Accelerated Graph Features A cache-accelerated method for storing and accessing graphs in C++

Table of Contents

| 1 | Intro | oduct | tion | 2 | |
|---|------------------|-------|---------------------------|------|--|
| | 1.1 | Ove | rview and motivation | 2 | |
| | 1.2 | Inst | allation | 3 | |
| 2 | User manual | | nual | 6 | |
| | 2.1 | Pack | kage structure | 6 | |
| | 2.2 | Cod | e assumptions | 7 | |
| 3 | Developer manual | | | 8 | |
| | 3.1 | Con | version pipeline | 8 | |
| | 3.1. | 1 | Pipeline description | 8 | |
| | 3.1.2 | | Converter function | 8 | |
| | 3.1.3 | | ConvertedGNXReciever | 9 | |
| | 3.1. | 4 | CacheGraph | 9 | |
| | 3.1. | 5 | Example | . 10 | |
| | 3.2 | Cacl | neGraph usage | . 10 | |
| | 3.3 | Writ | ting a Feature Calculator | . 12 | |
| | 3.3. | 1 | Calculator object | . 12 | |
| | 3.3.2 | | Wrapper function | . 14 | |
| | 3.3. | 3 | Python-side function | . 15 | |
| 4 | List of features | | | | |
| 5 | Benchmarks | | . 18 | | |
| | 5.1 Initia | | al Benchmarks | . 19 | |
| | 5.1. | 1 | Clustering Coefficient | . 20 | |
| | 5.1. | 2 | Node Page Rank | . 21 | |
| | 5.1. | 3 | K Core | . 22 | |
| | 5.1.4 | 4 | RFS Moments | 23 | |

1 Introduction

1.1 Overview and motivation

Calculating network features is an expensive business, consisting of calculations that often require traversing the entire network in order to compute the feature.

These days, the problem is magnified because of sheer size of the networks. The networks today can reach sizes of millions of nodes and billions of edges.

The networkx library, which is used to deal with large networks, is simply not fast enough to deal with those kinds of networks in an efficient manner.

To deal with those kinds of networks, a lower level solution must be used.

The solution described below is two-fold:

First, the code is written in C++, with an interface to python. The C++ language gives us the speed and flexibility we need to manage the large networks and expensive operations.

Second, the code is based around an extremely efficient data structure for storing large graphs, first devised by Dr. Lev Muchnik, a senior lecturer in the Hebrew University of Jerusalem. The data structure used arrays to store an adjacency list in such a way that the computer's cache is used extensively, accelerating the access time to the memory.

The manual below is split into two sections:

User manual: a description of the features that can be used in python code.

Developer manual: details of the C++ implementation for future developers using the same code structure.

1.2 Installation

The installation of the package requires several components:

- ➤ The Boost.Python library¹
- ➤ The shared library of C++ code (which needs to be built from source)
- > The python wrapper code

The Boost.Python library can be built by two different methods:

- 1. Installation from source
- 2. Installation using the conda² package manager

While the second method is considerably easier, it is also less flexible than the manual installation from source.

1.2.1 Installation from source

Disclaimer: these installation instructions were tested on linux only, use on other operating systems at your own risk.

An example of the makefile that can be used with this method is *makefile.dsi*.

Instructions:

- 1) Download a compressed version of the Boost libraries. The version used for testing was 1.67, but more advanced versions should also work. The compressed tar.bz2 archive can be downloaded from https://www.boost.org/users/history/version1670.html.
- 2) Unpack the archive into a local folder using the command (make sure to change the version number to the one you downloaded):

The archive will unpack into the current folder and usually create it's own subdirectory.

- 3) Install a version of python3. For testing, the python3 version that is downloaded with Conda was used, but any version should work.
- 4) Run ./bootstrap.sh from the main Boost folder that was just created with the following options:
 - --with-python=[main python path]/bin/python3
 - --with-python-version=3.7 (or whatever your version is)
 - --with-python-root=[main python path]/lib/python3.7 (make sure the version matches the one above).

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¹ https://www.boost.org/doc/libs/1 66 0/libs/python/doc/html/index.html

² https://conda.io/en/master/

Generally speaking, the main python path will be either the /usr directory or the main Conda path (e.g. ~/anaconda3/).

- 5) Run the actual installation:
 - a. Add the python you are using to the correct environment variable:

export CPLUS_INCLUDE_PATH=\$CPLUS_INCLUDE_PATH:[main python path]/include/python3.7m (or whatever directory exists and matched the version you are using.)

- b. Run ./b2
- 6) Update the following values in the makefile (it is recommended to create a copy of *makefile.dsi* and edit it).
 - → Python version (long and short)
 - → Python include and lib directories
 - → Boost include and lib directories

All of these variables can be found in the top of the makefile.

7) Add the same boost lib path the LD_LIBRARY_PATH is ~/.bashrc (which runs whenever you open a shell). Add the following line in ~/.bashrc:

export LD_LIBRARY_PATH=\$LD_LIBRARY_PATH:[main boost path]/stage/lib

8) Build the library by running

make -f your makefile name

From the features_algorithms/accelerated_graph_features/src directory.

1.2.2 Installation using Conda

Disclaimer: Again, these instructions were only tested on linux.

The makefile for these installation instructions is Makefile-conda.

Assuming you already have a version of Conda3 installed:

1) Create an environment from the configuration file env.yml that can be found at

features_algorithms/accelerated_graph_features/src

Move into the directory and run:

conda env create -f env.yml

You should now have an environment called "boost" with all the neccesary packages installed.

2) Build the library by running

conda activate boost

make -f Makefile-conda

1.3 Building the library for with GPU-accelerated features

Building the library with support for GPU features is done on top of a Conda installation.

On a machine where CUDA is installed, *Makefile-gpu* can be used to build the library with GPU features. The modifications needed on a per-basis system are the CUDA links in the makefile. The current configurations is a general case where the cuda libraries are in the /usr/local directory. Modifying the makefile requires changing the lines with the NV links on the top of the makefile.

In order to build the librarary, run the following:

conda activate boost

make -f Makefile-gpu

2 User manual

The code is exposed to the user in Python via a python package.

The C++ code is hidden behind pure python functions that handle the pre- and post-processing for the C++ code.

The exact processing pipeline is described in the next section – Developer manual.

In this section we'll describe the way the user will interact with the code, with examples.

There are two methods of interacting with the package. You can decide to interact with the base functions directly, which allows some flexibility, or you can use the FeatureCalculators which encapsulates the functions, which allows you to easily use the accelerated features in any existing code that already uses the graph measures package.

2.1 Package structure

As we said, the code is encapsulated in a pure python package.

The package can be imported like every other python package by using the following line:

import features algorithms.accelerated graph features.src as af

The package allows immediate access into the feature functions.

All the functions are of the format:

af.function_name(nx_graph, **kwargs)

Where:

- nx graph is a Graph object (or DiGraph) from the networkx³ package
- **kwargs are further keyword arguments needed for the feature calculation.
- **kwargs will always be able to receive the following arguments:
 - → with weights: a boolean specifying whether to add weights to the C++ code.
 - → cast_to_directed: a boolean specifying whether to force to graph into a directed form.
 - → converter_function: if given, replaces the internal converter function, the specifications for such a function are given in the developer section.

The function return either a list (of there is a value returned for each node) or a float (if there is only one number returned from the feature calculation.

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³ https://networkx.github.io/

2.2 Code assumptions

Since the code to convert the nx Graph into a format compatible with the C++ code is internally contained inside the python wrapper package, there are several assumptions made on the structure of the graph passed into the function.

If you wish to specify a different converter function, please look at the relevant section in the developer manual.

The assumptions are as followes:

- a) The nodes are labeled with numbers
- b) Those numbers are the sequence [0,...,num_of_nodes-1]
- c) Currently, the library does not support weighted graphs.

Note that there are no assumptions on whether the graph is directed/undirected, but there is an effect on the result of the code (for details, see the developer section).

3 Developer manual

This section dives into the internal structure of all sections of the code, and is meant both for debugging and extending the code.

3.1 Conversion pipeline

As we are dealing with code from both Python and C++, there is a conversion routine set in place that enables the use of the same graph in both Python and C++.

The conversion takes in (in Python) a Graph or DiGraph object (from the networkx library) and outputs (in C++) a CacheGraph object, which is the format used for cacheaware storage of the graph and access to it in C++.

3.1.1 Pipeline description

The pipeline is described in the diagram to the right.

The input of the conversion is a graph from the networkx package, and the output is a CacheGraph (the graph structure used in the C++ code).

Converter function ConvertedGNXReciever CacheGraph

3.1.2 Converter function⁴

The converter function is a pure python function that receives a nx.Graph as input and returns two or three lists: indices, neighbors [,weights].

A weights list is returned only if weights were requested (by setting with weights).

The default converter function is located in the *graph_converter.py* file.

For those who wish to write their own converter function, here are a few guidelines to follow:

- ✓ The function should be able to determine whether the graph is directed (or if a directed version is requested) and act accordingly.
- ✓ The function should handle the case for both weighted and unweighted graphs
- ✓ The function should not break for:
 - A node without neighbors
 - An unweighted graph where the user asked for weights
- ✓ The function should notify the user when working on a directed graph.

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⁴ Can be found in graph_converter.py

3.1.3 ConvertedGNXReciever⁵

This is a C++ object that has dual responsibilities:

- Convert a dictionary of lists (that was created by the Python converter) into a CacheGraph. A pointer to the created CacheGraph is saved in the Reciever.
- ❖ Memory management: the graph will be deleted alongside to converter at the end of the converter's scope. This will usually occur at the end of the wrapper function (that is exposed to Python) and so the graph will be cleaned automatically at the end of the calculation.

3.1.4 CacheGraph⁶

The CacheGraph object contains 3 private members:

- 1) NumberOfNodes: an unsigned int that holds the number of nodes in the graph
- 2) Graph: a vector of unsigned ints which holds a sorted neighbor list for each node.
- 3) Offsets: a vector of type **int64** (which is our shorthand for **unsigned long long** in linux) that holds at position *i* the offset of the first neighbor of node *i* in the *Graph* array.

Access to the graph is conducted by iterating both of the arrays (see the CacheGraph usage section below).

⁵ Can be found in the utils directory of the C++ code

⁶ Can be found in the *arch* directory of the C++ code

3.1.5 Example⁷

There follows an example of the conversion routine for a simple graph.

The given graph is the one composed of these edges: (0->1, 0->2,0->3,2->0,3->1,3->2).

The behavior of the conversion now depends on whether the graph is directed (either by passing in a nx.DiGraph or by specifying *cast_to_directed* as true) or undirected.

If the graph is directed, the result is as follows:

Indices: [0, 3, 3, 4, 6] Neighbors: [1, 2, 3, 0, 1, 2]

Else, the results for the undirected graph are:

Indices: [0, 3, 5, 7, 10]

Neighbors: [1, 2, 3, 0, 3, 0, 3, 0, 1, 2]

It is worth noting a few things:

- Notice that the last element in the indices list actually contains the number of edges in the graph (i.e. the length of *neighbors*). This fact will be used in the C++ code.
- The length of the *neighbor* list in the undirected version isn't double the length of the undirected one, and that is because a bidirectional edge (0<->2) already exists in the directed graph.
- Note the *indices* list of the directed version. Since node 1 has no neighbors (it doesn't point to any other node), it's corresponding neighbor list section is empty and so the index in the *indices* list doesn't increment).

3.2 CacheGraph usage⁸

The CacheGraph object is designed around the principle of cache-awareness. The most important thing to remember when accessing the graph in the C++ code is that we are aiming to accelerate the computations by loading sections of the graph into the cache ahead of time for quick access. When using the CacheGraph, this comes into effect when we iterate over the offset vector first and then access the blocks of neighbor nodes in the graph vector.

By doing this, we are pulling the entire list of a certain node's neighbors into the cache, allowing us to iterate over them extremely quickly.

A concrete example of a good use and bad use case of the CacheGraph are the two popular search strategies BFS and DFS.

⁷ This example can be found in the test python converter.py file.

⁸ Example code based on the code in the clustering coefficient feature calculator

When using BFS we are utilizing the full power of the cache-aware storage mechanism by pulling in the entire list of a node's neighbors to be processed at once, which is fast due to the fact that the neighbors are in the cache.

In contrast, when using DFS, we are not going over all of a node's neighbors at once but jumping from one node to the other, which completely nullifies the speed advantage that the cache can give us, as the contents of the cache need to be constantly swapped out by the proccessor.

There is a general code structure used when dealing when a CacheGraph that is designed as part of our cache-aware strategy:

```
// This snippet is taken out of the clustering coefficient feature calculator
    //There is usually no need to declare the CacheGraph manualy, as it is a
   member variable of the FeatureCalculator class.
   CacheGraph* mGraph;
    //Get the neighbors and offsets from the graph
    const unsigned int* neighborList = mGraph->GetNeighborList();
    const int64* offsetList = mGraph->GetOffsetList();
    // For readability, we save the begining and the end of the current
   neighbor list at each step.
    int64 begin offset, end offset;
    // We start by iterating over all the nodes in the graph
    for (unsigned int i = 0; i < numOfNodes; i++) {</pre>
              // Below are the edges of the neighbor list for the current
      node (including begin offset, excluding end offset)
       begin offset = offsetList[i];
       end offset = offsetList[i + 1];
        // Now we can iterate over all of the neighbors.
        // It is important to note that behind the scenes a large section of
        the neighbor list is already
        // put in the cache by the os, which means our access time is reduced
        dramatically.
        for (auto p = neighborList + begin offset; p < neighborList +</pre>
        end offset; ++p)
            //p is a pointer to the current node in the adjacency list.
            //Example usage: the offset of the current node is offsetList[*p]
            //Another example: check if there are neighbors inside the
            neighbor list of the current node.
            for (auto q = neighborList + begin offset; q < p; ++q) {</pre>
            //We need to go over all the pairs, but we don't want to count
            every edge twice, so we only iterate over q<p.
                if (mGraph->areNeighbors(*p, *q))
                    connectedNeighbors++;
        }
    }
```

3.3 Writing a Feature Calculator⁹

Writing a feature calculator consists of two parts:

- Creating a class that inherits from the FeatureCalculator¹⁰ class (or one of its subclasses, such as NodeFeatureCalculator, GPUFeatureCalculator, etc.).
- Creating a wrapper function that will be exposed to the user in Python
- Creating a pure python function that will perform the preprocessing necessary from the Python side.

3.3.1 Calculator object

The main body of code for the feature calculation will be held in this class.

As a class that inherits from FeatureCalculator, the skeleton of the code is already build, enabling the developer to focus on the calculation itself.

The FeatureCalculator class is a template class that gives you the following:

- 1. Access to a **const CacheGraph*** that is the CacheGraph given to the Calculator.
- **2.** The need to only implement the *Calculate* function, which will be called by the wrapper function.

Notice that when you extend the base FeatureCalculator class, you must specify the template argument for it.

In the next page there is an example of a skeleton version of a FeatureCalculator class, which includes the basic requirements needed in each FeatureCalculator.

https://www.boost.org/doc/libs/1_66_0/libs/python/doc/html/tutorial/tutorial/techniques.html#tutorial.techniques.reducing compiling time

⁹ Here we will be building the python side package as described in the Boost.Python manual:

¹⁰ Can be found in the *arch* directory of the C++ code

```
#ifndef EXAMPLEFEATURECALCULATOR_H_
#define EXAMPLEFEATURECALCULATOR H
#include "FeatureCalculator.h"
// Notice that we're returning a float from this feature calculation,
// it could just as easily have been a vector<float>
class ExampleFeatureCalculator: public FeatureCalculator<float> {
public:
   ExampleFeatureCalculator();
   // This is the only function you actually need to implement!
   virtual float Calculate();
   virtual ~ExampleFeatureCalculator();
   // You can add whatever private variables\methods you want!
};
#endif /* EXAMPLEFEATURECALCULATOR H */
#include "../includes/ExampleFeatureCalculator.h"
ExampleFeatureCalculator::ExampleFeatureCalculator() {
   // If there is any additional preprocessing you would like to perform,
   // this is the place for it.
}
* The actual calculation goes here
float ExampleFeatureCalculator::Calculate() {
   return (float) mGraph->GetNumberOfNodes();
ExampleFeatureCalculator::~ExampleFeatureCalculator() {
   // Don't forget to release any memory you may have allocated!
```

3.3.2 Wrapper function

The code for the wrapper function is usually written in a separate C++ file, under the wrappers directory. Truth be told, there are two wrapper functions – one which actually wraps the Feature Calculator, and another that defines it for the Boost.Python module.

An example of the two can be found below:

```
#ifndef CALCULATORWRAPPEREXAMPLE H
#define CALCULATORWRAPPEREXAMPLE H
#include <boost/python.hpp>
#include <boost/python/dict.hpp>
#include "../includes/ConvertedGNXReciever.h"
#include "../includes/ExampleFeatureCalculator.h"
using namespace boost::python;
void BoostDefExampleCalculator();
void ExampleCalculatorWrapper(dict converted graph);
#endif
#include "ExampleWrapper.h"
void BoostDefExampleCalculator() {
  def(ExampleCalculatorWrapper, "example feature");
float ExampleCalculatorWrapper(dict converted graph) {
  ConvertedGNXReciever reciever (convertedGraph);
  ExampleFeatureCalculator calc();
  calc.setGraph(reciever.getCachedGraph());
  return calc.Calculate();
  // At this point in time the graph is also deleted
}
//This is the file which is compiled into an .so
#include <boost/python.hpp>
#include "../wrappers/ExampleWrapper.h"
// ... other imports ...
BOOST PYTHON MODULE ( features)
   // ... other boost def wrappers ...
  BoostDefExampleCalculator();
}
```

Note that if the feature returns a vector, it is the responsibility of the wrapper function to convert it into a python list (boost/python/list.hpp).

The separation of the boost definition functions into different files is important. The compilation against Boost.Python is the most time-expensive compilation, and we don't want one change in a single calculator to cause us to recompile all of the wrapper functions.

3.3.3 Python-side function

The python side function is responsible for both pre- and post-processing for the feature calculator.

Pre-processing consists of converting the graph into a dictionary of lists (as done by the functions in the *graph_coverter.py* file) and of any other additional transformation needed to adapt the rest of the parameters to a format compatible with the C++ code.

Post-processing can include any action necessary to convert the results of the C++ code into a Python-friendly format.

Luckily for us, since most of the actions in the python side function are the same for all exposing functions, there is a python decorator¹¹ designed to save you the trouble of writing the same code over and over¹².

```
This code is in a pure python file which contains all the wrapper functions.
from feature wrapper decorator import FeatureWrapper
from test python converter import create graph
@FeatureWrapper
def example feature(graph, **kargs):
    # example feature is a C++ function exposed to python
    from features import example_feature
   print(graph['indices'])
   print(graph['neighbors'])
    # Here 0 is the default value for the argument
    example arg = kargs.get('example arg', 0)
    res = example feature(graph, example arg)
    # Any post-processing goes here
    return res
# Usage is as follows:
result = example feature (create graph(), example arg=42)
```

https://www.thecodeship.com/patterns/guide-to-python-function-decorators/https://www.artima.com/weblogs/viewpost.jsp?thread=240808

¹¹ The Feature Wrapper decorator can be found in the feature wrapper decorator.py

¹² For more on python decorators, see:

Notice that the python function must have **two and only two** arguments:

- → *Graph:* the user passes in the nx.Graph, but because of decorator magic, inside the function it is a converted graph dictionary
- → **kwargs: any other arguments must be keyword arguments. Among them may be with_weights, cast_to_directed and converter_function.

In this code example you can also see the name of the compile C++ shared library is _features, which in the context of the python package means that the code is invisible to any code outside the function.

All function which are exposed to the outside python code (i.e. the python wrapper functions) are visible only because they are imported in the __init__.py file of the package.

4 List of features

There follows a list of all the features available in the library.

Notes about the table:

- Name is the name of the python wrapper function (which is the one the user calls).
- **Arguments** are all the arguments given to **kwargs (in python) i.e. not including the graph itself.
- **Return type** is the python type returned (e.g. list and not vector)
- **Notes** contains any other assumption and/or conditions for the feature (for example, that the graph must be undirected).

| Name | Arguments | Return type | Notes |
|-----------------|---|---|------------------------|
| node_page_rank | dumping: float, iterations: int | List of floats | |
| k_core | | List of ints (unsigned) | |
| bfs_moments | | List of tuples (float,float) | |
| 3- and 4-motifs | level: int gpu: boolean, whether to use the GPU version | A matrix of size num_nodes X num_motifs | Also has a GPU version |
| flow | threshold:float | List of floats | |
| Attractor_basin | alpha:int | List of floats | |

5 Benchmarks

Since the entire point of creating this package is to enable faster computations of graph features, below we include some benchmark speed tests for the various features.

The purpose of showing the benchmarks is twofold:

- 1) Prove the necessity of having faster computation for large graphs
- 2) Allow the user to know when it is better to use the python version of the feature calculation, as using a feature calculator from this package involves some overhead (such as the conversion routine and access to the shared library).

The features will be timed for graphs of increasingly larger size (in both the number of nodes and the number of edges).

The graphs used will be random graphs (specifically, erdös-reyni graphs generated by the networkx package¹³). Since we are using random graphs, we will use several different graphs for each benchmark, taking the average of the time. The graphs will be generated by a predetermined set of seeds, so as to use the same graphs for all benchmarks.

The python features will be calculated either by internal functions in the networkx package or by the feature calculators from *graph-measures*.

¹³ https://networkx.github.io/documentation/networkx-

^{1.10/}reference/generated/networkx.generators.random graphs.erdos renyi graph.html#networkx.generators.random graphs.erdos renyi graph

5.1 Initial Benchmarks

As a first test of the viability of the project each feature was tested over a set of random graphs with an increasing number of nodes.

To maintain consistency between all tests, the graphs were erdös-reyni graphs generated by the NetworkX function¹⁴ using a constant random seed set to 123456.

The number of nodes was chosen from the following: [50, 100, 300, 500, 1000, 2000, 3000, 4000, 5000]. The p value for the graphs was a constant 0.8.

The results of the initial benchmarks are extremely promising.

The four features in this benchmark were: Clustering Coefficient, Node Page Rank, K Core and BFS Moment.

The first three were tested against the corresponding functions in the NetworkX package¹⁵, and the latter to the existing code in the *graph-measures* GitHub repository.

The calculation of the feature in C++ is made up of two distinct parts:

- 1) Conversion: converting the Graph object of the networkx package into the format used by the C++ code.
- 2) Feature calculation: the actual calculation of the feature in C++.

Each part is timed separately, as the conversion is only needed once per graph (and afterward the converted graph can be written to a binary file that can be loaded directly into the C++ code), and the feature needs to be calculated on its own.

On the other hand, the python code has no conversion of the graph, and so only has one time measurement.

The actual benchmarking code can be found in the *feature_benchmarks.py* file, and the timer used for timing the benchmarks is in the *graph_timer.py* file.

Clustering Coefficient: https://networkx.github.io/documentation/networkx-

1.9/reference/generated/networkx.algorithms.cluster.average_clustering.html

Node Page Rank: https://networkx.github.io/documentation/networkx-

1.10/reference/generated/networkx.algorithms.link analysis.pagerank alg.pagerank.html

K Core: https://networkx.github.io/documentation/networkx-

1.7/reference/generated/networkx.algorithms.core.k core.html

¹⁴ https://networkx.github.io/documentation/networkx-

 $[\]underline{1.10/reference/generated/networkx.generators.random\ graphs.gnp\ random\ graph.html\#networkx.generators.r}$ $\underline{andom\ graphs.gnp\ random\ graph}$

¹⁵ The functions' docs can be found in the following links:

5.1.1 Clustering Coefficient

We'll start with the worst results.

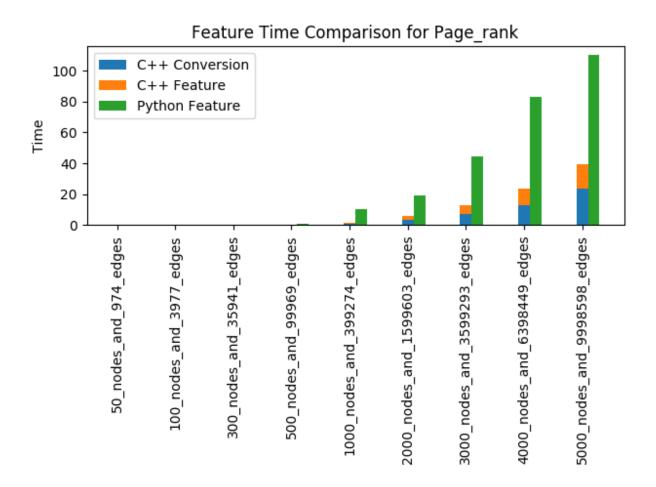
The clustering isn't the ideal feature for the cache-aware format we're using, and as such the time saved, while not negligible, is mainly due to the fact that the C++ language code runs faster than Python.



It can also be seen that in this case, as the feature time is $O(n^2)$, the conversion time is irrelevant.

5.1.2 Node Page Rank

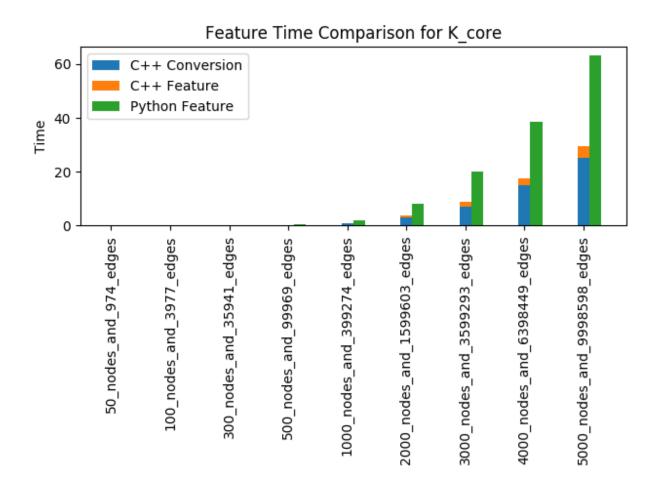
This feature and the ones following are linear by the number of nodes and utilize out cacheaware graph extremely well. As such, the gain of the C++ feature is much greater, especially when considering the face that the graph can be converted ahead of time, saving the conversion time in each feature.



In this case, as the feature itself takes less time, the cost of performing the conversion each time can be seen. This demonstrates a case where converting the graph ahead of time can constitute a large time saver.

5.1.3 K Core

This case is an even better improvement that the Page Rank, again if the graph is preconverted.

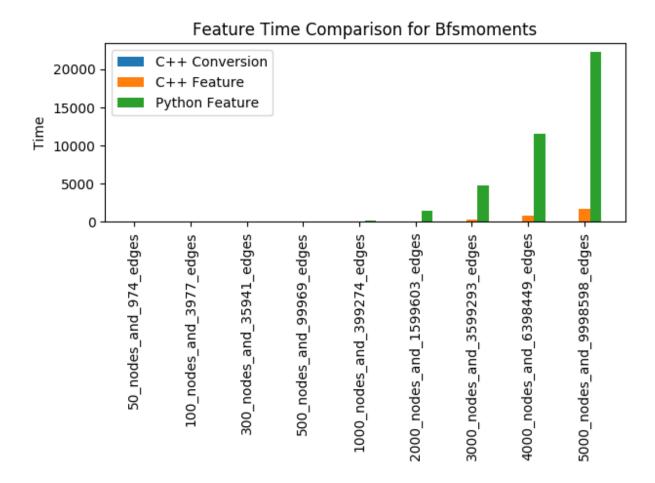


5.1.4 BFS Moments

In comparison to the last two features, this feature takes much longer.

This attribute has two consequences:

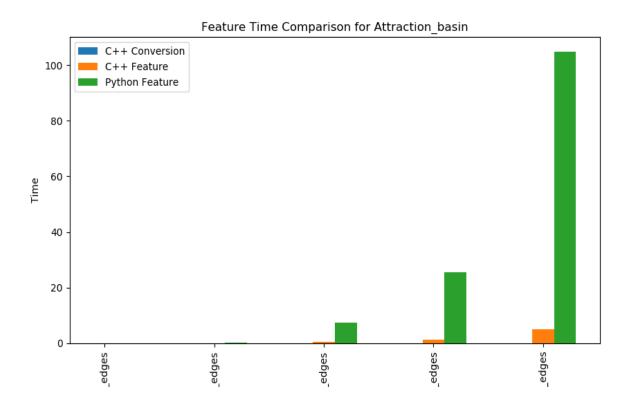
- 1) The conversion time, in comparison with the time it takes to calculate the feature, is again negligible.
- 2) The gain of the C++ code is much greater than the previous features, as it has a longer time to take advantage of the cache-aware graph format.



5.1.5 Other BFS based features

As with the previous feature, these two features (attraction basin and flow), give us a 10X boost over the corresponding python code.

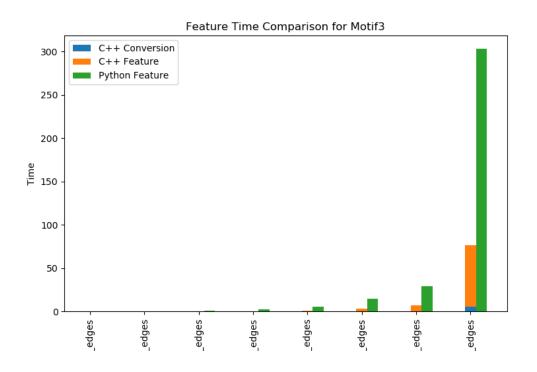
It should be noted that these features were tested on much more powerful servers than the previous features, and so are not in the same time scale as them. The speed improvement given by the C++ code, though, can still be clearly seen.

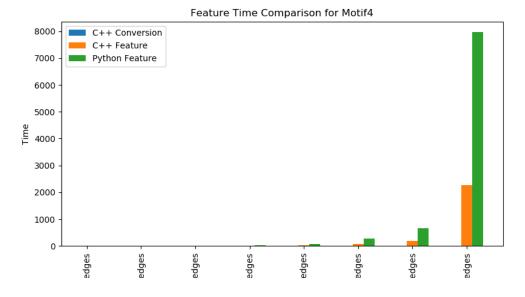


5.1.6 Motifs

The motif algorithm is one of the most commonly used algorithms in the package.

While the C++ code in this case does not give us the 10X boost the BFS features give us, it still gives us a respectable 3X-4X boost.





In addition to the usual C++ code, there also exists a GPU version for the motif code.

The GPU gives us an additional 10X boost **over the C++ code**, which results in a total of up to 50X over the python code. It can also be seen that the GPU code requires an overhead, making it unsuited for small graphs.

