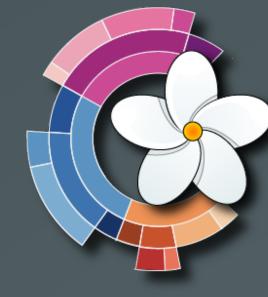
Introduction





Visualization of Understudied Genes

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Conclusion

This visualization is a catalogue of the publications of genes over time. This can serve to draw attention to gaps in knowledge and influence where further investigation is needed.

Further Research

- Automated data extraction from NCBI and cloud storage of data
- Text mining to identify sentences from PubMed abstracts
- Use natural language processing to transform unstructured text in documents into normalized, structured data suitable for analysis or to drive machine learning algorithms

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Rodriguez-Esteban, R., & Jiang, X. (2017). Differential gene expression in disease: A comparison between high-throughput studies and the literature. *BMC Medical Genomics*, 10(1), 1–10. https://doi.org/10.1186/s12920-017-0293-y

Methods



U.S. National Library of Medicine
Strategic Plan

Center for

Information

Biotechnology

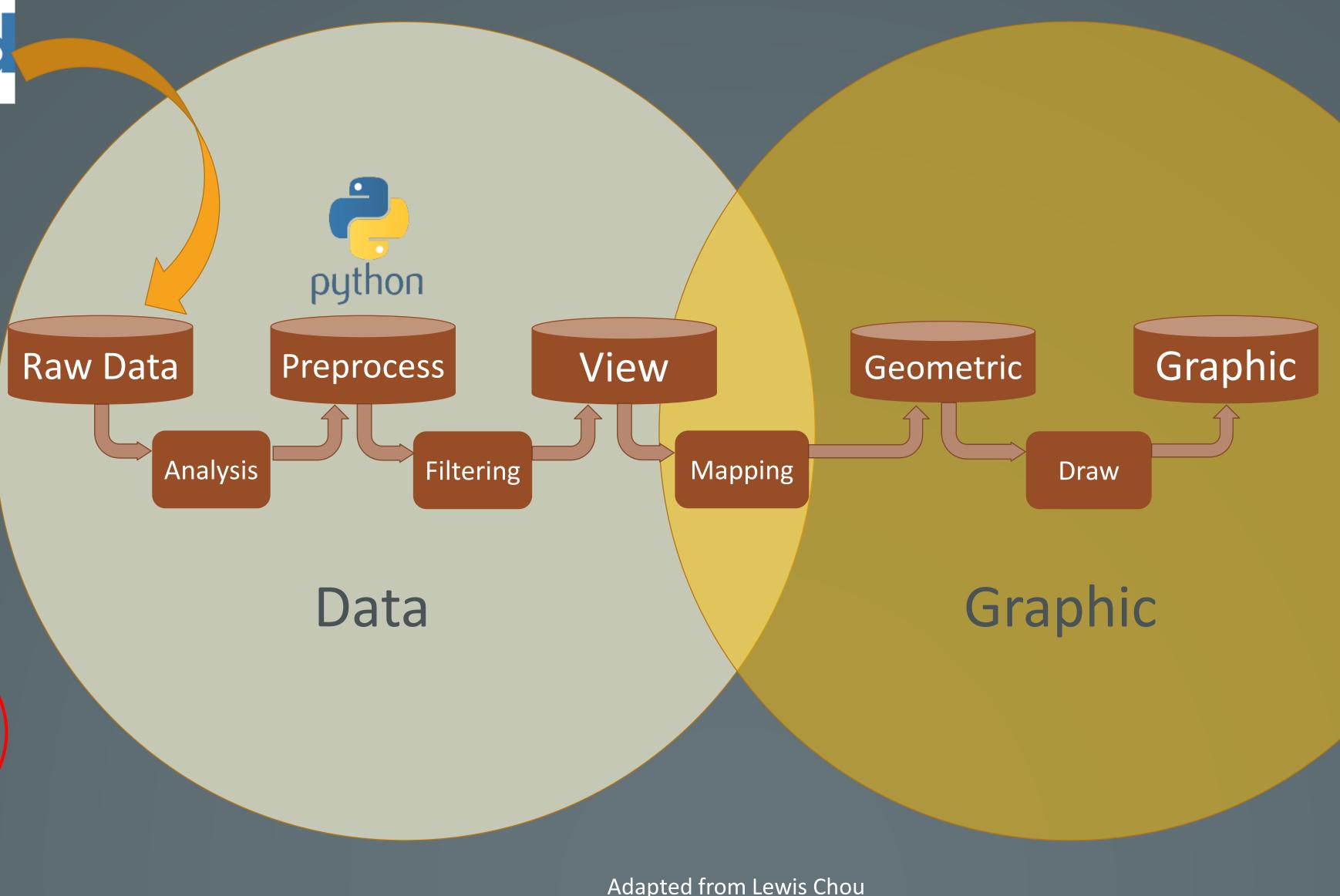
GOAL 1: Accelerate discovery and advance health by providing the tools for data-driven research

GOAL 2: Reach more people in more ways through enhanced dissemination and engagement pathways

GOAL 3: Build a workforce for datadriven research and health

Genes currently being studied 9%

Genes not being researched 91%



Query NCBI to extract list of gene IDs which are orthologs to the human gene

Use the list of IDs to retrieve synonyms differing across orthologs

Use synonyms to query PubMed and return PubMed IDs of articles about the genes

Use PubMed IDs of articles to return the date of publication

Graph the count of publications denoted by the ortholog along a timescale