

REPORT

Hyper clique Pattern Discovery

Following is the list of work performed for the Hyper clique pattern Discovery paper :

- Implemented Hyper clique Miner Algorithm (code attached)
- Proof of correctness using association pattern mining examples
- Executing algorithm on **PUMSB dataset**
- Examining the number of patterns generated by hyper clique miner for various minimum support threshold
- Examining the execution time of hyper clique miner for various minimum support threshold
- Examining the number of patterns for various number of attributes
- Examining the execution time for various number of attributes (hyper clique miners)
- Conclusions drawn

Hyper clique Miner Algorithm Implementation

Below is the list of functions used in the algorithm (Complete code has been attached):

def calc_hc(item) : This function calculates h-confidence for a given item set.

def calc_sup(item) : This function calculates the value of support for a given item.

def aprioriGen(Lk, k) : This function returns C_{k+1} from the given L_k .

def antimonotone() : This function returns patterns after pruning using anti-monotone property.

def cross_support() : This function returns patterns after pruning using cross-support property.

Correctness Proof for the algorithm implementation using association pattern mining example

I have taken the example given on page number 19 of the research paper.

TID	Items
1	1
2	2
3	3, 4
4	1, 2
5	1, 2
6	1, 2
7	1, 2, 3, 4, 5
8	1
9	2
10	3, 5

Minimum support = 0.0

h-confidence threshold = 0.6

myfunc(0.0,0.6)

https://colab.research.google.com/drive/1eSKi8AsraHwQ2qr5lSmWLPbeE-KSQI#scrollTo=tORv_EKImRDf&printMode=true

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Final_output_DM3.ipynb - Colaboratory

```
[ ] [frozenset({1}), frozenset({2}), frozenset({3}), frozenset({4}), frozenset({5})]
2  [frozenset({1, 2}), frozenset({3, 4}), frozenset({3, 5})]
3  []
   []
   =====
   /usr/local/lib/python3.6/dist-packages/ipykernel_launcher.py:22: FutureWarning: get_valu
8
```

Please refer **Example_execution.pdf** attached for the detailed implementation.

Executing Algorithm code on PUMSB dataset

Since PUMSB dataset was very big , I have done sampling (2k rows) of the given data and executed code for the same .

- Min_sup=0.4
- h-confidence_threshold=0.7
- Total number of patterns found after pruning =14644

Output Screen shot :

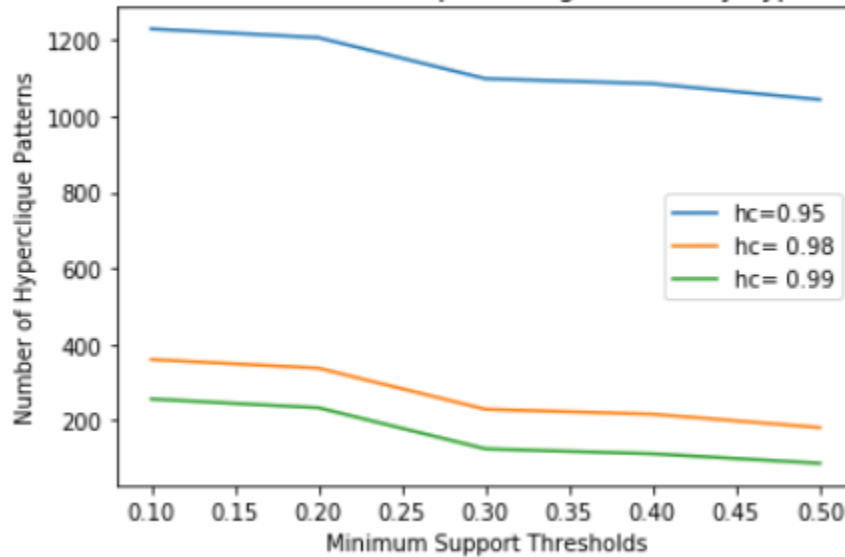
```
myfunc(0.4,0.7)

[ ]> [frozenset({14}), frozenset({15}), frozenset({17}), frozenset({66}), frozenset({84}), fr
2
/usr/local/lib/python3.6/dist-packages/ipykernel_launcher.py:23: FutureWarning: get_valu
{frozenset({3403, 4493}), frozenset({4499, 4414}), frozenset({4434, 7092}), frozenset({1
3
{frozenset({4680, 4786, 4518}), frozenset({180, 4430, 4428}), frozenset({168, 4499, 4436
4
set()
=====
14644
```

NOTE : Please refer **PUMSB_output.txt** file attached for the generated patterns and **Final_output_PUSMB.pdf** for the complete code execution.

Examining the number of patterns generated by hyper clique miner for various minimum support threshold

On the Pumsb data set Number of patterns generated by hyperclique miner

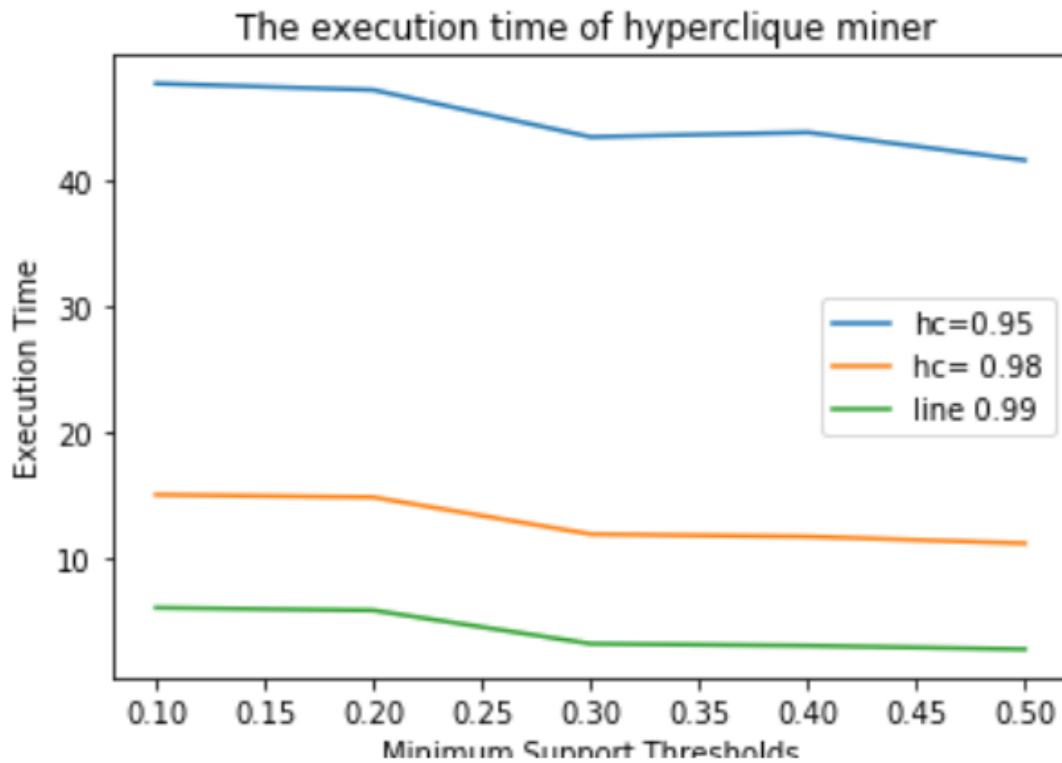


Inferences Drawn :

- Increasing the value of h confidence threshold results in lesser number of hyper clique patterns.
- *Number of hyper clique patterns decreases with increase of min support values for a constant h confidence threshold.*
- Number of hyper clique patterns significantly increases by slight decrement of h conf threshold value.

Please find the implementation details in **minsup_vs_patterns.pdf** attached.

Examining the execution time of hyper clique miner for various minimum support threshold

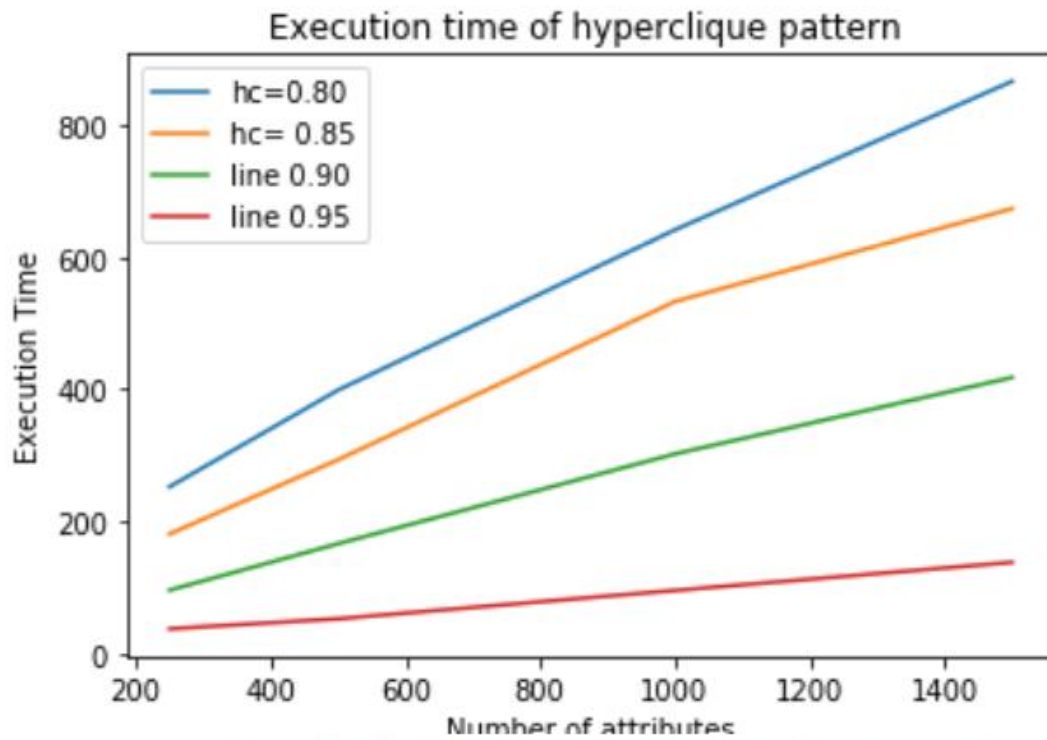


Inferences Drawn:

- Smaller value of H confidence threshold results in larger number of hyper clique patterns. Therefore execution time increases.
- Execution time decreases with increase of min support value for a constant value of h confidence.

Please refer **min_sup_vs executiontime.pdf** for necessary details.

Examining the execution time for various number of attributes (hyper clique miners)

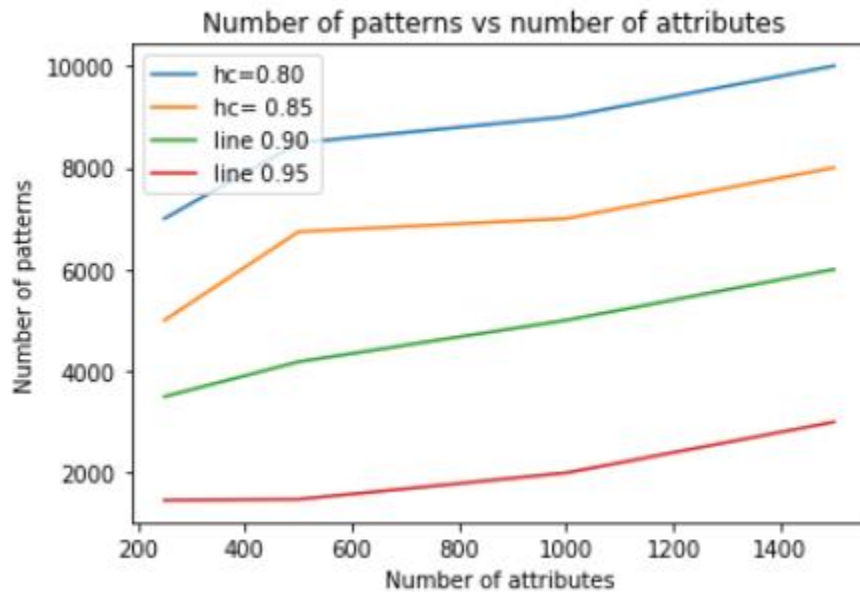


Inferences Drawn

- Execution time increases with increase of number of attributes for a constant h confidence value
- Execution time increases with decrease in h confidence value.

Please refer **number_of_attr_vs_execution_time.pdf** for more details.

Examining the number of patterns for various number of attributes



Inferences Drawn:

- Number of hyper clique patterns increases with increase of number of attributes for a constant h confidence threshold.
- High value of h confidence threshold results in lesser number of hyper clique patterns.

Please refer [attr_vs_number_patterns.pdf](#) for more details.

Conclusions

- Cross support property (Corollary 1) tested successfully and can be used to avoid generating spurious pattern involving items from different support levels.
- Combination of anti-monotone and cross support properties worked correctly for efficient discovery of hyper clique patterns at low levels of support.
- Using Apriori algorithm may take significant amount of time even for smaller data set and usually gets trapped in low memory issues, hyper clique miner algorithm successfully overcome this problem.
- Though hyper clique miner is efficient yet large datasets requires RAM of larger size hence had to run the code on the sample of the data set.