

**Title:** Technical Report

**Project Title:** Connectivity and coverage of underwater sensor networks

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## Abstract

## 1 Introduction

Interest of monitoring underwater environment is growing day by day. Sensor network is a promising tool for underwater environment monitoring because of its remote monitoring and control technology. Underwater Sensor networking is using for military surveillance[8], disaster prevention, assisted navigation, offshore exploration, tsunami monitoring and oceanographic data collection. Coverage and connectivity are two important aspect of underwater sensor network. One can determine the quality of surveillance of a underwater wireless sensor network from network coverage. Many to one data flow from a set of sources to a common sink over a tree based routing topology, is a fundamental communication architecture of a underwater sensor networks. Connectivity is an

Table 1: Physical Layer Parameters

Parameter Name	Value	Description
Standard	IEEE 802.11	Can be used as medium access control protocol (MAC) for underwater acoustic communication with modification[5].
Range	1m-10km [6]	Short range modem provides very high bandwidth (typically MHz or more) where long range modem provides low bandwidth( several bps)
Data Rates	bps - MBps	Varies from standard to standard from 5 bps to 19200 bps.
Energy Source	battery, external sources	Primarily battery is used as a power source but for most of the standard external power supply can be used as a power supply.
Well-known models	LinQuest Inc, Hydro International	UWM1000,UWM2000,S1510 Radio Modem,Digital Underwater Modem UM 30
Path loss model		Path loss of underwater acoustic communication channel depends on the distance between the transmitter and receiver and signal frequency [?].

important issue for UWSN in-order to perform localization [16], [17],[4], routing [12],[15],[9]. In this task we are interested to measure the connectivity of UWSN. We are considering a network consists of anchored sensor nodes or free moving sensor nodes. The connectivity we are interested, can be

- All node connectivity: every node is connected with sink.
- 2-terminal connectivity: is the connectivity that two given vertices, called the source and the sink, can communicate.

- k-terminal connectivity: compute the probability that every operational pair of sites in  $k$ -can communicate with each other.
- In [1] authors propose a distributed node deployment scheme which can increase the initial network coverage in an iterative basis. They assuming that the nodes are initially deployed at the bottom of the water and can only move in vertical direction in 3-D space. The idea is to relocate the nodes at different depths based on a local agreement in order to reduce the sensing overlaps among the neighbouring nodes. Redundancy is observe by one of the node called Leader. It utilizes vertex colouring problem formulation in-order to determine coverage overlap. The nodes continue to adjust their depths until there is no room for improving their coverage. They consider both tether and and untethered architecture for node deployment.
- The work of [17] highlighted the localization of a sensor node. According to their model, the moving speed of underwater node changes continuously and shows semi-periodic properties. They predict the future mobility pattern from the past mobility information. The nodes maintain connectivity in-order to provide accurate localization information.
- In [3] authors proposes Meandering Current Mobility Model (MCM) which is able to capture the physical movement of the sensor nodes with ocean currents. In MCM, nodes are moving by the effect of meandering sub-surface currents and vortices. According to MCM, there is a strong correlations between nearby sensors. Vertical movements in ocean are negligible with respect to horizontal ones. Thus, in their model they neglect vertical displacement which makes mobility in 2D. This model is more realistic than other mobility models [7][17] for UWSNs since nodes are drifted according to the movement of the ocean. They studied dynamic coverage and connectivity as a function of time under the MCM model. They also consider the effect of different deployment strategies on network coverage and connectivity.
- Connectivity and coverage are important issue for UWSN. In [2] authors tackling the problem of determining how to deploy minimum number of sensor nodes so that all points inside the network is within the sensing range of at least one sensor and all sensor nodes can communicate with each other, possibly over a multi-hop path. They used sphere-based communication and sensing model. They place a node at the center of each virtual cell created by truncated octahedron-based tessellation. They provide solutions for both limited and full communication redundancy requirements.
- The work of [10], [11] extends [3] and proposes a ring like motion to capture the basic of sea flow. According to this model, there are two types of mobility, uncontrollable mobility which breaks the coverage of sensor network and controllable mobility which restore the coverage of sensor network. In the ring-like model [10], they consider local variety and main circulation in-order to capture some characteristics of water bodies. They use probability in-order to determine the next position of a node.
- In [13], [14] author proposed a polynomial algorithm to solve Steiner tree problem. The Steiner tree problem in an undirected graph is the problem of finding a tree spanning a pre-specified set of vertices at minimum cost where the cost of a tree is equal to the sum of the cost of its edges.

We device a scheme where we approximate the UWSN by using a special types of graph which is called partial  $k$ -tree. According to our scheme each node can be located into a set of regions with certain probabilities which is known as probabilistic locality set. The locality set of a node at a

certain time instant is known from the initial location of the node and underwater current.

We introduce a problem called the Connectivity in Underwater Sensor Network (CUSN) problem for Underwater Wireless Sensor Network (UWSN). We have a set of nodes and we have a probabilistic locality set of each node. We would like to compute the probability that the network is connected.

## 2 System Model and Problem Formulation

In the section we are going to describe notation, connectivity and coverage model and node connectivity model. Finally we present problem formulation for  $Conn(G, \mathbf{R})$ .

### 2.1 Notation

In this section we are going to describe some of the notations, we used throughout this paper.

- $G = (V, E)$  : the underlying connectivity graph of a given UWSN.
- $R_{tr}(v)$  : the transmission range for node  $v$ .
- $R_v = \{r_{(v,1)}, r_{(v,2)}, \dots\}$  : the locality set for node  $v$ .
- $\mathbf{R} = \{R_v : v \in V\}$  : the set of all the locality sets of all node in  $G$ .
- $p(r_{(v,i)})$  : the probability that node  $v$  is in region  $r_{(v,i)}$ .
- $PES$  : an ordering  $(v_1, v_2, \dots, v_{n-k})$  of the nodes of  $G$  for every  $i \in \{1, 2, \dots, n-k\}$  :, the node  $v_i$  is simplicial in the subgraph of  $G$ .
- $\{K_{(v,1)}, K_{(v,2)}, \dots, K_{(v,k)}\}$  : are  $k$ -cliques associated with vertex  $v$  in a  $k$ -tree, where  $v$  is a simplicial vertex.
- $K_{(v,base)}$  : the base clique to which  $v$  is attached and it is formed by all adjacent vertices of  $v$ .
- $\{T_{(v,1)}, T_{(v,2)}, \dots, T_{(v,k)}\}$  : tables associated with clique  $\{K_{(v,1)}, K_{(v,2)}, \dots, K_{(v,k)}\}$  respectively.
- $V(G')$  : the nodes of a subgraph  $G'$ .
- $Reach(r_{(x,i)}, r_{(y,j)}) = 1$  if node  $x$  in region  $r_{(x,i)}$  can reach node  $y$  in region  $r_{(y,j)}$  else 0.
- $Conn(G, \mathbf{R})$  : The probability that the nodes of  $G$  are in a state where all nodes are connected  $= \sum Pr[S : S \text{ is a connected state of nodes in } G]$ .

### 2.2 Connectivity and Coverage Model

Underwater Wireless Sensor Network (UWSN) is modelled by a graph  $G$  of  $(V, E)$  where  $V = (v_1, v_2, v_3, \dots, v_n)$  is a set of vertices and  $E$  is a set of edges. The transmission range for node  $v \in V$  is  $R_{tr}(v)$ . One node can transmit data to other node if they are within transmission range of each other.

### 2.3 Location Probability Model

We are considering each sensor node  $v \in V$  can be located into a set of regions,  $R_v = \{r_{(v,1)}, r_{(v,2)}, \dots\}$  with certain probabilities,  $p(r_{(v,1)}), p(r_{(v,2)}), \dots$  which are called location probabilities. Mathematical sum of a node's location probabilities are 1 but in some cases we are considering a set of regions with high probable values where sum of locations probabilities can be less than 1. Location probability of a node in a specific region, is independent of location probability of that node located into other regions. Also location probabilities of one node is independent of the location probabilities of other nodes. Also we are considering the probable locations of a node can be contiguous as well

as non-contiguous.

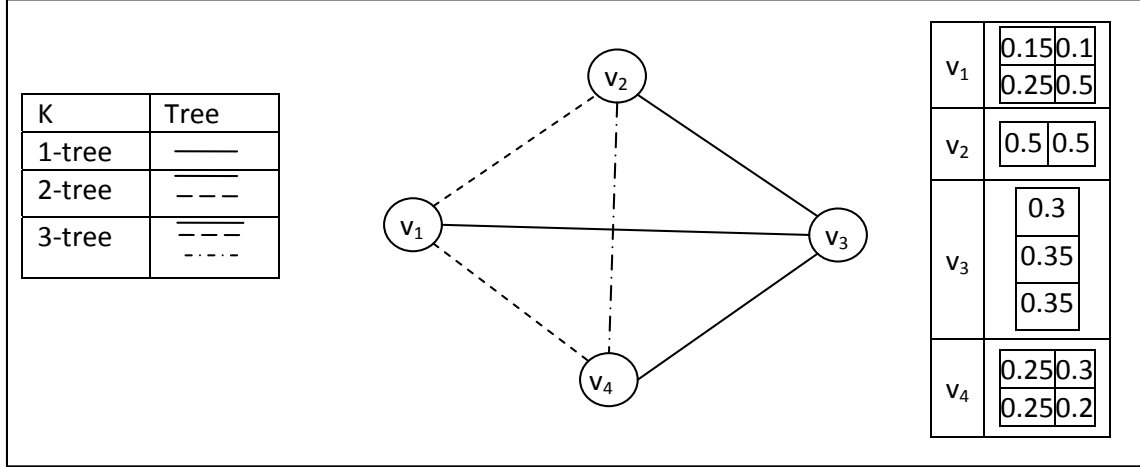


Figure 1: A partial 3-tree with 4 nodes.

**Example 2.1.** Fig. 1 illustrates a network of 4 nodes. Locality set of node  $v_1, v_2, v_3$  and  $v_4$  has 4, 2, 3 and 4 regions. We are also considering  $v_4$  as a sink node. Transmission range  $R_{tr} = 8.5$  unit. By using this transmission range node  $v_1$  from any of its probable regions can communicate with all probable regions of node  $v_2$  and  $v_3$  but not with node  $v_4$ . Similarly node  $v_2$  from all probable regions can communicate with all probable regions of node  $v_4$  and node  $v_1$  but for node  $v_3$  only the region with probability value 0.3. Node  $v_3$  can communicate with node  $v_4$  when  $v_4$  is located in the regions with probability values 0.25 and 0.2. ■

## 2.4 Node Connectivity Model

If one or more regions in the locality set of node  $x \in V$  is within the transmission radius  $R_{tr}(y)$  of one or more regions in the locality set of node  $y \in V$  and vice-versa then node  $x$  is reachable from node  $y$ .

A network state,  $S$  arises when each node is assigned to a specific region from its locality set. A connected state,  $S_{conn}$  is a network state where each node can communicate with others.

## 2.5 Problem Statement

Now we formally define the problem.

**Definition** (The  $Conn(G, \mathbf{R})$  Problem). Let  $G$  is a UWSN where each node  $v \in V$  can be located into a set of regions  $R_v = \{r_{(v,1)}, r_{(v,2)}, \dots\}$  with probability  $p(r_{(v,i)})$  where  $i = 1, 2, \dots$ . Also  $R_{tr}(v)$  is the transmission radius for node  $v \in V$ . we would like to find the probability  $Conn(G, \mathbf{R})$  that each node is connected. ■

An exhaustive approach to compute  $Conn(G, \mathbf{R})$  can be

- compute  $Reach(r_{(x,i)}, r_{(y,j)})$  for every pair of region  $r_{(x,i)}, r_{(y,j)}$  where  $(x, y) \in V$
- find all network states  $\mathbf{S}$  for the given graph  $G$ .
- find all connected states  $\mathbf{S}_{conn} \subset \mathbf{S}$

- calculate  $Conn(G, \mathbf{R}) = \sum Pr[S : S \in \mathbf{S}_{conn}]$  where  $Pr[S]$  can be computed by multiplying all nodes specific regional probability in state  $S$ .

Running time of the exhaustive algorithm can be calculated as follows

- let  $f(n, \mathbf{R})$  is the time to find all network states  $\mathbf{S}$  for the given graph  $G$ .
- let  $g(n, \mathbf{R})$  the time to find all connected states  $S \subset \mathbf{S}$
- in-order to calculate  $Conn(G, \mathbf{R})$  the upper bound can be  $|R_1| \times |R_2| \times \dots \times |R_n|$ . Also let  $v$  is the node located maximum number of regions then we can calculate it  $|R_1| \times |R_2| \times \dots \times |R_n| = \mathcal{O}(|R_v|^n)$
- so the total running time will be  $\mathcal{O}(|R_v|^n)$  as  $f(n, \mathbf{R})$  and  $g(n, \mathbf{R})$  minor in comparison.

### 3 k-trees

In this section, we will define clique,  $k$ -tree and perfect elimination sequence which is used in the main algorithm.

A clique is a set of vertices that induce a complete subgraph of a graph  $G$ . A clique with  $k$  vertices is considered to be a  $k$ -tree.

**Definition.** The class of  $k$ -tree can be defined recursively as follows:

- The complete graph on  $k$  vertices is a  $k$ -tree.
- Lets  $G_n$  is a  $k$ -tree with  $n$  vertices where  $n \geq k$ . Then we can construct a  $k$ -tree  $G_{n+1}$  of  $n + 1$  vertices by adding a vertex adjacent to exactly  $k$  vertices, namely all vertices of a  $k$  clique of  $G_n$ . ■

A partial  $k$ -tree is any subgraph of a  $k$ -tree. partial  $k$ -trees are rich class of graph. Forest is an example of partial 1-tree. Series-parallel graphs and chordal graphs are subfamily of partial 2-trees. Also Halin graphs, Nested SAT and IO-graphs are subclasses of partial 3-trees.

**Example 3.1.** Fig. 1 depict a partial 2-tree. The complete graph of 2 vertices namely  $v_1$  and  $v_2$  is a 2-tree. Then we added vertex  $V_3$  which is adjacent to both  $v_1$  and  $v_2$  is a 2-tree of 3 vertices. Finally a vertices  $s$  is added to the clique  $v_1v_2$  to form the 2-tree with 4 vertices.■

#### 3.1 Perfect Elimination Sequence

A perfect elimination sequence (PES) in a graph is an ordering of the vertices of the graph such that, for each vertex  $v$ ,  $v$  and the neighbors of  $v$  that occur after  $v$  in the order form a clique. In-order to find PES we need simplicial vertex.

**Definition** (Simplicial Vertex). A simplicial vertex of a graph  $G$  is a vertex  $v$  such that the neighbours of  $v$  form a clique in  $G$ . Clearly, if  $G$  has a PES, then the last vertex in it is simplicial in  $G$ .■

**Example 3.2.** In fig. 1 we show that there are two simplicial vertices,  $v_1$  and  $s$ . But  $s$  is our sink node so we are not eliminating  $s$ . So After elimination of  $v_1$ , we will be able to eliminate either  $v_2$  or  $v_3$ . So the PES will be either  $v_1, v_2$  or  $v_1, v_3$ .■

### 4 Main Algorithm

In this section we present an algorithm to compute the exact solution for  $Conn(G, \mathbf{R})$  problem. More specially, the algorithm takes as input a UWSN network  $G$  where each node has a probabilistic

locality set, a PES and compute Prob, a solution to the input  $Conn(G, \mathbf{R})$  instance. The function uses a dynamic programming framework to solve  $Conn(G, \mathbf{R})$  problem.

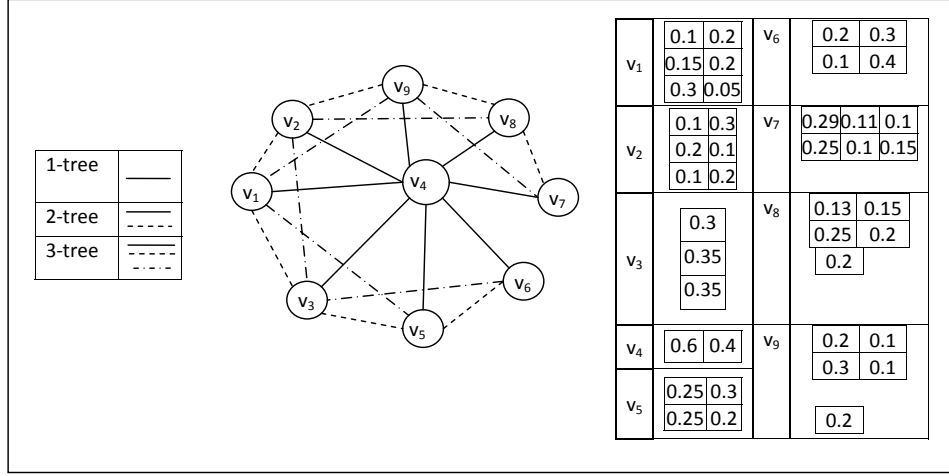


Figure 2: A UWSN modelled by a 3-tree.

#### 4.1 Key Data-structures

Each clique associated with a table. Each row in the table defines key-value mapping.

- A key is a set of sets namely partition(s) of nodes along with corresponding regions in that particular clique.
- A value is a probability which is the multiplication of regional probability of nodes for the clique.

**Example 4.1.** Fig. 2 illustrates a graph  $G$  which is a 3-tree with 9 nodes and their corresponding probabilistic locality set. There are 19 triangles so there are 19 cliques associated with this 3-tree. For each clique the algorithm maintain a table. For every table there are some rows as key-value mapping. For example the clique  $\langle v_1, v_2, v_3 \rangle$  can be partitioned 5 different ways including  $\{v_1, v_2, v_3\}$ ,  $\{v_1, v_2\}\{v_3\}$ ,  $\{v_1, v_3\}\{v_2\}$ ,  $\{v_1\}\{v_2, v_3\}$  and  $\{v_1\}\{v_2\}\{v_3\}$ . For each partition there are  $6 \times 6 \times 4 = 144$  rows as key-value mappings because the locality set of node  $v_1, v_2$  and  $v_3$  are 6, 6 and 4 respectively. One of the row is  $\{v_1, v_2, v_3\}^{(1,1,1)} 0.004$  where  $v_1, v_2$  and  $v_3$  are in same partition with region 1, 1 and 1 respectively and the probability 0.004 is multiplication of corresponding regional probabilities. ■

When the algorithm starts elimination node by the order of  $PES$  there will be table merging and details is presented in section 4.2.

#### 4.2 Algorithms Description

We now explain the main steps of function main, merge and merge partitions.

##### 4.2.1 Function Main

Main function eliminates every node according to the PES by merging all cliques associated with that node and updating the result to base clique. The last remaining clique associates with the sink

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**Algorithm 1:** Function Main( $G, \mathbf{R}, p(r_{(v,i)}), PES$ )

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**Input:** a UWSN  $G = (V, E)$  is a partial  $k$ -tree where each node,  $v \in V$  can be located into a set of regions  $R_v = \{r_{(v,1)}, r_{(v,2)} \dots\}$  with probability,  $\{p(r_{(v,1)}), p(r_{(v,2)}) \dots\}$  and  $(x, y) \in E$  if  $x \in V$  can be located one of it's locality set and  $y \in V$  can be located one of it's locality set, so that they reach each other.  $PES$  is a perfect elimination sequence  $(v_1, v_2, \dots, v_{n-k})$  of  $G$ .

**Output:** Prob, a solution to the input instance.

**Notation:**  $Temp$  is a map from keys to probabilities.

```
1 Initialize every clique by a table.
2 for  $i = 1, 2, \dots, n - k$  do
3   node  $v_i$  is associated with  $k$ -cliques,  $K_{(v_i,1)}, K_{(v_i,2)}, \dots, K_{(v_i,k)}$ 
   //  $T_{(v_i,1)}, T_{(v_i,2)}, \dots, T_{(v_i,k)}$  are the tables associated with cliques  $K_{(v_i,1)}, K_{(v_i,2)}, \dots, K_{(v_i,k)}$ 
   respectively
4    $Temp = T_{(v_i,1)}$ 
5   for  $j = 2, 3, \dots, k$  do
6      $Temp = merge(Temp, T_{(v_i,j)})$ 
   end
   // clique  $K_{(v_i,base)}$  is the base clique and table  $T_{(v_i,base)}$  is the base table of node  $v_i$ 
7    $Temp = merge(Temp, T_{(v_i,base)})$ 
8   remove node  $v_i$  from  $Temp$  and assign the result to  $T_{(v_i,base)}$ 
  end
9 return  $Prob = \sum (All\ probability\ for\ single\ partition\ in\ the\ remaining\ table)$ 
```

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node. Finally main function calculates the connectivity from the last clique by adding probabilities for those row which is associate with single partition.

In more details,

- Step 1 : initialize each clique by a table describe in subsection 4.1.
- Step 2 : processes each node  $v_i$  in PES. Every processing node,  $v_i$  is associated with  $k$  cliques.
- Step 3 : finds all those cliques,  $K_{(v_i,1)}, K_{(v_i,2)}, \dots, K_{(v_i,k)}$ .
- Step 4 : assign table  $Temp$  with table  $T_{(v_i,1)}$  associated with clique  $K_{(v_i,1)}$ .
- Steps 5-6 : iteratively merge all associated tables of node  $v_i$  and assign the result to table  $Temp$  by using *merge* function.
- Step 7 : merge  $Temp$  table with base table of node  $v_i$   $T_{(v_i,base)}$  by using *merge* function and assign the result to  $Temp$ .
- Step 8 : remove the vertex  $v_i$  and it's locality set from  $Temp$  and assign the result to  $T_{(v_i,base)}$ .
- Step 9 : calculate and return connectivity for the network.

**Example 4.2.** One of the PES of the 3-tree depict in fig. 2 is  $\langle v_1, v_6, v_5, v_7, v_8, v_9 \rangle$ . In-order to eliminate  $v_1$ , the algorithm merge three cliques  $\langle v_1, v_3, v_4 \rangle$ ,  $\langle v_1, v_2, v_3 \rangle$  and  $\langle v_1, v_2, v_4 \rangle$  to  $\langle v_1, v_2, v_3, v_4 \rangle$ . Next step the algorithm find the base clique  $\langle v_2, v_3, v_4 \rangle$  and merge with  $\langle v_1, v_2, v_3, v_4 \rangle$ . Finally it deletes  $v_1$  from merged clique and update the result to  $\langle v_2, v_3, v_4 \rangle$  ■

#### 4.2.2 Function merge

The primary task of merge function is to merge two table  $T_1, T_2$ , update keys and values of the newly created table  $T$  and return the table  $T$ .

$T_1(v_1, v_2, v_3)$			$T_2(v_1, v_3, v_4)$			$Temp(v_1, v_2, v_3, v_4)$	
.	.	$\times$	.	.	$\Rightarrow$	.	.
$\{v_1, v_2\}^{(1,1)}\{v_3\}^{(2)}$	0.002		$\{v_1\}^{(1)}\{v_3, v_4\}^{(2,1)}$	0.008		$\{v_1, v_2\}^{(1,1)}\{v_3, v_4\}^{(2,1)}$	0.0008
.	.		.	.		.	.
.	.		.	.		.	.
.	.		.	.		.	.

Figure 3 a). Merging two table into  $Temp$

$Temp(v_1, v_2, v_3, v_4)$			$Temp(v_2, v_3, v_4)$	
.	.		.	.
$\{v_1, v_2\}^{(1,1)}\{v_3, v_4\}^{(2,1)}$	0.0008	$\Rightarrow$	$\{v_2\}^{(1)}\{v_3, v_4\}^{(2,1)}$	0.0008
.	.		.	.
.	.		.	.
.	.		.	.

Figure 3 b). Deleting node  $v_1$  from  $Temp$

In more details,

- Step 1 : finds the common vertices between two tables and assign the common nodes probability 1.
- Step 2 : check if whether or not the common vertex set  $C$  is empty. If it is empty then the algorithm returns otherwise go to next step.
- Steps 3-5 : performs the row-wise merging by using function  $mpar$  and create another row.
- Steps 6-7 : updates the locality set for every vertex  $v$  of newly created row.
- Steps 8-9 : calculates the regional probability of common nodes.
- Step 10 : updates the newly created row probability by multiplying row-wise probability and dividing by the sum of common nodes regional probabilities.
- Step 11 : insert the row into table  $T$  and
- Step 12 : return table  $T$ .

**Example 4.3.** Fig 3 a). illustrates merging row  $\{v_1, v_2\}^{(1,1)}\{v_3\}^{(2)}$  0.002 of table  $T_1$  with row  $\{v_1\}^{(1)}\{v_3, v_4\}^{(2,1)}$  0.008 of table  $T_2$  which is fundamental operation for merging two tables. The function merge uses  $pMerge$  function which takes two partitions  $\{v_1, v_2\}\{v_3\}$  and  $\{v_1\}\{v_3, v_4\}$  and returned  $\{v_1, v_2\}\{v_3, v_4\}$  because  $\{v_1, v_2\} \cup \{v_1\} = \{v_1, v_2\}$  and  $\{v_3\} \cup \{v_3, v_4\} = \{v_3, v_4\}$ . The merge function updates the locality set of each node of the merged partition  $\{v_1, v_2\}^{(1,1)}\{v_3, v_4\}^{(2,1)}$  from the locality set of nodes in merging partitions. There are two common nodes  $v_1$  and  $v_3$  located in region 1 and 2 respectively with probability 0.1 and 0.2 respectively in the merging partitions. It update the probability of newly created partition 0.0008 by multiplying row-wise probabilities  $0.002 \times 0.008$  and dividing the probability by product of the common nodes corresponding regional probabilities  $0.1 \times 0.2$ .

Fig 3 b). shows the deletion of node  $v_1$  from  $Temp$ . The function main simply look for the node and remove it and it's locality set from key. ■



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**Algorithm 2:** Function merge( $T_1, T_2$ )

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**Input:** Two tables  $T_1$  and  $T_2$  that share at least one common vertex

**Output:** A table  $T$

**Notation**  $C$  is a set of vertices and  $Obj$  is a row of table  $T$  and  $Prob\_C$  is a double variable

```
1 set  $C$  = the set of common vertices between  $T_1$  and  $T_2$  , set  $Prob\_C = 1$ 
2 if  $C \neq \emptyset$  then
3   foreach row  $r$  in  $T_1$  do
4     foreach row  $s$  in  $T_2$  do
5        $Obj.par = pMerge(r.par, s.par)$ 
6       foreach vertex  $v_i$  in  $Obj.par$  where  $i = 1, 2, \dots, k + 1$  do
7          $Obj.loc[v_i] = r.loc[v_i] || s.loc[v_i]$ 
8       end
9       foreach vertex  $v \in C$  do
10         $Prob\_C = Prob\_C * s.loc[v]$ 
11      end
12       $Obj < Obj.par : Obj.loc > = \frac{Prob[r] \times Prob[s]}{Prob\_C}$ 
13      Insert  $Obj$  in  $T$  as a row.
14    end
15  end
16 end
17 return Table  $T$ 
```

---

#### 4.2.3 Function Merge Partitions

The merging partitions is mainly done using the union operation by mapr function. The mpar function takes as input two partitions  $P_1$ ,  $P_2$  and merge them into one partition  $P$ .

In more details,

- Steps 1-2 : adds all the sets of  $P_2$  to  $P_1$ .
- Steps 3-4 : selects two sets  $s^*$  and  $t^*$  from partition  $P_1$
- Step 5: checks whether or not they are disjoint . If they are not disjoint then it will go to step 6 otherwise it will return to step 4.
- Step 6 : delete  $s^*$  from  $P_1$ .
- Step 7 : computes the union of  $s^*$  and  $t^*$  and insert it at the beginning of partition  $P_1$ .
- Step 8 : delete  $t^*$  from  $P_1$ .
- Steps 9-10 : set the iterator to the beginning of  $P_1$  and return to step 5.
- Step 11 : assign  $P_1$  to  $P$  and finally the algorithm return  $P$ .

---

**Algorithm 3:** function pMerge( $P_1, P_2$ )

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**Input:** Two partitions  $P_1$  and  $P_2$

**Output:** A partition  $P$

**Notation:**  $s$  and  $t$  are two set iterators and their corresponding set are indicated by  $s^*$  and  $t^*$ .

```
1 foreach set  $s^*$  in  $P_2$  do
2    $P_1.push\_back(s^*)$ 
3 end
4 for ( $s = P_1.begin(); s \neq P_1.end(); ++s$ ) do
5   for ( $t = s.next(); t \neq P_1.end(); ++t$ ) do
6     if  $s^* \cap t^* \neq \emptyset$  then
7        $P_1.delete(s^*)$ 
8        $P_1.push\_front(s^* \cup t^*)$ 
9        $P_1.delete(t^*)$ 
10       $s = P_1.begin()$ 
11       $break$ 
12    end
13  end
14 end
15 set  $P = P_1$ 
16 return  $P$ 
```

---

### 4.3 Verification Cases

In this subsection we are going to present some cases by using these one can verify the correctness of our algorithm.

- For a partial  $k$ -tree, there should be more than one PES. In our algorithm, we tried all elimination sequence and we got same result.

### 4.4 Correctness

In this section we prove the correctness of our algorithm. It is an exhaustive algorithm which takes care all possible configuration. Our algorithm consists of two fundamental operation, partition merge and table merge. First we prove that partition merge and table merge are correct then we prove that our algorithm is correct.

The partition merge perform core operations of the algorithm. The partition merge is correct because it is merging two partitions into one given that they share at least one node. The table merge is correct because of the following reason. The table merge is merging each row of one table with each row of another table by holding the condition that they share atleast one node with same regional position. It is using partition merge in-order to merge two rows which is correct. The main algorithm is elimination a node by merging all clique associated with that node and updating the merged clique with base clique of that node. Each clique is associated with table so cliques are merged by using table merge operation which is correct. That concludes that the algorithm is correct.

## 4.5 Running time

## 5 Simulation Results

In this section we present simulation results that aim to investigate the following aspect

- (a) complexity of our algorithm increases with increasing the value of  $k$  for partial  $k$ -tree.
- (b) influence of  $k$  on accuracy where  $k = 1, 2, 3, \dots$
- (c) effect of choosing different  $k$ -trees.

Table 2: Running time (RT) with respect to  $k$

k	Network I	Network II	Network III
1	30	30	20
2	290	380	380
3	85000	875000	940000

Table 3: Accuracy with respect to  $k$

k	Network I	Network II	Network III
1	62	32.96	81.8
2	71.2	36.15	98.95
3	75	37.31	100

We used three networks in-order represent running time and accuracy. Network I consists of 7 nodes with locality set of each node range from 4 to 8. Network II consists of 9 nodes where each node can be located from 3 to 6 locations. Network III consists of 12 nodes with locality set of each node range from 2 to 8.

- a)  **$k$  versus running time** table 1 shows three scenarios running time with respect to  $k$  of. In every scenario the running time of the algorithm increases with  $k$ . We also observe that difference between the running time 1-tree and 2-tree are 10 to 15 times where the running time between 2-tree and 3-tree are more than 1000 times. This is because of complexity of 3-tree is much more than 2-tree.
- b)  **$k$  versus accuracy** table 2 illustrates accuracy increases with  $k$ . The increase of accuracy is large for 1-tree to 2-tree than for 2-tree to 3-tree.
- c1) **Effect of random edge selection.** Fig 6 illustrates connectivity versus transmission range when the algorithm selects randomly. We have the following observation from fig 6.
  - with increase of transmission range the connectivity is also increasing. This is due to, increase of transmission range the probability of an edge between two nodes also increasing.
  - connectivity for 2-tree is always greater than or equal to the connectivity of 1-tree for same transmission range. This is because in 2-tree there are more edges in comparison with 1-tree.
  - The above is also true for 2-tree to 3-tree.
- c2) **Effect of greedy edge selection.** Fig 7 illustrates connectivity versus transmission range when we are selecting edges by greedy techniques. In greedy techniques those edges are selected which has a high probable values. It is observable from figure that the network is fully connected with a smaller transmission range for all tree in comparison with random edge selection strategy.

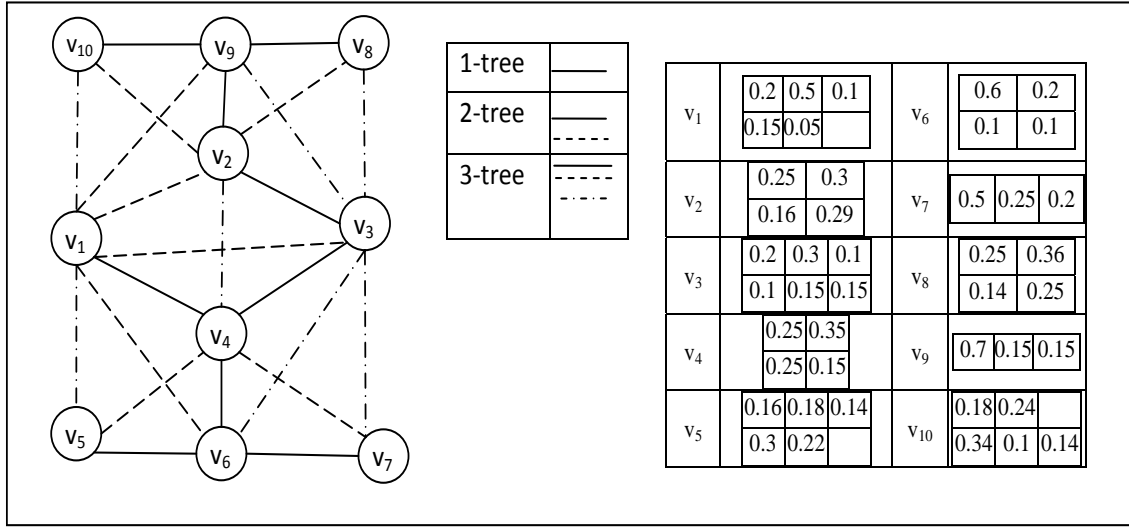


Figure 3: NetworkI

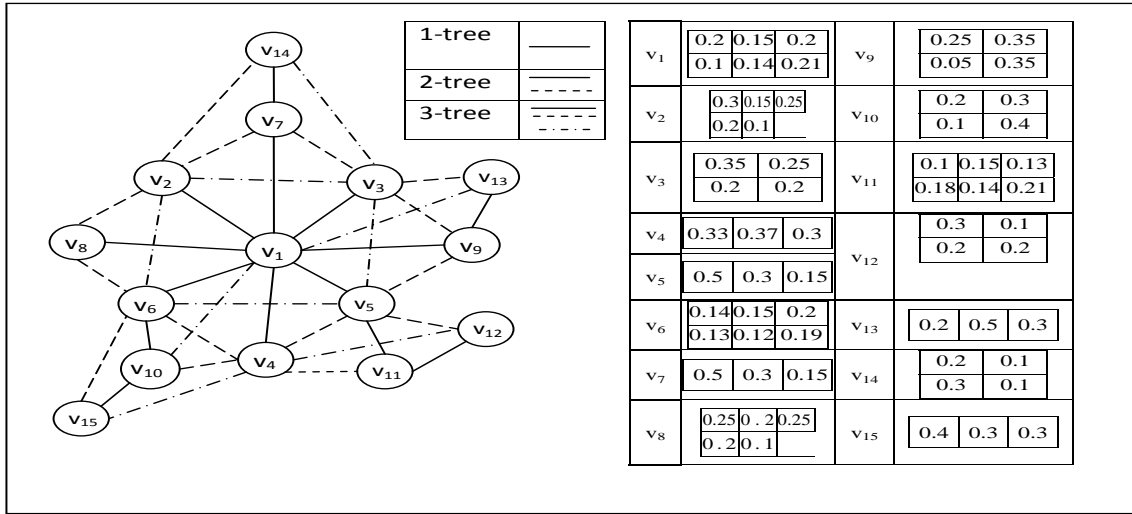


Figure 4: NetworkII

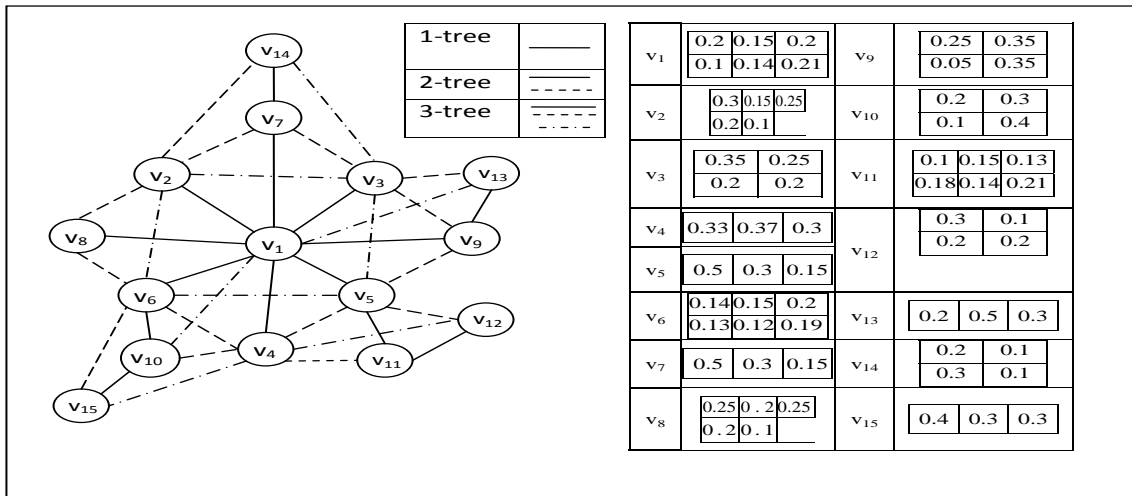


Figure 5: NetworkIII

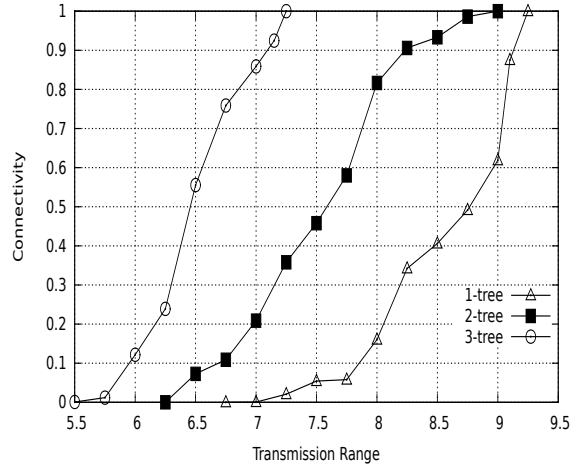


Figure 6: Connectivity versus transmission range with random edge selection.

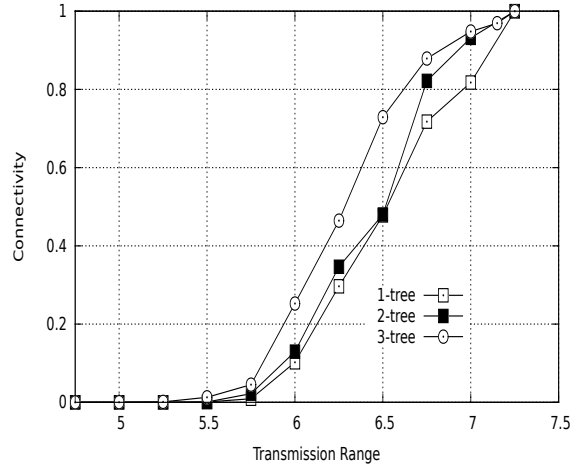


Figure 7: Connectivity versus transmission range with greedy edge selection.

## 6 Network with Relays

In this section we add relay nodes to our network. So our data structures changes as a result main algorithm and merge function also changes in some aspect. We describe the updates and show the results after adding relays.

We added relays,  $Rel \subset V$  to the network in addition to sensor nodes which is referring target  $Tar \subset V$  node in this section.

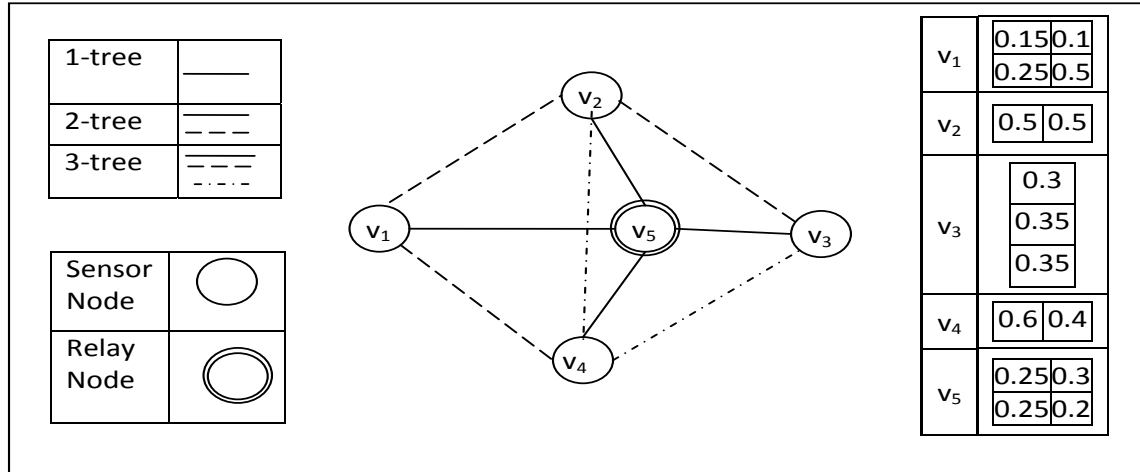


Figure 8: A partial 2-tree with 6 nodes.

**Example 6.1.** Fig ?? illustrates a network of 6 nodes where  $v_1, v_2, v_3$  and  $v_4$  are target nodes and  $v_5, v_6$  are relays.  $v_5$  is enhancing the connectivity of the network but  $v_6$  is not increasing the connectivity. So we can simply ignore  $v_6$  and measure the connectivity from  $v_1$  to  $v_5$

### 6.1 Problem Statement

In this section we define the problem.

**Definition** (The  $Conn(G, Tar)$  Problem). We are given,  $V$  the set of all nodes where each node  $v \in V$  is located in a set of regions,  $R_v = \{r_{(v,1)}, r_{(v,2)}, \dots\}$  with probability  $p(r_{(v,i)})$  where  $i = 1, 2, \dots$ .  $Rel \in V$  is the set of relay nodes and  $Tar \in V$  is the set of target nodes where  $V = Rel \cup Tar$ . Also, if  $x \in V$  and  $y \in V$  are two nodes where  $x$  can be located into one of it's locality set and  $y$  can be located into one of it's locality set that they can communicate with each other then there is a link between  $x$  and  $y$ , denoted by  $(x, y) \in E$ . Here we use relay nodes to enhance the performance of the network but we are interested to find out the probability  $Conn(G, Tar)$  that all target nodes  $Tar$  are connected.

## 6.2 Key Data Structures

We already know from section 4.1 that a typical row in a table is key-value mapping. A key consists of i). partitions ii). regional set iii). target node attach. We explain them this section.

- i) partitions: There can be more than one partition associated with each row. Each partition consists with one or more node. We use braces to distinguish each partition. Two or more nodes are in the same partition means they can communicate each other when they are in regions indicated by regional set. For example in the key  $\{v_1, v_2\}_{(1)}^{(1,1)} v_3_{(0)}^{(2)}$ , there are two partitions including  $\{v_1, v_2\}$  and  $\{v_3\}$ . Also node  $v_1$  can reach node 2 when they are both in region 1 of their corresponding locality set but neither node  $v_1$  nor node  $v_2$  from region 1 can reach node 3 when node  $v_3$  is in region 2 of it's locality set.
- ii). regional set: The regional set of a partition consists of the position of corresponding node in the locality set. The regional set is indicated as a superscript of partition and is surrounded by parentheses. For example in the key  $\{v_1, v_2\}_{(1)}^{(1,1)} v_3_{(0)}^{(2)}$ , there are two regional sets (1, 1) and (2) associated with two partitions  $\{v_1, v_2\}$  and  $\{v_3\}$  respectively. More specifically the partition-regional set pair  $\{v_1, v_2\}_{(1)}^{(1,1)}$  indicates that node  $v_1$  and  $v_2$  are both located in region 1 of their corresponding locality set.
- iii). Target node attach: The target node attach associates with every partition. If there is one or more target node in a partition then the value of target node attach is 1 otherwise it is 0. In the above example we indicate the target node attach as a subscript of the partition. First partition in the above key  $v_1$  and  $v_2$  are target nodes so as a subscript we put 1 and for second partition we simply put 0 to indicate that  $v_3$  is a relay node.

## 6.3 Main function

- Step 8: check whether or not the node  $v_i$ , we are going to eliminate is a relay node. If  $v_i$  is relay node, It goes to next step otherwise it goes to step 10.
- Step 9: search every partition in every row of the table  $Temp$  for node  $v_i$ . It deletes node  $v_i$  from every partition that contains node  $v_i$ . It also updates the regional sets of corresponding partitions from which  $v_i$  was deleted by removing the corresponding regions of  $v_i$ .
- Step 10: If  $v_i$  is sensor node then it goes to next step.
- Step 11: search every partition in every row of table  $Temp$  for  $v_i$ . If the partitions that contains  $v_i$  consists of more than one node then remove  $v_i$  from that partitions. It also removes the location information of  $v_i$  from the regional sets corresponding to those partitions. If a

$T_1(v_1, v_2, v_3)$			$T_2(v_1, v_3, v_4)$			$Temp(v_1, v_2, v_3, v_4)$
.	.		.	.		.
$\{v_1, v_2\}_{(1)}^{(1,1)} \{v_3\}_{(0)}^{(2)}$	0.002	$\times$	$\{v_1\}_{(1)}^{(1)} \{v_3, v_4\}_{(1)}^{(2,1)}$	0.008	$\Rightarrow$	$\{v_1, v_2\}_{(1)}^{(1,1)} \{v_3, v_4\}_{(1)}^{\{2,1\}}$ 0.0008
.	.		.	.		.
.	.		.	.		.
.	.		.	.		.

Figure 3 a). Merging two table into  $Temp$

$Temp(v_1, v_2, v_3, v_4)$			$Temp(v_2, v_3, v_4)$
.	.		.
$\{v_1, v_2\}_{(1)}^{(1,1)} \{v_3, v_4\}_{(1)}^{(2,1)}$	0.0008	$\Rightarrow$	$\{v_2\}_{(1)}^{(1)} \{v_3, v_4\}_{(1)}^{(2,1)}$ 0.0008
.	.		.
.	.		.
.	.		.

Figure 3 b). Deleting node  $v_1$  from  $Temp$

partition contains  $v_i$  itself this kind of partition is called bad partition. The algorithm simply ignore bad partition.

---

**Algorithm 4:** Function Main( $G, \mathbf{R}, Tar, Rel, p(r_{(v,i)}), PES$ )

---

**Input:** a UWSN  $G = (V, E)$  is a partial  $k$ -tree where each node,  $v \in V$  can be located into a set of regions  $R_v = \{r_{(v,1)}, r_{(v,2)} \dots\}$  with probability,  $\{p(r_{(v,1)}), p(r_{(v,2)}) \dots\}$  and  $(x, y) \in E$  if  $x \in V$  can be located one of it's locality set and  $y \in V$  can be located one of it's locality set, so that they reach each other. A set of target nodes  $Tar = \{v_1, v_2, \dots\}$  where  $Tar \subset V$ . A set of relay nodes  $Rel = \{v_1, v_2, \dots\}$  where  $Rel \subset V$ .  $PES$  is a perfect elimination sequence  $(v_1, v_2, \dots, v_{n-k})$  of  $G$ .

**Output:** Prob, a solution to the input instance.

**Notation:**  $Temp$  is a map from keys to probabilities.

```
1 Initialize every clique by a table.
2 for  $i = 1, 2, \dots, n - k$  do
3   node  $v_i$  is associated with  $k$ -cliques,  $K_{(v_i,1)}, K_{(v_i,2)}, \dots, K_{(v_i,k)}$ 
   //  $T_{(v_i,1)}, T_{(v_i,2)}, \dots, T_{(v_i,k)}$  are the tables associated with cliques  $K_{(v_i,1)}, K_{(v_i,2)}, \dots, K_{(v_i,k)}$ 
   respectively
4    $Temp = T_{(v_i,1)}$ 
5   for  $j = 2, 3, \dots, k$  do
6      $Temp = merge(Temp, T_{(v_i,j)})$ 
   end
   // clique  $K_{(v_i,base)}$  is the base clique and table  $T_{(v_i,base)}$  is the base table of node  $v_i$ 
7    $Temp = merge(Temp, T_{(v_i,base)})$ 
8   if  $v_i$  is a relay node then
9     remove node  $v_i$  from  $Temp$  and assign the result to  $T_{(v_i,base)}$  after updating the
     regional set.
   end
10  else
11    Remove  $v_i$  from all the partitions of  $Temp$  except the partition with  $v_i$  itself and
    assign the result to  $T_{(v_i,base)}$  after updating the regional set.
    // We simply ignore the row where there is a partition of  $v_i$  itself.
  end
end
12 return  $Prob = \sum (All\ probability\ for\ single\ partition\ in\ the\ remaining\ table)$ 
```

---



## 6.4 Merge Function

---

**Algorithm 5:** Function  $\text{merge}(T_1, T_2)$ 


---

**Input:** Two tables  $T_1$  and  $T_2$  that share at least one common vertex

**Output:** A table  $T$

**Notation**  $C$  is a set of vertices and  $Obj$  is a row of table  $T$  and  $Prob\_C$  is a double variable

```

1  set  $C$  = the set of common vertices between  $T_1$  and  $T_2$  , set  $Prob\_C = 1$ 
2  if  $C \neq \emptyset$  then
3      foreach row  $r$  in  $T_1$  do
4          foreach row  $s$  in  $T_2$  do
5               $Obj.par = \text{pMerge}(r.par, s.par)$ 
6              foreach vertex  $v_i$  in  $Obj.par$  where  $i = 1, 2, \dots, k+1$  do
7                   $Obj.locmap[v_i] = r.locmap[v_i] || s.locmap[v_i]$ 
8              end
9              foreach node  $v$  in every partition  $P$  in  $Obj.par$  do
10                 foreach node  $u$  in every partition  $Q$  in  $s.par$  do
11                     if  $v == u$  then
12                          $Obj.tAttach[P] = \max(Obj.tAttach[P], s.tAttach[Q])$ 
13                     end
14                 end
15             end
16             foreach node  $w$  in every partition  $T$  in  $r.par$  do
17                 if  $v == w$  then
18                      $Obj.tAttach[P] = \max(Obj.tAttach[P], r.tAttach[T])$ 
19                 end
20             end
21             end
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- Step 8: selects every node  $v$  of every partition  $P$  in the newly created row  $Obj$ .
- Step 9: iterates every node  $u$  of every partition  $Q$  in the row  $s$ .
- Step 10: check whether  $u$  and  $v$  are the same same node or not. If they are same then it goes to next step otherwise it goes to step 9.
- Step 11: updates the target attach entry of partition  $P$  by taking the max between the target attach entry of partition  $P$  and target attach entry of partition  $Q$ .
- Step 12: iterates every node  $w$  of every partition  $T$  in the row  $r$ .

Table 4: Accuracy with respect to  $k$

k	Network I	Network IR	Network II	NeworkIIR	NetworkIII	Network IIIR
1	30.23	86.96	5.23	22.27	15.99	28.44
2	53.72	96.72	17.55	59.43	80.23	89.3
3	60.20	97.60	20.96	60.5	83.3	92.3

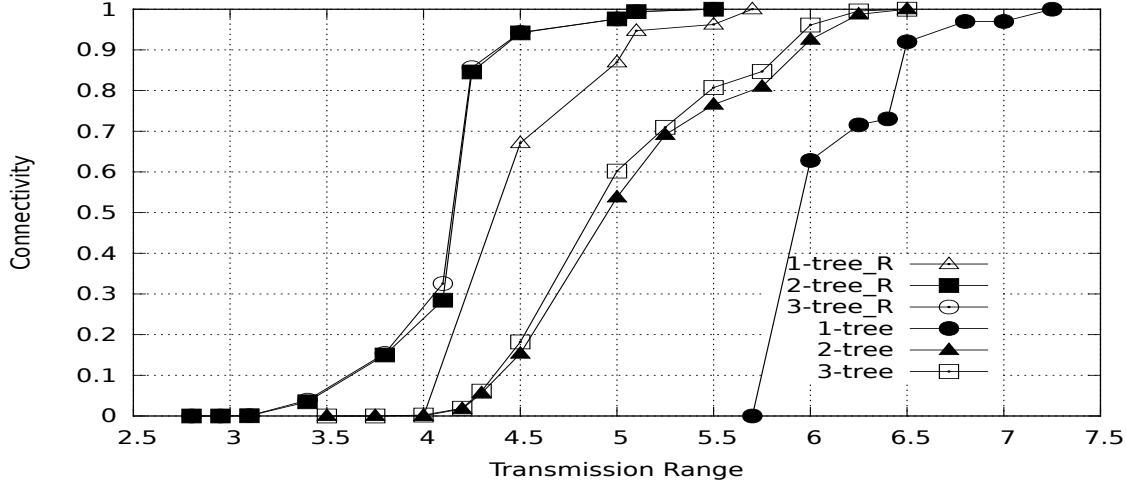


Figure 9: NetworkIII

- Step 13: check whether  $w$  and  $v$  are the same same node or not. If they are same then it goes to next step otherwise it goes to step 9.
- Step 11: updates the target attach entry of partition  $P$  by taking the max between the target attach entry of partition  $P$  and target attach entry of partition  $T$ .

## 6.5 Simulation Results

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