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Anomaly Detection in Dynamic Networks: A Survey

Stephen Ranshous\(^1,2\), Shitian Shen\(^1,2\), Danai Koutra\(^3\), Christos Faloutsos\(^3\), and Nagiza F. Samatova\(^1,2,\+)

\(^1\)Department of Computer Science, North Carolina State University
\(^2\)Computer Science and Mathematics Division, Oak Ridge National Laboratory
\(^3\)Computer Science Department, Carnegie Mellon University

\(+\)Corresponding author: samatova@csc.ncsu.edu

Abstract

Anomaly detection is an important problem with multiple applications, and thus has been studied for decades in various research domains. However, for dynamic networks, no comprehensive survey exists covering the richness of methods proposed and their differences, as well as the variety of problems they solve. In light of this, the current survey focuses on anomaly detection in time-evolving networks. The methods covered are classified into five categories based on their technical approaches, and subsequently broken down by their output, i.e., the type of anomalies they detect—nodes, edges, subgraphs, and events. Some of the methods discussed cannot easily be placed into a single category, leveraging multiple approaches. For such cases, the method has been classified based on the predominant approach utilized. We cover a total of 46 methods: 9 based on communities, 2 on compression, 15 on decomposition, 11 on distance metrics, and 9 probabilistic based. Within these categories we highlight the similarities and differences showing the wealth of different techniques derived from similar conceptual approaches.

Introduction

Anomaly detection, or outlier detection, is the well-known problem\(^9,37\) of “finding patterns in data that do not conform to expected behavior”\(^14\). This area has been well-covered for a variety of domains including sequence data\(^16\), discrete sequence data\(^15\), and temporal data\(^32\); see also a general comprehensive overview\(^14\). However, anomaly detection in dynamic networks\(^1\) has been barely touched in existing works\(^11,32\). No extensive survey exists, despite the popularity and the growing importance of the topic.

A surge for researching novel methods for real world networks that are constantly evolving has been motivated by a number of applications. Dozens of methods specifically designed for dynamic environments have emerged over the last five years (see

\(^1\)Throughout this paper we will use the terms networks and graphs interchangeably.
summary in Table 1 and Table 2. These techniques complement methods for static graphs [59, 27, 26, 6], as the latter cannot be easily adopted for dynamic networks. They find real world applications in numerous domains, including but not limited to:

- Detection of ecological disturbances such as wildfires [21, 20] and cyclones [18] by analyzing the ecological evolution.
- Intrusion detection for individual systems [87] and network systems [85, 29, 23] by observing past normal behaviors and comparing new actions to the expected model.
- Identifying abnormal users and events in communication networks [64, 84], such as days with an abnormally high number of messages.

We present in this survey techniques for performing anomaly detection in dynamic networks in hopes to bridge the gap between the increasing number of of emerging algorithms for this important graph mining task and the lack of their comprehensive analysis. We start by introducing four different types of anomalies that these algorithms detect, namely, anomalous nodes, edges, subgraphs, and events. We then continue with an extensive overview of the existing methods based on the proposed taxonomy that takes into account their underlying design principles, such as those based on graph communities, compression, decomposition, distance metrics, and probabilistic modeling of graph features. Finally, we subcategorize each taxonomic group further based on the types of anomalies detected.

Types of anomalies

In this section we identify and formalize four types of anomalies that arise in dynamic networks. These categories represent the output of the methods, not the implementation details of how they detect the anomalies, e.g., comparing consecutive time points, using a sliding window technique, etc.

Because graphs are assumed to be dynamic, their vertices and edges can be inserted or removed at every time step; however, for the sake of simplicity, we assume that the node correspondence and the edge correspondence across different time steps is resolved due to unique labeling of nodes and edges, respectively. We define a graph series as an ordered set of graphs with a fixed number of time steps, and a graph stream as an ordered set of graphs where new time steps are continuously added. The majority of techniques we discuss assume that the input graphs are simple, undirected, and unweighted. Hence, we assume that all graphs are of this type unless otherwise noted.

Type 1: Anomalous nodes

Problem 1. Given a fixed graph series $G$ or graph stream $\mathcal{G}$, find a subset of the vertices such that every vertex in the subset has an “irregular” evolution compared to
the other nodes in the graph. Optionally, identify the time point(s) where the nodes are determined to be anomalous.

Applications. Consider a graph series \( G \) where each graph \( G_t \) is a single day of network IP traffic. Assume that it is known that at time \( t \) a virus spreads throughout the system causing network congestion. By comparing the activity of each node on the days surrounding the attack to their activity on the day of the attack, one can detect the nodes from which the virus originated, as, at time \( t \), those nodes should have been more active than usual. Typical applications of this type of anomaly detection are identifying nodes that contribute the most to a discovered anomaly, such as in communication networks \[5\] (also known as attribution), and observing the shifts in community involvement \[28, 44\].

Example. An example of nodes that do not share the same evolution pattern as the community they initially belong to is shown in Figure 1. In the first time step, all of the nodes are found to be in the same community according to some extracted feature values, which is indicated by the blue circle. In time steps two and three, there is a shift in the values of the community as a whole, altering the distribution of the feature values. Some nodes, however, do not follow this trend and end up as outliers (anomalous nodes), indicated by the red crosses.

Type 2: Anomalous edges

Problem 2. Given a fixed graph series \( G \) or graph stream \( G \), find a subset of the edges such that every edge in the subset has an “irregular” evolution. Optionally, identify the time point(s) where they are determined to be anomalous.

There are two main types of irregular evolution for edges: (i) abnormal edge weight evolution, where the weight of a single edge fluctuates over time and has inconsistent spikes in value, and (ii) appearance of unlikely links in a graph \[66, 1\], where two nodes that are not typically connected or part of the same community have an edge added between them.

Applications. Consider a dynamic communication network where the nodes are people, the edges represent communication between two people, and the edge weight corresponds to the number of messages they exchange. Imagine persons \( a \) and \( b \) typically message each other 10 times each day, represented by the weighted edge \((a, b, 10)\). Then, suddenly, they have a day where they message each other 25 times. It is natural to assume that something happened between them because of the sharp increase in communication. This is the intuition behind edge detection. Anomaly detection of edges appears in many areas, such as vehicle traffic patterns \[52\] and finding unlikely social interactions \[39\].

Example. An example of edges that have abnormal weight evolution is shown in Figure 2, where the anomalous edge weights are highlighted. By examining the edge weights, one can observe that they tend to hover around a certain value, changing by \( \pm 0.05 \) at
most each time step. However, at time $t = 2$, edge $(3, 4)$ suddenly has a weight of 0.77, radically different from its average weight, 0.21, in the rest of the time points. Similarly, edge $(1, 4)$ at $t = 3$ has an edge weight of 0.93, which is roughly double its typical value (0.48).

**Type 3: Anomalous subgraphs**

**Problem 3.** Given a fixed graph series $G$ or graph stream $G$, find all subgraphs that have “unusual” behavior. Optionally, identify the time point(s) where they are determined to be anomalous.

**Applications.** Given a dynamic network representing vehicle traffic, where nodes are street intersections, edges are the streets, and the edge weights are the travel times for that particular road, how can traffic accidents be identified? Accidents typically block traffic on the roads surrounding the intersection where they occur, increasing the travel time by a substantial amount for those roads. If we were able to identify regions of the graph that exhibited this type of behavior, we would be able to identify such accidents automatically. This is just one possible application area [57], others include detecting the change in communities in the food web [19], and detecting changes and threats in social networks [59, 56].

**Example.** Figure 3 shows an example of two types of community based anomalous subgraphs. The shrunken community shown in Figure 3a is when a single community loses a considerable number of its members. Closely related is the idea of a split community shown in Figure 3b when a single community divides into several distinct communities.

**Type 4: Event and change detection**

**Problem 4 (Event detection).** Given a fixed graph series $G$ or graph stream $G$, find a time point at which the graph exhibits behavior sufficiently different from the others.

**Applications.** In the example scenario presented in the node anomaly section, we assumed that the time point that marked an event was known. However, in real world applications the event points are seldom known, requiring methods to discover them from the data. Event detection has been used to detect when the collective motions in molecular dynamics simulations suddenly change [65], to find frames in a video that are unlike the others [82, 71], and to detect disturbances in the ecosystem (e.g., wildfires) [20].

**Example.** A simple example is given in Figure 4a. At time $t = 3$, there is clearly a large structural change in the graph, increasing the connectivity to almost a clique. However, at the next time step, the graph is back to normal, having only sparse connections between the nodes.
Problem 5 (Change detection). Given a fixed graph series $G$ or graph stream $G$, find a time point at which a substantial change in the graph persists until the next point of change.

It is important to note the distinction between event and change detection. While events represent isolated incidents, change points mark a point in time where the entire behavior of the graph changes and the difference is maintained until the next change point.

Applications. Consider the DBLP network where authors are linked to the venues they publish per year. By profiling the authors, one can spot that most of them publish in specific conferences consistently, while few authors act as “bridges” and switch areas of interest and venues over the course of their career [48]. Change detection has been also applied to mobile communication graphs [5] and traffic networks [43].

Example. Figure 4b illustrates the concept of change detection. Time steps $t = 1, 2$ have approximately the same structure, with sparse connectivity between the nodes. However, at time $t = 3$ there is a large structural change, similar to that in Figure 4a, except the new structure is maintained.

Methods

Community based methods

Community based methods track the evolution of communities and their associated nodes in the graphs over time [77, 78, 79]. Various community-based approaches differ in two main points: (i) in the aspects of the community structure they analyze, e.g., the connectivity within each community vs. how the individual nodes are assigned to the communities at each time step; and (ii) in the definitions of communities they use, e.g., soft communities where each node has a probability of belonging to each community vs. disjoint communities where each node is placed into at most one community in the graph. Moreover, based on how the community evolution is viewed, it can be applied for the detection of different anomaly types. For example, a rapid expansion or contraction of a community could indicate that the specific subgraph for that community is undergoing drastic changes, whereas a drop in the number of communities by two corresponds to an abnormal event.

Node detection

Main Idea. A group of nodes that belong to the same community are expected to exhibit similar behavior. Intuitively this means that if at consecutive time steps one node in the community has a significant number of new edges added, the other nodes in the community would also have a significant number of new edges. If the rest of the nodes in the community did not have new edges added, the node that did is anomalous.
Based on this logic, using soft community matching and looking at each community individually, the average change in belongingness (the probability the node is part of the community) for each node can be found for consecutive time steps. Nodes whose change in belongingness is different from the average change of the nodes in the community are said to be evolutionary community outliers [35].

The changes in a node’s community belongingness may form a pattern over time, called a soft temporal pattern. In [70], a Non-Negative Matrix Factorization approach in combination with the Minimum Description Length principle are used to automatically detect node roles and build transition models. The community memberships are found slightly differently in [34], where Xmeans is used. However, in both cases, the patterns common across all of the nodes in the graph are extracted, then each node’s patterns are compared to the extracted ones. If a node’s patterns are not similar to any of the extracted common patterns, then the node is anomalous. Later extended to networks derived from heterogeneous data sources [33], the two step approach in [34] is modified to an alternating iterative approach. Instead of extracting patterns first and then identifying outliers, the patterns and outliers are found in an alternating fashion (pattern extraction → outlier detection → pattern extraction → …) until the outliers discovered do not change on consecutive iterations. By alternating back and forth between pattern extraction and outlier detection, the algorithm accounts for the affect the outliers have on the communities discovered [35].

If the edge weights are considered as the connection strength between two nodes, then the removal of a node with a high edge weight impacts the node it is connected to more than the removal of a node with a low edge weight [44]. The “closeness” of nodes is defined as a function of the weighted path between them instead of strictly the length of the path (e.g., one-hop paths). A node’s “corenet” is built based on the closeness values of nodes. Neighboring nodes within two hops and with a weighted path higher than a threshold value are part of the corenet. At each time step every node is given an outlier score based on the change in its corenet, and the top outlier scores are declared anomalous.

Subgraph detection

**Main Idea.** Instead of looking at individual nodes and their community belongingness, entire subgraphs that behave abnormally can be found by observing the behavior of communities themselves over time.

Six different types of community-based anomalies have been proposed [19]: shrink, grow, merge, split, born, and vanish. To detect such anomalies, time steps are compared using graph and community representatives. Conversely, instead of finding changes, communities that are conserved, or stable, can be identified. Constructing multiple networks at each time step based on different information sources, communities can be conserved across time and networks. Networks which behave similarly can be grouped using clustering or prior knowledge. If a community is conserved across time and the networks within its group, but has no corresponding community in any other group of
networks, then the community is anomalous [18]; two communities are considered corres-
dponding communities if they have a certain percentage of their vertices in common.
Unlike [18, 19] which consider only the structure of the network, in social networks there is often more information available. For example, in the Twitter user network, clusters can be found based on the content of tweets (edges), as well as the users (vertices) involved. If the fraction of the tweets (edges) added during the recent time window for a cluster is much larger than the fraction of tweets (edges) added anytime before the window, then this influx is declared as an evolution event for that cluster [3]. A cluster that experiences an evolution event is marked as an anomalous subgraph at the time when the evolution event occurs.

Change detection

Main Idea. Changes are detected by partitioning the streaming graphs into coherent segments. The beginning of each segment represents a detected change.

The segments are found online by comparing the node partitioning of the newest graph to the partitioning found for the graphs in the current segment. Node partitioning can be achieved with many methods, but in [25] it is done using the relevance matrix computed by random walks with restarts and modularity maximization. When the partitioning of the new graph is much different than the current segment’s, a new segment begins and the time point for the new graph is output as a detected change. The similarity of two partitions is computed as their Jaccard coefficient.

Compression based methods

The methods discussed in this section are all based on the minimum description length (MDL) principle [68]. The minimum description length principle, and compression techniques based on this principle, exploit patterns and regularity in the data to achieve a compact graph representation [31]. Applying this principle to graphs is done by viewing the adjacency matrix of a graph as a single binary string, flattened in row or column major order. If the rows and columns of the matrix can be rearranged such that the entropy of the binary string representation of the matrix is minimized, then the compression cost (also known as encoding cost [69]) is minimized. Anomalies can then be defined as graphs or edges that inhibit compressibility.

Edge detection

Main Idea. An edge is considered anomalous if the compression of a subgraph has higher encoding cost when the edge is included than when it is omitted.

Node partitioning can be done by rearranging the rows and columns in the adjacency matrix. In [13], a two step alternating iterative method is used for automatic partitioning of the nodes. In the first step, for a fixed number of partitions, the nodes are
assigned to them so that the encoding cost is minimized. The second step iteratively splits the partitions with high entropy into two. Any node whose removal from the original partition would result in a decrease in entropy is removed from that partition and placed into the new one. Once the method has converged, meaning steps 1 and 2 are unable to find an improvement, the edges can be given outlierness scores. The score for each edge is computed by comparing the encoding cost of the matrix including the edge to the encoding cost if the edge is removed.

Change detection

Main Idea. The main idea is that consecutive time steps that are very similar can be grouped together leading to low compression cost. Increase in the compression cost means that the new time step differs significantly from the previous ones, and thus signifies a change.

To detect changes in a graph stream, similar consecutive time steps can be grouped into segments. When considering the next graph in the stream, it can either be grouped with the graphs in the current segment, or it can be the beginning of a new one. The decision to start a new segment is made by comparing the encoding cost of the current segment without the next graph to the encoding cost of the segment if the next graph were included. If the node partitioning for the new graph is very similar to the node partitioning of the graphs in the segment then the encoding cost will not change much. However, if the partitions are very different, the encoding cost would increase because of the increase in entropy. Changes in the graph stream are the time points when a new segment begins [73]. This method is also parameter-free.

Decomposition based methods

These techniques represent the set of graphs as a tensor, most easily thought of as a multidimensional array, and perform its factorization or dimensionality reduction. The most straightforward method of modeling a dynamic graph as a tensor is to create a dimension for each graph aspect of interest, e.g., a dimension for time, source nodes, destination nodes, etc. For example, modeling the Enron email dataset can be done using a 4-mode tensor, with dimensions for sender, recipient, keyword, and date. The element \((i, j, k, l)\) is 1 if there exists an email that is sent from sender \(i\) to recipient \(j\) with keyword \(k\) on day \(l\), otherwise it is 0.

Similar to compression techniques, decomposition techniques search for patterns or regularity in the data to exploit. One of the most popular methods for matrices (2-mode tensors) is singular value decomposition (SVD) [30], and for higher order tensors \((\geq 3\) modes\) is PARAFAC [36], a generalization of SVD. The main differences between the decomposition based methods are whether they use a matrix or a higher order tensor, how the tensor is constructed, and the method of decomposition.
**Node detection**

**Main Idea.** Matrix decomposition is used to obtain activity vectors per node. A node is characterized as anomalous if its activity changes significantly between consecutive time steps.

Due to the computational complexity of performing principal component analysis on the entire graph, it is advantageous to apply it locally. One approach is to have each node maintain an edge correlation matrix $M$, which has one row and column for every neighbor of the node. The value of an entry in the matrix for node $i$, $M(j, k)$, is the correlation between the weighted frequencies of the edges $(i, j)$ and $(i, k)$. The weighted frequencies are found using a decay function, where edges that occurred more recently have a higher weight. The largest eigenvalue and its corresponding vector obtained by performing PCA on $M$ are summaries of the activity of the node and the correlation of its edges, respectively. The time series formed by finding the changes in these values are used to compute a score for each node at each time step. Nodes that have a score above a threshold value are output as anomalies at that time [86].

**Event detection**

**Main Idea.** There are two main approaches: (a) Tensor decomposition approximates the original data in a reduced dimensionality, and the reconstruction error is an indicator of how well the original data is approximated. Sub-tensors, slices, or individual fibers in the tensor that have high reconstruction error do not exhibit behavior typical of the surrounding data, and reveal anomalous nodes, subgraphs, or events. (b) Singular values and vectors, as well as eigenvalues and eigenvectors are tracked over time in order to detect significant changes that showcase anomalous nodes.

Using the reconstruction error as an indicator for anomalies has been employed for detecting times during molecular dynamics simulations where the collective motions suddenly change [65], finding frames in a video which are unlike the others [82], and identifying data that do not fit any concepts [8].

To address the intermediate blowup problem—when the input tensor and output tensors exceed memory capacity during the computation.—Memory-Efficient Tucker (MET) decomposition was proposed [46], based on the original Tucker decomposition [83]. The Tucker decomposition approximates a higher order tensor using a smaller core tensor (thought of as a compressed version of the original tensor) and a matrix for every mode (dimension) of the original tensor. Similarly, methods have been developed for offline, dynamic, and streaming tensor analysis [75], in addition to static and sliding window based methods [74]. These extensions allow the method to operate on continuous graph streams as well as those with a fixed number of time points. Compact Matrix Decomposition (CMD) computes a sparse low rank approximation of a given matrix. By applying CMD to every adjacency matrix in the stream, a time series of reconstruction values is created and used for event detection [76]. Colibri [81] and
ParCube [62] can be used in the same fashion and provide a large increase in efficiency. The PARAFAC decomposition has been shown to be effective at spotting anomalies in tensors as well [48].

A probabilistic model that represents the graph expected at a given time step can be created using the Chung-Lu random graph model [22]. Taking the difference between the real graph’s adjacency matrix and the expected graphs forms a residual matrix. Anomalous time windows are found by performing SVD on the residual matrices—on which a linear ramp filter has been applied—and by analyzing the change in the top singular values [54]. The responsible nodes are identified via inspection of the right singular vectors. More accurate graph models that also consider attributes are proposed in [55].

Performing PCA on the data, the calculated eigenvectors can be separated into “normal” and “anomalous” sets by projecting the data onto each eigenvector, and flagging as anomalous the projections that contain data points outside 3 standard deviations of the values. At each time step, the components are found by projecting the data onto its normal and anomalous subspace. Changes are detected when modifications in the anomalous components are above a threshold in traffic flow data [50]. Expanding on this method, joint sparse PCA and graph-guided joint sparse PCA were developed to localize the anomalies [45]. The responsible nodes are more easily identified by using a sparse set of components for the anomalous set. Nodes are given an anomaly score based on the values of their corresponding row in the abnormal subspace. As a result of the anomalous components being sparse, the nodes that are not anomalous receive a score of 0. Due to the popularity of PCA in traffic anomaly detection, a study was performed identifying and evaluating the main challenges of using PCA [67].

Change detection

**Main Idea.** The activity vector of a graph is the principal component, the left singular vector corresponding to the largest singular value obtained by performing SVD on the weighted adjacency matrix. A change point is when an activity vector is substantially different from the “normal activity” vector.

The normal activity vector is the left singular vector obtained by performing SVD on the matrix formed by the activity vectors for the last $W$ time steps. Each time point is given a score based on the difference between the normal vector and activity vector. Anomalies can be found online using a dynamic thresholding scheme, where time points with a score above the threshold are output as changes [42]. The nodes responsible are found by calculating the ratio of change between the normal and activity vectors. The nodes that correspond to the indexes with the largest change are labeled anomalous. Similar approaches have used the activity vector of a node-to-node feature correlation matrix [5], and a node-to-node correlation matrix based on the similarity between node’s neighbors [43].
Distance based methods

Using the notion of distance as a metric to measure change is natural. Two objects that have a small difference in a measured metric can be said to be similar. The metrics measured in graphs are typically structural features, such as the number of nodes. Once the summary metrics are found for each graph, the difference or similarity, which are inversely related, can be calculated. The variation in the algorithms lies in the metrics chosen to extract and compare, and the methods they use to determine the anomalous values and corresponding graphs.

Edge detection

**Main Idea.** *If the evolution of some edge attribute (e.g., edge weight) differs from the “normal” evolution, then the corresponding edge is characterized as anomalous.*

In [52], a dynamic road traffic network whose edge weights vary over time is studied. The similarity between the edges over time is computed using a decay function that weighs the more recent patterns higher. At each time step, an outlierness score is calculated for each edge based on the sum of the changes in similarity scores.

Viewing the network at a stream of edges, meaning the network does not have a fixed topology as the road traffic network did, the frequency and persistence of an edge can be measured and used as an indicator of its novelty. The persistence of an edge is how long it remains in the graph once it is added. Set system discrepancy [17] is one way to measure the persistence and frequency of the edges. When a new edge arrives, its discrepancy is calculated and compared to the mean discrepancy value of the active edge set. If the weighted discrepancy of the edge is more than a threshold level greater than the mean discrepancy, the edge is declared novel (anomalous) [1]. This method also allows anomalous nodes and subgraphs to be detected.

Subgraph detection

**Main Idea.** *A subgraph with many “anomalous” edges is deemed anomalous.*

Fixed subgraphs over a time interval are scored by the sum of their edge weights over that time interval. Significant anomalous regions (SARs) are the fixed subgraphs that have the highest scores. Every edge at every time stamp is given its own anomaly score, which is a function of the probability of seeing that particular edge weight on that particular edge given the distribution of weights for that edge over all graphs in the series. Mining SARs is analogous to finding the heaviest dynamic subgraphs (HDS) [57]. A similar approach mines weighted frequent subgraphs in network traffic, where the edge weights correspond to the anomaly contribution of that edge [38].
Event detection

Main Idea. Provided a function $f(G_i, G_j)$ that measures the distance between two graphs, a time series of distance values can be constructed by applying the function on consecutive time steps in the series. Anomalous values can then be extracted from this time series using a number of different heuristics, such as selecting the top $k$ or using a moving average threshold.

Extracting features from the graphs is a common technique to create a summary of the graph in a few scalar values, its signature. Local features are specific to a single node and its egonet (the subgraph induced by itself and its one-hop neighbors), such as the node or egonet degree. Global features are derived from the graph as a whole, such as the graph radius. The local features of every node in the graph can be agglomerated into a single vector, the signature vector, of values that describe the graph using the moments of distribution (such as mean and standard deviation) of the feature values.

In [10], the similarity between two graphs is the Canberra distance, a weighted version of the $L_1$ distance, between the two signature vectors. A similar approach is used in [51] to detect abnormal times in traffic dispersion graphs. Instead of an agglomeration of local features, it extracts global features from each graph, and any graph with a feature value above a threshold is anomalous.

As an alternative to extracting multiple features from the graph, the pairwise node affinity scores may be used. Pairwise node affinity scores are a measure of how much each node influences another node, and can be found using fast belief propagation [47]. In [49] the scores are calculated for two consecutive time steps, and the similarity between the two graphs is the rooted Euclidean distance (Matusita distance) between the score matrices. A moving threshold is set on the time series of similarity scores using quality control with individual moving range. The exponential weighted moving average has also been used as a way to dynamically set the threshold, tested on distribution features extracted from Wikipedia revision logs [7].

Complementary to feature similarity, one can look at the structural differences between graphs to identify the magnitude of change. These methods focus on the function that defines the distance between graphs instead of finding the optimal features to use as summaries. Many metrics have been developed and tested to quantify the differences between graphs. Ten different distance functions (e.g., weight distance, maximum common subgraph (MCS) weight difference, graph edit distance) were evaluated on TCP/IP traffic graphs with known anomalies (ground truth). Box-Jenkins autoregressive moving average (ARMA) modeling [12] was used to set the threshold and identify anomalies. Of the ten distance functions tested, the MCS based methods performed the best [63]. In [61], five different distance scoring functions were tested on web graphs with specific types of anomalies: missing subgraph, missing vertices, and connectivity change. The best method was the signature similarity, which is done by extracting a signature vector from each graph and finding the distance between them. The features used were each vertex and its PageRank value, and each edge $(u, v)$ with a weight of $u$’s PageRank divided by the number of outlinks from $u$. A fixed threshold was set to find graphs with abnormally low similarity scores.
Instead of finding the difference between two consecutive graphs, events can be detected using the time series of robustness values for each graph. Robustness is a measure of the connectivity of the graph. A graph with a high robustness will retain the same general structure and connectivity even when some nodes or edges are removed. Finding events is then finding when the robustness value changes significantly [53].

Probabilistic model based methods

With a foundation in probability theory, distributions, and scan statistics, these methods typically construct a model of what is considered “normal” or expected. Deviations from this model are flagged as anomalous. The type of model used, how it is constructed, what it is modeling, and the method for determining outliers is what differentiates these approaches.

Anomalous node detection

Main Idea. There are three main approaches: (a) Modeling some node features using probabilistic approaches, and tracking the nodes that do not fit the models; (b) Building scan statistics time series and detecting points that are several standard deviations away from the mean; (c) Node classification.

One way to model the relationship between nodes is considering the communication between them as a Bayesian discrete time counting process. A model is fit to the expected number of communications (weight) for each edge and is continuously updated as new graphs are considered. At any time point when the actual number of communications is different from the expected number by a statistically significant amount both nodes are added to the set of anomalies [39]. Statistical significance is determined by a predictive \( p \)-value dropping below a threshold.

Scan statistics are often called “moving window analysis,” where the local maximum or minimum of a measured statistic is found in specific regions of the data. In a graph a scan statistic can be considered as the maximum of a graph invariant feature, such as the number of edges, found for each node and its neighborhood in the graph. The local statistic for each node is normalized using the mean and standard deviation of its recent values. The scan statistic of a graph is the maximum of all of the local scan statistics. Normalizing the values accounts for the history of each node, meaning the statistic for each node is relative to its own past instead of the values of the other nodes. Building a standardized time series of the scan statistic values, any value that is five standard deviations away from the mean of the series is considered an event. The node most responsible is identified as the one that was chosen for the scan statistic value for the entire graph [64].

The Markov Random Field Model (MRF) is used to uncover the states for nodes and infer the maximum likelihood assignment by belief propagation algorithm. In [60], anomalies (fraudsters) are uncovered in an online auction network by discovering bipartite cores, which are posited to be the interaction behavior between fraudsters and...
accomplices. It incrementally updates the model as new edges are added, taking advantage of the fact that an edge insertion or removal will affect only a local subgraph. Nodes that are assigned the label of fraudster are anomalous.

Anomalous subgraph detection

**Main Idea.** Fixed subgraphs (e.g., like paths and stars), multigraphs, and cumulative graphs are used to construct models on the expected behaviors. Deviations from the models signify an anomalous subgraph.

To identify hacker behaviors in a network, scan statistics are combined with a Hidden Markov Model (HMM) for edge behavior. Unlike [39] that used neighborhoods, the local scan statistics are based on two graph shapes, the $k$-path and star. Comparing the scan statistics of a sliding window to its past values, and using an online thresholding system, local anomalies are identified [58]. The local anomaly is the subgraph representing the $k$-path or star used to derive the statistic.

Another method to model a dynamic network, instead of using a series of graphs, is to have a single multigraph where parallel edges correspond to communication between nodes at two different time steps. The initial multigraph is decomposed into telescoping graphs for each time window [24]. Telescoping graphs that are large in size, but have a low probability of appearing, are output as anomalies.

Likewise, a cumulative graph, which included every edge seen up until the current time step, could be used. Edge weights are calculated using a decay function where more recent edges weigh more. The connected components in the cumulative graph for the entire time series are found and monitored for unusual behavior. Unusual behavior is defined as a deviation from the model constructed using the cumulative graph for expected behavior at a given time step [80].

Event detection

**Main Idea.** Deviations from the models of the graph likelihood or the distribution of the eigenvalues reveal when an event occurs.

Maintaining a structural summary of the network by keeping multiple partitionings of the graph can be used to detect graph outliers online [4]. The structural summary and partitionings allow an edge likelihood value to be calculated for every edge in the new graph. After calculating the edge likelihoods, the graph likelihood is derived by taking the geometric mean of the edge likelihoods. If the graph likelihood is several standard deviations below the average of all the graphs up until that point, the graph is an outlier.

In [41], a similar edge likelihood approach is to construct an edge probability matrix for each time window in the series, where each element in the matrix $(i,j)$ is the probability of having an edge between node $i$ and $j$. Using Expectation Maximization to
calculate a potential score for every sender-recipient pair, the probability of all possible source-destination node tuples can be estimated. Tuples represent an email, where there is one sender and multiple recipients, while edges represent a link between two individual nodes. Time windows that are anomalous are found by comparing the average log-likelihood scores for the windows. Individual emails that are abnormal can also be found by looking at the emails with the lowest likelihood.

Based on the assumption that each node has a time series of feature values, a node-to-node correlation matrix can be generated for each time step. The eigen equation of the correlation matrix is compressed by keeping the largest eigenvalues and a set of low dimension matrices (one matrix for each node). By learning the distributions of the eigenvalues and matrices, both events and the nodes responsible can be identified. When the eigenvalues deviate from the expected distribution an event has occurred. The nodes whose matrices deviate from the matrix distribution the most are considered responsible [40].

**Conclusion**

In this survey we first introduced four types of anomalies that occur in a dynamic graph (node, edge, subgraph, and events) and an example of each. Individual nodes in the graph are considered anomalous if their evolution is unlike the rest of the nodes, see Figure 1 for an example. Anomalous edges typically relate to irregular weight evolution or the insertion/removal of an unlikely edge, Figure 2 shows an example of an irregular edge weight. Entire subgraphs that behave unlike the rest of the graph, for example, in terms of community evolution or communication patterns, are anomalous. Figure 3 shows two types of anomalous community evolutions. The last type of anomaly is points in time which represent a large change in the graph or an isolated abnormal fluctuation in the graph behavior (event and change detection, respectively, see Figure 4).

We partitioned the methods into five main categories derived from the intuition behind each: community, compression, decomposition, distance, and probabilistic model based. Community based methods track the evolution of communities over time, using the changes in the communities to identify anomalies. Different methods have different definitions of a community and may look at the evolution of a node’s community belongingness instead of the behavior of communities themselves. Decomposition based methods try to find a low dimensional approximation of the data represented as a tensor or a sequence of matrices. When the approximation is poor and there is a high reconstruction error it indicates a lack of regularity, hence abnormal behavior. Performing the decomposition online and the target of the decomposition is the major difference between the methods presented. Distance based methods use graph invariant summaries and find the difference between consecutive values. They differ mainly in their choice of summary values and method of declaring a value to be abnormal. Probabilistic methods create a model of “normal” or “expected” behavior and declare deviations from this model anomalous. The type of model used and the method for determining how closely the model is
being followed is what separates these methods. [Table 1] and [Table 2] show a summary of the methods presented in this survey, the type of anomalies they detect, and the year the paper was published. The field of anomaly detection in dynamic graphs is relatively young and is rapidly growing in popularity, as indicated by the number of papers published in the past five years. For further reading on anomaly detection in graphs and other domains we direct the reader to [2, 14, 15, 16, 32, 72].

Acknowledgements

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Figure 1: Three snapshots of a dynamic graph, where the nodes are plotted in a 2-dimensional space based on features extracted. At time step 1, all of the nodes belong to the same community, indicated by the blue circle, and are hence not anomalies (blue points). At time steps 2 and 3, the distribution of feature values for the community shifts, however, some nodes do not obey this pattern and are marked as anomalous (red crosses).

Figure 2: An illustration of anomalous edges that occur due to an irregular pattern of their weight over time, with anomalous edges highlighted. At each time step a node’s weight typically shifts by $\pm 0.05$ at most. However, edge $(3, 4)$ has a spike in its weight at time step 2, unlike any other time in the graph. Similarly, at time step 3, edge $(1, 4)$ spikes in value. These spikes cause the edges to be considered anomalous.
<table>
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Table 2  Summary of methods and the type of problems they solve (continued).

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Figure 3: Two different types of anomalous subgraphs. A shrunken community is shown in Figure 3a, where a community loses members from one time step to the next. A split community is shown in Figure 3b, where a single community breaks into several distinct smaller communities.

Figure 4: An example of an event, a change, and the difference between them. In both Figure 4a and Figure 4b the graphs initially have only a few edges. At time step 3, the graphs become highly connected, almost forming a clique. The major difference is observed at the time step following the insertion of the edges. In (a) the edges are then removed and the graph returns to a state similar to the one before the insertion, representing an isolated event. In (b), however, the graph maintains its new state indicating the sustained shift, or change, in the graph.