# Working Title

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## 1 Model

We work with the stochastic model of [3], which considers the evolution over time of a repairable dependability system operating in a randomly changing environment. The system consists of a collection  $\Omega = \{1, 2, ..., N\}$  of  $N < \infty$  component types. Each component type  $i \in \Omega$  has a redundancy  $1 \le r_i < \infty$ , and the  $r_i$  components of type i are assumed to be identical. A component can be either operational (up) or failed (down).

The environment changes randomly within a set  $\mathcal{E} = \{0, 1, 2, ..., L\}$ . For example, we can think of the environment as representing the current load on the system, and if there are two possible environments, 0 and 1, then 0 (resp., 1) may represent a low (resp., high) load. Once the environment enters  $e \in \mathcal{E}$ , it remains there for an exponentially distributed amount of time with rate  $\nu_e > 0$ , after which the environment changes to e' with probability  $\delta_{e,e'} \ge 0$ , where  $\delta_{e,e} = 0$  and  $\sum_{e' \in \mathcal{E}} \delta_{e,e'} = 1$ . We assume the matrix  $\delta = (\delta_{e,e'} : e, e' \in \mathcal{E})$  is irreducible; i.e., for each  $e, e' \in \mathcal{E}$ , there exists  $k \ge 1$  and a sequence  $e_0 = e, e_1, e_2, \ldots, e_k = e'$  with each  $e_i \in \mathcal{E}$  such that  $\prod_{i=0}^{k-1} \delta_{e_i,e_{i+1}} > 0$ . In other words, it is possible to eventually move from environment e to environment e'.

The components in the system can randomly fail and then be repaired. When the environment is  $e \in \mathcal{E}$ , the failure rate and repair rate of each component of type i are  $\lambda_{i,e} > 0$  and  $\mu_{i,e} > 0$ , respectively. If there is only one environment e, i.e.,  $|\mathcal{E}| = 1$ , then the lifetimes and repair times of components of type i are exponentially distributed with rates  $\lambda_{i,e}$  and  $\mu_{i,e}$ , respectively. Exponential distributions are frequently used to model lifetimes of hardware and software components; e.g., see [5]. We assume that all operating components of a type i have the same failure rate  $\lambda_{i,e}$  in environment e. Thus, in a system with redundancies for which not all components of a type are needed for operation of the system, the extras are "hot spares" since they fail at the same rate as the main components.

Our model includes probabilistic instantaneous cascading failures, which occur as follows. The ordered set  $\Gamma_i$  specifies the types of components that a failure of a type-i component can cause to simultaneously fail. When a component of type i fails, it causes a single component of type  $j \in \Gamma_i$  to fail at the same time with probability  $\phi_{i,j}$  (if there are components of type j up), and we call  $\phi_{i,j}$  a component- affected probability. The events that the individual components of types  $j \in \Gamma_i$  fail immediately are independent. Thus, when a component of type i fails, there are independent "coin flips" to determine which components in  $\Gamma_i$  fail, where the coin flip for  $j \in \Gamma_i$  comes up heads (one component of type j fails) with probability  $\phi_{i,j}$  and tails (no components of type j fail) with probability  $1 - \phi_{i,j}$ .

We allow for a cascading failure to continue as long as there are still components operational in the system. For example, the failure of a component of type i may cause a component of type j to fail (with probability  $\phi_{i,j}$ ), which in turn makes a component of type k fail (with probability  $\phi_{j,k}$ ), and so on. As noted in [3], the SAVE package [1] allows for only one level of cascading, but the unlimited cascading in our model makes it significantly more difficult to analyze.

We can think of a cascading failure as a tree of simultaneously failing components. The root is the component, say of type i, whose failure triggers the cascade. The root's children, which are from  $\Gamma_i$ , are those components whose immediate failures were directly caused by the root's failure. At any non-root level of the tree, these components' failures were directly caused by the failure of their parents at the previous level. Although all the failing components in a cascade fail at the same time, we need to specify an order in which they fail for our problem to be well-defined, as we explain later in Section 2.1. We assume the components in a tree fail in breadth-first order.

There is a single repairman who fixes failed components using a processor- sharing discipline. Specifically, if the current environment is e and there is only one failed component, which is of type i, then the repairman fixes that component at rate  $\mu_{i,e}$ . If there are b components currently failed, then the repairman allocates 1/b of his effort to each failed component, so a failed component of type i is repaired at rate  $\mu_{i,e}/b$ .

## 2 Markov Chain

We want to analyze the behavior of the system as it evolves over time. Because of the processor-sharing repair discipline and the exponential rates for the event lifetimes, it will suffice to define the state of the system as a vector containing the number of failed components of each type and the current environment. Thus, let  $S = \{x = (x_1, x_2, \dots, x_N, x_{N+1}) : 0 \le x_i \le r_i \ \forall i \in \Omega, x_{N+1} \in \mathcal{E}\}$  be the state space, and let  $Z = [Z(t) : t \ge 0]$  be the continuous-time Markov time (CTMC) living on S keeping track of the current state of the system. (If we had instead considered a first-come-first-served repair discipline, then the state space would need to be augmented to keep track of the order in which current set of down components failed.) We assume that Z starts in environment  $0 \in \mathcal{E}$  with no components failed, i.e., state  $(0,0,\ldots,0)$ . As noted in [3] the CTMC is irreducible and positive recurrent.

We now describe the CTMC's infinitesimal generator matrix  $Q = (Q(x,y): x, y \in S)$ , where Q(x,y) is the rate that the CTMC Z moves from state  $x = (x_1, \ldots, x_N, x_{N+1})$  to state  $y = (y_1, \ldots, y_N, y_{N+1})$ . If  $y_i = x_i$  for each  $i \in \Omega$  and  $y_{N+1} \neq x_{N+1}$ , then (x,y) is an environment transition with  $Q(x,y) = \nu_{x_{N+1}} \delta_{x_{N+1},y_{N+1}}$ . If  $y_i = x_i - 1$  for one  $i \in \Omega$ ,  $y_j = x_j$  for each  $j \in \Omega - \{i\}$ , and  $y_{N+1} = x_{N+1}$ , then (x,y) is a repair transition corresponding to the repair of a component of type i, and  $Q(x,y) = x_i \mu_{i,x_{N+1}}/(\sum_{j\in\Omega} x_j)$ . If  $y_i \geqslant x_i$  for all  $i \in \Omega$  with  $y_j > x_j$  for some  $j \in \Omega$  and  $y_{N+1} = x_{N+1}$ , then (x,y) is a failure transition in which  $y_i - x_i$  components of type i fail,  $i \in \Omega$ . Any other (x,y) with  $x \neq y$  not falling into one of the above three categories is not possible, so Q(x,y) = 0. The diagonal entry  $Q(x,x) = -\sum_{y\neq x} Q(x,y)$ , as required for a CTMC; e.g., see Chapter 5 of [4].

We now determine the rate Q(x,y) of a failure transition (x,y). First consider the case when cascading failures are not possible, i.e.,  $\Gamma_i = \emptyset$  for each type i. Then the only possible

failure transitions (x, y) have  $y_i = x_i + 1$  for one  $i \in \Omega$ ,  $y_j = x_j$  for each  $j \in \Omega - \{i\}$ , and  $y_{N+1} = x_{N+1}$ , and this corresponds to a single component of type i failing. Then  $Q(x, y) = (r_i - x_i)\lambda_{i, x_{N+1}}$ .

Cascading failures complicate the computation of Q(x,y) for a failure transition (x,y). As mentioned before, a cascading failure is modeled as a tree T built from the multiset B of simultaneously failing components, where B has  $y_{\ell} - x_{\ell}$  failing components of type  $\ell$ ,  $\ell \in \Omega$ . A tree T has a rate

$$R(T) = (r_i - x_i)\lambda_{i,x_{N+1}} \rho \eta, \tag{1}$$

where

- $(r_i x_i)\lambda_{i,x_{N+1}}$  is the failure rate of the root (assumed here to be of type i),
- $\rho = \rho(T)$  is the product of the  $\phi_{j,k}$  terms for a parent node of type j causing a child of type  $k \in \Gamma_j$  to fail, and
- $\eta = \eta(T, x)$  is the product of the  $1 \phi_{j,k}$  terms from a node of type j not causing a component of type  $k \in \Gamma_j$  to fail when there are components of type k up.

A difficulty arises since there can be many such trees corresponding to the multiset B of components failing in (x, y), and calculating Q(x, y) requires summing R(T) over all possible trees T that can be constructed from B. The number of such trees grows exponentially in the number of failing components in the cascade; see [3].

## 2.1 Example of Computing a Tree's Rate

We now provide an example of computing the rate R(T) of a tree T. Let  $\Omega = \{A, B, C\}$ , with  $r_A = r_B = r_C = 4$ . Also, let  $\Gamma_A = \{B, C\}$ ,  $\Gamma_B = \{A, C\}$ , and  $\Gamma_C = \{A, B\}$ . Suppose that  $\mathcal{E} = \{0\}$ , and consider the failure transition (x, y) with x = (2, 2, 3, 0) and y = (4, 4, 4, 0). Thus, (x, y) corresponds to 2 components each of types A and B failing and a single component of type C failing. One possible tree T corresponding to (x, y) is shown in Figure 1. We assume the nodes in T fail in breadth-first order.

The nodes depicted as double circles form the tree of failing components. The single circles correspond to components in some  $\Gamma_i$  but did not fail. A component type j in some  $\Gamma_i$  could have not failed because either there are components of type j up at this point but its coin flip came up tails (with probability  $1 - \phi_{i,j}$ ), or there were no more components of type j up at this point. Each node has a label of the form i-ID, where i denotes the type of the component for that node, and ID is the position of the node in a breadth-first ordering of all the nodes (single and double circles). We include the IDs just to simplify the discussion here. We call the tree of all nodes the supertree corresponding to the tree T of failing nodes. The supertree is used to compute R(T) of T as follows. Let  $u_i$  be the number of components of type i available in the system. Since the root is a component of type I and there are I and the cascade is I and I are root then causes a component of type I to fail at node ID I, and this occurs with probability I and the node at ID I did not fail, and at this point there are I are I and I are components of type I still up, so this non-failure occurs

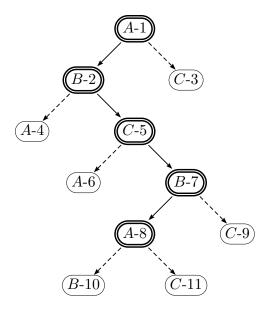


Figure 1: An example of a supertree.

with probability  $1 - \phi_{A,C}$ . Instead of stepping through the rest of the supertree one node at a time, we notice that all the  $\phi_{i,j}$  terms must be included if T is to correspond to (x,y). Thus, in (1) we have  $\rho = \phi_{A,B} \phi_{B,C} \phi_{C,B} \phi_{B,A}$ . We notice the following when calculating  $\eta$  in (1):

- $1 \phi_{i,j}$  terms are included if and only if there are still components of type j up.
- Each time we encounter a node of type j that has failed in the breadth- first enumeration of T, we decrement  $u_j$  by 1.

Keeping these observations in mind, we now calculate  $\eta$ . For component type A we have  $u_A=2$  before we traverse through T. As we do a breadth-first traversal through T, at ID 1,  $u_A=1$ . We see that  $u_A>0$  until ID 8. So  $\eta$  includes terms  $1-\phi_{B,A}$  and  $1-\phi_{C,A}$  from IDs 4 and 6, respectively. For component type B we have  $u_B=r_B-x_B=2$  before we traverse through T. Since components of type B have already been exhausted at ID 7, we do not include  $1-\phi_{A,B}$  at ID 10 in  $\eta$ . For component type C we see that  $u_C=1$  before we traverse through T, and  $u_C=0$  at ID 5. Hence, the only contribution to  $\eta$  from a component of type C not failing is  $1-\phi_{A,C}$  from ID 3. Taking the product over all component types yields  $\eta=(1-\phi_{B,A})(1-\phi_{C,A})(1-\phi_{A,C})$ . Therefore, R(T) is  $2\lambda_{A,0} \rho \eta$ . In our implementation  $\eta$  is calculated through a data structure called BFHist, which we describe later in Section 3.1.

We previously stated that the order in which the components fail in a tree must be specified for the tree's rate to be well defined. To see why, suppose instead that the components in Figure 1 fail in depth-first order. The depth-first traversal of T is A-1, B-2, A-4, C-5, A-6, B-7, A-8, B-10, C-11, C-9, C-3. Initially, the number of components up of each type are  $u_A = 2$ ,  $u_B = 2$  and  $u_C = 1$  as before. In this traversal a component of type C fails at ID 5, which makes  $u_C = 0$ .

Thus,  $\eta$  for the depth-first traversal does not include the terms  $1 - \phi_{A,C}$ ,  $1 - \phi_{B,C}$  and  $1 - \phi_{A,C}$  from the subsequent type-C nodes at IDs 11, 9 and 3. In contrast, the breadth-first traversal includes one  $1 - \phi_{A,C}$  term at ID 3. Hence, it is necessary to define the order in which the components fail, even though they fail simultaneously.

# 3 Algorithms

We now provide efficient algorithms for generating all possible trees and constructing the Q-matrix. A tree corresponds to a multiset of particular components failing, and cascading failures starting from different states can have the same multiset of components failing. Hence, a particular tree may correspond to several different transitions. Our algorithm generates each possible tree only once and determines all the transitions to which this tree corresponds. This avoids generating the same tree numerous times for each corresponding transition, as was originally done in [3]. Moreover, rather than building each new tree from scratch, as was done in [3], our current algorithm builds larger trees from smaller ones already considered, leading to additional savings in the overall computational effort.

Computing the rate (1) of a given tree depends on the state from which the cascading failure began and the multiset of components that fail in the cascade. We do not actually construct the supertree in our algorithm to compute a tree's rate, but instead build a breadth-first history to keep track of the information necessary to compute  $\eta$  in (1). This breadth-first history concisely recounts the creation of the tree and thus allows us to obtain  $\eta$  without having to build supertrees per transition as was done in [3]. This is all done in Algorithms 1 (SeedTrees), 2 (AddTreeLevel) and 3 (ComputeTreeRate).

SeedTrees starts the tree generation and initializes the necessary data structures for AddTreeLevel. AddTreeLevel adds a new level to an existing tree in a recursive fashion, updates  $\rho$  in (1) to include the component-affected probabilities, and builds the tree's breadth-first history. ComputeTreeRate computes the rate of a completed tree for all the transitions it corresponds to using  $\rho$  and  $\eta$  computed from the breadth-first history populated in AddTreeLevel. We will now discuss each algorithm in detail. References to line numbers in the algorithms are given within angled brackets  $\langle \ \rangle$ .

### 3.1 SeedTrees

SeedTrees loops through all component types to choose a root for the tree to be constructed  $\langle 1 \rangle$ . One of the data structures initialized in SeedTrees is *BFHist*, which stores the breadth-first history using an array of linked lists, where the array is indexed by component type. Each linked list stores the respective parents of nodes exclusive to the supertree and stores the symbol @ for the nodes that did fail in the tree.

The dynamic array level contains the current bottom level of the tree that we are building, and it initially contains the root, which is of type  $rootC.type \langle 5 \rangle$ . We add the symbol @ to the list at BFHist[rootC.type] to denote that one component of the root's type has failed. We update nFailed by setting the counter for rootC.type to  $1 \langle 5-7 \rangle$ . We initialize  $\rho$  in (1) to 1 since no components so far have been caused to fail by the failure of other components  $\langle 8 \rangle$ .

If the component type rootC.type at the root cannot cause any other components to fail, only the trivial tree of one node can be made with this type of root. We then evaluate this single-node tree's rate in ComputeTreeRate because it cannot be grown further. If  $\Gamma_{rootC.type} \neq \emptyset$  (i.e., it can cause other types of components to fail), then we call AddTreeLevel to proceed with building taller trees by adding another level to the current tree  $\langle 9-13 \rangle$ .

### **Algorithm 1** SeedTrees( $\Gamma$ )

where  $\Gamma$  is an array of ordered sets that describes which components can cause which other components to fail

```
1: for rootC.type \in \Omega do
      level = []; {Dynamic array of failed components at tree's current bottom level}
3:
      nFailed = [0, 0, \dots, 0]; {Array that counts failed components of each type in the tree}
      BFHist = [(), (), ..., ()]; {Array of linked lists that keeps a history of parent compo-
4:
      nent types in breadth-first order; BFHist is indexed by component types}
      add rootC.type to level;
5:
      nFailed[rootC.type] = 1;
6:
      add @ to BFHist[rootC.type]; {Signifies one component of type rootC.type has failed}
7:
      \rho = 1; {initialize product of component-affected probabilities}
8:
     if \Gamma_{rootC.tupe} == \emptyset then
9:
        ComputeTreeRate(nFailed, BFHist, \rho, rootC.type);
10:
11:
12:
        AddTreeLevel(level, nFailed, BFHist, \rho, rootC.type);
13:
      end if
14: end for
```

### 3.2 AddTreeLevel

Given the bottom-most tree level, AddTreeLevel determines all the possibilities for the next level that can be added to the current tree by taking the Cartesian product of the power sets of the nonempty  $\Gamma_i$  for each of the failed components at the current bottom level. For each possible next level, we recursively call AddTreeLevel. We use  $\mathcal{P}$  to denote the power set-like operation on an ordered set. In our realization of the power-set operation, components in each subset maintain their relative ordering from the original set. Note that by only maintaining one bottom-most level at a time, as opposed to passing the entire tree per recursive call, we save some memory.

We implement this power set-like operation by generating all possible binary numbers with  $\sum_{i=1}^{|level|} \Gamma_{level[i]}$  bits. A total of  $\prod_{i=1}^{|level|} 2^{\Gamma_{level[i]}}$  such binary numbers are generated. In the binary number, 1 denotes a failed node and 0 denotes a node that is included in a  $\Gamma$  set but did not fail. If we have a tree where none of the leaf nodes at the current bottom level can cause any other components to fail (i.e., have empty  $\Gamma_i$ s), we get no nextLevelPossibilities

 $\langle 1 \rangle$ .

Otherwise, we choose one possible choice for the failed components in the next level from the nextLevelPossibilities  $\langle 2 \rangle$ . Each node in T has three attributes: a component type i; an ID, which is its breadth-first ID in the supertree; and a parentID, which is the breadth-first ID of this node's parent in the supertree. For each node parent C in the current bottom level that acts as a parent, potentially causing other nodes to fail, we iterate through all of its possible children, i.e., through  $\Gamma_{parentC.type}$ . If any of these children actually fail, then they will be members of the set oneNextLevelPossibility  $\langle 4-6 \rangle$ . Now if the redundancy of the component type we just tried to add as a child has already been exhausted, it cannot fail and hence our tree is invalid and we move on to the next possibility  $\langle 10-12 \rangle$ .

If a child of type childC.type has failed (i.e., it exists within nextLevelPossibilities), the relevant data structures are updated to signify this occurrence. We increment the number of failed components of type childC.type, add the symbol @ to BFHist[childC.type] (to mark that a failure of this type has occurred at the current location in the tree), and update  $\rho$  by multiplying it by  $\phi_{i,j} \langle 11-13 \rangle$ . If the child does not actually fail, then we add parentC.type to BFHist[childC.type]. Later in ComputeTreeRate for each transition (x,y), we evaluate which of these children could have failed but did not, and which could not have failed because their type's redundancy was exhausted.  $\langle 14-16 \rangle$ .

We do the above updates to the data structures for each potential parent node in level and each of its children in oneNextLevelPossibility. If at least one child has been added, we recursively add another level to the current tree  $\langle 19-23 \rangle$ . Otherwise, this tree has not changed in this pass through AddTreeLevel and we call ComputeTreeRate to compute the rate of the finalized tree  $\langle 24-26 \rangle$ . We make sure that trees are not double counted because once a tree passes through AddTreeLevel unmodified, it is processed and discarded. We avoid making duplicate trees because each oneNextLevelPossibility is unique.

## **Algorithm 2** AddTreeLevel(level, nFailed, BFHist, $\rho$ , rootC.type)

where *level* is the current level of failed components, nFailed counts failed components by type in the tree, BFHist is breadth-first history,  $\rho$  is a cumulative product of component-affected probabilities, root C. type is the root component's type in the current tree

1: 
$$nextLevelPossibilities = \underset{\substack{i=1:\\\Gamma_{level[i]} \neq \varnothing}}{\overset{|level|}{\times}} \mathcal{P}(\Gamma_{level[i]});$$

{Builds all possibilities for the next level (given the current level) by taking a Cartesian product of the power sets of non-empty  $\Gamma$  sets of failed nodes in the current level

- 2: for  $oneNextLevelPossibility \in nextLevelPossibilities$  do
- addedAChildFlag = False;3:
- 4: for  $parentC \in level$  do
- 5:
- $\begin{array}{l} \textbf{for } i \in \Gamma_{parentC.type} \ \textbf{do} \\ \textbf{if } \exists \ childC \in oneNextLevelPossibility : childC.type == i \ \&\& \ childC.parentID == \\ \end{array}$ 6: parentC.ID then

```
addedAChildFlag = True;
7:
             if nFailed[childC.type] == r_{childC.type} then
8:
               goto line 2; {Invalid Tree, it requires more components than available}
9:
             end if
10:
             nFailed[childC.type] = nFailed[childC.type] + 1;
11:
             add @ to BFHist[childC.type]; {One component of type childC.type has failed}
12:
13:
             \rho = \rho * \phi_{parentC.type, childC.type};
             {Update rate with appropriate component-affected probability}
14:
          else
             add parentC.type to BFHist[childC.type]; {One component of type childC.type
15:
             has not failed, but was present in \Gamma_{parentC.type}
          end if
16:
        end for
17:
      end for
18:
      if addedAChildFlag then
19:
        AddTreeLevel(oneNextLevelPossibility, nFailed, BFHist, \rho, rootC.type);
20:
        {Tree can be grown further}
      else
21:
        ComputeTreeRate(nFailed, BFHist, \rho, rootC.type);
22:
        {Current tree is complete because it cannot be grown further}
23:
      end if
24: end for
```

## 3.3 ComputeTreeRate

In ComputeTreeRate, for a given tree, we determine all of the transitions (x, y) in the Qmatrix that use this tree, and then update the total rates of each of those transitions (x, y)by adding in the rate of the current tree to the current rate for (x, y). However, even if
several transitions use the same tree, the rate of the tree may differ for those transitions
since the failure rate of the root and  $\eta$  are transition-dependent.

Let  $x' = (x_1, x_2, ..., x_N)$  and  $S' = \{(x_1, x_2, ..., x_N) : 0 \le x_i \le r_i, \forall i \in \Omega\}$ . In line 1 of ComputeTreeRate, we only loop through  $x' \in S'$  and not  $x \in S$  because the structure of a tree does not depend on the environment; only the tree's rate does. Also, as we explained in the example tree-rate calculation in Section 2.1, computing  $\eta$  in (1) requires keeping track of the number  $u_i$  of components of each type i still up as we perform a breadth-first traversal through the supertree.

Each time we encounter the symbol @ in the *BFHist*, we reduce the number available by one  $\langle 5-7 \rangle$ . If there are components still available, we update  $\eta \langle 8-9 \rangle$ . As soon as  $u_i$  reaches zero, we break because that type has then been exhausted and can no longer fail  $\langle 10-12 \rangle$ .

Once we have finished calculating  $\eta$ , we loop through all environments since the same tree can be used for transitions occurring in different environments  $\langle 15 \rangle$ . We convert  $x' \in S'$  into a state  $x \in S$  by appending an environment  $e \langle 16 \rangle$ . We generate a potential state y by

adding nFailed to x' and appending environment  $e \langle 17 \rangle$ . Invalid states y will be generated in some instances because simply adding nFailed will cause some component types' number failed to exceed their redundancies  $\langle 18 - 20 \rangle$ . We finally compute the rate of the tree for the transition (x, y) using (1), and add this to the current cumulative rate Q(x, y) for the transition  $\langle 21 \rangle$ .

### **Algorithm 3** ComputeTreeRate(nFailed, BFHist, $\rho$ , rootC.type)

where level is the current bottom level of failed components, nFailed counts the failed components of each type in the tree, BFHist is breadth-first history,  $\rho$  is a cumulative product of component-affected probabilities, rootC.type is the root's type in the tree

```
1: for x' \in S' do
      \eta = 1; {Cumulative product of complement probabilities of components that could
      have failed but did not
      for i \in \Omega do
 3:
         u_i = r_i - x_i';
 4:
         for parentC.type in BFHist[i] do
 5:
           if parentC.type == @ then
 6:
              u_i = u_i - 1;
 7:
           else if u_i > 0 then
 8:
              \eta = \eta * (1 - \phi_{parentC,tupe,i});
 9:
           else
10:
              break; {need u_i > 0 or else there cannot be any more failed nodes of type i}
11:
12:
           end if
         end for
13:
      end for
14:
      for e \in \mathcal{E} do
15:
         x = (x', e);
16:
         y = (x' + nFailed, e);
17:
         if y is not a valid state then
18:
19:
           continue;
         end if
20:
         Q(x,y) = (r_{rootC.type} - x[rootC.type]) * \rho * \eta;
21:
      end for
22:
23: end for
```

## 3.4 Computing a Tree's Rate Revisited

We now use our example in Section 2.1 to show how to compute  $\eta$  from (1) using the data structure BFHist for Figure 1. We assume the tree in Figure 1 was first constructed from several recursive calls to AddTreeLevel, which also built BFHist as follows.

1	4	@-1	<i>B</i> -2	C-5	@-8
I	3	@-2	@-7	A-8	
(	7	A-1	@-5	<i>B</i> -7	A-8

We then proceed to ComputeTreeRate to compute  $\eta$ . As in Section 2.1, prior to iterating through *BFHist*, we have  $u_A = 2$ ,  $u_B = 2$  and  $u_C = 1$ . Iteration through *BFHist* occurs as specified in ComputeTreeRate  $\langle 3 - 14 \rangle$ .

- Iteration through the linked list at index A is as follows: The @-1 means a component of type A has failed at ID 1, so we then decrement  $u_A$  to 1. Then for the next two entries, since  $u_A$  is still positive,  $\eta$  includes  $(1 \phi_{B,A})$  and  $(1 \phi_{C,A})$  from B-2 and C-5 respectively in its product. Finally @-8 means a component of type A failed at ID 8, so we decrease  $u_A$  to 0.
- Iteration through the linked list at index B is as follows: @-2 and @-7 mean components of type B have failed at IDs 2 and 7 respectively, so we decrement  $u_B$  from 2 to 1 and then from 1 to 0. Since  $u_B$  is now 0, we cannot include any terms from this row in  $\eta$ .
- Iteration through the linked list at index C is as follows: Since  $u_C$  starts out positive, we include  $(1 \phi_{A,C})$  from A-1. @-5 means a component of type C has failed at ID 5, so we decrement  $u_C$  to 0. Since  $u_C = 0$ , again we cannot include any terms from this row in  $\eta$ .

Multiplying the contributions from each index results in  $\eta = (1 - \phi_{B,A})(1 - \phi_{C,A})(1 - \phi_{A,C})$ .

# 4 Performance Analysis

**TODO** Discuss MTTF and SSU computations, and how we utilize Up-state condition to parition S into U and U'.

When computing dependability measures such as the mean time to failure (MTTF) and steady-state unavailability (SSU), we rely on third-party libraries to perform the matrix operations [2]. We can easily swap out libraries for these tasks, so there is no need to discuss the performance of the Q-matrix operations. Note that the models with larger numbers of states do not necessarily correspond to larger numbers of trees. The time to generate trees (TGT) increases as the number of trees generated increases, and the number of trees increases as the number of cascading failures increases. When models have a state-space larger than 1000 states, the overhead from the matrix operations significantly effects the total computation time (CT).

TODO Introduce Models, Show Total Run times, TGT, MTTF and SSU for each model

States	Comp. Types	Redudancies	$\Gamma_{sets}$	Env.
81	A, B, C, D	$r_A = 2, r_B = 2,$	$\Gamma_A = \{B, C\}, \ \Gamma_B = \{A, D\},$	1
		$r_C = 2, r_D = 2$	$\Gamma_C = \{\}, \ \Gamma_D = \{A, B, C\}$	
288	A, B, C, D	$r_A = 3, r_B = 3,$	$\Gamma_A = \{B, C\}, \ \Gamma_B = \{A, C\},$	2
		$r_C = 2, r_D = 2$	$\Gamma_C = \{B, D\},  \Gamma_D = \{C\}$	
640	A, B, C, D	$r_A = 4, r_B = 3,$	$\Gamma_A = \{B, C\}, \ \Gamma_B = \{A, C\},$	2
		$r_C = 3, r_D = 3$	$\Gamma_C = \{B, D\},  \Gamma_D = \{B\}$	
125	A, B, C	$r_A = 4, r_B = 4,$	$\Gamma_A = \{B, C\}, \ \Gamma_B = \{A, C\},$	1
		$r_C = 4$	$\Gamma_C = \{A, B\}$	
1944	A, B, C, D, E, F	$r_A = 2, r_B = 2,$	$\Gamma_A = \Gamma_B = \{C, D\},$	2
		$r_C = 2, r_D = 2,$	$\Gamma_C = \{A, E\}, \ \Gamma_D = \{B, F\},$	
		$r_E = 2, r_F = 2$	$\Gamma_E = \{\}, \Gamma_F = \{\}$	

Table 1: Relation of TGT vs number of trees across various models

States	Trees	TGT	СТ
81	978	0.12	0.58
288	4507	0.26	0.78
640	27746	1.64	2.48
125	321372	6.22	6.69
1944	6328	1.42	21.21

Table 2: Relation of TGT vs number of trees across various models

The number of trees can grow exponentially in the number of components in the cascade (see [3]), and this limits the size of the models that our algorithm can handle. To address this issue, we explored heuristic approaches that reduce the computational effort by halting the generation of trees based on a stopping criterion. We insert the necessary breaking condition in AddTreeLevel as line  $\langle 18.1 \rangle$ , which will cause the program to skip over to the next enumeration of the bottom-most tree level. By skipping over the bottom-most tree level we don't generate a particular tree and thus neither evaluate its rate nor update the Q-matrix with said rate. The goal is to generate trees with relatively higher rates to get an approximate MTTF and SSU while saving time by skipping trees with relatively lower rates.

#### New lines inserted as <18.1, 2, 3>

- 1: **if** stopping criterion satisfied **then**
- 2: goto line 2;
- 3: end if

### Stopping Criteria

#### 1. $height > \tau_h$

We first consider the height threshold, where we stop generating trees after the height of the current tree reaches  $\tau_h$ . It is necessary to create a global variable height to keep track of the height. Our implementation requires a slight modification to AddTreeLevel, where we introduce the height as a method parameter. We also modi-

fied line  $\langle 20 \rangle$  to increment the *height* on every successive call.

2. 
$$\sum_{i=1}^{N} nFailed[i] > \tau_n$$

We also considered the *node threshold*, so constructed trees will contain a maximum of  $\tau_n$  failed components.

3.  $\rho < \tau_r$ 

We also used a rate threshold, where we stop generating trees after  $\rho$  in (1) drops below  $\tau_r$ .

Increasing  $\tau_h$  and  $\tau_n$  leads to a monotonically increasing number of generated trees, whereas the number of generated trees decreases monotonically in  $\tau_r$ . Setting  $\tau_h = \tau_n = \infty$  or  $\tau_r = 0$  leads to generating all trees, in other words, we do not approximate MTTF and SSU and Q = Q'. We can also get the exact values for the dependability measures by choosing thresholds

$$\tau_{h_0} = \tau_{n_0} = \sum_{i=1}^{N} r_i \text{ and } \tau_{r_0} = \max_{i,j \in \Omega: j \in \Gamma_i} \phi_{i,j}.$$

Although  $\tau_r$  starts at zero, we must provide an upper bound so that we can analyze and compare all of the thresholds with the same control variable. Since we halt tree generation prematurely, the resulting matrix Q' (that sums the rates over trees that are actually generated) can differ from the matrix Q that results from summing over all trees. Solving for the MTTF and SSU with Q' rather than Q leads to inaccuracies in the values for these dependability measures.

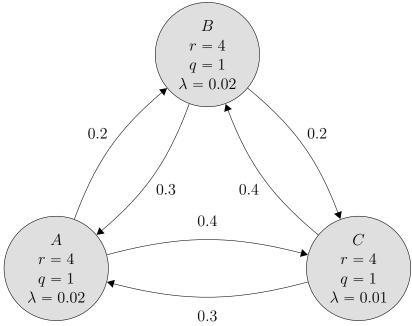


Figure 2: Graph representation of the 125-state model **TODO** Discuss the utility of giving model details

## 4.1 Enumerating thresholds

For the model shown in Figure 2, we try setting different values for the thresholds to examine the trade-off in the TGT and errors in the resulting MTTF and SSU. Starting from i = 0, we can calculate the next subsequent value of a threshold based on the index i. Here are governing recursive equations for choosing our thresholds:

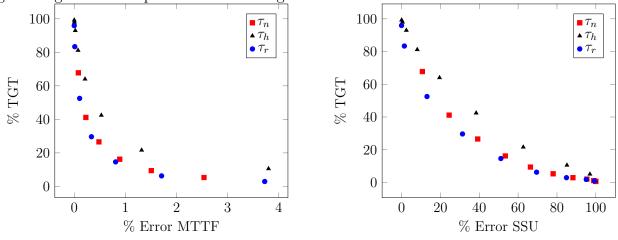


Figure 3: Two graphs showing the relation between % TGT and %Error in MTTF/SSU

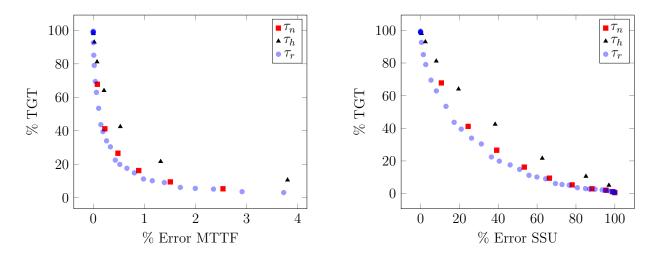


Figure 4: This graph shows an alternative view with smaller stepping in  $\tau_r$ 

Figure 4 plots the (TGT, MTTF) pairs for the different stopping criteria considered using different values for each threshold. Decimal values for  $\tau_n$  and  $\tau_h$  are not valid, since we do not define a fraction of a component. On the other hand,  $\tau_r$  can be sliced into more points with a slight modification to our recursive equation. The true value of the MTTF, obtained when all trees are generated, is 44.124, and this defines the horizontal asymptote to which the approximate MTTFs for each stopping criterion are approaching as TGT increases.

Points closer to the origin of the graph represent more efficienct thresholds. For a fixed TGT, the stopping criterion based on  $\tau_r$  leads to an approximate MTTF with lower error

than the other thresholds. Also, for a fixed approximate MTTF value, the stopping criterion based on  $\tau_r$  again leads to a smaller TGT than the other thresholds. Thus, the points plotted from using  $\tau_r$  define the efficient frontier.

## 4.2 Choosing an appropriate $\tau_r$

Use Djikstra's algorithm to find the shortest path with the largest rate.

### 5 Remarks

## 5.1 MTTF Computation

We remove the down-states from the matrix to speed up MTTF calculation.

### 5.2 Bookmarks

We do not store all failed components *BFHist*, we simply multiply until we hit a not-failed component. OR We cache BFH between succesive @ symbols to get complement cumulative component affected probabilities in constant time.

## References

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