

Friendship prediction and homophily in social media

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Social media have attracted considerable attention because their open-ended nature allows users to create lightweight semantic scaffolding to organize and share content. To date, the interplay of the social and topical components of social media has been only partially explored. Here we study the presence of homophily in three systems that combine tagging of social media with online social networks. We find a substantial level of topical similarity among users who lie close to each other in the social network. We introduce a null model that preserves user activity while removing local correlations, allowing us to disentangle the actual local similarity between users from statistical effects due to the assortative mixing of user activity and centrality in the social network. This analysis suggests that users with similar interests are more likely to be friends, and therefore topical similarity measures among users based solely on their annotation metadata should be predictive of social links. We test this hypothesis on several datasets, confirming that social networks constructed from topical similarity capture actual friendship accurately. When combined with topological features, topical similarity achieves a link prediction accuracy of about 92%.

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The present paper is an extended version of a previously published conference paper [Schifanella et al. 2010].

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1. INTRODUCTION

Online social networking systems, together with online systems for content organization and sharing, entangle cognitive, behavioral, and social aspects of a user community through an underlying technological platform. The resulting “ecosystems” provide new possibilities to mine and investigate the various processes at play in the interactions of individuals, and to study the ways in which users relate with the information they share.

Key open questions deal with understanding the concepts of similarity and influence, tracking the emergence of shared semantics, and determining the interplay between social proximity and shared topical interests among users. The emergence, spreading, and stability of any shared concept depend critically on these factors.

Social media supporting tagging are especially relevant to a data-driven investigation of these questions, because they stimulate users to provide light-weight semantic annotations in the form of freely chosen terms [Golder and Huberman 2006]. Social annotations based on tags are valuable for research because they externalize the three-way relation between users, items of interest (resources), and metadata (tags). Usage patterns of tags can be employed to monitor interests, track user attention, and investigate the emergence and spread of shared concepts through a user community. Moreover, several social media platforms support explicit representations of social links between users, making an objective definition of social proximity available. They also combine several aspects of user activity, such as exposing resources, tagging items, belonging to discussion groups, and relating to other users.

In this paper, we consider three different online social systems: Flickr, Last.fm, and aNobii. In these systems, users share resources (pictures, songs/artists, and books, respectively), form social networks, and tag items to produce social classifications of data commonly called *folksonomies*.

The three systems differ significantly by size, category of resources, and the precise ways in which users tag resources and relate to each other. Taking advantage of datasets built from the three systems, we address the following question: *How does the similarity between user profiles relate to their proximity on the social network?* More precisely, are neighboring users more similar, both in the amount of activity they devote to the system, and in the content of their activity, than users who lie further apart in the social network? If so, how does this local similarity fade when the distance on the social graph increases? And can we predict the existence of social links from knowledge of the similarity among user profiles?

In the remainder of this paper, after an overview of related work in Section 2 and of our datasets in Section 3, we provide in Section 4 a thorough analysis along several axes. This data analysis highlights the heterogeneity of user activities and the correlations in the various metrics measuring the different activities of a single user. We also show the existence of non-trivial mixing patterns: the amounts of different user activities are correlated across neighbours. Section 4 exposes the

substantial levels of topical similarity that exist among users who are close to each other in the social network. In Section 5, we evaluate the performance of predictors of online social links based on the similarity of user profiles. We consider a number of topical similarity measures from the literature. Scalable similarity measures, such as Maximum Information Path, proposed by some of the authors, are among those achieving the best predictive performance. The role of language communities in these predictions is investigated in Section 6.

2. RELATED WORK

In prior work we explored the correlation between proximity in an online social network and topical similarity and we analyzed the extent to which similarity among users based only on tagging information can be an accurate predictor of social ties [Schifanella et al. 2010]. We analyzed samples from the Flickr and Last.fm social networks. The present paper expands on our prior work both in breadth and in depth.

Similarity between the members of social groups, or between individuals sharing a social link, is known as homophily in the social networks literature, and has long been observed and studied [McPherson et al. 2001]. Homophilic phenomena can be present due to selection mechanisms or social influence, two effects that are often confounded and difficult to disentangle [Leenders 1997; Aral et al. 2009; Shalizi and Thomas 2010]. The increasing availability of data from online social networks has created ideal laboratories for testing and quantifying such social phenomena and theories (see, e.g., Crandall et al. [2008], Aiello et al. [2010], Szell et al. [2010], and Leskovec et al. [2010]).

We find in the literature several studies on the evolution of online social systems and correlations between different user features. Leskovec and Horvitz [2008] present a study on the Microsoft Messenger network, showing correlation between user profile information and communication patterns. Evolutionary patterns in the Flickr social network have been studied by Kumar et al. [2006] and Mislove et al. [2007; 2008]. Marlow et al. [2006] perform a quantitative study on tag usage in Flickr. They discuss the heterogeneity of tagging patterns and perform a preliminary analysis of vocabulary overlap between pairs of users. Their analysis shows that neighbors in the social graph have a higher vocabulary overlap, on average. However, no assessment is made of biases that could be responsible for the reported observation: here we explore such biases. The role of groups as coordination tools in Flickr is investigated by Prieur et al. [2008]. They point out a strict relation between the density of the social network and the density of the network of tag co-usage among group members. Leskovec et al. [2008] perform a comparative study on the microscopic evolutionary dynamics between several social networks, in which a special emphasis is placed on the arrival process of new nodes and on the dynamics of attachment. Influence of social contacts on browsing patterns in Flickr has been analyzed by Lerman and Jones [2007] and van Zwol [2007], who provide insights into the activity patterns of users. Correlation between topical overlap among user interests in tagging systems and other indicators of social behavior is explored by Santos-Neto et al. [2009]. They consider CiteULike and Connotea systems, which both lack an explicit social network component, so they look at collaboration rela-

tions determined by the participation in the same discussion group.

Predicting the presence of a link between two entities in a network is one of the major challenges in link mining [Getoor and Diehl 2005]. A common approach to the link prediction problem is to infer ties from the structural properties of the social network. Liben-Nowell and Kleinberg [2003] discuss several notions of node similarity based on social graph structural features for link prediction. Prediction of future links in a question-answering bulletin board service is performed by Murata and Moriyasu [2007], showing network proximity scores calculated from local topological information to be accurate predictors of future links. Huan [2006] defines a cycle formation model for social graphs that relates the probability of the presence of a link with its ability to form cycles. The parameters of the model are estimated using the generalized clustering coefficients of the network. The power of the model is evaluated on the Enron email dataset. Another probabilistic network evolution model aimed at link prediction is proposed by Kashima and Abe [2006]. The idea is that links appear in the network due to a copying process where status labels associated to edges are copied from one node to another with a probability that is dependent on the relative topological position of the two nodes. Clauset et al. [2008] present a hierarchical decomposition algorithm for network clustering which can also be applied to predict missing interactions in networks. The generated dendrograms determine the probability of connection for every pair of vertices. Links are predicted between pairs that have high probability of connection within the hierarchical random graphs but that are unconnected in the observed network. This technique yields positive results on several small networks.

Another line of research focuses on link detection through supervised learning methods trained on the topological features of graphs [Popescul et al. 2003; Hasan et al. 2006]. Prediction of the sign of an existing link in friends-foes social networks (e.g., Slashdot Zoo) is presented by Leskovec et al. [2010], who use a logistic regression classifier trained with two classes of topological features: node degree and triadic closure.

Link prediction can also be based on features that describe user profiles, based on the principle that people with similar tastes are more likely to establish social contacts. Caragea et al. [2009] propose an ontology-based classification of user features and show that the semantics captured by the ontology can effectively improve the performance of a topology-based machine learning classifier for social link prediction. Li et al. [2008] cluster Delicious users extracting implicit relations based on the similarity of their tag vocabulary. Leroy et al. [2010] leverage group membership information from Flickr to build a probabilistic graph useful to detect the hidden social graph. Mislove et al. [2010] explore the complementary question: can we predict topical similarity from the social network? Again, here we discuss the role of global correlation in biasing such prediction.

Even if the majority of papers is focused on link prediction on simple (directed or undirected) graphs, techniques have also been developed for different kinds of networks like weighted networks [Lü and Zhou 2009], bipartite networks [Dunlavy et al. 2010; Benchettara et al. 2010; Kunegis et al. 2010], and signed social graphs [Leskovec et al. 2010]. Finally, some approaches based on probabilistic models such as relational Markov networks [Taskar et al. 2003] and probabilistic relational

models [Getoor et al. 2003] deserve to be cited. However, these approaches have not been extensively tested on real-world datasets.

In a survey of link prediction techniques, Lü and Zhou [2010] compare several structural similarity metrics in terms of accuracy and computational efficiency. In our own previous work [Markines et al. 2008; Markines et al. 2009; Markines and Menczer 2009], we made a systematic analysis of a broad range of semantic similarity measures that can be applied to the three-dimensional folksonomy space to extract similarity networks of tags, resources, or users. Here, we use such measures to perform link prediction based on the folksonomy information.

3. DATASETS

In the following, we report on the main features of our datasets and we describe the data retrieval methods we used to build them. For each dataset, we collected at least the information about the social network, the tag assignments, and the group affiliations. Table I summarizes the sizes of our three dataset.

Table I. Dataset statistics					
Dataset	Users	Triples	Tags	Tagged items	Groups
Flickr	130,840	90,723,412	1,420,656	20,599,583	92,301
Last.fm	90,049	6,971,166	194,763	894,615	69,306
aNobii	86,800	5,378,190	143,182	918,181	3,581

3.1 Flickr

We collected the tagging information about the pictures uploaded in Flickr between January 2004 and January 2006 by means of API methods (`flickr.com/api`). The crawling effort was distributed by splitting the above time interval into smaller time windows to be crawled independently. A global tag knowledge base, initialized with a minimal set, was shared between parallel crawlers. Crawlers issued search queries limited to their specific time interval to retrieve information about photos marked with the tags stored in the common database. New tags were added to the shared database as they were discovered by individual crawlers.

Separate crawls were made to explore group affiliations and the social network. In Flickr jargon, social ties are called *contacts*; they are *directed* and do not require acceptance by the linked user. The overall crawl was performed during the first half of 2007.

Our analysis will focus on the network of about 130 thousand users for whom we have tag, group, and contact information.

3.2 Last.fm

In Last.fm, each user is linked to *friends* through undirected links that are established given the consent of both endpoints. Users also have a public list of *neighbors*, computed by the system as recommendations for potential new friendship contacts. An affinity value, ranging from 0 to 1, is also assigned to each member of the neighbor set. Users can annotate songs, artists or albums with tags, and can create or join groups. Users also have a public *library*, i.e., a list of the artists they have

listened to. User profile information includes an optional geographic specification at the country level.

We used both API calls (`last.fm/api`) and web crawling methods to build the dataset. The API can be used to retrieve user profiles, friendships and neighborhood relationships and a list of the 50 top artists in the user library (i.e., those with the highest playcount). The API does not allow for the collection of a user's complete activity and group affiliation information, so we extracted the $(user, item, tag)$ triples and the group membership relations via web crawling and scraping. The user set we consider was selected by a BFS crawl of the friendship network. The crawls took place in January 2010. We started from three randomly chosen users and for each of them we performed a crawl up to those nodes that resided 4 hops away. The corresponding snapshots include approximatively 100 thousand users each, with an overlap of about 20% between them. Since we found that the results of our analysis are consistent across the three samples, we report the findings for a single representative one.

Recently, the Last.fm API was extended with a similarity function, called *tasteometer*, which, given in input a pair of users or artists, returns an affinity score ranging from 0 to 1. This value is different from the one provided by the neighborhood score and, most of all, it can be computed for *any* pair of users or artists. Jointly with the crawling activity, we retrieved the tasteometer values for a large set of user pairs to compare the performance of our tag-based similarity functions in the link prediction task with the performance of the tasteometer. Further details are given in Section 5.

3.3 aNobii

Users in aNobii (`anobii.com`) fill their digital book collections with titles selected from the public aNobii book database, which contains the metadata (such as publication year, authors etc.) of about 20 millions different publications, written in 49 different languages. Each personal book collection is partitioned into a *library*, which is a set of titles that the user is reading or has already read, and a *wishlist* that lists the books that the user wants to read in the near future. Books in the user collection can be annotated with arbitrary tags. Since books in libraries form the vast majority of the overall book collections, here we focus mainly on books from libraries.

Users can also provide public information about their profile, such as gender, age, marital status, and a detailed specification of their geographic location including country and hometown. Affiliation with thematic, user-generated groups is also possible.

Two different types of social ties can be established between users: *friendship* and *neighborhood*. The aNobii website suggests that people should be friends if they know each other in real life. Users should establish neighborhood ties with people who have a library they consider interesting. Surprisingly, although these two types of social links are formally different, they are equivalent from a structural point of view. In fact, both are *directed* and can be created without any consent of the linked user, who is not even notified when a new incoming tie is established. Furthermore, both links activate a monitoring on the linked user's library that triggers notifications on library updates. Given this strong structural similarity,

and since the two types of links are *mutually exclusive*, in the following we deal with the *union* between friendship and neighborhood networks and we generically refer to the union network as the aNobii social network.

We crawled the aNobii network in December 2009 starting from a random seed of users and following the social links in a forward BFS fashion. We explored the entire giant strongly connected component and the out component of the social network, for a total of 86,800 users. We collected each user’s profile information, group affiliations, library, and tag assignments through web scraping.

4. DATA ANALYSIS

In most folksonomies, the activity of users has many facets. In Flickr, for instance, users can upload pictures and tag them, participate in groups, and comment on photos. In Last.fm, users can listen to music and tag songs according to their characteristics and personal tastes. In aNobii, users can add books to their libraries, tag them, join groups, and create a list of books they wish to read.

Since social networks are explicitly built by users, we can also consider the number of friendship relations to be a measure of activity in each folksonomy we consider. When links are directed, we consider out-degree as a proxy for activity, while in-degree measures popularity.

In this section, we first analyze the activity patterns of individual users, and show their considerable heterogeneity. We also investigate the correlations between various activity indicators.

4.1 Heterogeneity and Correlations

Figures 1 and 2 show the distributions of the number of neighbors in the social network and the probabilities of finding a user with a given number n_t of distinct tags in her vocabulary, a total tagging activity a , belonging to n_g groups, and having (for aNobii) n_b and n_w books in her library and wishlist, respectively.

All these distributions are broad, spanning multiple orders of magnitude, showing that the activity patterns of users are highly heterogeneous. For each activity measure most users have little activity while some are extremely active, and all intermediate values are represented. No characteristic or “typical” value of the activity can be sensibly defined as evident from a standard deviation that is orders of magnitude larger than the average, for each activity measure.

Given this high level of disparity between users, a natural question arises about the correlations between the different types of activity: do users who have many neighbors also use many tags, belong to many groups, and so on? The simplest way to examine this issue is to compute the average activity of a type for users having a certain value of another activity type. For instance, we can measure the average number of distinct tags for users having k neighbors in the social network:

$$\langle n_t(k) \rangle = \frac{1}{|u : k_u = k|} \sum_{u: k_u = k} n_t^u, \quad (1)$$

where n_t^u is the number of distinct tags of user u . As shown in Fig. 3, all types of activity have a clearly increasing trend for increasing values of the out-degree; users who have more contacts in the social network tend also to be more active in terms

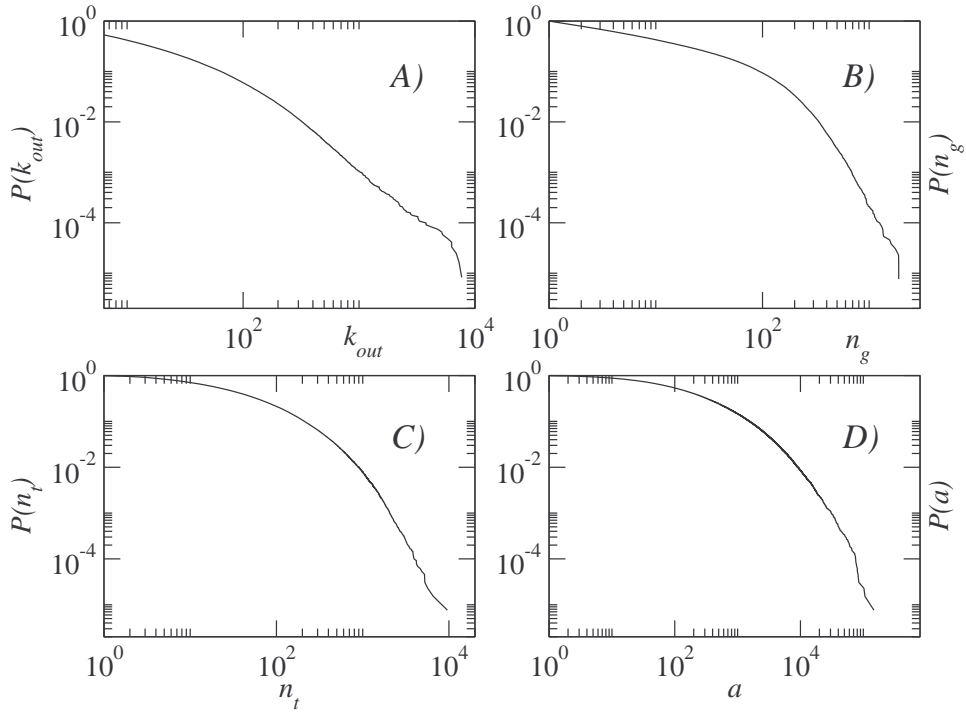


Fig. 1. Flickr complementary cumulative distributions of (A) the number k_{out} of neighbors of a user, (B) the number n_g of groups of which a user is a member, (C) the number n_t of distinct tags per user, and (D) the number a of tag assignments per user.

of tags and groups. Overall, the various activity metrics are all positively correlated with one another. For instance, in Flickr, the Pearson correlation coefficients are: 0.349 between k and n_t , 0.482 between k and n_g , 0.268 between k and a , 0.429 between n_t and n_g , 0.753 between n_t and a , and 0.304 between n_g and a .

Despite these correlations, large fluctuations are still present. First, the strong fluctuations at large degree values are due to the smaller number of highly-connected nodes over which the average is performed. Notably, users with a large number of social contacts but using very few tags and belonging to very few groups can be observed. We can investigate in more detail the degree of correlation between activity types through the conditional probabilities of the type $P(n_t|k)$, i.e., the probability for a user to have n_t tags, knowing that she has k neighbors in the social network, where the average $\langle n_t(k) \rangle$ is simply the first moment of this conditional distribution. As shown for some examples in Fig. 4, these distributions, although narrower than the distributions shown in Figures 1 and 2, remain broad. This shows that, despite the strong correlations observed, users with a given activity level in the social network remain quite heterogeneous.

4.2 Mixing Patterns

While the previous analysis concerns the correlations between the diverse activity levels of a single user, another important question concerns the correlations between

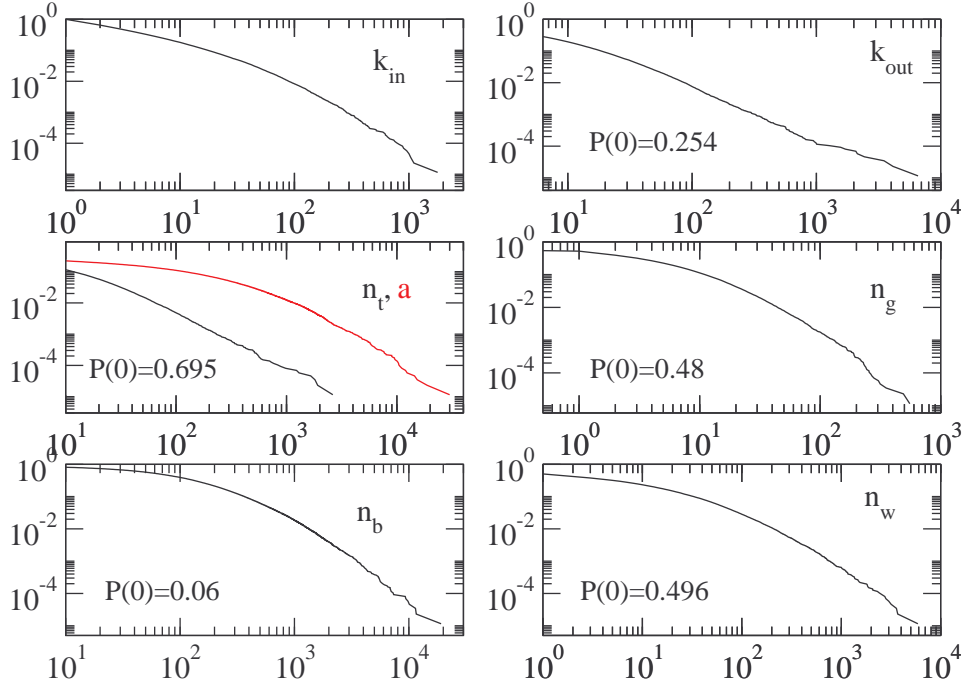


Fig. 2. Complementary cumulative distributions of the measures of activity of aNobii users: in-degree k_{in} and out-degree k_{out} in the social network, number of distinct tags n_t and tagging activity a (total number of annotations by a user), number of group memberships n_g , number of books in a user library n_b and in a user wishlist n_w .

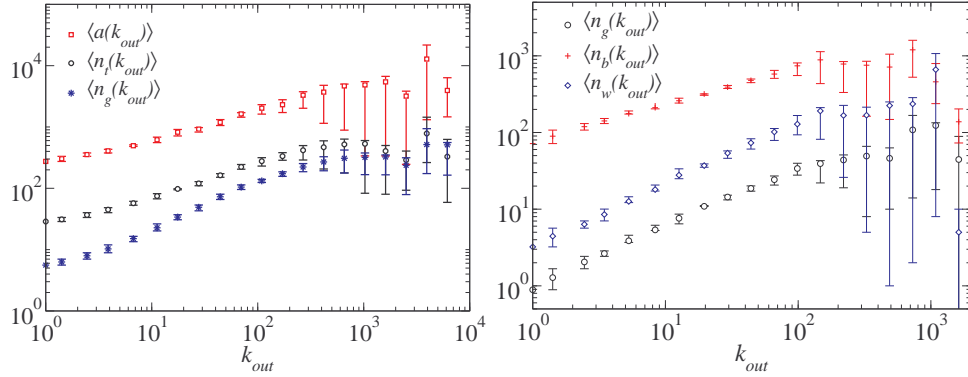


Fig. 3. Left: Average number of distinct tags ($\langle n_t \rangle$), of groups ($\langle n_g \rangle$), and of tag assignments ($\langle a \rangle$) of users having k_{out} out-neighbors in the Flickr social network. Right: Correlations between the activity of aNobii users and their number of declared friends and neighbors: group memberships $\langle n_g \rangle$, library $\langle n_b \rangle$ and wishlist sizes $\langle n_w \rangle$, averaged over users with k_{out} out-links, vs k_{out} . The data has been log-binned: the symbols indicate the average, and the errorbars the 25 and 75 percentiles for each bin.

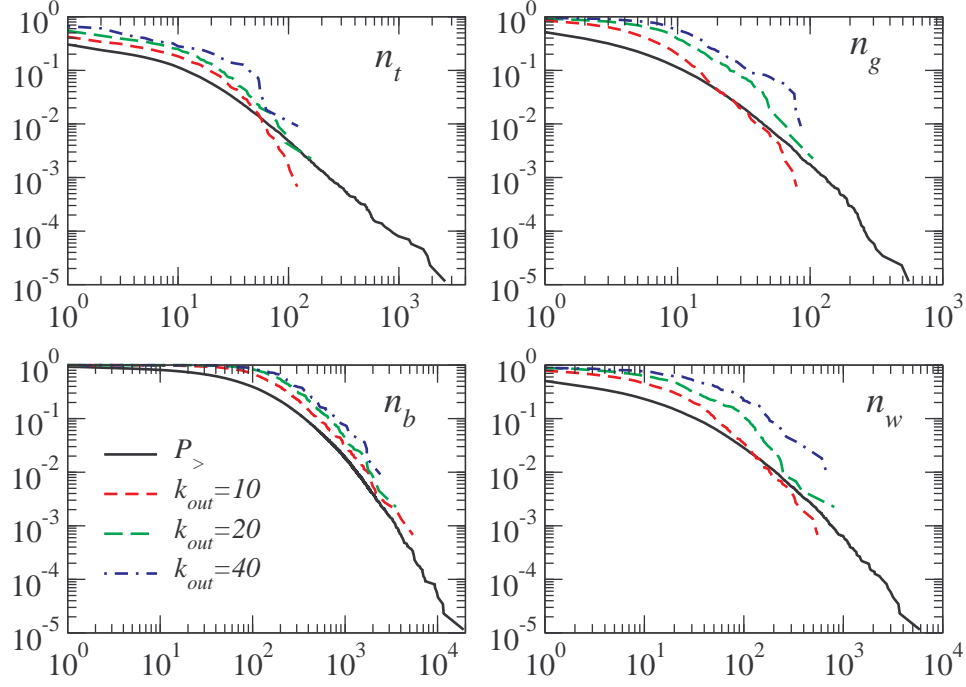


Fig. 4. Complementary cumulative conditional distributions $P(n|k_{out})$ in aNobii, where n is the number of tags n_t , of groups n_g , of books n_b , and wishlist size n_w , compared with the global cumulative distributions $P_>$ (black lines). Even among the subset of users with a given k_{out} , a strong disparity is still observed in the amount of activity.

the activity metrics of users who are linked in the social network. This is a long-standing problem in the social sciences, ecology and epidemiology: a typical pattern, referred to as “assortative mixing,” describes the tendency of nodes of a network (here, the users), to be linked to other nodes with similar properties [Newman 2003; 2002; Newman and Park 2003]. This property is also commonly called “homophily” [McPherson et al. 2001].

Mixing patterns can be defined with respect to any property of the nodes. In the case of social media, since each user is endowed with several properties characterizing his activity and social connectivity, it is interesting to characterize mixing patterns with respect to each of these properties.

One of the most investigated mixing pattern involves the degree of nodes. A commonly used measure is the average nearest neighbors degree of a user u ,

$$k_{nn}^u = \frac{1}{k_u} \sum_{v \in \mathcal{V}(u)} k_v, \quad (2)$$

where the sum runs over the set $\mathcal{V}(u)$ of neighbors of u . To characterize mixing patterns with respect to node degree, a convenient measure can be built on top of k_{nn}^u by averaging over all nodes u that have a given degree k [Pastor-Satorras et al.

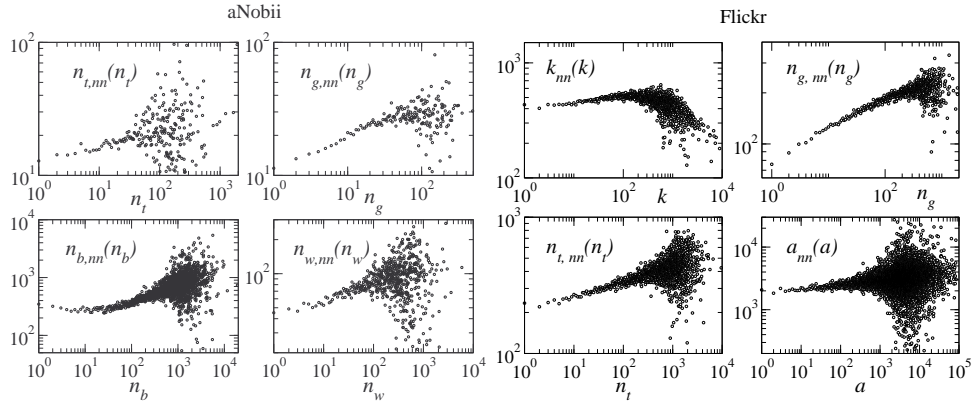


Fig. 5. Mixing patterns in the aNobii and Flickr dataset: average number $n_{t,nn}(n_t)$ of distinct tags of the nearest neighbors of users having n_t distinct tags; average number $n_{g,nn}(n_g)$ of groups of the nearest neighbors of users belonging to n_g groups; average number $n_{b,nn}(n_b)$ of books of the nearest neighbors of users who have read n_b books; average wishlist size $n_{w,nn}(n_w)$ of the nearest neighbors of users who have a wishlist of size n_w ; average out-degree $k_{nn}(k)$ of the nearest neighbors of users having out-degree k ; and average number $a_{nn}(a)$ of distinct triples of the nearest neighbors of users having a distinct triples. The quantitative differences between the various mixing patterns are not relevant to the present discussion, therefore we omit a more detailed analysis here.

2001; Vázquez et al. 2002]:

$$k_{nn}(k) = \frac{1}{|u : k_u = k|} \sum_{u: k_u = k} k_{nn}^u. \quad (3)$$

We can generalize the average nearest neighbors degree presented above for the case of any profile feature. We define the average number of tags of u 's nearest neighbors $n_{t,nn}^u = \frac{1}{k_u} \sum_{v \in \mathcal{V}(u)} n_t^v$, and, similarly, the average number of tag assignments used by her nearest neighbors, $a_{nn}^u = \frac{1}{k_u} \sum_{v \in \mathcal{V}(u)} a^v$, the average number of groups to which her nearest neighbors participate, $n_{g,nn}^u = \frac{1}{k_u} \sum_{v \in \mathcal{V}(u)} n_g^v$, and, in the case of the aNobii dataset, the average number of books read by her nearest neighbors, $n_{b,nn}^u = \frac{1}{k_u} \sum_{v \in \mathcal{V}(u)} n_b^v$ and the average wishlist size of her nearest neighbors, $n_{w,nn}^u = \frac{1}{k_u} \sum_{v \in \mathcal{V}(u)} n_w^v$.

By analogy with the case of $k_{nn}(k)$, we can compute the average number of distinct tags of the nearest neighbors *for the class of users having n distinct tags*,

$$n_{t,nn}(n) = \frac{1}{|u : n_t(u) = n|} \sum_{u: n_t(u) = n} n_{t,nn}^u, \quad (4)$$

and the average number of tag assignments used by the nearest neighbors *for the class of users with a tag assignments*,

$$a_{nn}(a) = \frac{1}{|u : a(u) = a|} \sum_{u: a(u) = a} a_{nn}^u. \quad (5)$$

Analogous definitions can be used with respect to all other profile features.

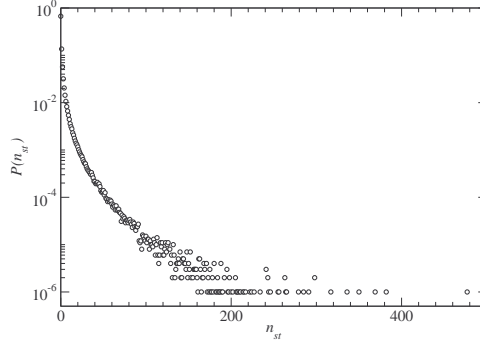


Fig. 6. Probability distribution of the number of shared tags for two randomly chosen Flickr users. The probability to have no tags in common is $P(0) \approx 0.67$, but the overall distribution is broad.

Fig. 5 shows clear assortative trends for several measures for both the aNobii and Flickr datasets, as in other social networks [Newman and Park 2003; McPherson et al. 2001]. Similar results are obtained for Last.fm (not shown). As before, large fluctuations are observed for large activity values, because of the small number of very active users. But we see for all activity measures that the average activity of the neighbors of a user increases with the user's own activity — the activities of socially connected users are correlated at all levels.

4.3 Topical Similarity

The previous analysis has focused on the amount of user activity, as quantified by several metrics, and on the corresponding correlations and mixing patterns. To understand the interplay between the social network and user activities, it is necessary to take into account not only the amount, but also the nature and content of the user activities. To compare users in detail, we therefore focus here on the *topical similarity* between user profiles as measured by the shared features — tags, groups, books, songs, and so on — in their profiles.

A first natural question regards the possible existence of some amount of *global* similarity between the users of a given folksonomy. For instance, in the context of tags, a simple test for the existence of a globally-shared vocabulary can be performed by selecting pairs of users at random and measuring the number of tags they share, n_{st} .

In the case of Flickr, this measure shows that there is actually no shared tag vocabulary; this is not very surprising, given that Flickr is a narrow folksonomy (see Section 5) and the broad range of interests of the users. The average number of shared tags is only about 1.6 in Flickr, and the most probable case is the absence of any tags shared by the selected users (Fig. 6).

Despite the lack of a globally shared profile, a number of mechanisms may however lead to *local* similarity of user profiles, in terms of shared tags, groups membership, books, musical tastes, and so on [McPherson et al. 2001]. The presence of a social link suggests some degree of shared context between the connected users, who are likely to have some interests in common, or to share some experiences, and who are moreover exposed to each other's content and annotations. As an ex-

Table II. Tags most frequently used by three Flickr users; A and B are friends.

User A	User B	User C
green	flower	japan
red	green	tokyo
catchycolors	kitchen	architecture
flower	red	bw
blue	blue	setagaya
yellow	white	reject
catchcolors	fave	sunset
travel	detail	subway
london	closeupfilter	steel
pink	metal	geometry
orange	yellow	foundart
macro	zoo	canvas

ample, Table II shows the 12 most frequently used tags for three Flickr users with comparable tagging activity. User A and user B have marked each other as friends, while user C has no connections to either A or B on the Flickr social network. All of these users have globally popular tags in their tag vocabulary. In this example, the neighbors A and B share various interests (expressed by the tags in bold).

As often discussed in the social sciences, homophily can emerge for different reasons, which are summarized in two scenarios: link selection and social influence [McPherson et al. 2001; Leenders 1997; Shalizi and Thomas 2010]. The former scenario considers that social links are preferentially created between individuals who are already similar and choose each other for establishing the social link precisely because they share some degree of similarity. In the latter scenario, individuals become more similar over time because they influence each other. Disentangling these scenarios is a delicate matter that requires longitudinal data sets, as social influence implies a temporal evolution of a relationship [Crandall et al. 2008; Aral et al. 2009; Aiello et al. 2010; Shalizi and Thomas 2010]. Regardless of the distinction between these possible mechanisms driving homophily, it is important to understand how to measure this effect, and how to relate it to the social network structure, in particular with the distance between users in the social network.

From this perspective, it is necessary to define robust measures of profile similarity between two users u and v , regarding the various types of activity. The first and simplest measure is given by the number of shared items for each activity: the number of shared tags n_{st} of the tag vocabularies of u and v , the number of shared groups n_{sg} to which both u and v belong, the number of common books in their libraries or wishlists for aNobii, and the number of common songs for Last.fm. These measures may however be affected by the amounts of activity of the users; two users who apply many tags may have more tags in common than two less active users, as it is more probable to find common items in two long lists than in two short ones. Therefore we need to normalize profile similarity measures to compensate for heterogeneous activity.

Let us first consider the case of the tags. Following Cattuto et al. [2008] we regard the vocabulary of a user u as a *feature vector* W whose elements correspond to tags and whose entries are the tag frequencies for that specific user's vocabulary, i.e., w_{ut} is the number of resources tagged with t by u . To compare the tag feature

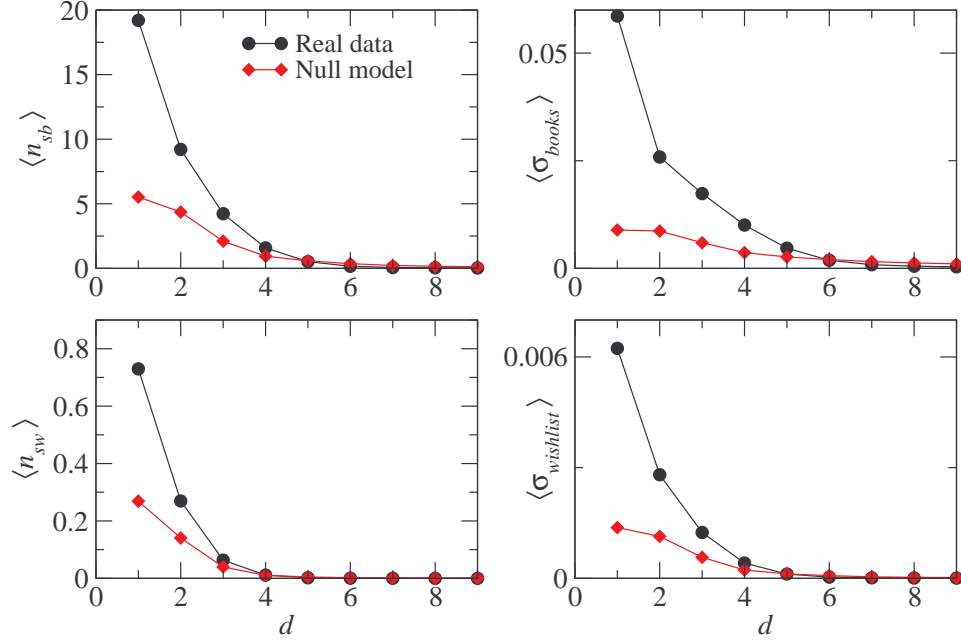


Fig. 7. Average library and wishlist similarity as a function of the distance on the aNobii social network. Top: average number of shared books in the libraries of users at distance d , and corresponding cosine similarity. Bottom: average number of shared items in the wishlists of users at distance d , and corresponding cosine similarity. The diamonds correspond to the null model discussed at the end of Section 4.3.

vectors of two users, we use the standard cosine similarity [Salton 1989] defined as

$$\sigma_{tags}(u, v) = \frac{\sum_t w_{ut} w_{vt}}{\sqrt{\sum_t w_{ut}^2} \sqrt{\sum_t w_{vt}^2}}. \quad (6)$$

This quantity is 0 if u and v have no shared tags, and 1 if they have used exactly the same tags, in the same relative proportions. Because of the normalization factors in the denominator, $\sigma_{tags}(u, v)$ is not directly affected by user activity.

Similarly, we can define the cosine similarities for group memberships and for books. Since a user belongs at most once to one group, and adds a book only once to her library, the elements of the group and book vectors are binary, and the similarities reduce to

$$\sigma_{groups}(u, v) = \frac{\sum_g w_{ug} w_{vg}}{\sqrt{n_g(u)} \sqrt{n_g(v)}}; \quad \sigma_{books}(u, v) = \frac{\sum_b w_{ub} w_{vb}}{\sqrt{n_b(u)} \sqrt{n_b(v)}}, \quad (7)$$

where w_{ug} is 1 if u belongs to group g and 0 otherwise, and w_{ub} is 1 if u has book b in her library and 0 otherwise.

Figures 7 and 8 give an indication of how the similarity between users depends on their shortest path distance d on the social network, by showing the average similarity of two users as a function of d . In aNobii (Fig. 7), the average number of shared books is rather large for neighbors (close to 20), but it drops rapidly as d increases, and is close to 0 for $d \geq 4$. Similar results are obtained for the number

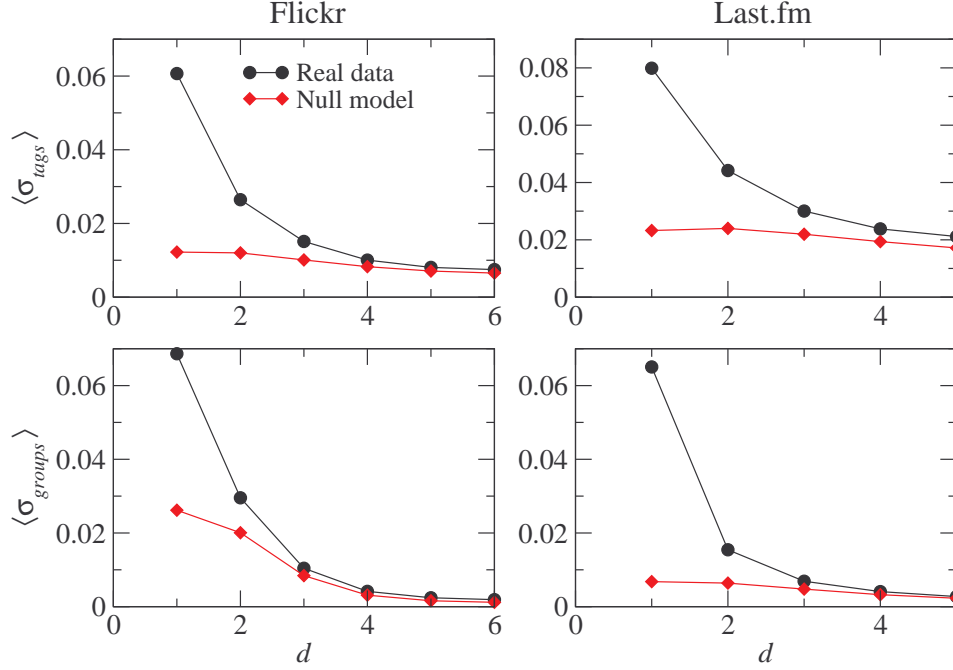


Fig. 8. Average tag and group similarity as a function of the distance on the Flickr and Last.fm social networks. Top: average cosine similarity between the tag vocabularies. Bottom: average cosine similarity between the groups participation vectors. The diamonds correspond to the null model discussed at the end of Section 4.3.

of common groups and tags, and hold for Last.fm and Flickr as well (Fig. 8). The cosine similarities display the same decreasing trend as the distance along the social network increases.

The shortest path distance between two users gives the minimum number of steps to navigate from one user to the other on the online social network. However, this measure of topological proximity between users can be sensitive to the addition or removal of a single link, and does not take into account the fact that more than one path can connect the users. To overcome this issue, the personalized PageRank [Haveliwala 2003] of one user v with respect to another user u can be considered. This algorithm essentially gives the probability, for a random walker starting from the profile page of user u , to visit the profile page of v , and therefore is a more robust measure of social network proximity. As shown in Fig. 9, the topical similarity between users increases with their social proximity, which is consistent with the trend of Fig. 7.

To gain more insight into the entanglement between similarities and distance on the social network, we present in Figures 10 and 11 the probability distributions of the selected similarity measures for pairs of users at social distance d . The figures clearly shows the dependence of all distributions upon the distance of the users along the social network: for users who lie at small distances on the social network, rather broad distributions are observed for the number of shared tags, groups, or books.

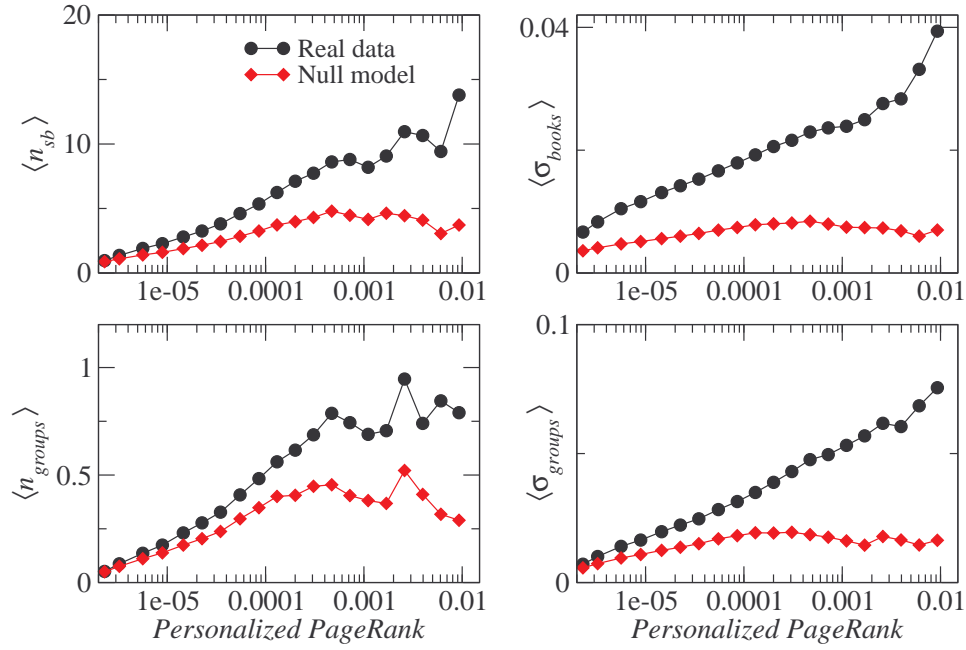


Fig. 9. Average library (top) and group (bottom) similarity between two aNobii users as a function of the personalized PageRank of one user with respect to the other user. The diamonds correspond to the null model discussed at the end of Section 4.3.

As the distance d along the network increases, the distributions become narrower. Two comments are in order: first, the distributions of n_{st} at short distances reach much larger values of n_{st} than in Fig. 6 (the same is observed for the number of shared groups or books). The reason is that, when choosing a random pair of nodes (as in Fig. 6), one is unlikely to select two neighboring nodes. Second, at any distance, the most probable value of n_{sg} or n_{st} is 0, even if the distributions are broad, and this probability increases with d . For instance, for Flickr users, the probability that two users do not share any tag is $P(n_{st} = 0) = 0.1$ if the users are neighbors ($d = 1$), 0.17 if they are at distance $d = 2$, 0.37 at $d = 3$. For groups, $P(n_{sg} = 0) = 0.17$ at $d = 1$, 0.4 at $d = 2$, 0.74 at $d = 3$. The distributions of cosine similarities between users at distance d show similar features: they mostly span the whole interval of possible values, but the probability of high similarity becomes smaller when d increases. The trends are similar for Last.fm (not shown).

The presence of assortative mixing patterns in the social network, with respect to the intensity of users activity, makes it necessary to investigate in more detail the observed local similarity of profiles. It could indeed be the case that such assortativity, by a purely statistical effect, yields an *apparent* local similarity between the tag vocabularies of users. For example, even in a hypothetical case of purely random tag assignments, it would seem more probable to find tags in common between two large tag vocabularies than between a small one and a large one. Furthermore, as we have shown, users who are more active have more friends, and their friends are also more active, therefore similarity with their friends may depend on their

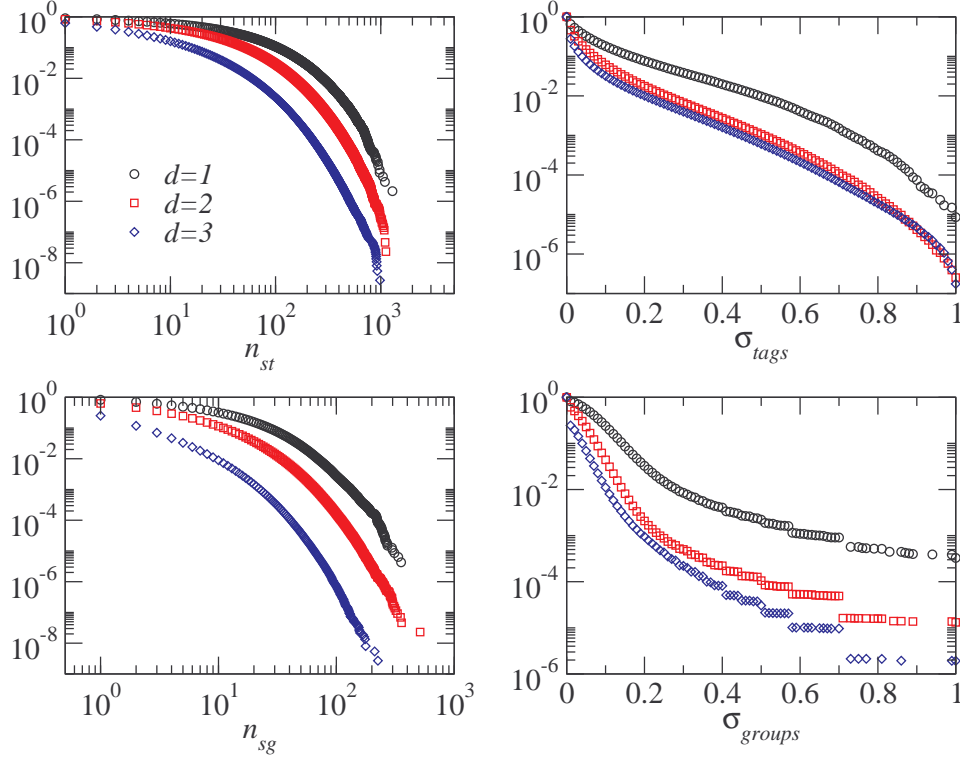


Fig. 10. Left: Complementary cumulative probability distributions of the number of shared tags and groups for two Flickr users lying at distance d on the social network, for different values of d . Right: Complementary cumulative probability distributions of the cosine similarity between the tag vocabularies and group memberships of two Flickr users.

greater activity alone.

To discriminate between effects simply due to the assortativity and those due to actual profile similarity, one has to construct a proper *null model*, i.e., an artificial system that retains the same social structure as the one under study, but lacks any feature similarity other than the one that may result from purely statistical effects. This is done by keeping fixed the social network and its assortativity pattern for the intensity of the activity, but destroying socially-related feature similarity by means of a random permutation of profile items. For instance, we proceed in the following fashion for the tags: (i) we keep the social network unchanged, preserving each user's degree k ; (ii) we shuffle the tags among users in such a way as to preserve each user's number of tag assignments a as well as number of distinct tags n_t . This guarantees that the distribution of frequencies of tags is left unchanged. Such a procedure is in the spirit of null models for detecting the importance of patterns in networks [Maslov et al. 2004] and of random models of networks with given degree distributions or correlation patterns [Molloy and Reed 1995; Catanzaro et al. 2005; Serrano and Boguñá 2005].

Using the null model defined above, we measure the similarity between users at

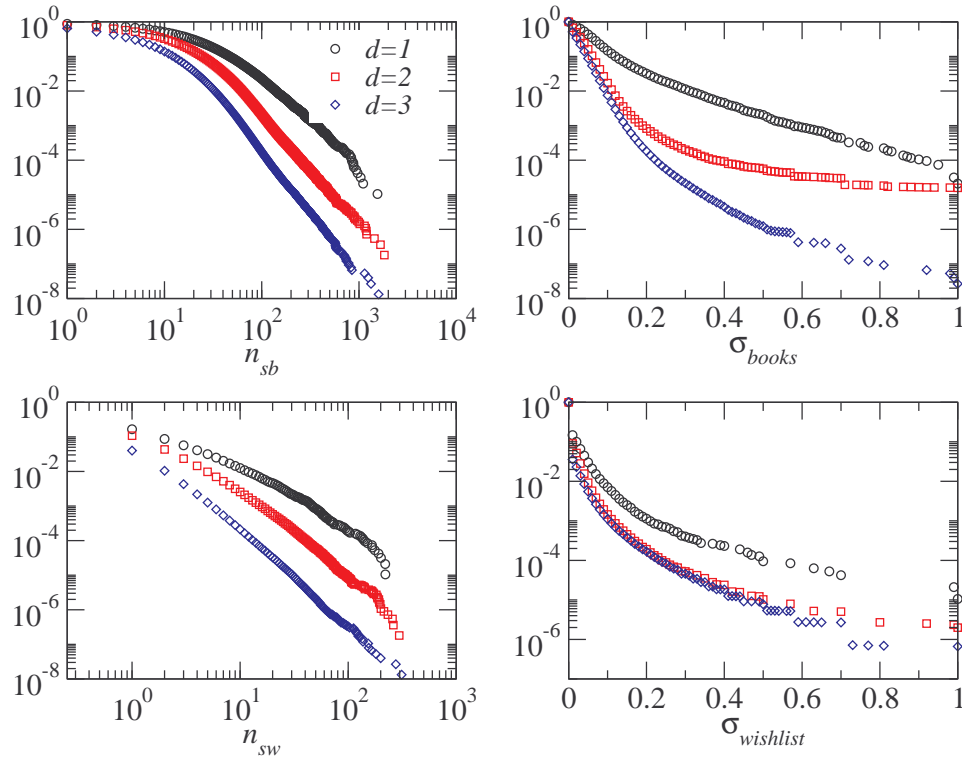


Fig. 11. Complementary cumulative distributions of the number of shared books and (left), and of the similarities in the lists of books (right), in the libraries (top) and in the wishlists (bottom) of aNobii users lying at distance d on the social network, for various values of d .

distance d on the social network in the same way as for the original data. As Fig. 7 shows, the average number of shared books in libraries and wishlists, as a function of the distance d , shows a similar trend to the original (non-shuffled) data. Similar curves are obtained for the number of shared tags and groups. For neighboring users, and also for next-to-nearest neighbors, the average numbers of shared tags or groups are generally significantly lower in the null model, but the distributions are very similar, as shown in Fig. 12(top). The assortative mixing between the amount of activity of neighboring users is therefore enough to yield a strong topical similarity *as simply measured by the number of shared tags, groups or books*. The case of cosine similarity is different: as shown in Figures 7 and 8, the average cosine similarity in the null model does not depend as strongly on distance in the social network. Analogously in Fig. 9 we see that the overlap measures in the null model are affected by social proximity, unlike cosine similarity. Fig. 12(bottom) also shows that the distributions of σ_{tags} are very different for the original and shuffled data, and do not depend on distance in the case of the shuffled data.

We conclude that the homophily measured by the cosine similarity is a genuine non-random effect and it is not only due to the assortative mixing.

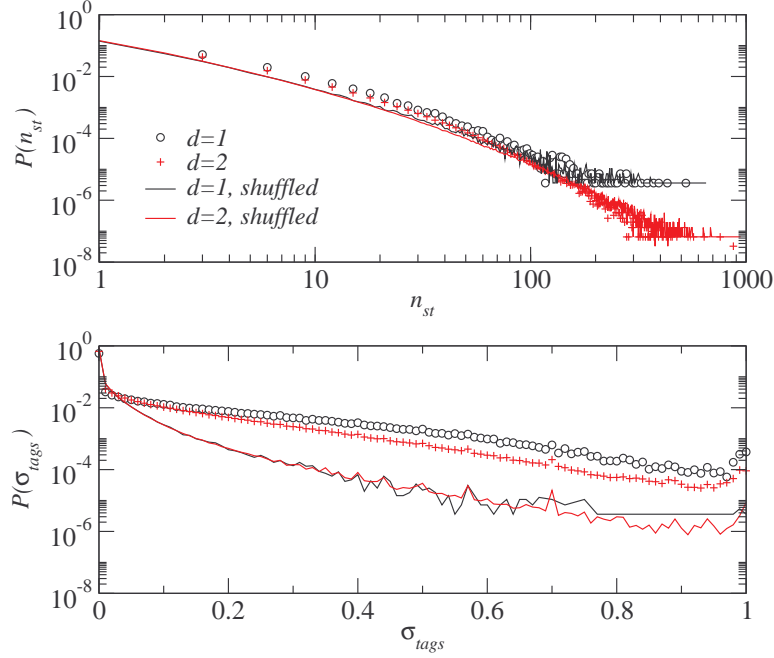


Fig. 12. Top: probability distributions of the number of shared tags of two Last.fm users lying at distance d on the social network, for $d = 1$ and $d = 2$ (symbols), and for the same network with shuffled tags (lines). Bottom: same for the distributions of the cosine similarities of the tag vocabularies.

5. SOCIAL LINK PREDICTION

The above findings on homophily naturally lead us to the hypothesis that the presence of a social tie could be predicted relying only on the topical similarity between users. In this section, we test this hypothesis focusing on the Last.fm and aNobii datasets. We do so for several reasons. First, in these two cases, the data include information on the user libraries, in addition to groups and annotations. Second, a prediction based on the tagging information is more meaningful in a *broad* folksonomy, in which users can label the same (global) set of items and pick from this global set to form their libraries. This condition allows us to deal with similarity based on shared content as well as shared vocabulary. Last.fm and aNobii are broad folksonomies, as any user can tag any artist or book. This is not the case for Flickr, which is considered a narrow folksonomy, as users normally tag only the pictures they upload themselves. Third, the Last.fm and aNobii datasets have peculiarities that allow to draw interesting conclusions regarding link prediction task. They both have a specification of users' mother tongues; as we shall see, language is a feature that can considerably affect prediction, and that should thus be taken into account when accuracy is measured. Additionally, Last.fm provides the *tasteometer* score, a user-to-user similarity metric computed by the system. While the tasteometer algorithm is not public, and therefore we do not have precise information of how its values are computed, we have verified empirically that it is largely independent from

social information and is based only on listening patterns. Consequently we can fairly compare the prediction accuracy achieved by user-to-user topical similarity measures to the one obtained by the system-provided similarity metric.

5.1 Methodology

Our link prediction problem can be defined as follows. Given a sample of users $U_s \subseteq U$, we want to predict the presence or absence of a social link for every pair $(u, v) \in \{U_s \times U_s | u \neq v\}$. Of course, the information about the social network topology is not known, but we are given the full information about the *features* that describe the user profiles. We deal in particular with four different features: *groups*, *library*, *tags*, and tagged *items*. Note the difference between *items* and *library* features. In aNobii, tagged items are a subset of the whole set of books in the user's library, while in Last.fm tagged items can be tracks, albums or artists and the library is composed only of the top 50 artists in the user's global playlist. Tags and items can be directly extracted from the three-dimensional triple space through aggregation (details are provided in Section 5.2).

We take into account each feature separately, so each user is described with a single *feature vector*. For each pair of users in U_s , we compute a similarity value between their feature vectors using the metrics defined in Section 5.2. In the case of Last.fm we also have the system-provided tasteometer similarity. Next, we sort the node pairs in decreasing order of their similarity score. The pairs with the highest topical similarity are those that we suppose are the most likely to be connected with a social link. For this reason, we predict the presence of a tie for every user pair whose similarity value is greater than or equal to a threshold value σ . To evaluate the accuracy of our predictions, we check the presence of each predicted link in the real social network and we count the number of true positives and false positives. As the value of σ decreases, a higher number of links is predicted, leading to an increase in the number of both true positives and false positives. We test the accuracy of our predictor for all the significant values of σ . The similarity measure that performs best for the prediction task is the one that achieves the best ratio between true positives and false positives, across all the possible threshold values. To quantitatively measure the prediction performance for the whole set of threshold values we consider ROC curves [Fawcett 2006] and we compare the area under the curve (AUC) achieved by the different features and similarity metrics considered. ROC curves are commonly used in the machine learning community for link prediction [Clauset et al. 2008].

Given this setting, it is important to select a significant sample of users U_s . Intuitively, one could choose $U_s \equiv U$. The problem is that, since the social graph's density is very low, the full social matrix $U \times U$ is extremely sparse. The problem of studying greatly biased datasets is well known by data miners and it is a common issue also in social link prediction, due to the intrinsic sparsity of social graphs [Liben-Nowell and Kleinberg 2003; Getoor and Diehl 2005; Leskovec et al. 2010; Lü and Zhou 2010]. It has been shown that the AUC is a good measure for performance evaluation when there is a strong class skew [Stäger et al. 2006]. However, in the first part of our evaluation we want to minimize the sparsity problem in order to compare the predictive power of different features in a less biased setting.

We thus restrict our initial analysis to several smaller subsets, each composed of

500 users sampled on the basis of one of two criteria. First, we extracted a *Most Connected* set for each feature, composed of the nodes with the highest out-degree and that have at least one element for the considered feature. Second, we sampled a distinct *Most Active* set for each feature, containing the 500 users with the largest number of elements for that feature. More in detail, we chose the sets of users with the highest number of groups, with the highest number of objects in their libraries, and, for both item and tag features, we chose the set of the 500 taggers with the highest number of triples.

The Most Active sampling provides the best scenario in which to explore the effectiveness of link prediction based on topical similarity. Furthermore, given the correlation between user activity and social connectivity (see Fig. 3), the Most Active nodes typically have a rather high degree, thus ensuring a relevant number of intra-sample social connections. As a result, the density of our social network samples ranges from 0.02 to 0.07, which is three–four orders of magnitude higher than the full networks.

In a second phase of the evaluation, we expand our observations with a sensitivity analysis to show how much the prediction accuracy is affected by the density, user activity, and size of the sampled subgraph, thus disentangling the evaluation from possible skew due to the narrower sample of active and connected users.

5.2 Similarity Metrics

To model the task of predicting social links we need to define measures of profile similarity between users. In particular, we have to select a robust similarity metric for the features that characterize the activity of users. In relation to group membership and library features we follow the approach in Section 4.3 that computes similarities by way of the standard cosine similarity as formalized in Equation 7.

For the remaining features we adopt the framework by Markines et al. [2009] that represents the system as a *tripartite* graph involving users, tags, and resources (e.g., books, songs, photos, etc.). A *triple* is a ternary relation between a user u , a tag t , and a resource r , and a set of triples is what we call a folksonomy F . We then define similarity measures $\sigma(u, v)$ where u and v can be two resources, tags, or users. Here we focus on similarity functions where u and v are two users.

Since measures for similarity and relatedness are not well developed for three-mode data such as folksonomies, we consider various ways to obtain two-mode views of the data. In particular, we consider two-mode views in which the two dimensions considered are dual — for example, users can be represented as sets of tags or resources. The process of obtaining a two-mode view from a folksonomy is called *aggregation*. Next we present different aggregation strategies and the set of similarity functions we adopted.

5.2.1 Aggregation Methods. In reducing the dimensionality of the triple space, we necessarily lose correlation information. Therefore, the aggregation method is critical for the design of effective similarity measures; poor aggregation choices may negatively affect the quality of the similarity by discarding informative correlations. Focusing on user similarity, we can aggregate across one of the tag or resource dimensions, obtaining a description of a user as a vector of, respectively, resources or tags. We consider four approaches to aggregate user information: *projection*,

distributional, *macro*, and *collaborative* aggregation. To simplify our exposition, in the following definitions we will adopt an aggregation across resources, meaning that a user will be represented as a vector of tags; analogous mechanisms apply when tags are selected as the aggregation dimension. An extensive discussion on these aggregation approaches can be found in our prior work [Markines et al. 2009].

Projection. The simplest aggregation approach corresponds to the projection operator $\pi_{u,t}(F)$ in relational algebra, assuming the triples are stored in a database relation F . Another way to represent the result of aggregation by simple projection is a matrix with binary elements where rows correspond to users (as binary vectors, or sets of tags) and columns corresponds to tags (as binary vectors, or sets of users).

Distributional. A more sophisticated form of aggregation stems from considering distributional information associated with the set membership relationships. One way to achieve distributional aggregation is to make set membership fuzzy, i.e., weighted by the Shannon information (log-odds) extracted from the annotations. Intuitively, a tag shared by two users may signal a weak association if it is very common. For example, let U be the set of users and U_t the users that annotate with t . We will use the information of tag t defined as $-\log p(t)$ where

$$p(t) = \frac{|U_t|}{|U|}. \quad (8)$$

Another approach is to define a set of *frequency-weighted* pairs (u, t, w_{ut}) where the weight w_{ut} is the number of resources tagged with t by u . Such a representation corresponds to a matrix with integer elements w_{ut} , where rows are user vectors and columns are tag vectors. We will use both of the above distributional aggregation approaches as appropriate for different similarity measures.

Macro. To compute an average function in class-partitioned datasets (e.g., documents partitioned into categories), micro- and macro-averaging approaches are possible. Micro-averaged scores are calculated considering the contribution from each element in each class. In contrast, macro-averaged values are obtained by first calculating the function for each class and then taking the average of the results. Micro-averaging gives equal weight to every element while macro-averaging gives equal weight to every class. Both approaches are broadly used in text mining [Feldman and Sanger 2006]. By analogy, distributional aggregation can be viewed as *micro-aggregation* if we think of resources as classes. Each annotation is given the same weight, so that a more popular resource would have a larger impact on the weights and consequently on any derived similarity measure. In contrast, macro-aggregation treats each resource's annotation set independently first, and then aggregates across resources. This will allow the similarity calculation to be *incremental*, breaking the dependency on global frequencies. In relational terms, we can select the triples involving each resource r , and then project, yielding a set of pairs for r : $\{(u, t)_r\} = \pi_{u,t}(\sigma_r(F))$. This results in per-resource binary matrices of the form $w_{r,ut}$. These matrix representations $w_{r,ut} \in \{0, 1\}$ are used to compute a local similarity $\sigma_r(u, v)$ for each resource r . When defining the Shannon information of a feature, the feature probability $p(t)$ must be replaced by a conditional probability $p(t|r)$. Finally, we macro-aggregate by voting, i.e., by summing across resources to obtain the global similarity. Macro-aggregation does not have a bias

toward resources with many annotations. However, in giving the same importance to each resource, the derived similarity measures amplify the relative impact of annotations of less popular resources.

Collaborative. Macro-aggregation lends itself to the exploration of collaborative filtering in folksonomies while the computation remains incremental. Thus far, we have only considered feature-based representations when working with a tripartite representation. That is, a user is described in terms of its tag or resource features. If two users share no feature, all of the measures defined on the basis of the aggregation schemes will yield a zero similarity. In collaborative filtering, on the other hand, the fact that one or more users vote for (or in our case annotate) two objects is seen as implicit evidence of an association between the two objects, even if they share no features. The more users share a pair of items, the stronger is the association. We want to consider the same idea in the context of user similarity in folksonomies. If many resources have been annotated by the same pair of users, even with different tags, the two users might be related. Likewise, if two users apply the same tags, even to annotate different resources, the two users might be related. We can capture this by adding a feature-independent local similarity to every pair (u, v) of users in macro-aggregation. In practice we can achieve this by adding a special “resource tag” t_r to all users that tagged r . This way all of r ’s users have at least one annotation in common. However, the information of such special tag would be $-\log(p(t_r|r)) = -\log(1) = 0$. To ensure that the special tag makes a non-zero contribution to the local similarity $\sigma_r(u, v)$, let us redefine the odds of tag t for resource r as

$$p(t|r) = \frac{|U_{t,r}|}{|U_r| + 1} \quad (9)$$

which is always less than 1 so that $-\log(p(t_r|r)) > 0$.

5.2.2 Similarity Measures. We wish to explore several information-theoretic, statistical, and practical similarity measures. Each of the aggregation methods requires revisions and extensions of the definitions for application to the folksonomy context. Having shown in our prior work [Markines et al. 2009] that distributional aggregation yields better accuracy than projection, and collaborative aggregation yields better accuracy than macro-aggregation, with the same computational complexity, we focus on distributional and collaborative aggregations. For brevity, we show definitions only for the similarity measures in the distributional case. These definitions are based on feature probabilities $p(x)$ defined in Equation 8. The definitions of the local similarities for collaborative aggregation are similar except that the feature probabilities are replaced by the conditional probabilities defined in Equation 9. We suppose that $u, v \in U$ represent users and X_u, X_v are their vector representations. Of course, the attributes of X depend on the aggregation dimension. In the following formulas we consider the case of aggregation across resources, i.e., the users are denoted by vectors of tags with tag elements w_{ut} .

Matching. The distributional version of the matching similarity is

$$\sigma(u, v) = - \sum_{t \in X_u \cap X_v} \log p(t). \quad (10)$$

Overlap: Distributional overlap is given by

$$\sigma(u, v) = \frac{\sum_{t \in X_u \cap X_v} \log p(t)}{\max(\sum_{t \in X_u} \log p(t), \sum_{t \in X_v} \log p(t))}. \quad (11)$$

Jaccard: Distributional Jaccard similarity is defined as

$$\sigma(u, v) = \frac{\sum_{t \in X_u \cap X_v} \log p(t)}{\sum_{t \in X_u \cup X_v} \log p(t)}. \quad (12)$$

Dice: Distributional version of Dice is defined as

$$\sigma(u, v) = \frac{2 \sum_{t \in X_u \cap X_v} \log p(t)}{\sum_{t \in X_u} \log p(t) + \sum_{t \in X_v} \log p(t)}. \quad (13)$$

Cosine: For the distributional version of the cosine, it is natural to use the frequency-weighted representation

$$\sigma(u, v) = \frac{X_u}{\|X_u\|} \cdot \frac{X_v}{\|X_v\|} = \frac{\sum_t w_{ut} w_{vt}}{\sqrt{\sum_t w_{ut}^2} \sqrt{\sum_t w_{vt}^2}}. \quad (14)$$

This formula is equivalent to Equation 6.

Maximum Information Path: The last measure we consider is *Maximum Information Path* (MIP) [Markines and Menczer 2009]. The MIP similarity is an extension of traditional shortest-path based similarity measures and Lin's similarity measure [Lin 1998]. MIP differs from traditional shortest-path similarity measures by taking into account Shannon's information content of shared tags (or resources). Lin's similarity measure only applies to hierarchical taxonomies, such as the case when bookmarks are organized in folders and subfolders. However, when the folksonomy includes non-hierarchical annotations, Lin's measure is undefined while MIP similarity is well defined and captures the same intuition. The association between two objects is determined by the ratio between the maximum information they have in common (most informative shared feature) and the information they do not share. Because of the dependency on log-odds, maximum information is not defined for projection aggregation. We define MIP for the distributional case as

$$\sigma(u, v) = \frac{2 \log(\min_{t \in X_u \cap X_v} [p(t)])}{\log(\min_{t \in X_u} [p(t)]) + \log(\min_{t \in X_v} [p(t)])}. \quad (15)$$

In prior work we have also explored mutual information and found it to be competitive but expensive [Markines et al. 2009], therefore we exclude it from the present analysis.

5.2.3 Computational Complexity. When computing similarity in large systems, the issue of scalability becomes crucial. The most important factor that affects scalability in the computation of the similarity matrix is the aggregation method adopted. The major difference between distributional and collaborative is *incrementality*. With distributional aggregation, similarities must be recomputed from scratch whenever new triples are added to the system, as frequency weights must be updated. Conversely, collaborative aggregation allows for incremental computation because each new triple affects only the contribution that the incoming tag or resource (depending on the aggregation dimension) gives to the overall similarity matrix.

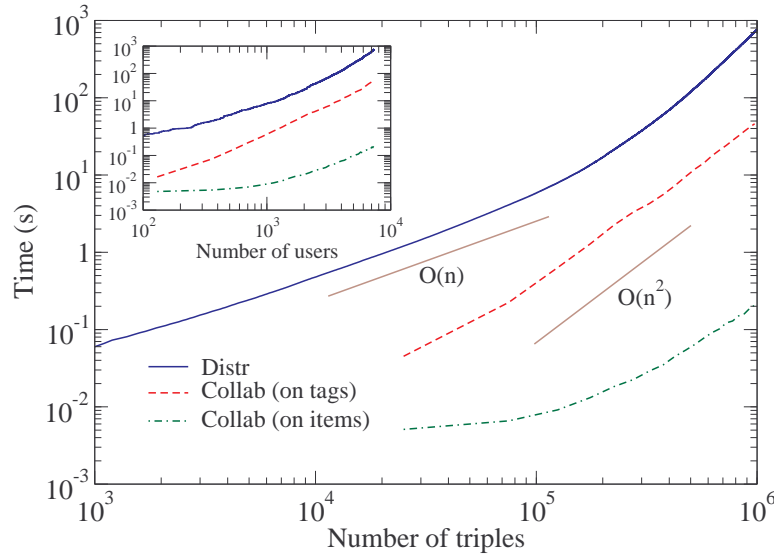


Fig. 13. Scalability of the MIP similarity computation for distributional and collaborative aggregations. Computation time of collaborative aggregation is shown for two different aggregation dimensions (items and tags). The figure reports the CPU time (in seconds) against the number n of triples (and users, in the inset). Two guides to the eye representing $O(n)$ and $O(n^2)$ complexity are shown in gray.

From a practical point of view, we can consider as scalable those measures that can be updated as a stream of incoming annotations is received. However, since the update time clearly depends on how many user pairs' similarity scores are affected by the new triples, we should study how the update time changes as the system size grows. We resort to an empirical analysis to examine how the update complexity scales with the number of users and triples in the system. Fig. 13 shows the complexity for the two different flavors of collaborative aggregations and a single representative case for distributional aggregation, since in this case the aggregation dimension does not impact performance. The experiment is performed on the aNobii dataset, using the MIP similarity.

We observe that up to around 10^5 triples the computational time is roughly linear with the system size; when the size of the system grows further the time becomes quadratic. Nevertheless, the computational advantage in using the collaborative paradigm instead of the distributional one is evident. Moreover, it is clear that collaborative aggregation over items outperforms aggregation over tags; this result is basically due to the different distributions of items and tags over users. When a new triple is added, the contribution of the value of the aggregation dimension (tag or item) of that triple to the overall similarity matrix should be recomputed. The greater the number of users who have that tag or item in their triple sets, the higher the number of pairs whose similarity score must be updated. In the considered sample the number of resources in the triple set is one order of magnitude larger than the number of distinct tags. Similar ratios hold in any big folksonomy. For this reason, it is more likely that the addition of a triple containing a very popular

tag will affect many more users (and consequently the similarity between them and others) than a triple with a very popular item.

5.3 Prediction with Single Features

We computed the similarity metrics on the Most Active and Most Connected sets of users for the four features considered, on the aNobii and Last.fm datasets. For the two folksonomy-related features, items and tags, we combined all the aggregation methods with all the similarity metrics defined above (except for the projection-MIP combination, which is not defined). For groups and library features we calculated the similarity using matching, overlap, Dice, Jaccard, and cosine metrics; note that these features do not require any aggregation and MIP is not defined for them.

We also considered two additional metrics as baselines. First, we queried the Last.fm API service for the tasteometer scores related to the same most active and Most Connected samples used for items and tags. Second, we computed the similarity in terms of number of common neighbors (CN) for the Most Connected samples, which, in this case, overlap with the Most Active samples since the number of connections is the feature considered. We introduce this widely used metric to compare the performance of network-based and feature-based similarities. Among all the known network-based metrics we opted for CN because, despite its simplicity, it has been shown to be an effective predictor of social ties [Newman 2001] and it is a local measure, whose computation is scalable.

Altogether we obtained 133 user similarity networks that we used to evaluate as many social link predictions. For brevity, we next report only on a selection of representative cases, restricting our evaluation to the best-performing instances. AUC values are summarized in Table III. Note that since the Last.fm library provided via the API has a size bounded to 50 artists, we cannot identify the Most Active users for this feature, therefore we omit the Most Active sample for the library feature. Not surprisingly, most of the highest AUC values are achieved for the Most Active samples, due to the greater amount of information available.

We note that the predictive potential is affected by the aggregation process, but without a clearly interpretable pattern. Regarding the folksonomy-based features, we find that the MIP similarity often outperforms the cosine metric, as well as the other measures (not shown). For the Most Active Last.fm users represented through items, MIP similarity outperforms the tasteometer baseline.

Predictions based on groups and libraries perform even better. For groups, we note a lower accuracy in the aNobii case compared to Last.fm, due to the relatively low cardinality of the group set; in fact, in aNobii we have about 3,000 groups, against about 70,000 groups in Last.fm. Inevitably, a lesser range of choice corresponds to a greater uniformity in the group affiliation behavior, thus making it more difficult to infer social connections. Lastly, the library results suggest that this is the best feature for prediction purposes, when applicable. Since in Last.fm the library feature vectors have at most just 50 elements each, the implicit social information carried by the elements in the library is very high. When the cardinality of the feature vector is unbounded, like in the Most Active scenario in aNobii, the AUC values become even higher. Of course, for aNobii, we expected the library-based prediction to be more accurate than the item-based ones, simply because the set of books in the library is a superset of the tagged books that can be retrieved

Table III. AUC values for Last.fm and aNobii social link predictions calculated for the four user features. The user samples considered are the most active with reference to the considered feature and the most connected users that have at least one element for that feature. The Last.fm results refer to one of our three snapshots; results for the other snapshots are consistent. The tasteometer similarity is calculated for the same most active and most connected sets used for items and tags. The feature vectors for items and tags are obtained through distributional or collaborative aggregation over the folksonomy. Shown in bold are the best results for each combination of dataset, sampling method, and feature.

Feature	Similarity	Last.fm		aNobii	
		Active	Connected	Active	Connected
Baselines	Tasteometer	0.734	0.759	-	-
	Common neighbors	0.927	-	0.854	-
Items	Distrib cosine	0.663	0.560	0.915	0.655
	Distrib MIP	0.749	0.559	0.878	0.649
	Collab MIP	0.589	0.613	0.652	0.561
Tags	Distrib cosine	0.579	0.625	0.652	0.554
	Distrib MIP	0.697	0.618	0.651	0.560
	Collab MIP	0.698	0.559	0.916	0.648
Groups	Cosine	0.810	0.677	0.662	0.690
Library	Cosine	-	0.769	0.923	0.768

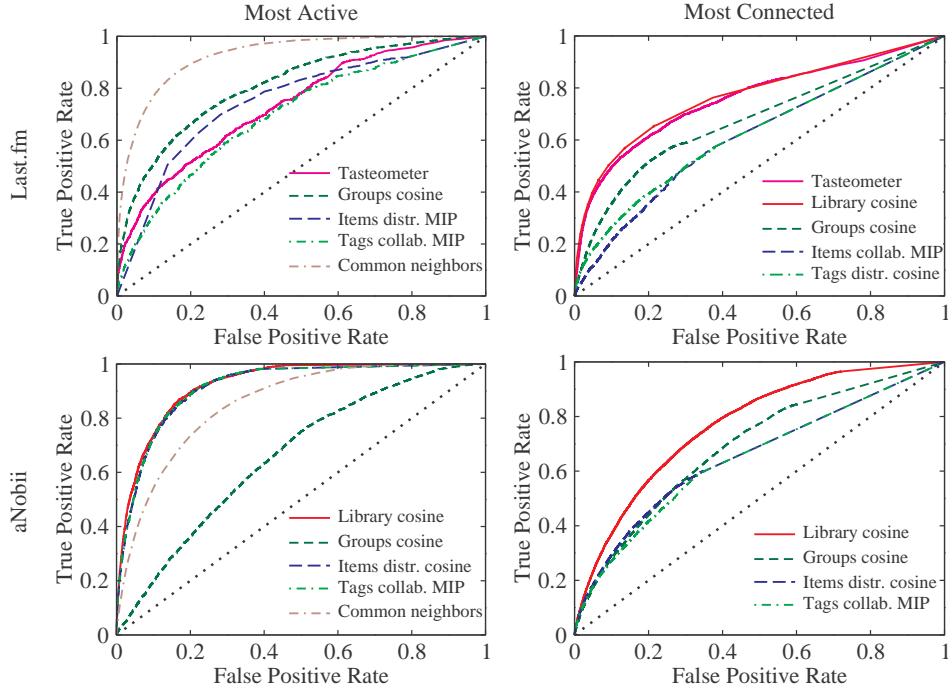


Fig. 14. ROC curves comparing the accuracy for the best feature-based predictions in both datasets. The tasteometer baseline curve is also shown for Last.fm.

from the folksonomy. Nevertheless, we observed that peak AUC values in aNobii are in part determined by the particularly strong geographically-biased clustering of its social network, which we discuss in detail in Section 6. Finally, we observe that the prediction based on common neighbors can be very accurate when a lot of information about social contacts is available. Indeed, in Last.fm’s Top Active sample, the CN metric performs best, while in aNobii the library and folksonomic features are more accurate.

In Fig. 14 we depict a summary comparison between the ROC curves of the best performing prediction measures, i.e., those shown in bold in Table III.

5.3.1 Sensitivity Analysis. The analysis on small user samples is useful for comparing the effectiveness of different metrics under different boundary conditions of connectivity and activity. However, to show that the results are not biased by this sampling procedure, we performed a sensitivity analysis for predictions made on the aNobii dataset, using the library feature. (Similar results hold for other features.) We first looked at the effect of sample size on accuracy. Top plots in Fig. 15 show the AUC values obtained for sample size up to 5000 users. Together with AUC, we also measure the precision at top N , i.e., what fraction of the pairs which have the N highest similarity values are actually connected; this is a metric which is commonly used to complement the AUC [Backstrom and Leskovec 2011]. The prediction accuracy is stable and does not depend significantly on sample size.

Second, to study how the prediction accuracy depends on the activity and connectivity values of the samples, we collected several samples of the same size (500 users) but with decreasing activity (starting from the Most Active sample) and connectivity (starting from the Most Connected sample). Results are depicted in bottom plots of Fig. 15. Since in many samples the number of connected users is very low, and the classic definition of precision at top N is not meaningful when the number of links is less than N , we present only the AUC values. We observe that even when activity and connectivity are greatly reduced, the prediction accuracy is stable and remains high.

5.3.2 Discussion. The overall picture that emerges from the experiments reveals some interesting results. First of all, the strong correlation between social linking and user activity, resulting in a noticeable homophily phenomenon, can be profitably exploited to accurately infer the structure of the social network given only information on user features. Considering various features results in quite different prediction performance, as seen in Fig. 14.

The only disadvantage of social link prediction based on folksonomic information is that in many systems a considerable portion of users does not use tags: 50% of users are taggers in Last.fm, only 30% in aNobii. However, when tagging information is available, results are encouraging. For folksonomy-based features (items and tags), our prediction methodology based on the MIP similarity metric tends to work better than the other similarity measures considered. This result holds across different collaborative tagging systems. When users are active (described by a high number of triples), we observe that good results are achieved with a distributional approach when aggregating over tags and with a collaborative approach when aggregating over items. In other words, from a collaborative filtering perspective,

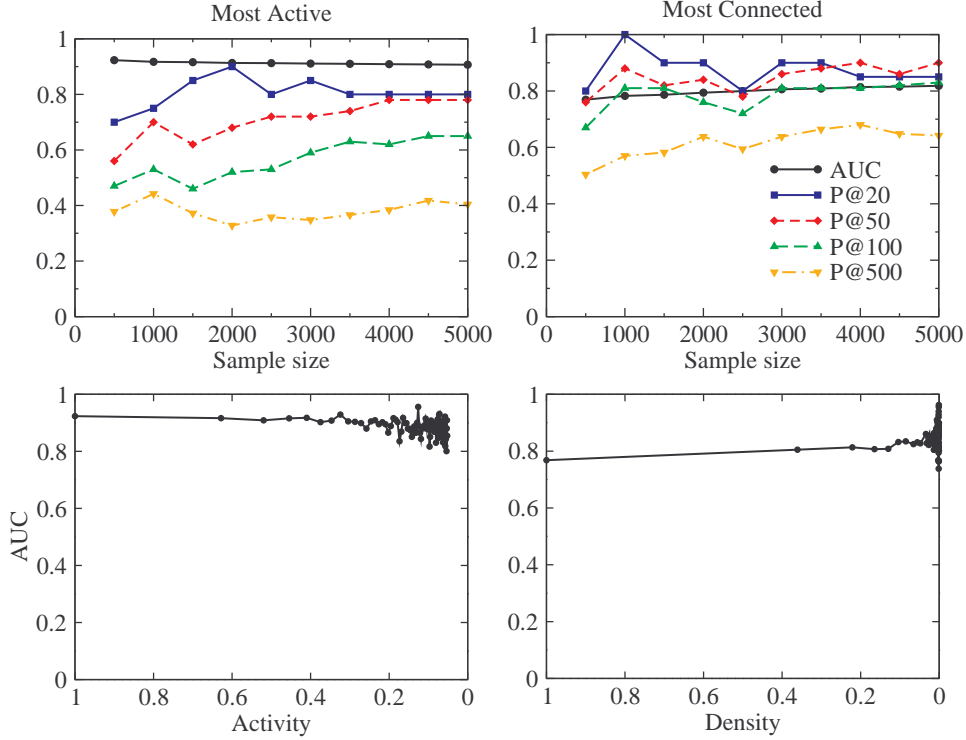


Fig. 15. Sensitivity analysis of link prediction based on the library feature in aNobii; the Most Active (left) and Most Connected (right) cases are considered. Top plots shows how AUC and precision at N change as the sample size is increased up to 5,000 users. Bottom plots show how AUC in different samples of size 500 but with decreasing values of density and activity (normalized on the highest value); the noisy patterns at the end of the lines are observed because the prediction on very sparse samples is much more sensitive to slight variations in the connectivity patterns.

knowing that two users share a tag is more informative for predicting their social link. In the distributional scenario with item representation, the accuracy compares favorably with that achieved by the Last.fm tasteometer.

It is interesting to notice that the ranking of aggregation methods by prediction accuracy is not consistent across datasets. For example, in the Most Connected scenario, the distributional and the collaborative approaches behave differently in the two datasets considered (see Table III). This means that even folksonomies with the same macro-structural properties (broad folksonomies, with similar numbers of users, triples and tags) can be characterized by inequivalent tagging patterns that lead to different performance of the prediction techniques.

Predictions made from the group feature can lead to even more accurate results. Groups behave very well if the total number of groups in the system is not too small with respect to the user population; furthermore, compared to the number of taggers, a larger portion of users in social systems take part in thematic groups, thus allowing the prediction for a wider user set. Even if we do not focus here on the causality aspects linking social connections with homophily, the high AUC values

obtained for the group feature could reasonably lead us to conjecture that groups are effective means of socialization, i.e., people know each other through groups.

The best performing profile feature is the library. Aside from the surprisingly high accuracy obtained for the Most Active set, where the information is maximal, the most important outcome is that the analysis of library feature vectors is very significant also in cases when considerably less information is available. From this viewpoint, the Most Connected scenario in Last.fm is particularly revealing because the accuracy is very high even if users are described with feature vectors containing at most 50 artists from their libraries. In a nutshell, the prediction task performs best if it relies on the main feature that denotes the social network topic; in our case study, books for aNobii and artists for Last.fm.

Finally, the common neighbors baseline seems to be a very good predictor of social links, sometimes performing even better than all other profile features. This is in part expected because the common neighbors measure captures the probability to form a *triadic closure*, which is a relevant phenomenon of attachment in social networks [Newman 2001]. Next, we explore a hybrid approach that combines profile features and network-based features to achieve even better link prediction accuracy.

5.4 Combining Features for Prediction

The common approach to link prediction based on multiple features relies on machine learning techniques. Prediction is seen as a binary classification problem that can be solved with a classifier trained on the features that describe the nodes. Positive and negative examples are chosen among pairs of nodes that are connected or disconnected, respectively [Lü and Zhou 2010].

Here we adopt this approach by selecting 10,000 positive samples and as many negative samples from the set of aNobii users who have at least one instance of each feature in their profile and with at least one outgoing edge. The features considered are simply the similarity scores, computed as described previously in this section. We use the J48 decision tree from WEKA [Hall et al. 2009] as a binary classifier and perform a 10-fold cross validation on our sample. We run the classifier for each profile feature and for the common neighbors separately, and then combine together all the profile features exclusively, before finally adding the information on common neighbors. We consider the AUC together with accuracy, False Positive and False Negative rates.

The results are shown in Table IV. The main observation is that combining different features results in a noticeable boost in predictive power. In particular, the combination of different profile features gives about a 14% improvement in AUC over the best performing profile feature taken individually, while adding topological information, like the number of common neighbors, leads to an additional 4% improvement. Thus, even if link prediction from network-based features can lead to accurate results, the best performance is achieved by their combination with profile features.

6. LANGUAGE COMMUNITY ANALYSIS

The very high accuracy of our social link predictions in aNobii motivated us to further inspect the reasons behind such a strong performance. We suspected that the results were somehow affected by the strongly clustered structure of the aNobii

Table IV. Predictive power of single and combined features using a decision tree on a balanced set of 10,000 positive and negative samples extracted from the aNobii dataset.

	Tags collab MIP	Groups	Library	CN	Profile feaures	All features
AUC	0.785	0.807	0.811	0.844	0.924	0.963
Accuracy	0.786	0.809	0.812	0.846	0.877	0.915
FP Rate	0.177	0.143	0.335	0.031	0.109	0.072
FN Rate	0.251	0.240	0.041	0.277	0.137	0.099

social network.

In fact, aNobii is split among two main groups: the Italian community (about 60% of users), and the Far East community, representing Hong Kong and Taiwan (about 20% of users). Since the type of literary items consumed by the great majority of people is strictly entangled with their mother tongue, the intersection of topical interests between the two communities is very small and prevalently limited to few worldwide best sellers. In addition to this, aNobii allows the insertion of annotations with any language character set. As a result, users fill their libraries with books written in their own language and they are motivated to annotate them using terms from their mother tongue vocabulary. Given the topical similarity between neighbors (see Section 4), and given that the two communities have very different topical interests, these two main groups turn out to be almost disconnected in the social network.

We show that this scenario directly affects the performance of our folksonomy-based prediction method, considering as a representative example the Most Active sample of users. The same qualitative considerations hold for other samples and features. We split the Most Active set of taggers into clusters on a country level (within the set of top 500 taggers, 249 are Italian and 137 are Taiwanese) and we calculate some basic measures for pairs of users that reside in the same country or in different countries (we neglect users who do not specify a location in the profile).

For every user pair, we compute the cosine similarity between their vocabularies and between their item sets, and we measure the portion of inter- and intra-cluster links. To show that the portion of links residing inside a language community is not simply due to statistical properties caused by the size imbalance between different communities, we repeated the same measure on a shuffled version of the social graph, where each node keeps its out-degree but rewires its links at random. Statistics are summarized in Table V. We notice that in aNobii the portion of inter-community links in the real network is considerably smaller than in the shuffled network, meaning that the clusters are nearly disconnected from each other due

Table V. Statistics on language communities that compose the Most Active users who declare country of origin in aNobii and Last.fm. We report on the portion of links that reside inside a cluster or, conversely, connect users belonging to different clusters in the real social network vs. its shuffled version (in parenthesis). The average cosine similarity for tag and item sets computed between pairs of users residing in the same or in different communities is reported as well.

	aNobii			Last.fm		
	Links (shuffl.)	Tags σ	Items σ	Links (shuffl.)	Tags σ	Items σ
Intra	85% (38%)	$3.4 \cdot 10^{-2}$	$2.2 \cdot 10^{-2}$	17% (9%)	$1.4 \cdot 10^{-1}$	$1.7 \cdot 10^{-2}$
Inter	15% (62%)	$4.7 \cdot 10^{-3}$	$1.8 \cdot 10^{-3}$	83% (91%)	$1.4 \cdot 10^{-1}$	$1.5 \cdot 10^{-2}$

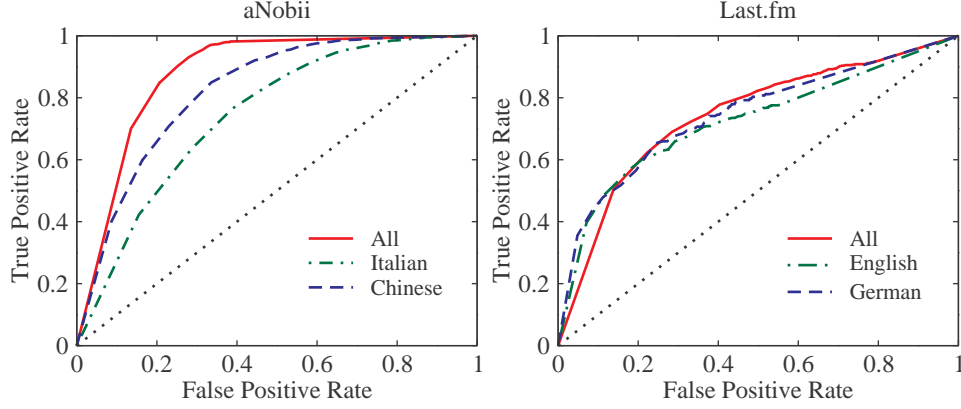


Fig. 16. ROC curves comparing the link prediction within different language communities in aNobii and Last.fm. The user samples considered are composed by the top 500 taggers in the whole system (All) or considering a single language community (Italian, Chinese, English, German). In all cases we used the MIP similarity metric using a distributional aggregation over tags.

to language homophily. Furthermore, on average, the similarity between pairs of users, calculated for both tags and items, is much lower for users belonging to different language clusters compared to pairs of users that reside inside the same cluster. The average inter-community topical overlap is thus very small. Hence, the prediction task is simpler with respect to a more homogeneous setting because both the social network and the folksonomy features are clustered according to language.

The effect of language on facilitating link prediction can be verified by focusing on communities with homogeneous language. We performed the prediction task on subsets of 500 active users within language communities. The comparison between the ROC curves is depicted in Fig. 16 for the MIP metric, using the aggregation on tags (the result is qualitatively the same for the other cases). The predictions are significantly more accurate in the mixed community than in the homogeneous communities: the prediction task is simpler in the former case because links between the two communities will almost never be predicted, thus considerably decreasing the number of false positives. Language thus plays a key role in the prediction task for multi-language communities.

To further confirm this observation, we performed the same tests on Last.fm, taking into account the most active users from the two largest language groups in our snapshot: the German community and the English-speaking community, composed by the union of users from USA, UK and Australia. As Fig. 16 shows, in this case the prediction accuracy is not clearly affected by the language. Such a different result is well explained by the statistics reported in Table V. Compared to aNobii, the language communities are fuzzier in Last.fm since the level of language homophily is considerably lower and close to that of the random shuffled network; furthermore, there is no clear difference between inter- and intra-cluster feature similarities. Therefore, language does not play a substantial role in the link prediction accuracy.

The impact of language clustering on the tagging behavior in the two social

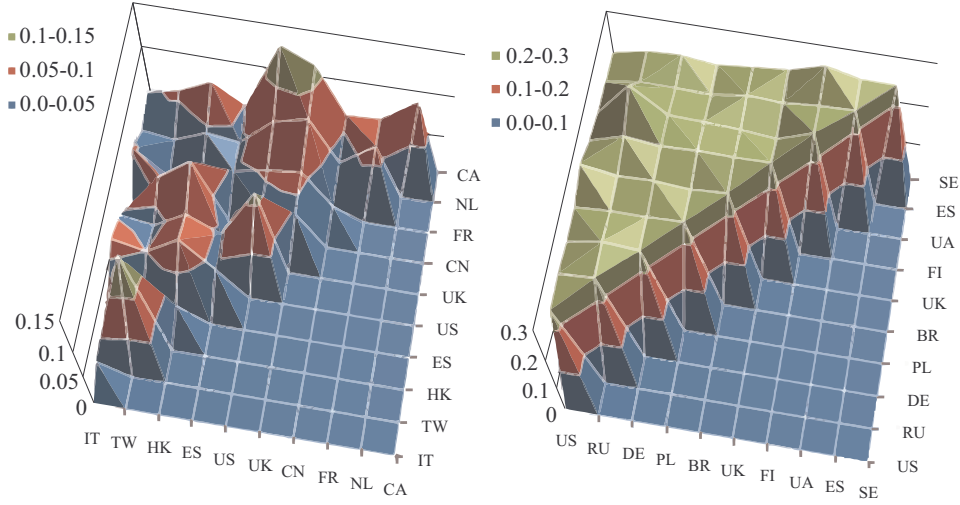


Fig. 17. Cosine similarity between the tag vocabularies used by the 10 biggest geographic communities in aNobii (left) and Last.fm (right). A tag vocabulary is the set of tags obtained by merging all the tag sets of all the users in the community. Color intensity denotes different classes of similarity values.

networks considered is also shown in the heat maps in Fig. 17. Here, for both aNobii and Last.fm, we computed the cosine similarity between the tag vocabularies in use by the 10 most populated geographic communities. The results clearly show an overall low similarity value between the aNobii communities, except for those countries that share the same language (e.g., Canada, United States, and United Kingdom or Taiwan and Hong Kong). On the contrary, the Last.fm geographic groups are more homogeneous, with a substantially higher similarity and no clear distinction on the basis of their official languages. A very similar scenario is found for similarity on item sets (not shown).

Linguistic constraints are more tangible in aNobii than in Last.fm, or in books compared to music. The tagging behavior and the literary tastes of aNobii users are strongly affected by their language, while in Last.fm, users across different languages tend to have more tags and more music items in common. In synthesis, language can strongly impact the tagging behavior; the level of homophily in a social network depends on the extent to which language is correlated with topical interests, which in turn depends on the nature of the objects shared between users.

7. CONCLUSIONS

In this paper we studied the correlation between social and topical features in three popular online social networks: Flickr, Last.fm, and aNobii. Interesting patterns that are common to all the considered social environments, despite their different magnitudes and semantic orientations, emerge from the study.

We first inspected the interplay between individual user features and social interaction. We observed a strong correlation between the social connectivity and the intensity of explicit user activities like tagging and participation in groups. Assorta-

tive mixing patterns between neighbors can be found for all the examined features as well. We showed a dependency between the topical similarity between pairs of users and their shortest-path distance on the social graph, using a null model built to discern purely statistical assortativity from actual homophily. We found a clear topical alignment trend between neighbors for all the examined features.

The results obtained for homophily led us to investigate the possibility of predicting the presence of a social link based on user profile features in Last.fm and aNobii. We built a user similarity network for each feature available (tags, tagged items, groups, and library items) using several similarity metrics. We used the similarity rankings of user pairs to perform predictions of social links and we evaluated prediction accuracy using ROC curves.

Results show that all the user profile features have substantial predictive power. Regarding the folksonomy-derived features, the similarity metric that we introduced to compare objects in the folksonomy space, namely the MIP similarity, produced the most accurate results. Moreover, the aggregation technique used to extract folksonomic features from the three-dimensional folksonomy space is very relevant to the prediction results. Overall, the library feature, i.e., the collection of user items (books in aNobii and artists in Last.fm) turned out to be the most predictive, and was very accurate even when users have a small number of library items. Finally, combining profile features with network-based features through a machine-learning approach leads to a boost in link prediction accuracy.

We also studied the effect that confounding aspects like spoken language and nationality have on the prediction task. We showed that the link prediction task is easier in social networks that are strongly clustered by language, because users in different clusters tend to have very different topical interests and to establish very few social ties. For this reason, we suggest that a preliminary analysis on the language community structure should be carried out before testing any feature-based link prediction algorithm on real social network data.

The present findings suggest a number of possible future directions. First, the causal relationships of homophily deserve to be explored in detail. More generally, given longitudinal snapshots of the social environment, evolutionary patterns of the interplay between connectivity and similarity could be studied. Using temporal datasets would also allow better testing of the validity of our link prediction approach by checking if predicted links actually appear in the near future.

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