

### $\textbf{PART} \ I$

### 1. Introduction & Motivation

- Overview & Contributions
- 2. Software Tools
  - Algorithms Provided
  - Advantages And Disadvantages
  - Metrics Calculations and Results
    - Case Studies
    - Practical Examples
    - Results Processing Time
    - Example Results

### PART II

- 1. Algorithm Developments
  - Green-Marl Language
  - Community Detection Algorithm
  - Similarity Ranking Algorithm
  - Metrics Calculations and Results
    - Case Studies
    - Practical Examples
    - Results Modularity & Processing Time
- 2. Summary & Conclusions



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### Introduction & Motivation

Generic Problem:

Nowadays, the huge amounts of data available pose problems for analysis with regular hardware and/or software.

Solution:

Emerging technologies, like modern models for parallel computing, multicore computers or even clusters of computers, can be very useful for analyzing massive network data.

### **Tutorial Overview & Contributions**

#### **1.** Aggregation of information:

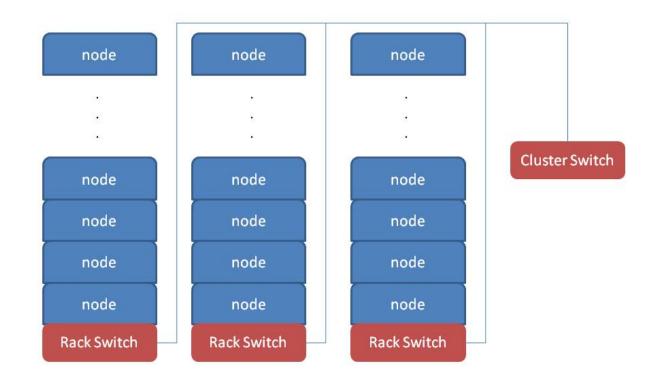
- a. What tools to use for analyzing large social networks
- b. What algorithms are already implemented with these tools
- c. Several Tools Advantages and Disadvantages
- **2. Implementation Example** of algorithms for large scale Social Network analysis and some results:
  - a. Community Detection algorithm implementation with Green-Marl language
  - b. Similarity Ranking algorithm implementation also with Green-Marl language

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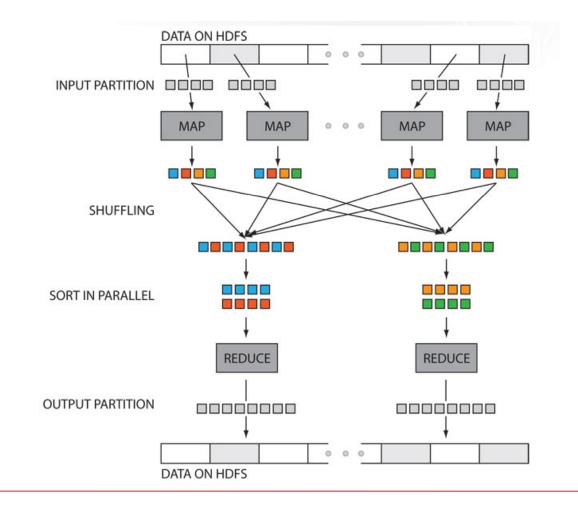
### - To list a few:

- 1. Hadoop Map/Reduce
- 2. Giraph
- 3. Graphlab
- 4. Pegasus
- 5. Green-Marl

#### Hadoop HDFS – Architecture of Compute Nodes



#### Hadoop Map-Reduce



#### Hadoop MapReduce Example – Counting terms in documents

Let start with something really simple. The code snippet below shows Mapper that simply emit "1" for each term it processes and Reducer that goes through the lists of ones and sum them up:

```
class Mapper
  method Map(docid id, doc d)
    for all term t in doc d do
      Emit(term t, count 1)
class Reducer
  method Reduce(term t, counts [c1, c2,...])
    sum = 0
    for all count c in [c1, c2,...] do
      sum = sum + c
    Emit(term t, count sum)
```

#### Hadoop MapReduce Advantages & Disadvantages

ТооІ	Hadoop MR
Advantages	<ul> <li>Ability to write MapReduce programs in Java, a language which even many non computer scientists can learn with sufficient capability to meet powerful data-processing needs</li> <li>Ability to rapidly process large amounts of data in parallel</li> <li>Can be deployed on large clusters of cheap commodity hardware as opposed to expensive, specialized parallel-processing hardware</li> <li>Can be offered as an on-demand service, for example as part of Amazon's EC2 cluster computing service Washington (2011)</li> </ul>
Disadvantages	<ul> <li>One-input two-phase data flow rigid, hard to adapt - Does not allow for stateful multiple-step processing of records</li> <li>Procedural programming model requires (often repetitive) code for even the simplest operations (e.g., projection, filtering)</li> <li>Map Reduce nature is not specially directed to implement code that presents iterations or iterative behavior</li> <li>Opaque nature of the map and reduce functions impedes optimization from Zinn (2010)</li> </ul>

Hadoop Map-Reduce Algorithms (Online Resources):

Highly Scalable Blog

•Log Analysis, Data Querying

- •Graph Analysis, Web Indexing
- •Text Analysis, Market Analysis

atbrox.com website

•Ads Analysis

- •Bioinformatics/Medical Informatics
- •Information Extraction and Text Processing
- •Artificial Intelligence/Machine Learning/Data Mining
- •Statistics
- •Numerical Mathematics
- •Graphs

#### **Algorithms Provided – Other tools**

Software	Pegasus	Graphlab	Giraph	Snap
Algorithms available from software install Parallel computing	<ul> <li>Degree</li> <li>PageRank</li> <li>Random Walk with Restart (RWR)</li> <li>Radius</li> <li>Connected Components</li> </ul>	<ul> <li>approximate diameter</li> <li>kcore</li> <li>pagerank</li> <li>connected component</li> <li>simple coloring</li> <li>directed triangle count</li> <li>simple undirected triangle count</li> <li>format convert</li> <li>sssp</li> <li>undirected triangle count</li> </ul>	<ul> <li>Simple Shortest Path (available from )</li> <li>Simple In Degree Count</li> <li>Simple Out Degree Count</li> <li>Simple Page Rank</li> <li>Connected Components</li> </ul>	<ul> <li>cascades</li> <li>centrality</li> <li>cliques</li> <li>community</li> <li>concomp</li> <li>forestfire</li> <li>graphgen</li> <li>graphhash</li> <li>kcores</li> <li>kronem</li> <li>krongen</li> <li>kronfit</li> <li>maggen</li> <li>magfit</li> <li>motifs</li> <li>ncpplot</li> <li>netevol</li> <li>netinf</li> <li>netstat</li> <li>mkdatasets</li> <li>infopath</li> </ul>
Can user configure number of	YES	YES	YES	NO
cores or machines?				

#### **Advantages & Disadvantages**

Tool	Pegas		Graphla		Gira		Snap	
Advantages	•	Similar positive points to Hadoop MR	•	Algorithms can be described in a node-centric way; same computation is repeatedly performed on every node. Significant amounts of computations are performed on each node. Can be used for any Graph as long as their sparse.	•	Several advantages over Map Reduce: - it's a stateful computation - Disk is hit if/only for checkpoints - No sorting is necessary - Only messages hit the network as mentioned from Martella (2012)	•	Optimized for Graph processing. Written with C++ which is intrinsically considered a fast language
Disadvantages	•	Similar negative points to Hadoop MR	•	Programmability: user must restructure his algorithm in a node centric way. There is an overhead of runtime system when the amount of computation performed at each node is small. Small world graphs: Graphlab lock scheme may suffer from frequent conflicts for such graphs.	•	Still in a very immature phase of development Lack of a complete offered algorithm library	•	Not developed to take advantage of parallel or distributed processing of tasks Some algorithms can be time consuming even for relatively small graphs due to the number of graph characteristics covered (eg. "centrality" algorithm)

#### **Metrics Calculations and Results – Use Case Studies**

•Network A – Relationships Between Tech. Companies and Financial Institutions.

>16.339 vertexes and 30.313 edges.

► Retrieved from Crunchbase API

•Network B – Relationships Between Personalities and Companies.

▶107.033 vertexes and 128.746 edges.

► Retrieved from Crunchbase API

•Network C – Amazon co-purchased products.

➤ 334.863 vertexes and 925.872 edges.

Retrieved from Stanford Large Network Dataset Collection

•Network D – Youtube online social network.

▶1.134.890 vertexes and 2.987.624 edges.

Retrieved from Stanford Large Network Dataset Collection

•Network E – Live Journal online social network.

> 3.997.962 vertexes and 34.681.189 edges.

Retrieved from Stanford Large Network Dataset Collection

#### **Practical Example with Graphlab – Triangle Counting**

😰 110414015@hpcgrid-centos6:~/graphlabapi/release/toolkits/graph_analytics
[110414015@hpcgrid-centos6 graph_analytics]\$ ./undirected_triangle_countgraph
=/home/110414015/Dados/com-lj.ungraph.tsvformat=tsv
This program counts the exact number of triangles in the provided graph.
<pre>INFO: mpi_tools.hpp(init:63): MPI Support was not compiled.</pre>
TCP Communication layer constructed.
<pre>INFO: metrics_server.cpp(launch_metric_server:219): Metrics server now liste</pre>
ning on http://hpcgrid-centos6:8090
INFO: distributed_graph.hpp(load_from_posixfs:1823): Loading graph from file
: /home/110414015/Dados/com-lj.ungraph.tsv
INFO: distributed_ingress_base.hpp(finalize:166): Finalizing Graph
INFO: distributed_ingress_base.hpp(exchange_global_info:493): Graph info:
nverts: 3997962
nedges: 34681189
nreplicas: 3997962
replication factor: 1
Number of vertices: 3997962
Number of edges: 34681189
Counting Triangles
INFO: synchronous_engine.hpp(start:1248): Iteration counter will only output
every 5 seconds.
INFO: synchronous_engine.hpp(start:1263): 0: Starting iteration: 0
INFO: synchronous_engine.hpp(start:1312): Active vertices: 3997962
INFO: synchronous_engine.hpp(start:1361): Running Aggregators
INFO: synchronous_engine.hpp(start:1263): 0: Starting iteration: 1
INFO: synchronous_engine.hpp(start:1312): Active vertices: 0
INFO: synchronous_engine.hpp(start:1373): 1 iterations completed.
Updates: 3997962
Counted in 14.4743 seconds
177820130 Triangles
Metrics server stopping.
[110414015@hpcgrid-centos6 graph_analytics]\$

#### **Case Studies - Metrics and their practical use**

•Triangles – involved in the computation of one of the main statistical property used to describe large graphs met in practice and that is the clustering coefficient of the node.

•K-Core – The concept of a k-core was introduced to study the clustering structure of social networks from and to describe the evolution of random graphs. It has also been applied in bioinformatics and network visualization.

•Friends of Friends – this algorithm is of good application in the commercial data networks where the results could serve as basis for a recommender system.

•Centrality Measures – The centrality measures algorithms have large application in several areas including Psychology, Anthropology, Business and communications, Ecology among many others.

#### **Processing Time**

Processing	Hadoop MR	Pegasus	Graphlab	Snap	
Time       "Friends of Friends"		Degree Measures	Triangles Counting	Centrality Measures	
Network A	rk A 16,040s 5,380s 0,048s		0,048s	374s (06m14s)	
Network <b>B</b> 23,880s		7,070s	0,103s	17400s(4h50m)	
Network C	138,980s	11,050s	0,305s	-[1]	
Network <b>D</b>	430,420s	23,330s	1,211s	-[1]	
Network <b>E</b>	1516,257s	35,680s	16,211s	-[1]	

[1] Value too high

#### **Example Results**

#### 1. Pegasus Degree

#### 2. Friends of Friends

10077	8507:2,17745:1,11077:1,24814:1,85008:1,24937:1,2569:1,2599:1,15721:1,26176:1
1008	73285:1,1469:1,35600:1,247:1,213:1,58475:1,51474:1,7522:1,1991:1,1010:1
1009	14833:1,35600:1,2050:1,11160:1,184:1,2474:1,7313:1,142:1,247:1,73285:1
10099	7613:1,7466:1,109:1,2474:1,12:1,357:1,27658:1,15:1,1135:1,26915:1
101	36:8,15:3,7293:3,26:2,7434:2,513:2,53:2,87:2,6:1,6319:1
1010	7490:4,1875:2,607:2,247:1,35509:1,100:1,1:1,57:1,1008:1,1009:1
1011	939:3,15:3,54:2,7279:2,7377:2,51820:1,5136:1,507:1,5:1,483:1
10116	55775:2,2870:2,39005:2,18924:2,72017:2,26185:1,25966:1,25866:1,25794:1,24768:1
1012	10996:1,1523:1
10120	35585:1,3192:1,31255:1,30752:1,30748:1,30663:1,27754:1,26857:1,26789:1,2665:1
10121	13289:1,11617:1,671:1,18956:1
10127	81082:1,9417:1,813:1,7542:1,7541:1,7227:1,27141:1,24898:1,15759:1,12134:1
10128	59502:1,5822:1,5739:1,56896:1,5344:1,4746:1,4410:1,43497:1,43350:1,4314:1

#### **Example Results**

#### 3. Centrality Measures with Snap

#NodeId	Degree	Closeness	Betweennes	EigenVector	Network Constraint	Clustering Coefficient	PageRank	HubScore	Authority Score
3	80.00	0.233747	1139257.1923 83	0.000461	0.016776	0.000633	0.001181	0.000094	0.029831
843	14.00	0.193071	164648.96552 8	0.000028	0.083915	0.00000	0.000798	0.00000	0.000021
844	16.00	0.207691	287289.05030 9	0.000061	0.071393	0.00000	0.000907	0.00000	0.001772
9	33.00	0.213657	310964.72449 0	0.000223	0.039056	0.00000	0.000361	0.00008	0.015517
1352	9.00	0.181062	96242.573356	0.000015	0.118590	0.00000	0.000539	0.00000	0.000147



### **1. Algorithm Developments**

- Green-Marl Language
- Community Detection Algorithm
- Similarity Ranking Algorithm
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- 2. Summary & Conclusions

#### **Green-Marl Language**

•Green-Marl, a DSL in which a user can describe a graph analysis algorithm in a intuitive way. This DSL captures the high-level semantics of the algorithm as well as its inherent parallelism.

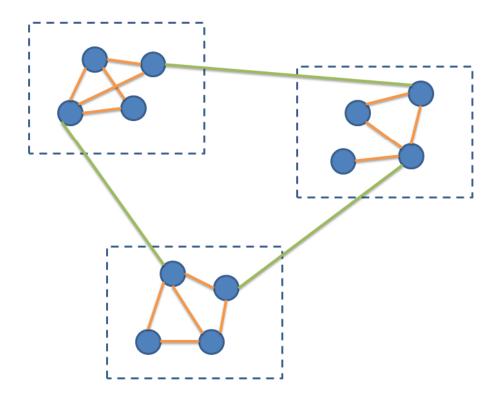
•The Green-Marl compiler which applies a set of optimizations and parallelization enabled by the high-level semantic information of the DSL and produces an optimized parallel implementation targeted at commodity SMP machines.

•An interdisciplinary DSL approach to solving computational problems that combines graph theory, compilers, parallel programming and computer architecture.

#### **Green-Marl Language** - Available Algorithms

Green-Marl Software Algorithms	Brief Description	OpenMP C++ compatible	Giraph/GPS compatible YES	
avg_teen_count	Computes the average teen count of a node	YES		
bc	Computes the betweenness centrality value for the graph	YES	NO	
bc_random	Computes an estimation for the betweenness centrality value for the graph	YES	YES	
communities	Computes the different communities in a graph	YES	NO	
kosaraju	Finds strongly connected components using Kosaraju's Algorithm	YES	NO	
pagerank	Computes the pagerank value for every node in the graph	YES	YES	
potential-friends	otential-friends Computes a set of potential friends for every node using triangle closing		NO	
sssp	SSP Computes the distance of every node from one destination node according to the shortest path		YES	
sssp_path	Computes the shortest paths from one destination node to every other node in the graph and returns the shortest path to a specific node.	YES	NO	
triangle_counting	Computes the number of closed triangles in the graph	YES	NO	

#### **Community Detection**



Simple Graph with 3 communities surrounded with dashed squares.

#### **Community Detection**

•Community detection is known to be a NP-complete problem.

•Community detection can be related to graph partitioning and there are good parallel algorithms for graph partitioning but for community detection it is a usual problem that relies on parallelism achievable from sequential algorithms.

•The top-down approach (divisive approach) or bottom-up approach (agglomerative approach) have inherent sequential flow with possibility of being parallelized on a higher amount on the first stages than the later stages.

•Because of the high computational overhead of community detection algorithms one cannot usually apply such algorithms to networks of hundreds of millions of nodes or edges. Thus, an efficient and high quality algorithm (modularity) for community detection is hard to achieve and a challenging problem as mentioned by Soman and Narang (2011).

**Similarity Ranking Algorithm** 

•SimRank proposed by Jeh and Widom (2002) has become a measure to compare the similarity between two nodes using network structure.

•Although SimRank is applicable to a wide range of areas such as social networks, citation networks, link prediction and others, it suffers from heavy computational complexity and space requirements.

•The basic recursive intuition behind SimRank approach is "two objects are similar if they are referenced by similar objects."

•Being an algorithm with  $O(n^2)$  time complexity where n is the number of nodes in the graph, it is a good choice to develop it in distributed computing environments.

#### **Results – Case Studies**

1. Community Detection Algorithm

#### Networks for Algorithms Modularity Comparison

•Zachary's Karate Club with 34 vertexes and 78 edges.

•Dolphin Social Network with 62 vertexes and 159 edges.

•American Colleague Football with 115 vertexes and 615 edges.

#### Networks for Algorithms Processing Time Comparison

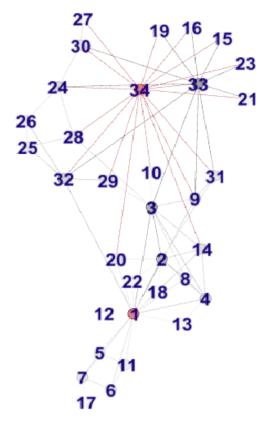
Network A with 16.339 vertexes and 30.313 edges.
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Network C with 334.863 vertexes and 925.872 edges.

2. Similarity Ranking Algorithm

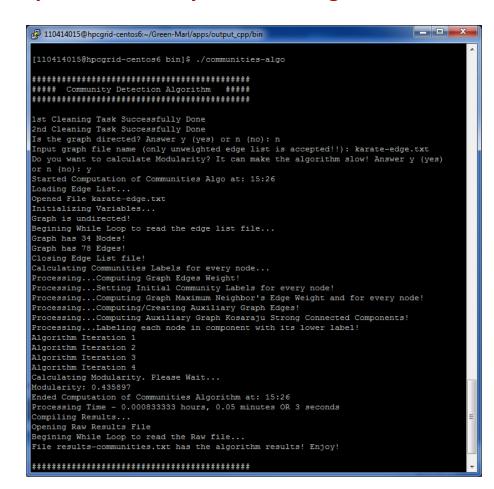
#### **Networks for Sequential vs Parallel Comparison**

•Network F with 471 vertexes and 250 edges.

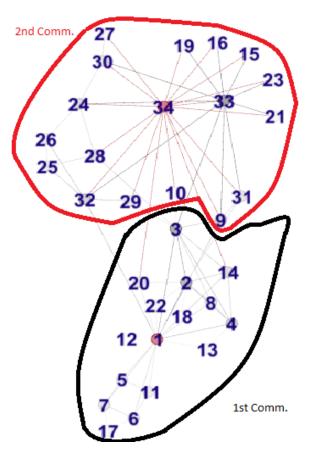
- •Network G with 892 vertexes and 500 edges.
- •Network H with 1.659 vertexes and 999 edges.



Zachary's Karate Club with 34 vertexes and 78 edges.

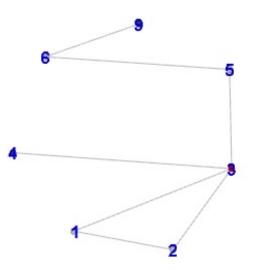


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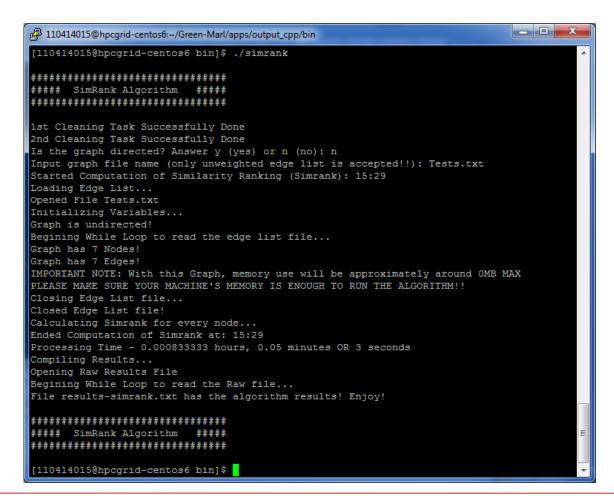
Zachary's Karate Club with 34 vertexes and 78 edges, divided in 2 Communities by the developed algorithm.

#### **Practical Example - Similarity Ranking Algorithm**



Test Network used in the development of the similarity algorithm.

#### **Practical Example - Similarity Ranking Algorithm**



#### **Practical Example - Similarity Ranking Algorithm**

	1	2	3	4	6	5	9
1	1.000000	0.235798	0.168164	0.350434	0.051199	0.209529	0.068624
2	0.235798	1.000000	0.168164	0.350434	0.051199	0.209529	0.068624
3	0.168164	0.168164	1.000000	0.066980	0.177689	0.043468	0.019956
4	0.350434	0.350434	0.066980	1.000000	0.018981	0.353290	0.106580
6	0.051199	0.051199	0.177689	0.018981	1.000000	0.012027	0.005073
5	0.209529	0.209529	0.043468	0.353290	0.012027	1.000000	0.353290
9	0.068624	0.068624	0.019956	0.106580	0.005073	0.353290	1.000000

#### **Community Detection Algorithm – Sequential vs Parallel**

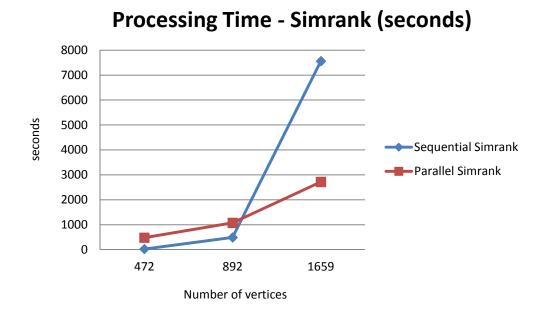
Modularity	Girvan – Newman Algorithm with Snap	Clauset-Newman-Moore Algorithm with Snap	Developed Algorithm with GM
Zachary's Karate Club	0.401	0.381	0.436
Dolphin Social Network	0.519	0.515	0.333
American College Football	0.599	0.549	0.339

Processing Time	Girvan – Newman Algorithm with Snap	Clauset-Newman-Moore Algorithm with Snap	Developed Algorithm with GM
Network A	288 (hours)	бs	4s
Network <b>B</b>	300+ (hours)	53s	133s
Network C	400+ (hours)	*	45659s

#### **Similarity Ranking Algorithm – Sequential vs Parallel**

Processing Time	Parallel Simrank with Green-Marl	Sequential Simrank with R
Network F	480s	25s
Network G	1073s	491s
Network <b>H</b>	2716s	7560s
Network A	26851s	1022000+ s

**Similarity Ranking Algorithm – Sequential vs Parallel** 



# Outline

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### 2. Summary & Conclusions

One of this part of the tutorial goals was to expose which tools to look for when dealing with big graphs studies.

We made the introduction to the tools used nowadays for distributed graph analysis

✤We wrote some practical examples of computing algorithms that leverage the tools potential for big scale graphs studies

Other tutorial goal was to prove the utility and diversity of the tools and algorithms available for graph studies.

✤We learned also that the increasing number of SDLs for big graph analysis make the choice of languages for programming tasks between two generic languages, C++ and Java.

✤The Green-Marl language was also a great tool in the set of tools available and some implementation results are given in this tutorial.

#### **Support Documents**

•"Large Scale Social Networks Analysis" – Thesis

•Document available for download on:

•http://www.ruisarmento.com/uploads/Large\_Scale\_Social\_Networks

<u>Analysis - 2013 - Aftermath.pdf</u>

•Code available for download:

•http://www.ruisarmento.com/uploads/Code.zip

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