# Multi-layer bipartite structural features to analyze YouTube Social Network 

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Heraklion, February 2022

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#### Abstract

This work investigates interactions on YouTube, concerning predicting missing or unseen interactions on multi-layer bipartite networks. More precisely, given a set of own interactions between YouTube users and videos, we measure how accurately we can predict comment interactions. We propose structural bipartite features, which enhance the performance of simple prediction models, to find missing or unseen links. Experimental validation of the proposed approach is carried out on multi-layer networks formed on YouTube. We have crawled an extensive dataset of YouTube videos, the channels that own them, and the authors of their comments. Using a machine learning framework, we find that we can predict future and unseen comment interactions on YouTube videos with precision $99 \%$. We also show that to predict a day's comment interactions it suffices to account network information generated 1 day prior. Our set-up is implemented on the MapReduce model. We propose two MapReduce algorithms, one that counts the bitruss number of an edge and one that clusters edges into blooms in a bipartite network.


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## Chapter 1

## Introduction

In this study, we investigate the comment interaction on the YouTube social network. The goal is to predict which YouTube user is commenting on what video. We approach this comment prediction task as an edge prediction problem for multi-layer bipartite graphs. We propose multi-layer structural features and compare their predictive power over simple structural features, on the YouTube bipartite networks.

YouTube is an online video sharing and social media platform. It is the second most visited website, with more than one billion monthly users [1]. YouTube users can watch, like, share, comment, put into a playlist, and upload their videos. Prior research studies on the social media platform have focused primarily on analyzing existing content and providing statistical insights for the platform [9, 21, 27, 28]. There is very little work related to predicting interactions on the platform solely.

The task of predicting interactions fits naturally in a graph framework. For that reason, we use bipartite graphs (bigraphs) to model relations between users and videos on YouTube. More precisely, we represent in a bigraph the relation 'own', which is user owns video and the relation 'comment', which is user comments video. Following, with the use of the topological features emerging from the 'own' and 'comment' graph, we answer the question 'which user is going to comment on what video'.

To make comment predictions on YouTube, we translate the problem to a link prediction task. Given a network at a specific period of time, the link prediction problem is considered as a task of discovering unseen links or predicting new ones that will occur in a future time. In this work, we use the link prediction as a binary classification problem, where first-order structures such as the node degree, edge existence, and the high-order structures such as the butterfly, the bloom, and the multi-layer topological structures are used as features.

A butterfly, also known as rectangle [2, 19, 20, 25, 24], is a complete $2 \times 2$ biclique and is considered as an analogue of a triangle in unipartite graphs. Figure 1.1 shows a user-video bipartite network with 2 butterflies formed, marked in dotted lines. We also use the bloom [26] as a structural feature to infer information about the bipartite network. Blooms can be considered clusters of butterflies. An example of bloom construction out of a bipartite graph is in figure 3.2. As high-order structural features, we also consider the topological features emerging when combining the 'own' and 'comment' interactions. These two YouTube interactions characterize nodes and their connections on two different relations. This setting is also known as multi-layer network, and we take it into account to make more sophisticated predictions.


Figure 1.1: A user-video bipartite network with 2 butterflies (dotted edges).
Overall, this study makes the following contributions:

- We compare the predictive power of first-order to high-order structures on the missing link task on a multi-layer bipartite network.
- We compare the predictive power of first-order to high-order structures on the formation of future links on a multi-layer bipartite network.
- We show that to predict a day's comment interactions suffices to consider data generated one day prior.
- We implement a MapReduce algorithm that counts the bitruss number of an edge in a bipartite network.
- We implement a MapReduce algorithm that clusters edges of a bipartite network into blooms.


## Chapter 2

## Related work

Given a network at a specific time, the link prediction problem is a task of discovering unseen links or predicting new ones that will occur in a future time. The link prediction has been extensively studied in unipartite and bipartite networks $[17,4,18,15,22,6,5,16,13]$. Al Hasan et al. [3] introduce the link prediction problem as a binary classification task, where various similarity metrics and data outside the graph topology scope were used as features. The same supervised classification approach has also been used by [10, 23, 7].

In unipartite graphs, link prediction algorithms make the assumptions of Triangle closure and Clustering [14], which cannot be in use on bipartite graphs. Technically, unipartite link prediction algorithms can apply to bipartite graphs, but they will not perform well because they are based on the triangle closure. For that reason, the link prediction algorithms need to be fine-tuned to match the structure of bipartite graphs.

In this study, we approach the link prediction problem as a binary classification task. We adopt multi-layer higher-order topological features that have application only on bipartite graphs. In contrast, authors of [6, 8] adapt some topological measures used in unipartite graphs for predicting links in bipartite graphs. They also transform bipartite graphs into unipartite and perform link prediction. The work of Allali et al. [5] propose a link prediction method that is based on the notion that if two nodes have a common neighbor in the graph, they will probably acquire more in the future.

We expand the idea of the triangle closure and build upon the analog structure for bipartite graphs, which is the butterfly. A butterfly (also known as rectangle), is a complete $2 \times 2$ biclique [24, 19, 25]. Besides the butterfly structure, we use the notion of the Bloom structure and bitruss number [26] of a link for our bipartite graph embedding. To enhance the predictive features, we also leverage structural features emerging when considering multi-layer graphs.

Multi-layer networks characterize multiple types of interactions not possible
to represent by using a traditional monolayer network approach. Figure 2.1 shows an example of a multi-layer bipartite network. Each layer represents a bipartite network, whose vertices are divided into two disjoint and independent sets, such that every edge connects a vertex from one set to the other [11]. In figure 2.1, there is a set with circle vertices and a set with square vertices represented on two networks, on Layer 1 and on Layer 2. For example, Layer 1 can represent the 'own' relations and Layer 2 can represent the 'comment' relation.


Figure 2.1: Multi-layer bipartite graph with 2 layers.

Close to our work lies the framework proposed in [22], where they study user interactions on Twitter. They use multi-layer directed networks and show that features involving triads turn out to be important for accurate predictions. There is little work regarding multi-layer bipartite link prediction. The study of [12] propose a community detection-base measures for link prediction.To the best of our knowledge, we are the first to address the multi-layer bipartite link prediction while taking into account multi-layer features on YouTube. We create a graph embedding with first-order and high-order structural features, which consists of multi-layer butterfly motif structures on the YouTube comment and own network.

## Chapter 3

## Design

In this section, we describe the set-up for the multi-layer link prediction in bipartite graphs, to predict the comment action on YouTube. We model the bipartite link prediction problem as a binary classification task, where each data point corresponds to a pair of vertices in the network. Vertices are divided into two disjoint and independent sets U and V , such that they compose an edge connecting a vertex in U to one in V .

The link prediction problem corresponds to two tasks, one to predict future links and one to predict missing links in a network. In our set-up we explore both prediction tasks. We need to create a graph embedding to perform both link prediction tasks. For this study, the graph embedding for these tasks is the same and is a vector that contains structural information for every edge/link in the network. Specifically, the vector contains first-order and high-order structural features of the multi-layer bipartite network. The first-order structural features are the degree of a node and the existence of an edge in the different layers. The high-order structural features we use in this study emerge from the butterfly motif and are the blooms, the bitruss number, and the multi-layer butterflies.

## 1 Definitions

Our problem is defined as an undirected bipartite graph $G(U, V, E)$, where $U(G)$ denotes the set of vertices in partition U and $V(G)$ denotes the set of vertices in partition V with $U(G) \cap V(G)=\varnothing$, and $E(G) \subseteq U(G) \times V(G)$ denotes the edge set. An edge between two vertices a and b in G is denoted as $(a, b)$ or $(b, a)$.

Butterfly (Rectangle): Given a bipartite graph G (single layer) and four vertices $(a, b, \in V(G)$ and $c, d \in U(G))$, a butterfly induced by the vertices a,
$\mathrm{b}, \mathrm{c}, \mathrm{d}$ is a (2,2)-biclique of G ; that is, a and b are both connected to c and d, respectively, by edges $(a, c),(a, d),(b, c),(b, d) \in E(G)$. For example in figure 3.1 there are 2 butterflies. One butterfly is supported by the edges((U1,V1), (U1,V2), (U2,V1), (U2,V2)) and the other is supported by the edges((U2,V2), (U2,V3), (U3,V2), (U3,V3)).

Bitruss number: Given a bipartite graph G, the bitruss number of an edge e, denoted as $\mathrm{b}(\mathrm{e})$, is a number k that indicates how many butterflies e supports. Figure 3.1 shows that the dotted edge(U2,V2) has bitruss number equal to 2, because this edge supports 2 butterflies, the solid edges((U1,V1), (U1,V2), (U2,V1), (U2,V3), (U3,V2), (U3,V3)) have bitruss number $=1$, and the dashed edges((U4,V3), (U4,V4), (U4,V5)) have bitruss number $=0$ because they do not support any butterfly.


Figure 3.1: A bipartite network with 2 butterflies and the bitruss count of each edge.

Bloom: Given a bipartite graph G, a bloom [26] is a cluster of butterflies. Blooms concentrate a set of edges that support butterflies that share at least one edge. Each edge in G can only be part of one bloom. The bipartite network in figure 3.2 has only one bloom with id V1U1. The bloom V1U1 is supported by the edges of butterflies made by the vertices [U1, U2, V1, V2] and [U2, U3, V2, V3].


Figure 3.2: Bloom construction out of a bipartite network.

Multi-layer Butterfly: Given two bipartite graphs $G_{1}$ (Layer 1), $G_{2}$ (Layer 2) and 4 vertices connected with 4 edges. A multi-layer butterfly extends the criteria of the single-layer butterfly and has two properties regarding the butterfly edges. First property is to have 1 edge at one layer and the remaining 3 at the other layer, and the second property is to have 2 edges at one layer and the remaining 2 at the other layer. In total, we define six structural types of multi-layer butterflies shown in figure 3.3. We set Layer $1\left(G_{1}\right)$ to be the relation 'own', marked in dotted lines and Layer $2\left(G_{2}\right)$ to be the relation 'comment', marked in solid lines. For example the Type 3 multi-layer butterfly structure from the figure 3.3 indicates that the two users own a video and that these two users have commented on each other's video.


Figure 3.3: The 6 structural types of multi-layer butterflies.

## 2 MapReduce Algorithms

For the bipartite link prediction implementation, we use the MapReduce programming model to scale out the problem. More specifically, we use Apache Spark, which uses the MapReduce distributed computing framework as its foundation. Spark is a multi-language engine for executing data engineering, data science, and machine learning on a single-node machine or clusters.

Following. we present the MapReduce pseudocode to describe the algorithms we use for the Edge Bitruss Count (algorithm 1) and the Edge Bloom Identification (algorithm 2) in a bipartite graph.

Given a bipartite network, the Edge Bitruss Count Algorithm 1 computes the bitruss number of an edge $b(e)$, for $b(e) \geq 1$. The bitruss number of an edge represents the number of butterflies the edge supports. The algorithm 1 returns a list of edges with their bitruss number.

For example, given as input to the algorithm 1 the Graph of figure 3.1, in line 1 we gather all nodes from set V of the graph G , so we get src_set $\leftarrow$ [V1,V2,V3,V4,V5], also shown in figure 3.4(i). In line 2, we declare a function
named $\mathrm{N}(\mathrm{n})$, which for every given node n it returns a list with its neighbors. In line 3, we declare the function count_butterflies(list), that given a list it returns its size minus 1 . This is because 2 source nodes can have butterfly support equal to the number of their common neighbors minus 1 . In line 4, we performe a cartesian product to all the node of set V of src_set where $V_{i} \neq V_{j}$, so src_cartesian $\leftarrow[(\mathrm{V} 1, \mathrm{~V} 2),(\mathrm{V} 1, \mathrm{~V} 3),(\mathrm{V} 1, \mathrm{~V} 4),(\mathrm{V} 1, \mathrm{~V} 5),(\mathrm{V} 2, \mathrm{~V} 3)$, (V2,V4), (V2,V5), (V3,V4), (V3,V5), (V4,V5)], also shown in figure 3.4(ii).

In lines 5-7, we calculate the potential butterflies for each pair created in src_cartesian, if they have more or equal to 2 neighbors. This condition is because two source nodes must have at least two common neighbors to form a butterfly. So butterflies $\leftarrow[((\mathrm{V} 1, \mathrm{~V} 2),(\mathrm{U} 1, \mathrm{U} 2)),((\mathrm{V} 2, \mathrm{~V} 3),(\mathrm{U} 2, \mathrm{U} 3))]$, also shown in figure 3.4 (iii). In lines $\mathbf{8 - 1 1}$, we select the first source node from the pair of source nodes, with all the destination nodes in the list of common neighbors, so as to construct an edge, and compute the bitruss number of them with the function count_butterflies. So for our example we get bitruss_src $_{1} \leftarrow[(1,(\mathrm{~V} 1, \mathrm{U} 1)),(1,(\mathrm{~V} 1, \mathrm{U} 2)),(1,(\mathrm{~V} 2, \mathrm{U} 2)),(1,(\mathrm{~V} 2, \mathrm{U} 3))]$, shown in figure $3.4(\mathrm{iv})$. Next we do the same for the second source node and we get bitruss_src $c_{2} \leftarrow[(1,(\mathrm{~V} 2, \mathrm{U} 1)),(1,(\mathrm{~V} 2, \mathrm{U} 2)),(1,(\mathrm{~V} 3, \mathrm{U} 2)),(1,(\mathrm{~V} 3, \mathrm{U} 3)]$, also shown in figure $3.4(\mathrm{v})$. Following in lines 12-13, we join the bitruss_src $c_{1}$ and bitruss_src $_{2}$ and we get a set that has a pair of source and destination node, which is an edge of G and its bitruss number. So bitruss $\leftarrow[((\mathrm{V} 1, \mathrm{U} 1), 1)$, ((V1,U2),1), ((V2,U1),1), ((V2,U2),1), ((V2,U3),1), ((V2,U2),1), ((V3,U2),1), ((V3,U3),1)]. Last in line 14, for each edge in the bitruss set we sum (reduce) their bitruss number and we get the result set that contains edges and their total bitruss number. For our example in this step we get edge_bitruss $\leftarrow$ [((V1,U1),1), ((V1,U2),1), ((V2,U1),1), ((V2,U2),2), ((V2,U3),1), ((V3,U2),1), ((V3,U3),1)], as shown in figure 3.4(vi).

Following we descrive the space complexity of the Edge Bitruss Count Algorithm 1. For input we have a graph $\mathrm{G}=(\mathrm{U}, \mathrm{V}, \mathrm{E})$, where V are the nodes in set $\mathrm{V}, \mathrm{U}$ are the nodes in set V and E are the edges between nodes of set U and set V . In line 1 we have $\mid$ src_set $|=|V|$. In line $\mathbf{2}$ we declare a function N that gets a node n and returns its neighbors. In our case we give as input the nodes of set V , so we have $N(n) \subseteq U$. We also have that $\sum_{n \in \text { src_set }}|N(n)|=|E|$. In line 4 we get $\mid$ src_cartesian $\left|=|V|^{2}-|V|=O\left(|V|^{2}\right)\right.$. For lines 5-7, we have $\mid$ butterflies $|\leq|$ src_cartesian $\mid+O\left(|V|^{2}\right)$. We get bitruss_src $c_{1} \leq O\left(|V|^{2} *|E|\right)$, in lines 8-9. We have the same complexity for bitruss_src $_{2} \leq O\left(|V|^{2} *|E|\right)$, in lines 10-11. In lines 12-13 we get bitruss $\leq O\left(|V|^{2}\right) *|E|+O\left(|V|^{2}\right) *|E|$, since is the union of bitruss_src $_{1}$ and bitruss_src $_{2}$. Last step in line 14 we have edge_bitruss $\leq|E|$.


Figure 3.4: Illustration of the Algorithm 1 with input the Graph of figure 3.1.

```
Algorithm 1 Edge Bitruss Count
Input: \(\operatorname{Graph} G=(U, V, E)\)
    src_set \(\leftarrow V(G) \quad / * V(G)\) are the source nodes */
    \(N(n) \quad / *\) Returns the neighbor set of node n */
    count_butterflies(list) /* Gets a list of destination nodes and returns list.size-1 */
    /* The number of butterflies 2 source nodes support, is equal to the number of their
    common neighbors - 1 */
    src_cartesian \(\leftarrow\) src_set \(\times\) src_set where \(V_{i} \neq V_{j}\)
    butterflies \(\leftarrow\left(\forall\left(s r c_{1}, s r c_{2}\right) \in \operatorname{src}\right.\) _cartesian \() \cdot \operatorname{map}\left(\left(s r c_{1}, s r c_{2}\right)\right.\),
                                    \(\left.N\left(s r c_{1}\right) \cap N\left(s r c_{2}\right)\right)\).
                                    filter \(\left(N\left(s r c_{1}\right) \cap N\left(s r c_{2}\right) \geq 2\right)\)
    /* To compose a butterfly 2 source nodes must have at least 2 common neighbors */
    bitruss_src \(c_{1} \leftarrow\) butterflies.map \(\left(\right.\) count_butterflies \(\left(N\left(s r c_{1}\right) \cap N\left(s r c_{2}\right)\right)\) :
                                    BitrussNumber, \(\left.s r c_{1} \times N\left(s r c_{1}\right) \cap N\left(s r c_{2}\right)\right)\)
    bitruss_src \(c_{2} \leftarrow\) butterflies.map(count_butterflies \(\left(N\left(\operatorname{src}_{1}\right) \cap N\left(s r c_{2}\right)\right)\) :
                                    BitrussNumber, \(\left.\operatorname{src}_{2} \times N\left(s r c_{1}\right) \cap N\left(s r c_{2}\right)\right)\)
    bitruss \(\leftarrow\) bitruss_src \(c_{1} \cup\) bitruss_src \(c_{2}\).
                        \(\operatorname{map}((s r c, d s t): E d g e\), BitrussNumber \()\)
4: edge_bitruss \(\leftarrow\) bitruss.reduce \((\) Edge, BitrussNumber \()\)
15: return edge_bitruss
```

Given a bipartite network, the Bloom Identification Algorithm 2 finds and constructs the blooms that exist in the network. Blooms concentrate a set of edges that construct butterflies that share at least one edge. We give an id to the bloom based on the vertices it concentrates.

For example, given as input to the algorithm 2 the Graph of figure 3.2, in line 1 we gather all nodes from set V of the graph G, so we get src_set $\leftarrow$ [V1,V2,V3,V4,V5], also shown in figure 3.5(i). In line 2, we declare a function named $\mathrm{N}(\mathrm{n})$, which for every given node n it returns a list with its neighbors. In line 3, we performe a cartesian product to all the node of set V of src_set where $V_{i} \neq V_{j}$, so src_cartesian $\leftarrow[(\mathrm{V} 1, \mathrm{~V} 2)$, (V1,V3), (V1,V4), (V1,V5), (V2,V3), (V2,V4), (V2,V5), (V3,V4), (V3,V5), (V4,V5)], also shown in figure 3.5(ii). In lines 4-6, we calculate the potential butterflies for each pair of src_cartesian, if they have more or equal to 2 neighbors. This condition is because two source nodes must have at least two common neighbors to form a butterfly. So butter flies $\leftarrow[((\mathrm{V} 1, \mathrm{~V} 2),(\mathrm{U} 1, \mathrm{U} 2)),((\mathrm{V} 2, \mathrm{~V} 3),(\mathrm{U} 2, \mathrm{U} 3))]$, also shown in figure 3.5(iii). In lines $\mathbf{7 - 9}$, we create the potential blooms. We construct a potential bloom id by selecting the minimum id between the two
source node ids and the minimum id from the neighbor' nodes of these source nodes and append it to the set of butterflies. So we have potential_bloomids $\leftarrow$ [(V1U1,(V1,V2),(U1,U2)), (V2U2,(V2,V3),(U2,U3))], also shown in figure 3.5(iv). Following in lines 10-13, we select the bloom id and the source nodes with all the destination nodes in the list of common neighbors and create all the potential edges with the bloom id they are given in the previous step. First we take the first node from the two source node, so we get edge_src $c_{1}$ bloomids $\leftarrow$ [(V1U1,(V1,U1)), (V1U1,(V1,U2)), (V2U2,(V2,U2)) (V2U2,(V2,U3))], which is shown in figure $3.5(\mathrm{v})$. Next we do the same for the second node from the two source node pair and we get edge_srcc_bloomids $\leftarrow[(\mathrm{V} 1 \mathrm{U} 1,(\mathrm{~V} 2, \mathrm{U} 1))$, (V1U1,(V2,U2)), (V2U2,(V3,U2)), (V2U2,(V3,U3))], shown in figure 3.5(vi). In lines 14-15, we take edge_src $c_{1}$ bloomids and edge_src $c_{2}$ bloomids, and construct the union of these two and as a result we get a set that has a bloom id and an edge of G. So in our example we get edge_bloomids $\leftarrow[((\mathrm{V} 1, \mathrm{U} 1), \mathrm{V} 1 \mathrm{U} 1)$, ((V1,U2),V1U1), ((V2,U2),V2U2), ((V2,U3),V2U2), ((V2,U2),V1U1), ((V3,U2),V2U2), ((V3,U3),V2U2),((V2,U1),V1U1)]. In the edge_src $c_{2}$ bloomids set in our example, the edge (V2,U2) has two bloom ids assigned the V1U1 and the V2U2. To distinguish which bloom id to assign to similar cases we do the following steps. In lines lines 16-17 we group by edge the edge_src $c_{2}$ bloomids set. We have edge_belong_blooms $\leftarrow[((\mathrm{V} 1, \mathrm{U} 1),[\mathrm{V} 1 \mathrm{U} 1]),((\mathrm{V} 1, \mathrm{U} 2),[\mathrm{V} 1 \mathrm{U} 1])$, ((V2,U2),[V1U1,V2U2]), ((V2,U3), [V2U2]), ((V3,U2),[V2U2]), ((V3,U3),[V2U2]), ((V2,U1),[V1U1])], as shown in figure 3.5(vii) In lines 1819, we now as last step we group by the bloom ids of the edge_belong_blooms set. From the result of grouping we select from the bloom ids set the minimum id. The minimum id is now the unique identifier of the bloom and the set of edges that belong to it. So we have blooms $\leftarrow[\mathrm{V} 1 \mathrm{U} 1,((\mathrm{~V} 2, \mathrm{U} 2),(\mathrm{V} 3, \mathrm{U} 2)$, (V2,U3), (V3,U3),(V1,U1), (V2,U1), (V1,U2))], as shown in in figure 3.5(viii).

Following we descrive the space complexity of the Bloom Identification Algorithm 2. For input we have a graph $\mathrm{G}=(\mathrm{U}, \mathrm{V}, \mathrm{E})$, where V is the nodes in set V , U is the set of nodes in set V and E are the edges between the nodes of U set and V set. In line $\mathbf{1}$ we have $\mid s r c_{-}$set $|=|V|$. In line $\mathbf{2}$ we declare a function N that gets a node n and returns its neighbors. In our case we give as input the nodes of set V , so we have $N(n) \subseteq U$. We also have that $\sum_{n \in \text { src_set }}|N(n)|=|E|$. In line 3 we get $\mid$ src_cartesian $\left|=|V|^{2}-|V|=O\left(|V|^{2}\right)\right.$. For lines 4-6, we have $\mid$ butterflies $|\leq|$ src_cartesian $\mid+O\left(|V|^{2}\right)$. We get edge_src $c_{1}$ bloomids $\leq$ $O\left(|V|^{2} *|E|\right)$, in lines 10-11. We also have edge_src__bloomids $\leq O\left(|V|^{2} *|E|\right)$, in lines 12-13. In lines 14-15 we get edge_bloom $\leq O\left(|V|^{2}\right) *|E|+O\left(|V|^{2}\right) *$ $|E|$, since is the union of $e d g e_{-} s r c_{1}$ bloomids and edge_src $2_{2}$ bloomids. Next in lines 16-17 we have edge_belong_blooms $\leq|E|$. Last step in line 18-19 we have blooms $\leq \mid$ butterflies in the G|.

(i) Algorithm 2 - line 1

(iii) Algorithm 2 - lines 4-6

(v) Algorithm 2 - lines 10-11

(ii) Algorithm 2 - line 3

(iv) Algorithm 2 - line 7-9

(vi) Algorithm 2 - line 12-13

(vii) Algorithm 2 - lines 16-17

Figure 3.5: Illustration of the Algorithm 2 with input the Graph of figure 3.2.

```
Algorithm 2 Bloom Identification
Input: Graph \(=(U, V, E)\)
    src_set \(\leftarrow V(G) \quad / * V(G)\) are the source nodes */
    \(N(n) \quad / *\) Returns the neighbor set of node \(n\) */
    src_cartesian \(\leftarrow\) src_set \(\times\) src_set where \(V_{i} \neq V_{j}\)
    butterflies \(\leftarrow\left(\forall\left(s r c_{1}, s r c_{2}\right) \in\right.\) src_cartesian \()\). map \(\left(\left(\right.\right.\) src \(\left._{1}, s r c_{2}\right)\),
        \(\left.N\left(s r c_{1}\right) \cap N\left(s r c_{2}\right)\right)\).
    filter \(\left(N\left(s r c_{1}\right) \cap N\left(s r c_{2}\right) \geq 2\right)\)
    /* To compose a butterfly 2 source nodes must have at least 2 common neighbors */
    potential_bloomids \(\leftarrow\) butter flies.map \(\left(\left[\right.\right.\) [src \(\left.c_{1}, s c_{2}\right]\).min,
                                    \(\left(N\left(s r c_{1}\right) \cap N\left(s r c_{2}\right)\right)\). min \():\) BloomId,
                                    \(\left.s r c_{1}, s r c_{2},\left(N\left(s r c_{1}\right) \cap N\left(s r c_{2}\right)\right)\right)\)
    : edge_src1_bloomids \(\leftarrow\) potential_bloomids.map (BloomId,
                                    \(\left(s r c_{1} \times N\left(s r c_{1}\right) \cap N\left(s r c_{2}\right)\right)\). combinations (2) : Edge)
    edge_src \(2_{2}\) bloomids \(\leftarrow\) potential_bloomids.map(BloomId,
                                    \(\left(s r c_{2} \times N\left(s r c_{1}\right) \cap N\left(s r c_{2}\right)\right) \cdot\) combinations(2) : Edge)
4: edge_bloomids \(\leftarrow\left(\right.\) edge_src \(c_{1}\) bloomids \(\cup\) edge_src__bloomids \()\).
15: \(\quad \operatorname{map}(\) BloomId, Edge \()\)
16: edge_belong_blooms \(\leftarrow\) edge_bloomids.groupBy (Edge).
17: \(\quad \operatorname{map}(E d g e, \operatorname{Set}(\) BloomId \():\) BloomIds \()\)
18: blooms \(\leftarrow\) edge_belong_blooms.groupBy (BloomIds).
19: \(\quad \operatorname{map}(\) BloomIds.min, Set(Edges))
20: return blooms
```


## 3 Bipartite Graph Features

In this section, we describe the 34 bipartite graph features we use to perform the multi-layer link prediction. We perform two tasks of the link prediction in a bipartite network. The first task is to predict future interactions, and the second is to predict unseen interactions. Since link prediction is a typical binary classification task, we use a simple logistic regression model to make predictions.

For both tasks, we create a graph embedding that is composed of the 34 bipartite graph features described in table 3.1. Besides the simple first-order structural features from no 1 to 10 on table 3.1, we propose the high-order structural features of the multi-layer bipartite network that are from no 11 to 28 and features emerging from the butterfly motif that are from no 29 to 34
in the table. To the best of our knowledge, these high-order features have not been used before in the literature for the bipartite link prediction problem.

In our set up given 2 graphs, Layer $1\left(G_{1}\right)$ and Layer $2\left(G_{2}\right)$, we aim to make link predictions for both tasks on Layer 2. For every edge in Layer 1 and Layer 2 we calculate the features in table 3.1 and compose the multi-layer graph embedding. The multi-layer graph embedding is given as input to the Logistic regression model to make predictions for both tasks.

| Structural <br> Features | no. | Feature Name | Value | Description |
| :---: | :---: | :---: | :---: | :---: |
|  | 1 | EdgeinLayer1 | $1 \mid 0$ | $1 \mid 0$ |
|  | 2 | EdgeinLayer2 | $1 \mid 0$ | N |

Table 3.1: Multi-layer bipartite edge graph embedding features.

## Chapter 4

## Implementation

In this study, we measure the effectiveness of the simple first-order and the higher-order structural features (table 3.1), for the link prediction problem. We compare the structural features' predictive power over multi-layer bipartite graphs on YouTube.

To evaluate these features' predictive power, we use them in two settings. The first setting predicts the formation of future links. To predict a day's links on the YouTube network, we examine two cases. In case one, we gather one day's prior dense sample of the YouTube network, and in case two, a sample of two prior consecutive days. We then compare these two cases' predictions and find that to predict a day's interactions, it is better to take into account network information generated one day prior, i.e., most information can be found on the previous day. For the second setting, we predict missing links from the network. We use a YouTube network sample and hide $30 \%$ of its links. Then with the information of the rest of the network, we predict these hidden links. We found that the higher-order structural features can predict with $99 \%$ precision missing and future interactions in the comment network.

## 1 Dataset

We use the YouTube Data API to monitor YouTube traffic generated between 19 of July 2020 and 22 of September 2020 on videos related to the US 2020 elections. To gather the topic-specific videos, we first use the Twitter API to obtain tweets that contain hashtags and keywords related to the US elections. From the corpus of those tweets, we extract the YouTube video links and gathered a dense sample of the YouTube graph, instead of a sparse random sample of the whole graph. The dataset collection is explained in detail in [21].

The resulting dataset contains 12,538 videos. Those videos have $3,091,176$
unique commenters and 27,927,909 comments and replies. From this YouTube dataset, we extract 2 bipartite graphs. The first graph represents the relation 'own', is a directed bipartite graph of user owns video, we define this graph to be the Layer 1. The second graph represents the relation 'comment', is a directed bipartite graph of user comments video, we define this graph to be the Layer 2. YouTube video and comment objects returned by the YouTube API are dated.

In this work, we explore both link prediction tasks, the prediction of the formation of future links, and the missing links predictions in a network. For both tasks, the goal is to predict if a user is to comment on a video. Specifically, we predict links on the YouTube 'comment' network, the Layer 2 in our setting.

## 2 Future Link Prediction

To predict the formation of future links, we randomly selected five different days of the gathered YouTube dataset between the three months of $19 / 7 / 20$ and $22 / 9 / 20$. To evaluate the predictive power of the proposed first and higherorder structural features (table 3.1), for each day of the five days, we construct 3 data sub-sets:

- we extract all data gathered for the day and call it the PredictSet,
- we extract all data gathered for the previous day and call it the $\operatorname{TrainSet}_{1}$,
- we extract all data gathered for the two previous days and call it the TrainSet $_{2}$.

These 3 data sub-sets include Layer 1 and Layer 2 relations, for the 'own' and 'comment' network, respectively.

We use TrainSet $t_{1}$ to generate an embedding using the structural features shown in table 3.1. We then train a Logistic regression model on the embedding and evaluate the predictive power over the PredictSet. We repeat the same procedure for TrainSet ${ }_{2}$.

Table 4.1 shows the structural characteristics of the five sets of data for the task of predicting links in the future. The first column (No.) refers to the dataset id. The second column (Day (2020)) is the date of each dataset. The third column (Datset) is the name of each data sub-set. The fourth and seventh columns $\left(V(G) L_{1}, V(G) L_{2}\right)$ shows the number of nodes in Layer 1 and Layer 2, respectively in the set V . The fifth and eighth columns $\left(U(G) L_{1}\right.$, $U(G) L_{2}$ ), shows the number of nodes in Layer 1 and Layer 2, respectively, in the set U. The sixth and ninth column $\left(E d g e s L_{1}, E d g e s L_{2}\right)$ shows the number
of edges in Layer 1 and Layer 2, respectively. The tenth column Butterflies $L_{2}$ shows the number of butterflies in Layer 2. The eleventh column Blooms $L_{2}$ shows the number of blooms in Layer 2. There are no records in the table for butterflies in Layer 1 because the relation 'own' cannot form this motif in the network, as one video cannot be owned by multiple users. Since there are no butterflies in Layer 1 there are no Bloom structures either.

We train two Logistic Regression models, one embedding containing only the first-order structural features and we call it the 'base model', and one with both first-order and high-order structural features, and call it the 'enhanced model'. Following, we evaluate and compare the predictive power of the two models and found that the enhanced model performs better. Specifically, for the future link prediction task the base model's precision is $97 \%$, and the enhanced model's precision is $99 \%$.

| No. | Day (2020) | Dataset | $\mathrm{V}(\mathrm{G}) L_{1}$ | $\mathrm{U}(\mathrm{G}) L_{1}$ | Edges $L_{1}$ | $\mathrm{V}(\mathrm{G}) L_{2}$ | $\mathrm{U}(\mathrm{G}) L_{2}$ | Edges $L_{2}$ | Butterflies $L_{2}$ | Blooms $L_{2}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | 07/08 | PredictSet | 1,082 | 2,568 | 2,568 | 143,678 | 2,568 | 179,306 | 1,573,999 | 4,144 |
|  | 06/08 | TrainSet ${ }_{1}$ | 1,076 | 2,457 | 2,457 | 141,665 | 2,457 | 178,945 | 1,496,857 | 4,493 |
|  | 05/08 \& 06/08 | TrainSet ${ }_{2}$ | 1,286 | 2,999 | 2,999 | 263,575 | 2,999 | 365,717 | 8,734,962 | 10,439 |
| 2 | 30/08 | PredictSet | 1,316 | 3,500 | 3,500 | 111,391 | 3,500 | 136,837 | 2,097,625 | 2,827 |
|  | 29/08 | TrainSet ${ }_{1}$ | 1,284 | 3,439 | 3,439 | 112,213 | 3,439 | 133,618 | 338,131 | 2,862 |
|  | 28/08 \& 29/08 | TrainSet $_{2}$ | 1,582 | 4,240 | 4,240 | 227,028 | 4,240 | 302,026 | 3,537,029 | 9,199 |
| 3 | 19/09 | PredictSet | 1,451 | 4,005 | 4,005 | 149,229 | 4,005 | 4,005 | 1,929,368 | 5,884 |
|  | 18/09 | TrainSet ${ }_{1}$ | 1,433 | 4,031 | 4,031 | 166,709 | 4,031 | 219,055 | 2725,572 | 6,932 |
|  | 17/09 \& 18/09 | TrainSet ${ }_{2}$ | 1,785 | 5,136 | 5,136 | 286,263 | 5,136 | 416,425 | 10,365,662 | 16,070 |
| 4 | 04/09 | PredictSet | 866 | 501 | 866 | 10,059 | 866 | 10,296 | 35 | 15 |
|  | 03/09 | TrainSet ${ }_{1}$ | 454 | 795 | 795 | 12,655 | 795 | 12,916 | 247 | 24 |
|  | 02/09 \& 03/08 | TrainSet ${ }_{2}$ | 4,315 | 1,523 | 4,315 | 269,916 | 4,315 | 403,965 | 8,107,893 | 13,325 |
| 5 | 24/07 | PredictSet | 899 | 1,796 | 1,796 | 95,570 | 1,796 | 112,827 | 661,254 | 1,803 |
|  | 23/07 | TrainSet ${ }_{1}$ | 866 | 1,725 | 1,725 | 80,839 | 1,725 | 92,214 | 383,554 | 1,188 |
|  | $22 / 07 \& 23 / 07$ | TrainSet $2_{2}$ | 2,141 | 1,080 | 2,141 | 156,443 | 2,141 | 189,557 | 1,637,450 | 3,477 |

Table 4.1: Structural characteristics of the 5 dataset used for the link prediction task.

## 3 Missing Link Prediction

For the missing link task, we select from the 5 random days described above, the data of TrainSet $_{2}$ (table 4.1). We split each of the 5 datasets into $70 \%$ train-set and $30 \%$ predict-set. For both sets we generate an embedding using the structural features shown in table 3.1 and train a Logistic regression model.

Following, we compare the predicting performance of the model trained only with an embedding containing the first-order structural features ('base
model'), to the model ('enhanced model') trained with all the features from table 3.1. We found that the precision of 'base model' is $95 \%$, and the precision of the 'enhanced model' is $99 \%$. Having more information about the YouTube network with the 'enhanced model', is better to find missing link interactions on the YouTube network.

## Chapter 5

## Results

## 1 Future Link Prediction

For the future link prediction problem, we address the following two questions:

- 'How many prior days' (1 or 2) of data are needed to predict if a user is going to comment on a video tomorrow?'
- 'What is the extra predictive value of the higher-order structural features over the first-order structural features?'

Table 5.1 shows the results for the five experiments and their average (Avg) values for Precision, F1 score, and Execution time in seconds, which includes Train and Predict time. When we train with one day's data (TrainSets ${ }_{1}$ ) for the base model, which only includes first-order structural features, we get $97 \%$ precision and the execution time takes 55 seconds. For TrainSets ${ }_{2}$, which have two days' data on the base model we get an average precision of $97 \%$ and the execution time is 59 seconds. When we make predictions with the enhanced model, which includes first and high-order structural features, for the TrainSets ${ }_{1}$ average precision is $99 \%$, and the execution time is around 8 hours. For the TrainSets 2 , we get average precision equal to $99 \%$, and the execution time is around 18 hours.

We found that to make predictions for a day's interactions, suffice to account for network information generated 1 day prior. When making predictions with the enhanced model, which includes the higher-order structures, the average precision is $99 \%$ for the five experiments. It is more time-efficient to make predictions with 1 day's data. With 1 day's data, the prediction is on average $56 \%$ faster than with 2 days' data in the enhanced model. We also found that the higher-order structural features perform $2 \%$ better with $99 \%$
average precision, compared to the first-order features with average precision $97 \%$.

From the above, we infer for the YouTube network the following:

- for more precise future link predictions is better to use the enhanced model (all structural features) with precision $99 \%$,
- the base model (only first-order structural features) can make predictions with $97 \%$ precision, and
- using one additional day's worth of data does not increase precision while adding $128 \%$ of running time when training with two days' data.

| No. | Dataset | Precision | Base model <br> F1 score | Execution Time* | Precision | Enhanced model <br> F1 score | Execution Time* |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | TrainSet $_{1}$ | 0.98 | 0.98 | 58 | 0.99 | 0.99 | 42,963 |
|  | TrainSet $_{2}$ | 0.98 | 0.98 | 60 | 0.99 | 0.99 | 78,061 |
| 2 | TrainSet $_{1}$ | 0.97 | 0.97 | 62 | 0.99 | 0.99 | 30,019 |
|  | TrainSet $_{2}$ | 0.97 | 0.97 | 66 | 0.99 | 0.98 | 60,496 |
| 3 | TrainSet $_{1}$ | 0.98 | 0.99 | 58 | 0.99 | 0.99 | 51,787 |
|  | TrainSet $_{2}$ | 0.98 | 0.98 | 58 | 0.99 | 0.98 | 92,921 |
| 4 | TrainSet $_{1}$ | 0.98 | 0.98 | 45 | 0.98 | 0.98 | 844 |
|  | TrainSet $_{2}$ | 0.98 | 0.98 | 55 | 0.99 | 0.98 | 65,991 |
| 5 | TrainSet $_{1}$ | 0.96 | 0.97 | 54 | 0.99 | 0.99 | 19,788 |
|  | TrainSet $_{2}$ | 0.96 | 0.97 | 58 | 0.99 | 0.99 | 33,959 |
|  |  | Avg Precision | Avg F1 score | Avg Exec Time* | Avg Precision | Avg F1 score | Avg Exec Time* |
|  |  |  |  |  |  |  |  |
|  | TrainSets $_{1}$ | 0.97 | 0.98 | 55 | 0.99 | 0.99 | 29,080 |
|  | TrainSets $_{2}$ | 0.97 | 0.98 | 59 | 0.99 | 66,285 |  |

* Execution Time in seconds

Table 5.1: Future Link prediction results for the 5 dataset, when trained with base and enhanced model.

## 2 Missing Link Prediction

For the missing link prediction problem, we address the questions 'What is the extra predictive value of the higher-order structural features over the first-order structural features?

In this task, we select from the 5 random days described in section 2 , the data of $\mathrm{TrainSet}_{2}$ (table 4.1) and splitted each of the 5 datasets into $70 \%$ trainset and $30 \%$ predict-set. We selected the data of TrainSets 2 because it contains more information about the network to test the higher-order features to the first-order features.

Table 5.2 shows the results for the five experiments and their average (Avg) values for Precision, F1 score, and Execution time in seconds, which includes Train and Predict time. For this task, we only use the dataset TrainSet ${ }_{2}$ that includes 2 days' network information.

In this predicting task, we again compare the base model, which only includes first-order structural features, to the enhanced model, which includes all of the structural features. When predicting with the base model we get an average precision of $95 \%$ and the running time is 35 seconds. With the enhanced model, the average precision is $99 \%$ and the execution time is around 9 hours.

We found for this task on the YouTube comment network on two days' interactions, is better to use the enhanced model, which includes the higherorder structures. The higher-order structural features perform $4 \%$ better than the first-order structural features with an average precision $99 \%$.

From the above, we infer for the YouTube network the following:

- for more precise missing link predictions is better to use the enhanced model (all structural features) with precision $99 \%$,
- using the enhanced model increases precision while adding a $99 \%$ increase of running time.

| No. | Dataset | Precision | Base model <br> F1 score | Execution Time* | Precision | Enhanced model <br> F1 score | Execution Time* |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | TrainSet $_{2}$ | 0.94 | 0.96 | 34 | 0.99 | 0.99 | 59,165 |
| 2 | TrainSet $_{2}$ | 0.96 | 0.97 | 38 | 0.99 | 0.99 | 41,357 |
| 3 | TrainSet $_{2}$ | 0.95 | 0.96 | 36 | 0.99 | 0.99 | 27,399 |
| 4 | TrainSet $_{2}$ | 0.95 | 0.96 | 33 | 0.98 | 0.99 | 24,682 |
| 5 | TrainSet $_{2}$ | 0.95 | 0.97 | 34 | 0.99 | 0.99 | 21,649 |
|  |  | Avg Precision | Average F1 score | Avg Exec Time* | Avg Precision | Avg F1 score | Avg Exec Time* |
|  | TrainSets $_{2}$ | 0.95 | 0.97 | 35 | 0.99 | 0.99 | 34,850 |

* Execution Time in seconds

Table 5.2: Missing Link prediction results for the $5 \mathrm{TrainSet}_{2}$ dataset, when trained with base and enhanced model.

## 3 Set-up \& Tools

For our experiments, we used a cluster of 5 servers with 32 -core $\operatorname{Intel}(\mathrm{R})$ Xeon(R) E5-2630 CPUs and 256 GB of main memory each, configured as 1 Spark Driver and 4 Spark Workers containing 3 Executors each. Each Executor used 83 GB of memory and 10 cores, resulting in 120 total cores. Nodes connect with a 40 Gb network.

## Chapter 6

## Conclusion

In this study, we measure how accurately we can predict comment interactions on the YouTube bipartite network, concerning predicting missing or unseen interactions. We propose high-order structural bipartite features, which enhance the performance of simple prediction models, to find missing or unseen links, which have never been used prior in the literature. Through experimentation on multi-layer networks, we find that we can predict future and unseen comment interactions on YouTube videos with precision $99 \%$. We show that to predict a day's comment interactions it suffices to account network information generated 1 day prior. Finally, we implement 2 MapReduce algorithms, one that counts the bitruss number of an edge, and one that identifies blooms in bipartite networks.

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[^0]:    This work has been performed at the University of Crete, School of Sciences and Engineering, Computer Science Department.

    The work has been supported by the Greek GSRT through the project ETAK, with project ID T1EDK-01800

