

Report of *Process Mining* Project

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The purpose of this report is to present the *Project 1: How much variable is my event log?* carried out for the Process Mining course at University of Padua.

Introduction

The aim of this report is to sum up the Process Mining project. In this project we aim at measuring how much variable are event logs, in terms of variety of behavior. Computing the variability of a log can indeed be rather useful, for instance, to decide for (a procedural or a declarative) model discovery, or to decide which prediction technique to apply. The are a lot of different ways to measure the variability of an event log, three of them are described in the following sections.

A possible way to measure the variability of an event log is counting the number of variants that it contains. A second possibility is to average the edit distance between each pair of traces in the event log.

The report is organized as follows: in Sec. , ?? and ??, the three ways to compute variability of a log have been proposed. For each section there are two subsections that explain advantages and disadvantages of the metric described in the section.

First approach

The first and easiest way to measure variability of an event log is counting the number of variants that it contains. We recall the definition of event, traces, process variant and process log: an event is an occurrence of an activity in a particular process instance, a trace is a sequence of events of the same such instance, a process variant is a sequence of process activities and a log is a multiset of such traces. The following function compute this metric:

Code snippet 1: Function compute_variant_variability

```
def compute_variant_variability(log: lg.EventLog) -> int:
    """Compute the number of variants present in the event log

Args:
log (lg.EventLog): The log to examine

Returns:
int: The number of variants present in the event log
"""

# For each case of the log we construct a tuple with the names of events.
```

```
# Then we collect all of them in a set to remove duplicates.
# Eventually we compute the length of the set.
return len(set(tuple(event["concept:name"] for event in case) for case in log))
```

This is the simplest and intuitive way to compute log variability. However, this metric does not consider the size of the log, i.e. big size log are penalized with respect to small size log. There is also a second reason that this way is unhelpful to our goal: it does not consider how variants differ to one another.

Example: If we have a process $\log L_1 = [\langle a,b,c \rangle^{10}, \langle b,c,d \rangle^{20}]$ the function returns 2 because in the log there are two different variants. If the log is $L_2 = [\langle a,b \rangle, \langle b,c \rangle, \langle c,d \rangle]$ the function returns 3; but L_1 is bigger than L_2 . This example shows how poorly informative this metric.

Second approach

In order to obtain a more informative metric to compute log variability, the second approach that we developed consists in averaging edit distance between each pair of traces in the event log. For this purpose, we decided to use the Levenshtein distance. Levenshtein edit distance is the most well-known string edit distance metric and it is defined by the number of insertions, deletions and substitutions required to convert one string into another. The basic idea to compute average edit distance is, first compute the sum of edit distance of all possible combination of two pair of traces and then divide it by the number of possible combination that is size_of_log * (size_of_log - 1).

Code snippet 2: Function compute_edit_distance_variability

```
def compute_edit_distance_variability(log: lg.EventLog) -> float:
    """Compute the average edit distance (Levenshtein distance) between each
    pairs of traces.

This function uses function `eval` of module 'editdistance' because it's
    implemented in C++ and it is faster than the corresponding implementation
    in python.

Args:
```

```
log (EventLog): The log to examine
11
   Returns:
   float: the average edit distance between each pairs of traces
13
14
    # Create a dictionary which contains variants as keys and its number of
    # occurrences as values
16
   variants_and_counts = Counter(
   tuple(event["concept:name"] for event in case) for case in log
18
19
   size_of_log = len(log)
20
21
    # For each pair of distinct variants (obtained by 'combinations') we
22
    # compute the edit distance and we multiply it for number of
23
    # occurrences of the variants.
    # In the end we sum all of them.
25
   sum_of_distances = sum(
   num_of_items_1 * num_of_items_2 * editdistance.eval(variant1, variant2)
27
   for (variant1, num_of_items_1), (variant2, num_of_items_2)
28
   in combinations(variants_and_counts.items(), 2)
29
   )
30
31
    # We multiply the sum of distances by 2 because to add the sum of distances
32
    # of each inverted pairs of variants.
    # In the end we divide it by the number of possible combination of pair
34
    # of traces
   return float(sum_of_distances * 2) / (size_of_log * (size_of_log - 1))
```

Some important caveats are:

- we use variants instead of the traces because, instead of calculating edit distance between two traces n times, we calculate the edit distance of two different traces multiplying by the number of repetitions of a trace (line 26 of the snippet 2).
- combination method of the python library itertools provide all the combinations of length two of different variants (line 29 of the snippet 2); the only case not covered by combinations is when two traces have the same variant but the edit distance is 0, so it does not affect the result (because the sum of all distances does not change summing 0).

• To optimize the calculation of edit distance, we use a C++ library with API for python because the above function has high complexity and with very big log (e.g. BPIchallenge2011.xes) it takes very long time¹.

Advantages

The main advantage is that this metric solves the problems of the metric described in Sect. . In fact:

- it takes into account the size of the log. Unlike the previous metric, this doesn't penalize big size logs.
- it considers how variants differ one another.

Disadvantages

This method has also the following disadvantages:

- it is very time-consuming for big size logs due to the fact that it has a high complexity.
- it does not consider the length of each trace, that's why logs with long traces are penalized).

Example: Consider two logs $L_1=[\langle a,b \rangle, \langle c,d \rangle]$ and $L_2=[\langle a,b,c,d,e \rangle, \langle a,b,c,f,g \rangle]$. compute_edit_distance_variability returns 2 for both logs but L_1 has a higher variability w.r.t. L_2 that has the first three events equal in both traces.

Third approach

To address the weakness of the metrics based on edit distance, we developed a third way that consists in normalization of edit distance. We normalise edit distance by the greatest possible distance between the traces to reflect that a distance of one operation on two

¹We know that computing edit distance between two traces has complexity O(nm) where n and m are the lengths of the traces. The whole function has a complexity of $O(l^2mm)$ where l is the number of variants in the event log.

very long strings should be considered less significant than on very short strings. In this way, we are able to compare the result with logs with a different average length.

Code snippet 3: Function compute_my_variability

```
def compute_my_variability(log: lg.EventLog) -> float:
    """Compute the average of normalized edit distances (Levenshtein distance)
    between each pairs of traces.
    This function uses function `eval` of module 'editdistance' because it's
    implemented in C++ and it is faster than the corresponding implementation
    in python.
    Args:
            log (EventLog): The log to examine
10
11
    Returns:
12
            float: a number between 0 and 1 included. The higher the number the
13
            more similar the traces are
14
                    - 0 : if all traces has nothing in common
15
                    - 1 : all traces belongs to the same variant (are equals)
16
    11 11 11
17
    # Create a dictionary which contains variants as keys and its number of
    # occurrences as values
19
    variants_and_counts = Counter(
20
                            tuple(event["concept:name"] for event in case) for case in log
22
   size_of_log = len(log)
23
24
    # For each pair of distinct variants (obtained by 'combinations') we
25
    # compute the average edit distance normalized (divided by longest trace)
26
    # and we multiply it for number of occurrences of the variants.
27
    # In the end we sum all of them.
28
   sum_of_distances = sum(
29
            float(num_of_items_1 * num_of_items_2 * editdistance.eval(variant1, variant2))
            / max(len(variant1), len(variant2))
31
            for (variant1, num_of_items_1), (variant2, num_of_items_2)
            in combinations(variants_and_counts.items(), 2)
33
    )
34
35
    # We multiply the sum of distances by 2 because to add the sum of distances
```

```
# of each inverted pairs of variants

# In the end we divide it by the number of possible combination of pair

# of traces

# return 1 - (sum_of_distances * 2 / (size_of_log * (size_of_log - 1)))
```

Advantages

This metric produces a result in the range [0, 1], that is immediately interpretable because it represent the fraction of how traces are equals. In fact, the higher the number is, the more similar the traces are. When this number is 0, all traces of a log have nothing in common. When it is 1, all traces of a log have the same variant. In conclusion, this metric solve the problem .

Example: Consider once again the two log of the previous example ($L_1=[\langle a,b \rangle, \langle c,d \rangle]$ and $L_2=[\langle a,b,c,d,e \rangle, \langle a,b,c,f,g \rangle]$ reported for convinience). compute_my_variability function returns 0 for L_1 which traces are completely different and 0.6 for L_2 which traces are somehow similar.

Disadvantages

The main disadvantage of this metric is the that it does not consider cycles in traces, i.e. traces with a repeated number of event are wrongly penalized. See the following example.

Example: This metric penalize the following traces A = ABCBCBCE and B = ABCE.

BPI Challenge 2011

In this section, we report and discuss the results of the three different metrics realized. The following table summarize the results.

Functions	Results
$compute_variant_variability$	981
$compute_edit_distance_variability$	195.88194492325937
$compute_my_variability$	0.14258102346211654

Table 1: Table of overall results

As you can see in Table 1, the result obtained applying compute_variant_variability is 981. This number indicates that BPIChallenge2011 log has a lot of variants. We verified this data importing the log on Disco and w.r.t. size of log (1143), the only conclusion is that most of traces are different one another. Eventually, this metric is too simple too capture an interesting variability that involves how different are this variants.

The second result obtained applying compute_edit_distance_variability is 195.88194492325937. This number is more informative w.r.t the previous result and it tell us that there is a big average edit distance between pair of traces, hence, the log has an high variability. But you cannot understand how much traces have something in common.

The last result obtained by applying compute_my_variability is 0.14258102346211654. This number gives more information than the others, because it tells how much two traces, random picked, have in common. In this case, the result is 15% which means that traces are disequal (hanno poco in comune) and we can conclude that the log has high variability.

Instructions

- 1. Extract Progetto.zip
- 2. Open a terminal inside the folder
- 3. Install requirements using the following command: pip install requirements.txt
- 4. Run the tests using: pytest
- 5. In file main.py you can find an use example and you can run it: python main.py