Viral marketing without tears: Limiting the harm caused by diffusing information to vulnerable users

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Motivation (1/2): Social networks and viral marketing

- Social networks are powerful communication infrastructures
 - Facebook (1.94 billion monthly active users¹)
 - Twitter (313 million monthly active users²)
- They allow diffusing information quickly to many users through word-of-mouth effects
 - good for advertising products or events through viral marketing
- The success of a viral marketing campaign on a social network can be measured by the number of influenced users

²https://about.twitter.com/company

¹http://newsroom.fb.com/company-info/

• Influence maximization

- Find k users (*seeds*) that influence the largest number of users, according to a diffusion model
- **Drawback**: Some users (*vulnerable users*) may be harmed by information diffusion
 - Promoting alcoholic drinks to people with drinking problems
 - Promoting junk food to obese people

How to limit the influence to vulnerable users, while maximizing the influence to the non-vulnerable users (so that users and companies benefit from viral marketing)?

Influence measure to quantify the quality of a seed-set

- Additive Smoothing Ratio (ASR)
- Baseline Heuristics for finding an ASR-Maximizing seed-set
 - GR natural greedy heuristic
 - *GR_{MB}*: a variation of *GR* (more efficient)

• Approximation algorithm for finding an ASR-Maximizing seed-set

• *ISS* (Iterative Subsample with Spread bounds): an efficient approximation algorithm

Background (1/2): Set functions

Monotonicity

A function $f : 2^U \to \mathbb{R}$ is monotone, if $f(X) \le f(Y)$ for all subsets $X \subseteq Y \subseteq U$, and non-monotone otherwise

Submodularity, supermodularity, and modularity

- A function $f : 2^U \to \mathbb{R}$ is submodular, if $\forall S \subseteq T \subseteq U$ and $j \in U \setminus T$: $f(S \cup \{j\}) - f(S) \ge f(T \cup \{j\}) - f(T)$ (1)
- supermodular, if and only if -f is submodular [3]
- modular, if Eq. 1 holds with equality



• diminishing returns property

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Background(2/2): Graph representation and IC model

Social network as a graph

- Directed graph G(V, E) that models a social network (at a certain time)
- V is partitioned into $\mathcal{N}(\text{non-vulnerable nodes})$ and $\mathcal{V}(\text{vulnerable nodes})$ and we assume $(\mathcal{N} \neq \emptyset)$

Independent Cascade (IC) model [2]



- Seed nodes are influenced at initial time point 0.
- At each next time point, each newly influenced node u activates its out-neighbor v independently, with probability p((u, v)).
- The process stops when no new nodes are activated.
- The spread (expected number of influenced users) for a seed-set S in the IC model is denoted with σ(S).

Difference

The difference $\sigma_{\mathcal{N}}(S) - \sigma_{\mathcal{V}}(S)$ between the spread of non-vulnerable and vulnerable users

Limitations

• It does not consider what fraction of all influenced users are vulnerable

Example

It favors promoting an alcoholic beverage to 140 users out of whom **40** have drinking problems, instead of 59 users with no drinking problems, since (140 - 40) - 40 > 59 - 0.

• It cannot be used to find a seed-set S with approximately maximum $\sigma_N(S) - \sigma_V(S)$ [1]

Ratio

The ratio $\frac{\sigma_{\mathcal{V}}(S)}{\sigma_{\mathcal{N}}(S)}$ between the spread of vulnerable and non-vulnerable users

Limitations

• It does not favor a seed-set that influences many non-vulnerable users (i.e., is good for viral marketing), among seed-sets that do not influence vulnerable users (does not distinguish seed-sets with $\sigma_V(S) = 0$).

Example

 S_1 and S_2 do not influence users with drinking problems:

- S_1 : 59 users with no drinking problems: $\frac{\sigma_V(S_1)}{\sigma_V(S_1)} = \frac{0}{59} = 0$
- S_2 : 2 users with no drinking problems: $\frac{\sigma_V(S_2)}{\sigma_V(S_2)} = \frac{0}{2} = 0$

• It cannot be used to find a seed-set with small or zero $\sigma_{\mathcal{V}}(S)$ and large $\sigma_{\mathcal{N}}(S)$.

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Our influence measure and problem definition

Additive Smoothing Ratio (ASR)

•
$$ASR(S,c) = rac{\sigma_{\mathcal{N}}(S) + c}{\sigma_{\mathcal{V}}(S) + c}$$
, where S is a seed-set and $c > 0$ is a constant

Example

 S_1 : 59 users with no drinking problems, $ASR(S_1, 1) = \frac{\sigma_N(S_1) + 1}{\sigma_V(S_1) + 1} = \frac{60}{1}$

 S_2 : 2 users with no drinking problems, $ASR(S_2, 1) = \frac{\sigma_{\mathcal{N}}(S_2)+1}{\sigma_{\mathcal{V}}(S_2)+1} = \frac{3}{1}$

Problem definition

- Given G(V, E) and c > 0, find a seed-set $S \subseteq V$ of size at most k with maximum ASR(S, c)
- NP-hard
- Cannot be approximated using algorithms for submodular and/or supermodular maximization because *ASR* is **non-monotone** and **neither submodular nor supermodular**.

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Baseline heuristics (1/2)

GR (GReedy heuristic)

Input: $\mathcal{N} \subseteq V, \ \mathcal{V} \subseteq V$, graph *G*, parameter *k*, constant *c* **Output**: Subset $S \subseteq \mathcal{N}$ of size $|S| \leq k$ $S_0 \leftarrow \{\}; \ i \leftarrow 0$ **While** i < kFind a node $u \in \underset{v \in \mathcal{N} \setminus \{S_i\}}{\operatorname{arg\,max}} \frac{\sigma_{\mathcal{N}}(S_i \cup v) - \sigma_{\mathcal{N}}(S_i) + c}{\sigma_{\mathcal{V}}(S_i \cup v) - \sigma_{\mathcal{V}}(S_i) + c}$ $S_{i+1} \leftarrow S_i \cup \{u\}$ $i \leftarrow i+1$ **Return** the subset $S \in \{S_1, \dots, S_k\}$ with the largest *ASR*

Limitation: The computation of $\sigma_{\mathcal{N}}$ and $\sigma_{\mathcal{V}}$ is slow (all paths from S to \mathcal{N} or \mathcal{V} in the graph need to be considered)

Baseline heuristics (2/2)

GR_{MB}

- Differs from GR in that it estimates the spread efficiently using the MIA (Maximum Influence Arborescence) Batch-update method [6]
- two orders of magnitude faster on average than *GR*, but less effective in terms of *ASR*



- For any pair of nodes *u* and *v*, find the **maximum influence path** from *u* to *v*
- Estimate influence probability *P_S(u)* as the union of maximum influence paths from *S* to *u*

(a)

•
$$\sigma_{\mathcal{N}} = \sum_{u \in \mathcal{N}} P_{\mathcal{S}}(u)$$

• $\sigma_{\mathcal{V}} = \sum_{u \in \mathcal{V}} P_{\mathcal{S}}(u)$

The ISS approximation algorithm (1/3)

Main ideas

 We define submodular (easier to maximize) functions ASR^L and ASR^U that bound ASR from below and from above:

$$ASR_{Y,c}^{\mathbf{L}}(S) = \frac{\sigma_{\mathcal{N}}(S) + c}{\widehat{\sigma_{\mathcal{V},Y}}(S) + c} = \frac{\sigma_{\mathcal{N}}(S) + c}{\sigma_{\mathcal{V}}(Y) + \sum_{u \in S \setminus Y} \sigma_{\mathcal{V}}(\{u\}) - \sum_{u \in Y \setminus S} (\sigma_{\mathcal{V}}(Y) - \sigma_{\mathcal{V}}(Y \setminus \{u\})) + c}$$
$$ASR_{Y,\pi^{Y},c}^{\mathbf{U}}(S) = \frac{\sigma_{\mathcal{N}}(S) + c}{\widetilde{\sigma_{\mathcal{V},\pi^{Y}}}(S) + c} = \frac{\sigma_{\mathcal{N}}(S) + c}{\sum (\sigma_{\mathcal{V},Y,\pi^{Y}}(u)) + c}$$

because ASR(S, c) is non-monotone and non-submodular (difficult to maximize). The bounds are based on the modular bounds for submodular functions in [1].

 $u \in S$

- We select seeds from a sample of \mathcal{N} of size approximately $\frac{|\mathcal{N}|}{k}$.
- Iterative construction of a seed-set, until ASR cannot improve.

The ISS approximation algorithm (2/3)

Simplified description of ISS

Input: $\mathcal{N} \subseteq V$, $\mathcal{V} \subseteq V$, graph *G*, parameter *k*, constant *c* **Output:** Subset $S \subseteq \mathcal{N}$ of size $|S| \leq k$

 $S_{pr} \leftarrow \{\}; S_{cur} \leftarrow \mathcal{N}$

While true

$$i \leftarrow 0; S_0^{\mathbf{0}} \leftarrow \{\}; S_0^{\mathbf{L}} \leftarrow \{\}; S_0^{\mathbf{U}} \leftarrow \{\}\}$$

While $i < k$

Uniform random sample with approximately $\frac{|\mathcal{N}|}{k}$ nodes $S_{i+1}^{\mathbf{0}} \leftarrow \text{add into } S_i^{\mathbf{0}}$ the node with max. marginal gain in ASR $S_{i+1}^{\mathbf{L}} \leftarrow \text{add into } S_i^{\mathbf{L}}$ the node with max. marginal gain in $ASR_{S_{pr,c}}^{\mathbf{L}}$ $S_{i+1}^{\mathbf{U}} \leftarrow \text{add into } S_i^{\mathbf{U}}$ the node with max. marginal gain in $ASR_{S_{pr,\pi}}^{\mathbf{U}} S_{pr,\pi}^{\mathbf{U}}$ $i \leftarrow i + 1$

 $S_{cur} \leftarrow$ best seed-set w.r.t ASR among S_k^{O} , S_k^{L} , S_k^{U} If S_{cur} not better than S_{pr} w.r.t. ASR

break

 $S_{pr} \leftarrow S_{cur}$

Return S_{cur}

The ISS approximation algorithm (3/3)

• ISS constructs a seed-set with expected value of ASR no less than $\mathcal{M} \cdot 23\%$ of the optimal, where \mathcal{M} depends on the constants c and k and the ASR^L function.

Theorem

ISS constructs a seed-set S such that:

$$\mathbb{E}[ASR(S,c)] \ge \max\left(\frac{\sigma_{\mathcal{V}}(S^*) + c}{\widehat{\sigma_{\mathcal{V},S_{pr}}}(S^*) + c}, \frac{c}{c + k \cdot \max_{u \in \mathcal{N}} \widehat{\sigma_{\mathcal{V},S_{pr}}}(\{u\})}\right) \cdot \frac{1}{e} \cdot (1 - \frac{1}{e}) \cdot ASR(S^*,c)$$

where $S^* = \arg \max_{S \subseteq \mathcal{N}, |S| \leq k} ASR(S, c)$, $\widehat{\sigma_{\mathcal{V}, S_{pr}}}$ is the modular upper bound used in ASR^{L} , and the expectation is over every possible S constructed by ISS.

Experimental setup

Evaluation of GR, GR_{MB}, ISS

Competitors:

- TIM [5]: a heuristic for maximizing σ_N(S) − σ_V(S),
- *RB*: employs *Greedy* [4] to the subset of non-vulnerable nodes that influence no vulnerable nodes
- Effectiveness measures: $\sigma_{\mathcal{N}}$, $\sigma_{\mathcal{V}}$, ASR, $\frac{\sigma_{\mathcal{N}}}{|\mathcal{N}|}$, $1 \frac{\sigma_{\mathcal{V}}}{|\mathcal{V}|}$
- Efficiency measure: Runtime

Datasets

Dataset	# of nodes	# of edges	avg in-degree	max in-degree	# of vuln. nodes	θ
	(V)	(E)			(\mathcal{V})	
WI	7115	103689	13.7	452	100	0.01
TW	235	2479	10.5	52	25	0.01
POL	1490	19090	11.9	305	100	0.003
AB	840	10008	11.9	137	10	0.01

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Comparison to RB

 GR constructs seed-sets that influence at least 5.5 and up to 38 times more non-vulnerable nodes than those constructed by RB, for different values of c and k





ASR with c = 1

- All our algorithms substantially **outperform** *TIM*
- *ISS* outperformed all other method **3.5 times** on average over all datasets, *k* value and $|\mathcal{V}|$ values





Spread of Vulnerable and Non-vulnerable Nodes

• Each point (x, y) corresponds to the values $(1 - \frac{\sigma_V(S)}{|\mathcal{V}|}, \frac{\sigma_N(S)}{|\mathcal{N}|})$, referred to as *protection* and *utility* of a seed-set S



- All our algorithms substantially **outperformed** *TIM* in terms of σ_N and/or σ_V
- *ISS* outperformed *TIM* with respect to **both protection and utility**, achieving overall better protection than GR and better utility than GR_{MB}

Efficiency

- Our methods are faster than TIM by at least one order of magnitude
- *TIM* is too slow (10 hours for k = 50 and a dataset with 235 nodes, and more than 17 days for larger datasets)





- Introduced the problem of performing viral marketing while limiting the influence to vulnerable nodes
- Proposed an influence measure and defined an optimization problem based on the measure
- Proposed two greedy baseline heuristics and the *ISS* approximation algorithm
- Experimentally showed that *ISS* outperforms *TIM* [5] and our baselines in terms of effectiveness and efficiency

Forthcoming IEEE AINA paper:

https://kclpure.kcl.ac.uk/portal/files/104770966/VIM_paper_final.pdf

Background (3/5): Modular bounds

- We review two bounds for a submodular function that are used in our approximation algorithm.
- The bounds are computed for a given subset $Y \subseteq U$.
- The bounds are modular and thus easier than f to optimize efficiently.



Modular upper bound [1]

The modular upper bound f_Y(X) of a submodular function f : 2^U → ℝ is a modular function [1]

$$\widehat{f_Y}(X) = f(Y) + \sum_{u \in X \setminus Y} (f(\lbrace u \rbrace) - f(\lbrace \rbrace)) - \sum_{u \in Y \setminus X} (f(Y) - f(Y \setminus \lbrace u \rbrace))$$
(2)

where $Y \subseteq U$ is a given subset of U.

Modular lower bound [1]

• The modular lower bound $f_{Y,\pi^Y}(X)$ of a submodular function $f(X): 2^U \to \mathbb{R}$ is a modular function

$$\widetilde{f}_{Y,\pi^{Y}}(X) = \sum_{u \in X} f_{Y,\pi^{Y}}(u)$$
(3)

where $Y \subseteq U$ is a given subset of U, π^{Y} is a random permutation of the elements of Y, π^{Y}_{u} is the prefix of π^{Y} , $\pi^{Y}_{u^{-}}$ is π^{Y}_{u} except u, and

$$f_{Y,\pi^{Y}}(u) = \begin{cases} f(\pi_{u}^{Y}) - f(\pi_{u^{-}}^{Y}), & \text{if } u \in Y \\ 0, & \text{otherwise} \end{cases}$$
(4)

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