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## VIRGINIA COMMONWEALTH UNIVERSITY



# Parsing MetaMap Files in Hadoop

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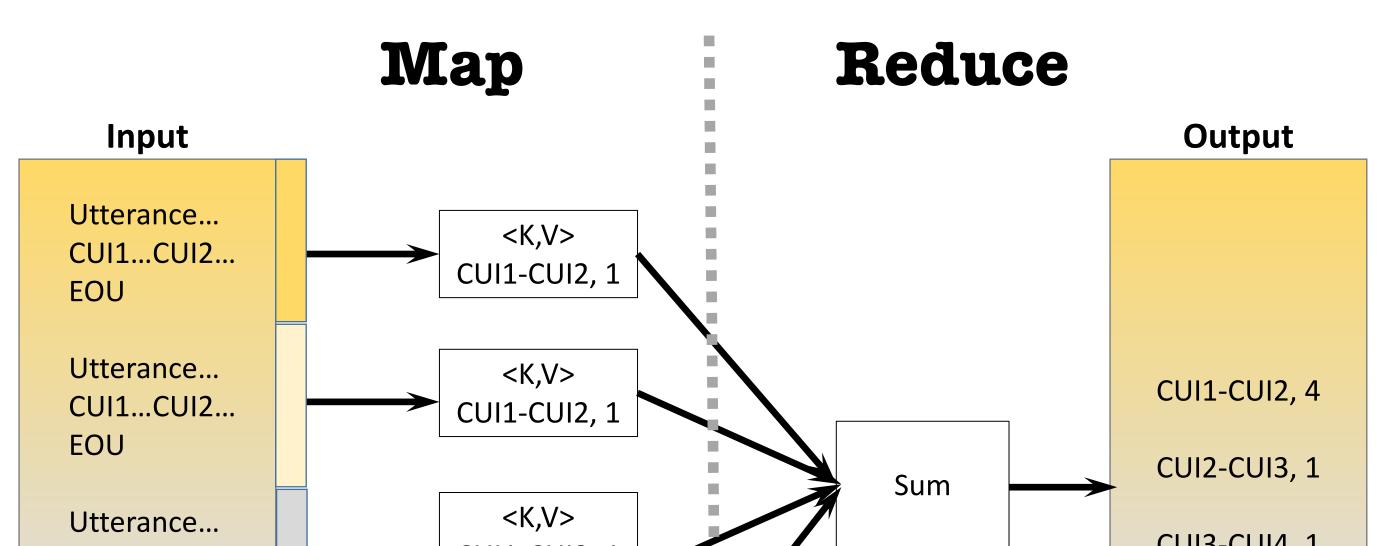


### Hadoop for the Desktop

The **deluge of data** in today's information-centric world requires bigger and better computing resources for processing. This can be a limiting factor in how much data labs with **limited computing resources** are able to handle. This project explores the pitfalls of a serial program that parses MetaMap files to identify UMLS CUI bigrams in a large set of scientific literature. The algorithm is re-implemented in Hadoop MapReduce to **overcome resource bottlenecks**.

## UMLS, CUIs, and MetaMap....Oh My!

### MapReduce in 30 seconds



UINISI	<ul> <li>Unified Medical Language System</li> <li>Repository of biomedical vocabularies</li> <li>Facilitates automated information retrieval (e.g. linking health information and billing codes across systems).</li> <li>Provides language normalization by linking similar terms to same concept/meaning.</li> </ul>	CUI1CUI2 CUI3CUI4 EOU Utterance CUI1CUI2 EOU Split input by record> identify <b>Key, Value</b> > pairs in parallel> sum values with same key		
CUI	<ul> <li>Concept Unique Identifier</li> <li>ID assigned to each unique concept in the UMLS.</li> <li>E.g. Headache and Cranial Pain are both assigned to the CUI <i>C0018681</i></li> </ul>	<ul> <li>MapReduce Advantages</li> <li>Inherent and scalable parallelization.</li> <li>Writes results to diskall intermediate and final.</li> </ul>		
	<ul> <li>Tool that identifies UMLS concepts in biomedical texts.</li> <li>Output mapped text to compressed MetaMap Machine Output (MMO) files.</li> <li>Parsed 23,343,329 citations to create the 2015 MedLine/PubMed Baseline dataset779 MMO files compressed to 132GB.</li> </ul>	<ul> <li>No MySQL communication.</li> <li>Results         <u>CuiCollectorMapReduce</u>         Extracts CUI bigrams using Hadoop MapReduce framework.         Desktop implementation (a single node Hadoop system).</li> </ul>		

### **UMLS::Association**

Dataset

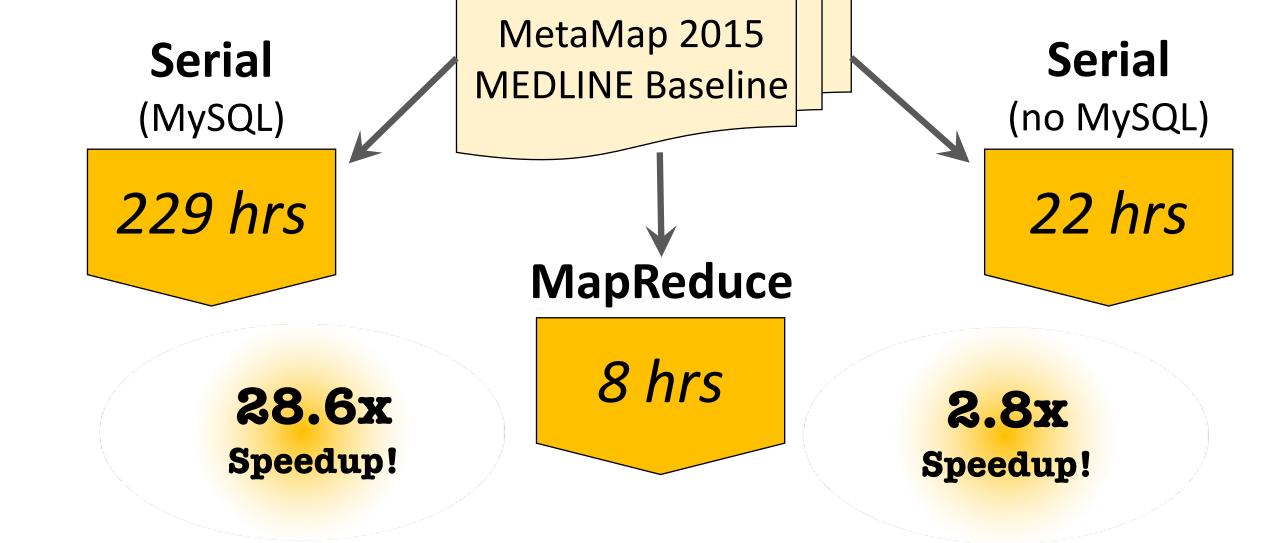
UMLS CUIs can be used to normalize biomedical and clinical text for use in natural language processing applications. By counting **CUI Bigram frequency**—the number of times two CUIs appear close to each other in text-the UMLS::Association package can **identify related concepts**. E.g. head ache and aspirin.

### MetaMap File Anatomy

#### Utterance

16691646.ti.1 "Statement of Cases of Gonorrhoeal and Purulent Ophthalmia..."

<b>Phrase 1</b> "Statement of Cases of <u>Gonorrhoeal</u> "					
Mapping	"Statement" "Cases" "Gonorrhoeal"		'Case (situation)'		
	C1710187	C0868928	C0018081	Case (situation)	
Mapping		" "Cases" " <u>Gon</u> C1533148		'Case unit dose'	
Phrase 2			<u>Adjacent CUI Bigrams</u>		
'Purulent Ophthalmia"			C1710187 - C0868928 C0868928 - C0018081		
Mapping "Purulent Ophthalmia" C0259800		ohthalmia"	C0018081 - C0259800		
		00	C1710187 - C1533148 C1533148 - C0018081		



### **Conclusion and Future Work**

Parsing CUI Bigrams in Hadoop on a desktop computer resulted in **significant speedup**, and enables **parsing larger datasets** that were not previously feasible. Algorithm improvements include a window size to collect distant CUI bigrams and crossing utterances to process full PubMed citations. Future work includes testing the scalability on a Hadoop cluster, resolving an issue with the compressed input file format to improve mapping efficiency, identifying optimal Hadoop settings for a desktop implementation, and re-implementing in SPARK to take advantage of its in-memory storage of intermediate results.

In conclusion, **desktop implementations** of Hadoop can **resolve computing resource problems** and **process data faster**, opening up **more research areas** in big data processing for smaller labs.



### **Serial Limitations**

- Processes one file/one utterance at a time with nested for loops.
   Regularly writes nested bigram hash table to MySQL database due to memory limitations, introducing DB communication latency.
- Perl code is **not paralellizable** due to limitations in sharing nested hashes across threads.

### References

[1] Bodenreider O. "The unified medical language system (UMLS): integrating biomedical terminology." *Nucleic acids research*, 2004, 32, D267-D270.
[2] Aronson AR and Lang F. "An overview of MetaMap: historical perspective and recent advances." *JAMIA*, 2010, 17:3, 229-236.

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