The DESQ Framework for Declarative and Scalable Frequent Sequence Mining

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- 1. Frequent Sequence Mining
- 2. Declarativity
- 3. Scalability
- 4. Summary

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Before and after



Anni wants to watch a movie. Anni loves LOTR1. But she does not want to see it. She had seen LOTR2 last week!

Movie streaming site



Let's look at some data

- Data from Netflix' online movie-streaming platform
 - 500k users, 18k movies, 100M ratings with timestamps
- ▶ 125k users rated both LOTR1 and LOTR2
- In which order?



105k users



20k users

Order matters!

- How to discover patterns in sequential data?

Frequent Sequence Mining

- Frequent sequence mining is a fundamental task in data mining
 - Data modeled as collection of sequences of items or events
 - Often items are arranged in a hierarchy
 - We seek frequent sequential patterns
- E.g., market-basket data
 - Sequence = purchases of a customer over time
 - Item = product (or set of products) + product hierarchy
 - Example pattern: DSLR Camera \rightarrow Tripod \rightarrow Flash
- E.g., natural-language text
 - Sequence = sentence or document
 - Item = word + syntactic/semantic hierarchy
 - Example pattern: person was born in location
- E.g., amino acid sequences
 - Sequence = protein
 - Item = amino acid
 - Example pattern: S L R

What constitutes a good pattern?

- Extensively studied
 - Interesting patterns should be new, surprising, understandable, actionable
 - No random patterns, common knowledge, redundancy
 - Details application-specific
- Many different variants, many algorithms
 - Constraints: length, positional/temporal, hierarchy, regex, ...
 - Scoring: frequency, utility, information gain, significance, ...
 - Pattern sets: all, top-k, maximality, closedness, MDL, ...
- Our research focuses on unifying frequent sequence mining
 - Study general properties instead of special cases
 - Avoid need for customized mining algorithms

DESQ

- DESQ = framework for declarative and scalable frequent sequence mining [TODS19, ICDM16, ICDE19]
 - Open source
- Key design goals are
 - 1. Usefulness
 - Can be tailored to application
 - Flexible constraints
 - 2. Usability
 - Describe pattern mining task in an intuitive, declarative way
 - Hide technical and implementation details
 - 3. Efficiency
 - Fast
 - Scalable
 - Competitive to specialized miners

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Special case: n-gram mining

An *n*-gram is a sequence of *n* consecutive words

- Extensively used in text mining and natural-language processing
- Web-scale n-gram models published by Google and Microsoft

Google books Ngram Viewer



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Going declarative

- ▶ If we simply mined all frequent *n*-grams, we may
 - 1. Produce many uninteresting patterns (low frequency threshold)
 - 2. Miss out on interesting patterns (high frequency threshold)
- DESQ allows data analysts to focus on what they consider relevant
 - Supports all traditional constraints (length, gap, hierarchy, ...)
 - Supports customized constraints that go beyond traditional constraints
- Based on a declarative pattern expression language
 - Describe relevant patterns, let DESQ take care of mining them
 - Syntax like regular expression
 - Adds capture groups and hierarchies

Some examples for text mining

1. Noun modified by adjective or noun

Ex: big country (110), green tea (337), research scientist (473) PE: ([ADJ|NOUN] NOUN)

2. Relational phrase between entities

Ex: lives in (847), is being advised by (15), has coached (10) PE: ENTITY (VERB⁺ NOUN⁺? PREP?) ENTITY

3. Typed relational phrases

Ex: ORG headed by ENTITY (275), PERS born in LOC (481) PE: (ENTITY[↑] VERB⁺ NOUN⁺? PREP? ENTITY[↑])

4. Google *n*-gram viewer data

Ex: a good day, a ADJ day, DET ADJ NOUN, have a good day PE: $(.^{\uparrow})(.^{\uparrow})?(.^{\uparrow})?(.^{\uparrow})?(.^{\uparrow})$

Pattern mining

- Under the hood, DESQ translates pattern expressions to finite state transducers (FST)
 - FST outputs all patterns that occur in a given input sequence
- Multiple sequential mining algorithms
 - Naive approach ("WordCount")
 - DesqCount ("WordCount" with frequency pruning)
 - DesqDfs (depth-first search)

Performance comparison (traditional constraints)





DESQ is competitive to state-of-the-art miners for traditional constraints.

Performance comparison (new constraints)



DesqDfs is method of choice and can be orders of magnitude faster than Naive or DesqCount.

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Distributed mining

Based on bulk synchronous parallel model

Key idea

- Partition data into smaller overlapping partitions using item-based partitioning
 - One partition for each frequent item
- Mine each partition locally
- Combine results

Key question

- What to communicate to partitions?
 - Inputs
 - Candidates



Communicate inputs

- Naïve approach: send each input sequence to all partitions for which it is "relevant"
- More efficient: send only relevant parts of input sequence
 - Example: only fantasy movies relevant for mining task



- Can reduce communication up to 100x

Communicate candidates

- Naïve approach: send each candidate subsequence to its corresponding partition
- More efficient: compress candidates
 - Shared structure
 - Non-deterministic finite automata (NFA)



- Can reduce communication by up to 100x

Performance comparison

Both approaches scale nearly linearly with number of input sequences. green: send inputs, blue: send candidates



- Up to 50x faster than naïve approaches
- Sending candidates is up to 5x faster for selective constraints
- 1-4x generalization overhead over specialized approaches

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Summary

DESQ: framework for declarative and scalable frequent sequence mining

- Find patterns in sequential data
- Declarative language to specify interest
- Item-based partitioning to scale to large datasets
- Open source: https://github.com/rgemulla/desq

[ICDM16] Beedkar, K.; Gemulla, R.: DESQ: Frequent Sequence Mining with Subsequence Constraints. In: ICDM, 2016.

[TODS19] Beedkar, K.; Gemulla, R.; Martens, W.: A Unied Framework for Frequent Sequence Mining with Subsequence Constraints. ACM Trans. Database Syst., 2019. [ICDE19] Renz-Wieland, A.; Bertsch, M.; Gemulla, R.: Scalable Frequent Sequence Mining With Flexible Subsequence Constraints. In: ICDE, 2019.

Thank you!