CSE 469: Assignment 2
Principle Association Analysis
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ASSIGNMENT 2: ASSOCIATION ANALYSIS

Frequent Itemsets – Gene Dataset

```
{gene 87}:
                                          sup = 0.67
L1 Frequent Itemsets
                              \{gene 89\}: sup = 0.59
{gene 1}: \sup = 0.83
                              {gene 9}:
                                          sup = 0.5
\{gene 12\}: sup = 0.54
                              {gene_90}:
                                          sup = 0.52
{gene 14}: \sup = 0.52
                              {gene 91}:
                                          sup = 0.65
{gene 17}:
            sup = 0.55
                              {gene 93}:
                                          sup = 0.53
{gene 21}:
            sup = 0.62
                              {gene 94}:
                                          sup = 0.62
{gene 22}:
            sup = 0.55
                              {gene 98}:
                                          sup = 0.51
{gene 23}:
            sup = 0.54
                              {gene 99}:
                                          sup = 0.56
\{gene 25\}: sup = 0.57
                             L2 Frequent Itemsets
\{gene 26\}: sup = 0.52
                                                  sup = 0.51
{gene 27}: sup = 0.51
                              {gene_5, gene_72}:
{gene 3}:
                              {gene 59, gene 5}:
            sup = 0.71
                                                  sup = 0.51
                              {gene 1, gene 59}:
\{gene 31\}: sup = 0.51
                                                  sup = 0.62
                                                  sup = 0.5
\{gene 36\}: sup = 0.61
                              {gene 47, gene 3}:
{gene 37}: \sup = 0.56
                              {gene 3, gene 72}:
                                                  sup = 0.53
{gene_39}:
                              {gene_47, gene_5}:
            sup = 0.51
                                                  sup = 0.53
{gene 4}:
            sup = 0.5
                              {gene 1, gene 67}: \sup = 0.55
\{gene 43\}: sup = 0.5
                              {gene 1, gene 3}:
                                                  sup = 0.63
\{gene 45\}: sup = 0.58
                              {gene 1, gene 8}:
                                                  sup = 0.53
\{gene 47\}: sup = 0.66
                              {gene 1, gene 84}: \sup = 0.5
\{gene 48\}: sup = 0.57
                              {gene 81, gene 1}: \sup = 0.51
                              {gene 87, gene 59}: \sup = 0.51
{gene 5}:
            sup = 0.73
                              {gene 1, gene 6}:
\{gene 50\}: sup = 0.5
                                                  sup = 0.59
\{gene_{53}\}: sup = 0.5
                              {gene_1, gene_89}:
                                                  sup = 0.52
\{gene 54\}: sup = 0.67
                              {gene 1, gene 72}: sup = 0.61
\{gene 55\}: sup = 0.55
                              {gene 87, gene 1}:
                                                  sup = 0.56
\{gene_{56}\}: sup = 0.51
                              \{\text{gene } 1, \text{ gene } \overline{2}1\}:
                                                  sup = 0.53
\{gene_{59}\}: sup = 0.76
                              {gene 87, gene 5}:
                                                  sup = 0.51
            sup = 0.66
                                                  sup = 0.5
{gene 6}:
                              {gene 91, gene 5}:
\{gene 60\}: sup = 0.54
                              {gene 1, gene 94}:
                                                  sup = 0.54
{gene 63}: \sup = 0.5
                              {gene 59, gene 3}: \sup = 0.56
                              \{gene_1, gene_{54}\}:
{gene 64}:
            sup = 0.5
                                                  sup = 0.58
\{gene 66\}: sup = 0.59
                              {gene 91, gene 1}: \sup = 0.55
{gene 67}:
            sup = 0.62
                              {gene 1, gene 47}:
                                                  sup = 0.59
                                                  sup = 0.59
{gene 71}: \sup = 0.58
                              {gene 5, gene 3}:
{gene_72}:
            sup = 0.74
                              {gene_59, gene_6}:
                                                  sup = 0.51
\{gene 75\}: sup = 0.57
                              {gene_1, gene_5}:
                                                  sup = 0.65
{gene 77}: \sup = 0.58
                              {gene 6, gene 5}:
                                                  sup = 0.52
\{gene_{78}\}: sup = 0.59
                              {gene 59, gene 72}: \sup = 0.62
{gene 8}:
            sup = 0.66
                             L3 Frequent Itemsets
\{gene 81\}: sup = 0.58
                              {gene 1, gene 72, gene 59}: \sup = 0.5
{gene 83}:
            sup = 0.5
                              {gene 1, gene 5, gene 3}: \sup = 0.52
{gene 84}:
            sup = 0.54
```

Length 3 Candidate Itemsets – Gene Dataset

```
{'gene_72', 'gene_3', 'gene_5'}
{'gene_47', 'gene_3', 'gene_5'}
{'gene_59', 'gene_5', 'gene_1'}
{'gene_1', 'gene_91', 'gene_5'}
{'gene_59', 'gene_5', 'gene_6'}
{'gene_59', 'gene_5', 'gene_5'}
{'gene_87', 'gene_5', 'gene_72'}
{'gene_87', 'gene_59', 'gene_1'}
{'gene_87', 'gene_1', 'gene_1'}
{'gene_72', 'gene_3', 'gene_1'}
{'gene_72', 'gene_3', 'gene_1'}
{'gene_72', 'gene_1', 'gene_5'}
{'gene_72', 'gene_1', 'gene_5'}
{'gene_59', 'gene_1', 'gene_5'}
{'gene_59', 'gene_3', 'gene_1'}
{'gene_59', 'gene_3', 'gene_5'}
{'gene_1', 'gene_47', 'gene_5'}
{'gene_59', 'gene_3', 'gene_1'}
{'gene_59', 'gene_3', 'gene_5'}
{'gene_59', 'gene_3', 'gene_5'}
{'gene_59', 'gene_3', 'gene_5'}
{'gene_59', 'gene_3', 'gene_72'}
{'gene_59', 'gene_87', 'gene_5'}
```

Apriori Algorithm Code

Below are the portions of code which I wrote to fill in the template functions provided. These have been nicely formatted to fit in this document. A full copy paste of the code is available in the appendix, read at your own risk.

Apriori Gen

```
def apriori_gen(freq_sets, k):
    n = len(freq_sets)
    if n<2: # Minimum 2 frequent itemsets needed to generate candidates
        return []
    # generate all possible candidate itemsets
    candidate_set = set()
    for i in range(0, n-1): # iterate through each element
        for j in range(i+1, n): # and try to combine it with every element after it
            commonElems = freq sets[i].intersection(freq sets[j])
            if len(commonElems) >= k-2: # if k-2 of the items in the sets match
                # combine the sets to make a length k itemset
                newCandidate = freq sets[i].union(freq sets[j])
                candidate set.add(newCandidate) # add that itemset to the list of candidates
    # find candidate itemsets which have k-1 length subsets that are infrequent
    invalidCandidates = set()
    for candidate in candidate set:
        for elem in candidate:
            k1subset = candidate.difference({elem}) # generate every possible k-1 subset
            subsetIsFrequent = False
            for freqItemset in freq_sets: # check that every k-1 subset is frequent
                if k1subset == freqItemset:
                    subsetIsFrequent = True
                    break
            if not subsetIsFrequent: # if one or more of the k-1 subsets was infrequent
                invalidCandidates.add(candidate) # prune the candidate itemset
    # prune the invalid candidates
    for candidate in invalidCandidates:
        candidate_set.remove(candidate)
    return list(candidate_set)
```

Get Freq

```
def get_freq(dataset, candidates, min_support, verbose=False):
    freq_list = []
    support_data = dict()
    for candidateSet in candidates:
        supportCount = 0
        for transaction in dataset:
            if candidateSet.issubset(transaction):
                supportCount += 1
        support = supportCount / len(dataset)
        if support >= min_support:
            freq_list.append(candidateSet)
        support_data[candidateSet] = support
```

Appendix

Full text of Apriori_template.py

```
from future import print function
import sys
def apriori (dataset, min support=0.5, verbose=False):
    """Implements the Apriori algorithm.
    The Apriori algorithm will iteratively generate new candidate
    k-itemsets using the frequent (k-1)-itemsets found in the previous
    iteration.
    Parameters
    dataset : list
       The dataset (a list of transactions) from which to generate
        candidate itemsets.
   min support : float
        The minimum support threshold. Defaults to 0.5.
   Returns
    F : list
       The list of frequent itemsets.
    support data : dict
       The support data for all candidate itemsets.
   References
    .. [1] R. Agrawal, R. Srikant, "Fast Algorithms for Mining Association
          Rules", 1994.
    C1 = create candidates (dataset)
    D = list(map(set, dataset))
    F1, support data = get freq(D, C1, min support, verbose=False) # get frequent 1-itemsets
    F = [F1] # list of frequent itemsets; initialized to frequent 1-itemsets
    k = 2 # the itemset cardinality
    while (len(F[k-2]) > 0):
       Ck = apriori gen(F[k-2], k) # generate candidate itemsets
        Fk, supK = get freq(D, Ck, min support) # get frequent itemsets
        support data.update(supK)# update the support counts to reflect pruning
        F.append(Fk) # add the frequent k-itemsets to the list of frequent itemsets
       k += 1
    if verbose:
        # Print a list of all the frequent itemsets.
        for kset in F:
           for item in kset:
                                            + "{"
               print(""
                                                                      + "".join(str(i) + ", " for i in
                                            + "}"
                                                                      + ": sup = " +
iter(item)).rstrip(', ')
str(round(support_data[item], 3)))
    return F, support data
def create candidates (dataset, verbose=False):
    """Creates a list of candidate 1-itemsets from a list of transactions.
```

Parameters

```
dataset : list
       The dataset (a list of transactions) from which to generate candidate
       itemsets.
   Returns
    The list of candidate itemsets (c1) passed as a frozenset (a set that is
    immutable and hashable).
    c1 = [] # list of all items in the database of transactions
    for transaction in dataset:
       for item in transaction:
            if not [item] in c1:
                cl.append([item])
   cl.sort()
    if verbose:
        # Print a list of all the candidate items.
       print(""
                            + "{"
                                               + "".join(str(i[0]) + ", " for i in iter(c1)).rstrip(', ')
+ "}")
    # Map c1 to a frozenset because it will be the key of a dictionary.
    return list (map (frozenset, c1))
def get freq(dataset, candidates, min support, verbose=False):
    This function separates the candidates itemsets into frequent itemset and infrequent itemsets based on
the min support,
        and returns all candidate itemsets that meet a minimum support threshold.
   Parameters
    dataset : list
       The dataset (a list of transactions) from which to generate candidate
       itemsets.
    candidates : frozenset
       The list of candidate itemsets.
   min support : float
       The minimum support threshold.
   Returns
    freq list: list
        The list of frequent itemsets.
    support data: dict
       The support data for all candidate itemsets.
    freq list = []
    support data = dict()
    for candidateSet in candidates:
        supportCount = 0
        for transaction in dataset:
            if candidateSet.issubset(transaction):
                supportCount += 1
        support = supportCount / len(dataset)
        if support >= min support:
            freq list.append(candidateSet)
            support_data[candidateSet] = support
```

```
return freq list, support data
def apriori gen(freq sets, k):
    """Generates candidate itemsets (via the F_k-1 \times F_k-1 \text{ method}).
    This part generates new candidate k-itemsets based on the frequent
    (k-1)-itemsets found in the previous iteration.
    The apriori gen function performs two operations:
    (1) Generate length k candidate itemsets from length k-1 frequent itemsets
    (2) Prune candidate itemsets containing subsets of length k-1 that are infrequent
    Parameters
    freq sets : list
        The list of frequent (k-1)-itemsets.
    k: integer
        The cardinality of the current itemsets being evaluated.
    Returns
    candidate list : list
       The list of candidate itemsets.
    n = len(freq sets)
    if n<2: # Minimum 2 frequent itemsets needed to generate candidates
        return []
    # generate all possible candidate itemsets
    candidate set = set()
    for i in range(0, n-1): # iterate through each element
        for j in range(i+1, n): # and try to combine it with every element after it
            commonElems = freq_sets[i].intersection(freq_sets[j])
            if len(commonElems) >= k-2: # if k-2 of the items in the sets match
                newCandidate = freq sets[i].union(freq sets[j]) # combine the sets to make a length k
itemset
                candidate set.add(newCandidate) # add that itemset to the list of candidates
    # find candidate itemsets which have k-1 length subsets that are infrequent
    invalidCandidates = set()
    for candidate in candidate set:
        for elem in candidate:
            klsubset = candidate.difference({elem}) # generate every possible k-1 subset
            subsetIsFrequent = False
            for freqItemset in freq_sets: # check that every k-1 subset is frequent
                if klsubset = freqItemset:
                    subsetIsFrequent = True
                    break
            if not subsetIsFrequent: # if one or more of the k-1 subsets was infrequent
                invalidCandidates.add(candidate) # prune the candidate itemset
    # prune the invalid candidates
    for candidate in invalidCandidates:
        candidate set.remove(candidate)
    return list(candidate set)
def loadDataSet(fileName, delim=','):
    fr = open(fileName)
```

```
stringArr = [line.strip().split(delim) for line in fr.readlines()]
    return stringArr
def run_apriori(data_path, min_support, verbose=False):
    dataset = loadDataSet (data path)
    F, support = apriori(dataset, min_support=min_support, verbose=verbose)
    return F, support
def bool transfer(input):
    ''' Transfer the input to boolean type'''
    input = str(input)
    if input.lower() in ['t', '1', 'true' ]:
       return True
    elif input.lower() in ['f', '0', 'false']:
       return False
    else:
        raise ValueError('Input must be one of {T, t, 1, True, true, F, F, 0, False, false}')
if __name__ = '__main__':
    if len(sys.argv) = \overline{3}:
       F, support = run_apriori(sys.argv[1], float(sys.argv[2]))
    elif len(sys.argv)=\overline{4}:
       F, support = run_apriori(sys.argv[1], float(sys.argv[2]), bool_transfer(sys.argv[3]))
       raise ValueError('Usage: python apriori templete.py <data path> <min support> <is verbose>')
    print(F)
   print(support)
    Example:
   python apriori_templete.py market_data transaction.txt 0.5
    python apriori templete.py market data transaction.txt 0.5 True
    111
```