

# ANGEL: efficient, and effective, node-centric community discovery in static and dynamic networks

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Our approach is primarily designed for social networks analysis and belongs to a well-known subfamily of Community Discovery approaches often identified by the keywords bottom-up and node-centric

## 1 GOALS

- we propose ANGEL , an algorithm that aims to lower the computational complexity of previous solutions while ensuring the identification of high-quality overlapping partitions

## 2 PRELIMINARIES

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## 3 CHALLENGES

- complex networks researchers agree that it is not possible to provide a single and unique formalization that covers all the possible characteristics a community partition may satisfy

## 4 PREVIOUS WORK / CITATIONS

- (Coscia et al. 2012): where the authors propose DEMON an approach whose main goal was to identify local communities by capturing individual nodes perspectives on their neighbourhoods and using them to build mesoscale ones
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- **This Work:**
  - Introduces a Label Propagation algorithm
    - \* Least complex kind of algorithm
    - \* Gives good quality results
  - In contrast to DEMON it focuses on lowering the time complexity while at the same time increasing the partition quality
  - Properties:
    - \* It produces a deterministic output
    - \* Allows for a parallel implementation

## 5 DEFINITIONS

- During each iteration, the label of  $v$  is updated to the majority label of its neighbours. As the labels propagate, densely connected groups of nodes quickly reach a consensus on a unique label

## 6 OUTLINE / STRUCTURE

- Node Labeling  $O(n + m)$  (Raghavan et al. 2007)
  - Initialize the labels at all nodes in the network. For a given node  $x$ ,  $C_x(0) = x$
  - Set  $t = 1$
  - Arrange the nodes in the network in a random order and set it to  $X$

- For each  $x \in X$  chosen in that specific order, let  $C_x(t) = f\left(C_{x_{i_a}}(t), \dots, C_{x_{i_m}}(t), C_{x_{i_{(m+1)}}}(t-1), \dots\right)$   
 $f$  here returns the label occurring with the highest frequency among neighbors and ties are broken uniformly randomly.
- If every node has a label that the maximum number of their neighbors have, then stop the algorithm. Else, set  $t = t + 1$  and go to (3)
- Community Matching:
  - Don't make use of the Jaccard similarity – a widely adopted strategy to address this kind of approaches
  - Each node has multiple labels
    - \* The ratio of nodes in it that already belongs to  $y$  w.r.t. the size of  $x$ :
      - Ratio is greater than (or equal to) a given threshold, the merge is applied and the node label updated
  - We assume that each node at time  $t$  carries three sets of labels
    - \* The identifiers of the communities it currently belongs to  $t$
    - \* The identifiers of the communities it was part of at  $t-1$
    - \* The identifiers of the communities it will be associated to at  $t + 1$
  - Event detection:
    - \* Birth (B): a community born at time  $t$  if there are no network substructures at  $t - 1$  that can be matched with it
    - \* Merge: two or more communities at time  $t$  merge iff they are matched to the same network substructure at  $t + 1$
    - \* Split (S): a community at  $t$  splits if it is matched to multiple network substructures at  $t + 1$
    - \* Continue (C): a community at  $t$  remains the same at  $t + 1$ ;
    - \* Death (D): a community dies at  $t$  if it is not matched with any network substructure at  $t + 1$ .

## 7 EVALUATION

- $\Psi(\mathcal{A}, \mathcal{B}) = \mathcal{A} \cap \mathcal{B} - (\mathcal{A} - \mathcal{B})$ : Quality metric to relate discovered events  $A$  to ground truth ones  $B$

## 8 CODE

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## 9 RESOURCES

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