$.
 - Finally, we define Div as the function that determines whether a subproblem is expandable by checking the result of Divide(subproblem).

Now that we have defined A_{\wedge} and Env, we only need to define our search control K_{\wedge} to complete our Process definition $P_{\wedge} = (A_{\wedge}, Env, K_{\wedge})$.

Our search control K_{Λ} chooses which state to transition to by using the following functions:

$$f_{eval}((pr,?)) := \begin{cases} 0 & if \ pr \ is \ solved \\ 1 + PartialEval(pr) & Otherwise \end{cases}$$

 $f_{leaf}(s)$ gives us a leaf with the minimal f_{eval} score in state s. In the event of a tie, choose the deepest leaf of the tied leaves. In the event of a second tie, randomly choose one of the tied deepest leaves to proceed with.

 $f_{trans}((pr,?))$ selects the only Div of pr that satisfies definition 4.

Whenever we come across a leaf (pr,?) such that $pr: GP \to Sl \land Contr(pr)$, we check our current environment $e = (Bestsol, assign) \in Env$. If Eval(pr) < Bestsol, then we take e' = (Eval(pr), pr) as our new environment. Once all leaves are solved, we take our current $e = (Bestsol, assign) \in Env$, and whatever assign is, will be our best solution. If assign = \$, then there are no valid solutions to the problem.

Now that we have defined out search process P_{\wedge} , we now need to define our search instance $Ins_{\wedge}(s_0,G_{\wedge})$. Using the natural definition of Ins_{\wedge} . We need to define our initial state s_0 and our Goal G_{\wedge} .

Our initial state is defined as follows:

$$s_0 = (partassign,?)$$

- Where partassign is an initial assignment as provided by a valid input.

Our Goal G_{Λ} is defined as:

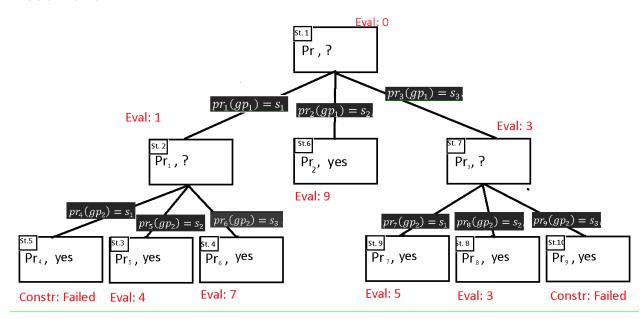
$$G_{\wedge}(s) = yes, if and only if$$

- 1.s = (pr', yes) or
 - o The root is sol-entry "yes", which means there's no valid solution.
- $2.s = (pr', ?, b_1, ..., b_n)$, $G_{\wedge}(b_1) = yes$ and the solutions to $b_1, ..., b_n$ meet our hard constraints, or
 - O All branches are sol-entry "yes" and are compatible, with the best solution stored in $e = (Bestsol, assign) \in Env$

3. We have explored all leaf nodes, they are all marked sol-entry "yes" by our definition, and they have no further possible transitions, again with our best solution stored in $e = (Bestsol, assign) \in Env$.

Now that we have defined our Process P_{Λ} , and Search Instance Ins_{Λ} , we have defined everything necessary for our And-Tree search model.

Visualization



Steps

- 1. We start with $s_0=(partassign,?)$, so [pr,?] which as we specified, has a default PartialEval=0. So, we divide it into its sub-solutions and assign gp_1 to s_1,s_2 , and s_3 respectively: $pr_1(gp_1)=s_1$, $pr_2(gp_1)=s_2$, $pr_3(gp_1)=s_3$.
- 2. Sub-solution s_1 derives $[pr_1,?]$ which has PartialEval=1 and can still be divided further into sub-solutions. Splitting into: $pr_4(gp_2)=s_1$, $pr_5(gp_2)=s_2$, $pr_6(gp_2)=s_3$, which we will find all to be full assignments and so they are given priority in order of evaluation.
- 3. pr_5 is solved (sol-entry "yes") and so we mark it as such: $[pr_5, yes]$, with it's Eval = 4. Currently our best (lowest) score, we store it in our Env.
- 4. pr_6 is solved (sol-entry "yes") and so we mark it as such: $[pr_6, yes]$, with it's Eval = 7, failing to beat our best solution, so we keep our environment unchanged.
- 5. pr_4 is violating a hard constraint, so we mark it as sol-entry "yes" so we can ignore it.
- 6. Back to pr_2 we find that it has a PartialEval = 9, which is worst than the solution we have stored in our environment and all other PartialEvals, so we needn't expand it and so mark it as sol-entry "yes" so we can ignore it.

- 7. pr_3 , like pr_1 and pr_2 , is derived from s_1 . We find that it can be divided further and since $PartialEval(pr_3)=3$ it's sub-solutions might be better than our current best one stored in the environment. So we mark it sol-entry "?": $[pr_3,?]$ and split from it into: $pr_7(gp_2)=s_1, pr_8(gp_2)=s_2, pr_9(gp_2)=s_3$, which we will find all to be full assignments.
- 8. pr_8 is solved (sol-entry "yes") and so we mark it as such: $[pr_8, yes]$, with it's Eval = 3, which beats our current best solution, so we replace the solution we stored in our environment with it.
- 9. pr_7 is solved (sol-entry "yes") and so we mark it as such: $[pr_7, yes]$, with it's Eval = 5, which does not surpass our best score, so we leave it.
- 10. pr_9 is violating a hard constraint, so we mark it as sol-entry "yes" so we can ignore it.
- 11. With that we are done searching as we have expanded our tree as much as possible and have stored the best solution among it's leaves in our environment, giving us the best solution of pr_8 .