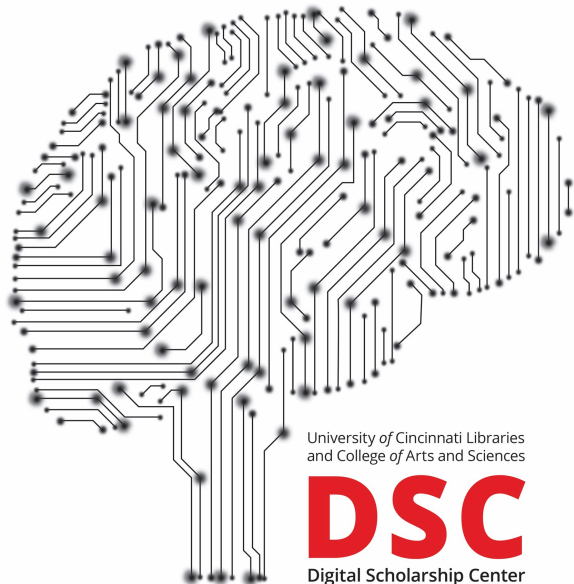


# The Digital Scholarship Center: Text Mining across Disciplines



James Lee, PhD  
Associate Vice Provost for Digital Scholarship  
Associate Dean of Research, University Libraries  
Director, Digital Scholarship Center



- What are Digital Scholarship Centers / Studios?
- The DSC has assembled research groups that genuinely span multiple disciplines, with people trained to think very differently about every step in the research process.
- Teams are composed of true partners across entire research lifecycle:
  - Formulation of research questions
  - Pitching grant proposals
  - Dataset cleanup and manipulation
  - Data analysis and visualization
  - Argument formation
  - Publication of findings
- Machine Learning and Data Visualization as discipline-agnostic methods: a “borrowing of techniques”.

# Andrew W. Mellon Foundation: Digital Integrator

- \$900,000 over 30 months from the Andrew W. Mellon Foundation.
- Specific goal: Advance our “catalyst” model of Digital Scholarship.
- The long view: Mellon supports culture change.
- Goals:
  - Model new transdisciplinary strategies and practices for translating between disciplines that rarely interact – connect domain experts with technical innovations in machine learning.
  - We provide resources and infrastructure to nurture research questions and collaborations that slip between the cracks of funding agencies.
  - Transdisciplinary computational tools / Human interpretable research products.

# Model: Catalyst vs. Concierge

- Six Digital Centers/Studios supported by Mellon initiative.
  - Leaders defining a new type of “digital integrator” collapsing silos.
  - Peers: UIUC, UConn, Brown, Emory, UC’s DSC, Duke.
  - Spectrum from “concierge” to “catalyst.”
- Digital integrators as hubs in the academic research ecosystem.
- Phase 2:
  - Digital Futures Building
  - \$3.1 Million in grant funding
  - Mellon Renewal

# Not Just a Feel-Good Story

- Multiple publications in a range of fields – content area, methodology, popular press.
- Grants from multiple bodies – agencies, foundations, industry.
  - Example: “Iowa Digital Bridges” – Mellon and Gates Foundations, NIH, NSF, NEH, NEA.
- A new perspective on teaching and doctoral training.

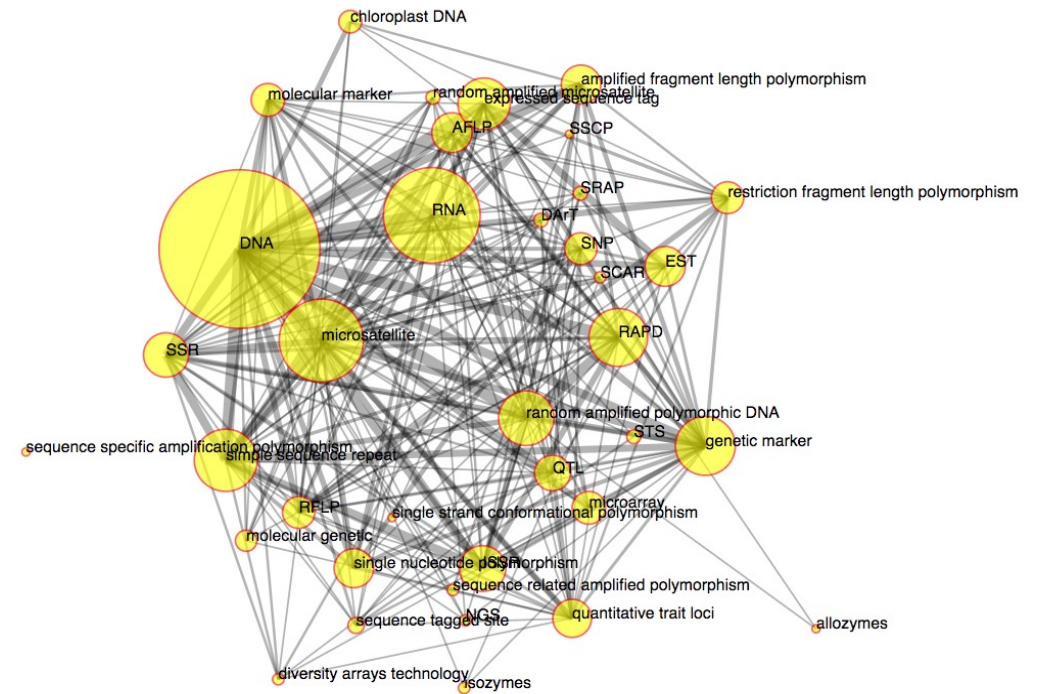
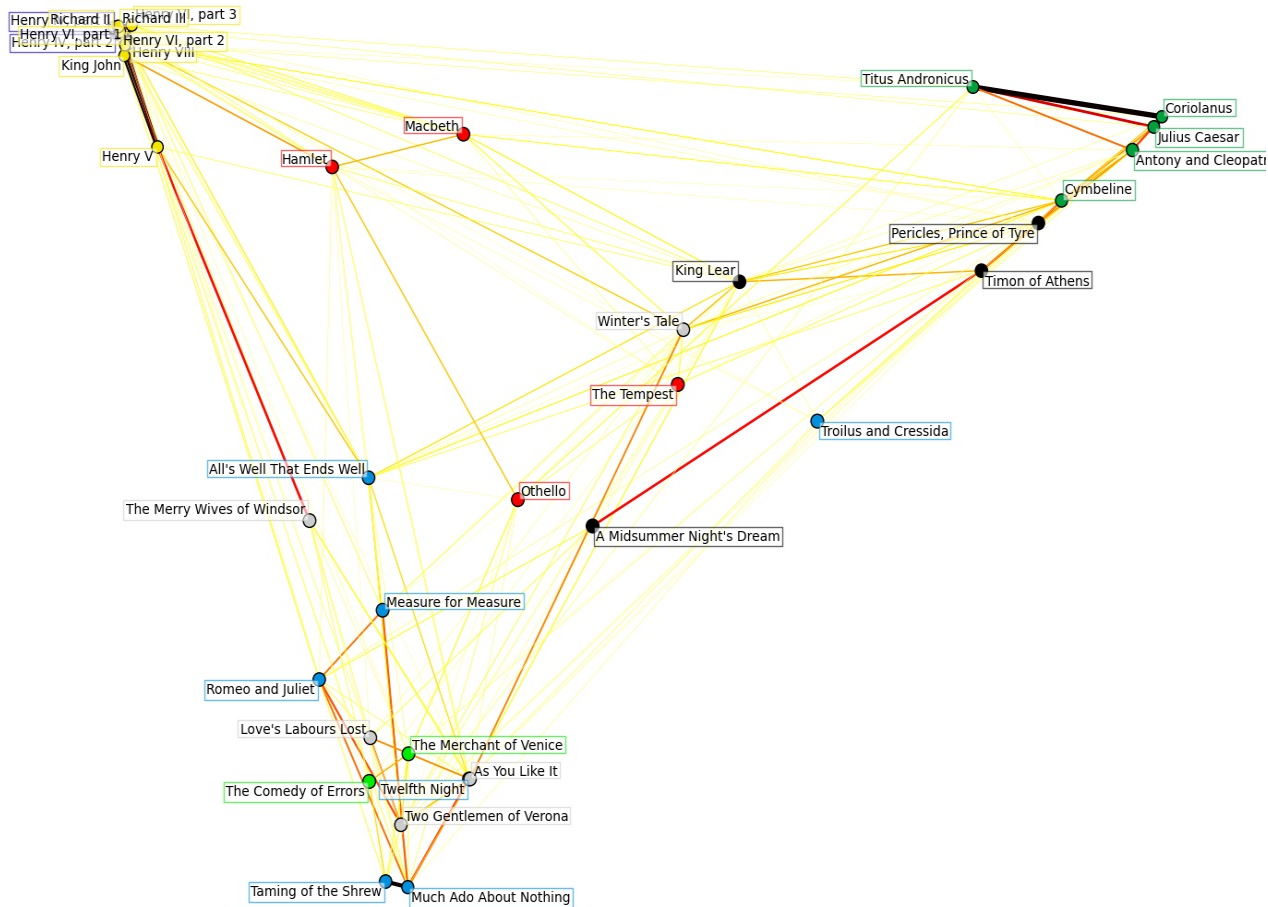
# Building Bridges: Data Structures

- The Shakespearean and the Astrophysicist.

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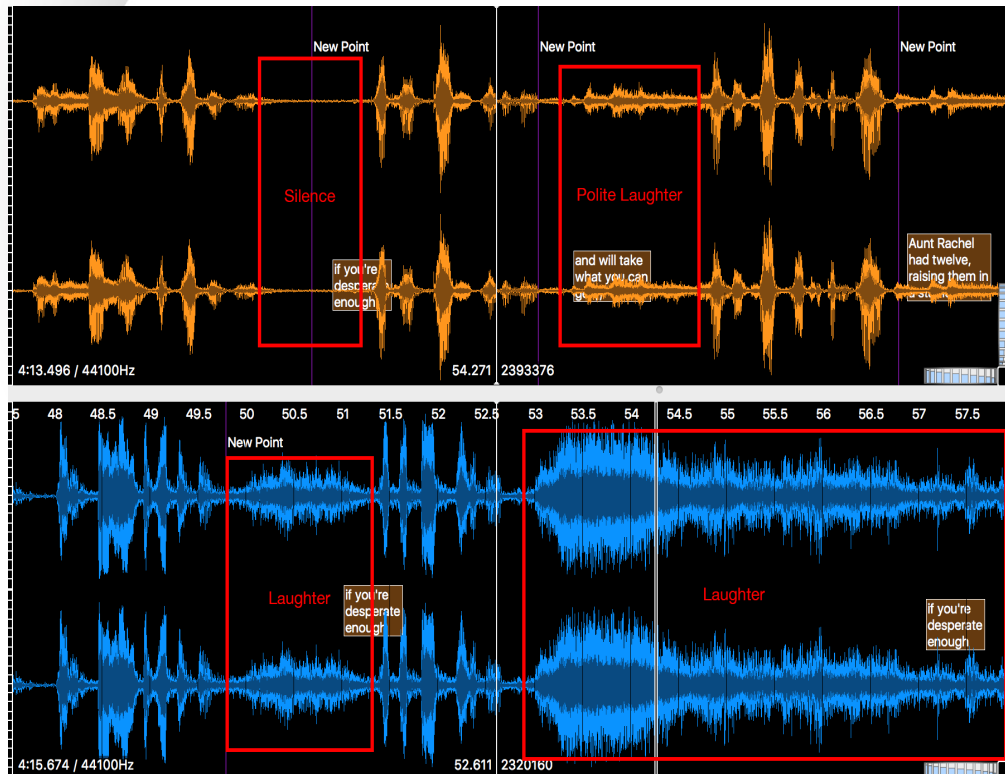
# Building Bridges: Data Structures



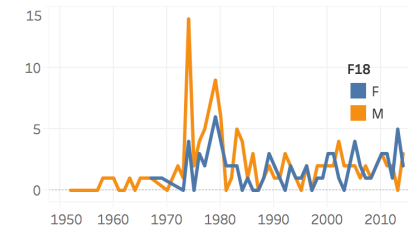


# Building Bridges: Data Visualization

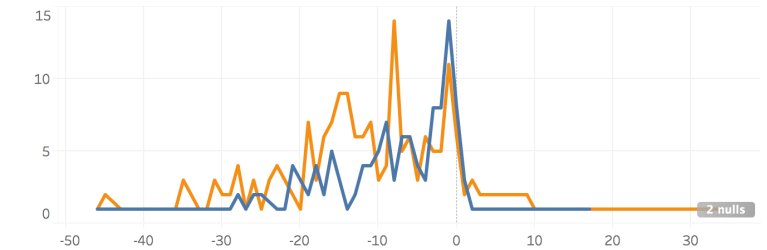
- Different visual frameworks for analysis and interpretability



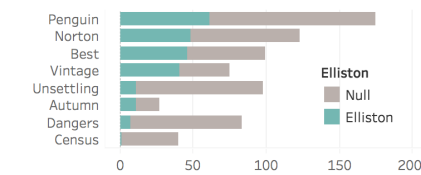
Trend of Featured Poets in Elliston



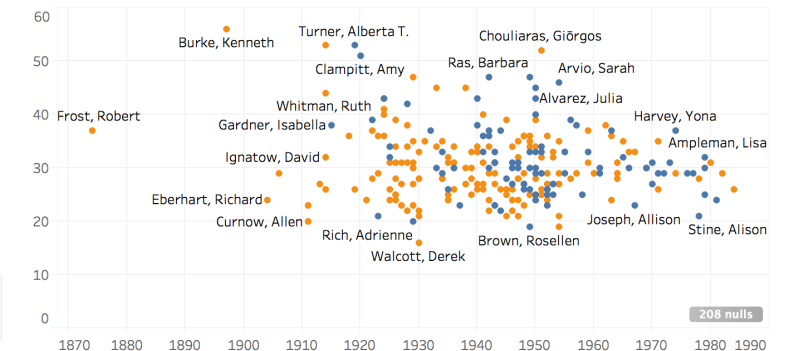
Difference in First book released to Elliston Invitation



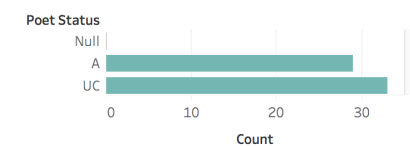
Number of Poets Featured in Elliston vs other



Age when first book is released

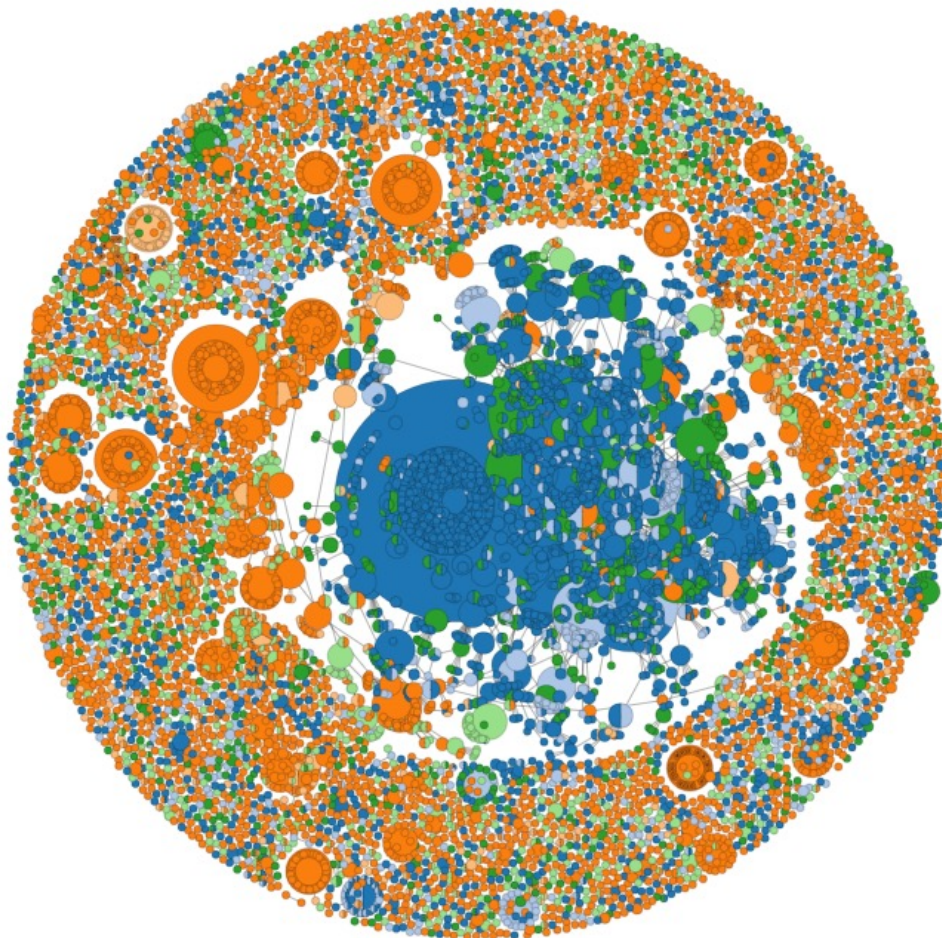


Poet from UC?

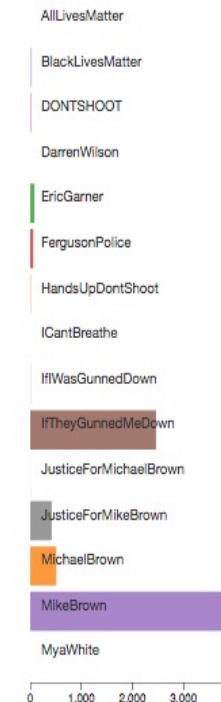


# Shared Motivations, Different Languages

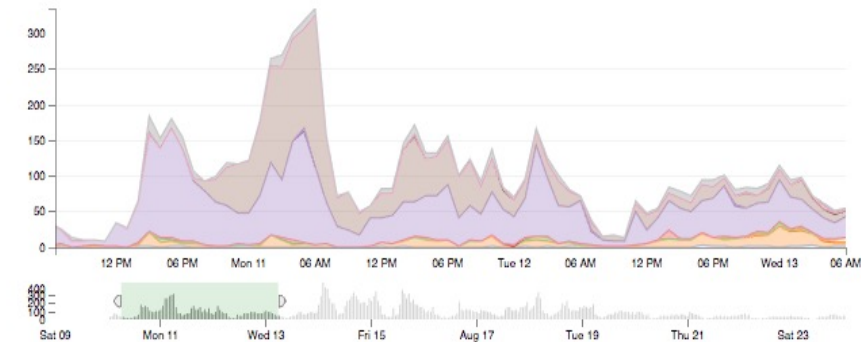
- Social Networks: Social network analysis and visualization.
- Team: DSC, A&S, Law, DAAP, CCHMC, CEAS, CECH.



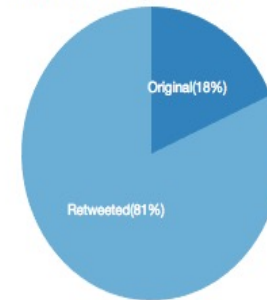
Hashtag Frequency



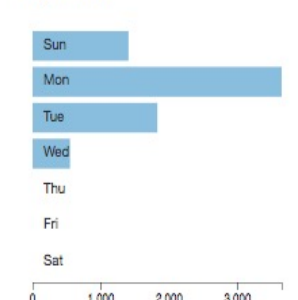
Hashtag Frequency/Time chart range: [08/10/2014 -> 08/13/2014] reset



Retweeted vs Original Tweets

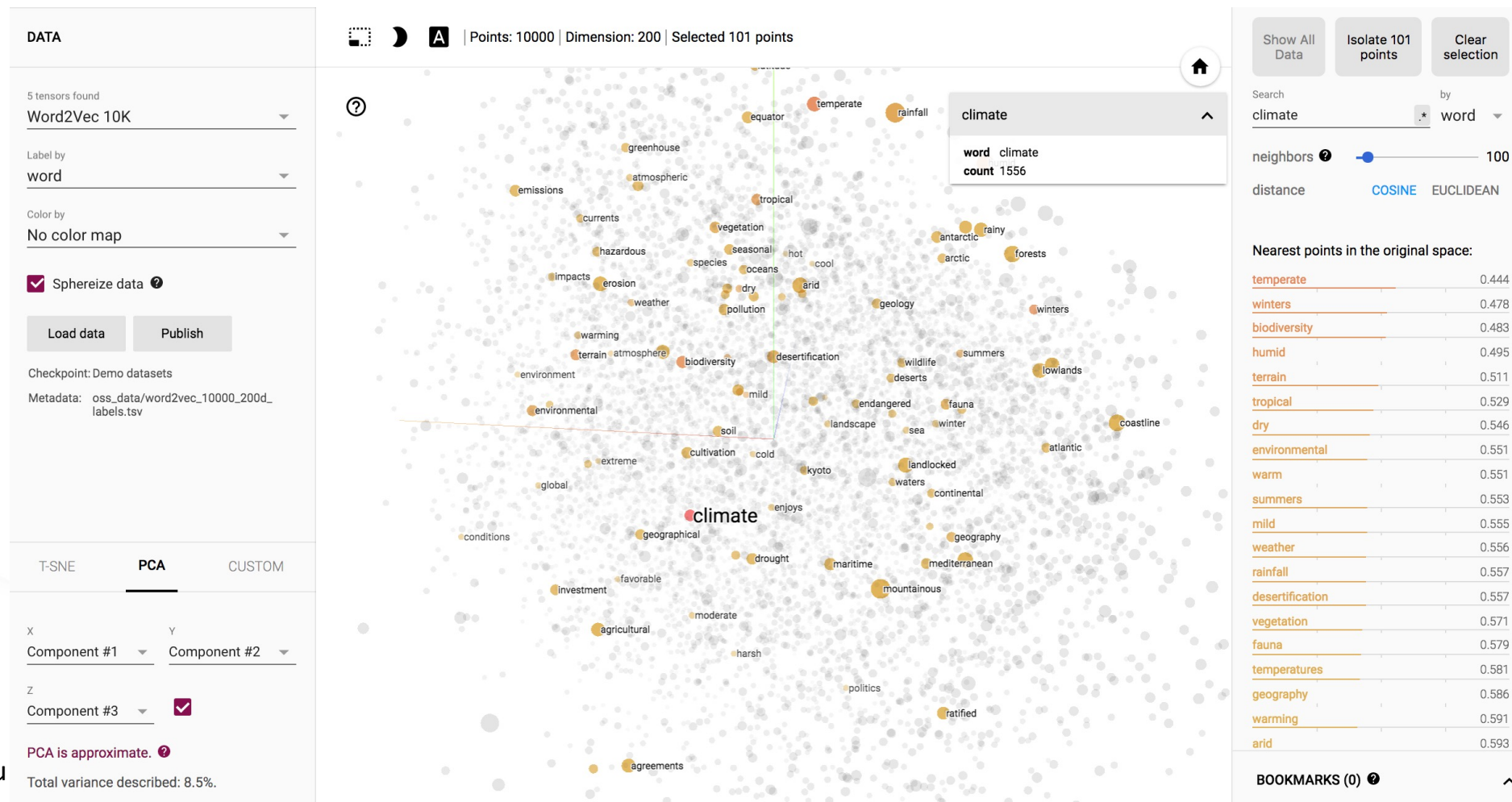


Day of Week



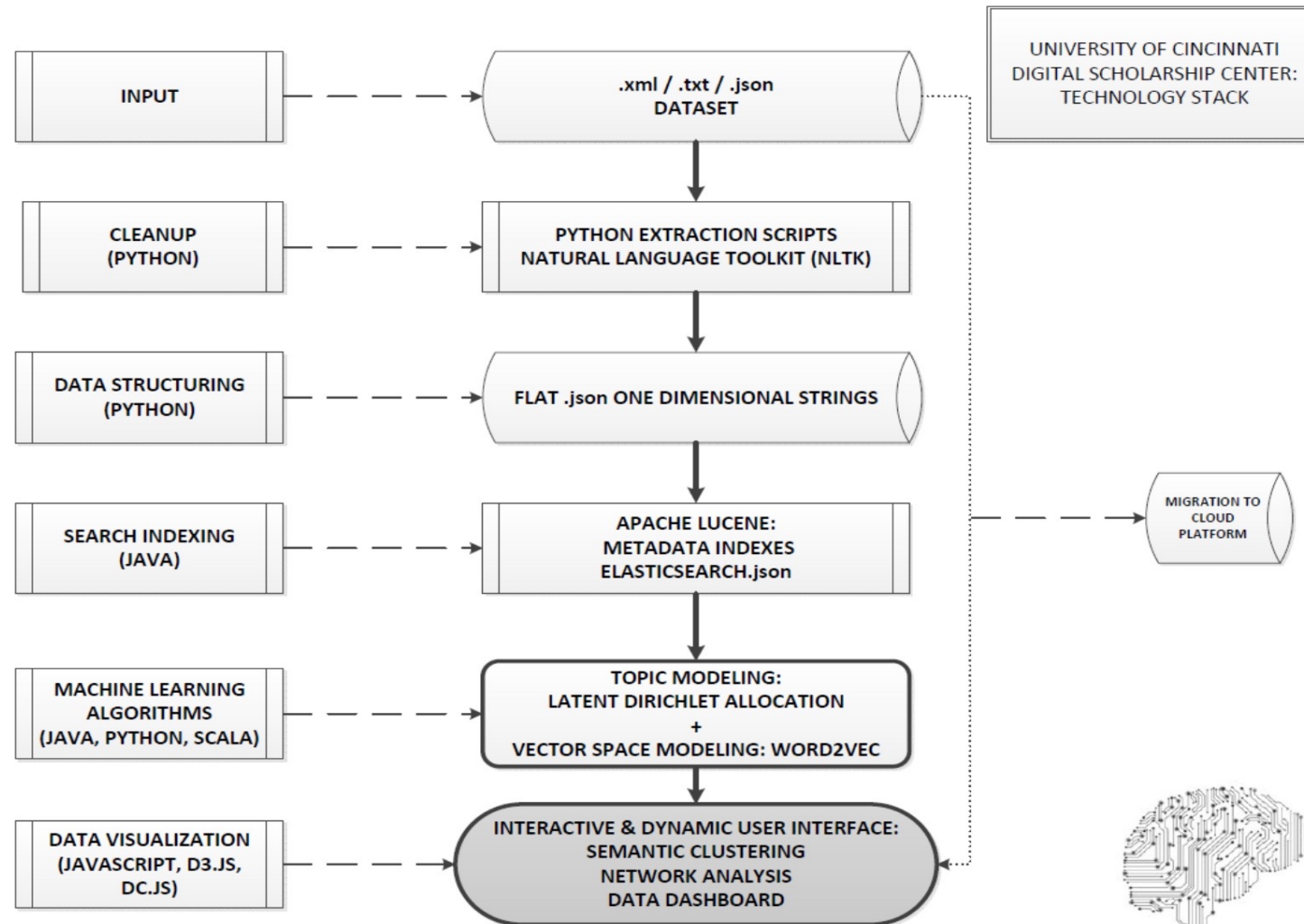
# Shared Motivations, Different Languages

- Using machine learning to extract salient features, cluster, and classify free text in EHR and literature.
- Team: A&S Biological Sciences, UC CoM, CCHMC Hospital Medicine, among others.



# Cloud Platform for Collaboration

## University of Cincinnati Digital Scholarship Center Technology Stack



Topic model:  | Term:

Topic models are trained on selected searches from the [Pubmed corpus](#). Visualization based on the [Python port](#) of the [LDAvis package](#).

Select a topic to see the related documents.

| Topic 6   |      |  |
|---|------|--|
| [19723248] Phylogenetic origin of Phyllolobium with a further implicati | 0.64 |  |
| [23071663] Genetic diversity and population structure of cucumber (Cucu | 0.54 |  |
| [16622792] Biodiversity of Streptomyces of high-mountainous ecosystems  | 0.39 |  |
| [26467618] Phylogeography of Phytophagous Weevils and Plant Species in  | 0.35 |  |
| [23962409] Phylogeography sheds light on the central-marginal hypothesi | 0.34 |  |
| [24065181] Contemporary pollen-mediated gene immigration reflects the h | 0.31 |  |
| [17191876] Biologically active ibogan and vallesamine derivatives from  | 0.30 |  |
| [18206283] Reduced nitrogen has a greater effect than oxidised nitrogen | 0.28 |  |
| [27886271] Speciation and genetic diversity in Centaurea subsect. Phalo | 0.28 |  |
| [22081412] Establishing the phylogenetic origin, history, and age of th | 0.26 |  |
| [22546007] Phylogeographic analysis reveals significant spatial genetic | 0.26 |  |
| [24498103] Genetic differentiation and genetic diversity of Castanopsis | 0.26 |  |
| [17638329] Topoisomerase-II-inhibitory principles from the stems of Spa | 0.25 |  |
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| [23478944] Post-Boreotropical dispersals explain the pantropical disjun | 0.24 |  |
| [27974324] The evolutionary history of Eugenia sect. Phyllocalyx (Myrta | 0.24 |  |
| [25072783] Temperate pine barrens and tropical rain forests are both ri | 0.23 |  |
| [22182994] A phylogeny of Delphinieae (Ranunculaceae) shows that Aconit | 0.22 |  |
| [23418542] Phylogeographic evidence for a link of species divergence of | 0.22 |  |
| [24028582] Large-scale pattern of genetic differentiation within Africa | 0.22 |  |
| [22730022] Range expansion of a selfing polyploid plant despite widespr | 0.21 |  |
| [23629053] Diversification of plant species in arid Northwest China: sp | 0.21 |  |
| [23560070] Tertiary origin and pleistocene diversification of dragon bl | 0.20 |  |

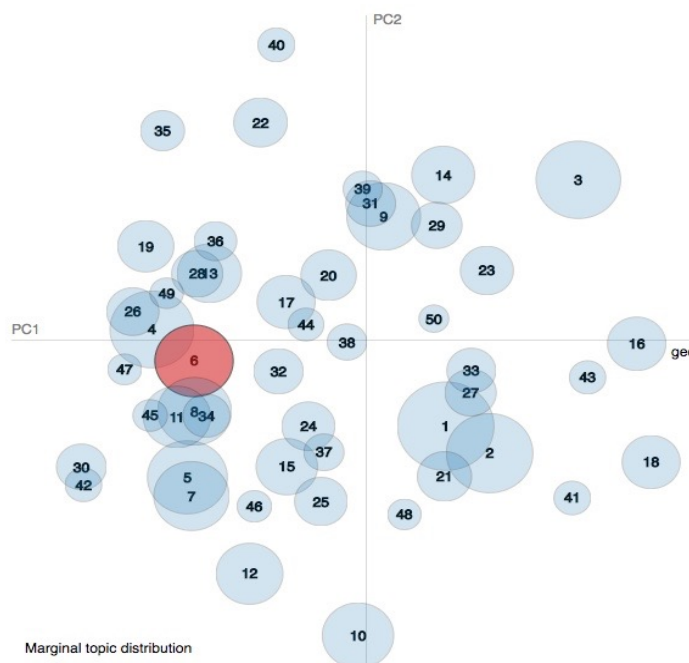
|               |   |
|---------------|---|
| PMID          | 20854478  |
| Article Title | Forest refugia revisited: nSSRs and cpDNA sequences support historical isolation in a wide-spread African tree with high colonization capacity, Milicia excelsa (Moraceae). |
| Pub Year      | 2010  |
|               | Africa [D000349]  |
|               | Bayes Theorem [D001499]   |
|               | Cluster Analysis [D016000]  |
|               | DNA, Chloroplast [D018742]  |
|               | genetics [Q000235]  |
|               | DNA, Plant [D018744]  |
|               | genetics [Q000235]  |
|               | Evolution, Molecular [D019143]  |
|               | Gene Flow [D051456]   |
|               | Gene Pool [D005788]   |
| Mesh          | Genetics, Population [D005828]  |
| Headings      | Microsatellite Repeats [D018895]  |
|               | Models, Genetic [D008957]   |
|               | Moraceae [D029586]  |
|               | genetics [Q000235]  |
|               | Pollen [D011058]  |

+plant +biodiversity-bow [50t,100p,2017-08-05,03.18.18] (7036 articles)

Selected Topic: 6 Previous Topic Next Topic Clear Topic

Slide to adjust relevance metric:<sup>(2)</sup>  $\lambda = 1$  0.0 0.2 0.4 0.6 0.8 1.0

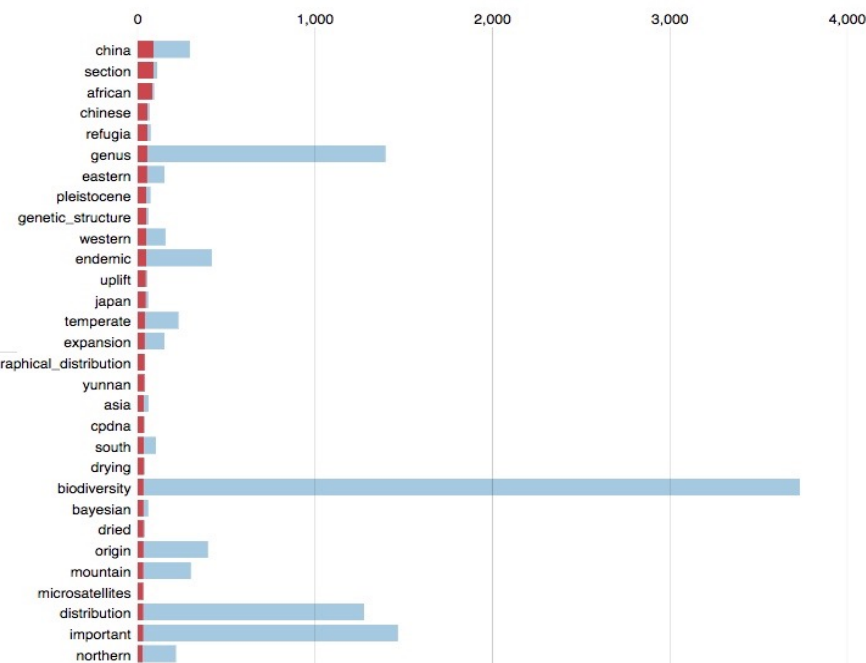
Intertopic Distance Map (via multidimensional scaling)



Marginal topic distribution



Top-30 Most Relevant Terms for Topic 6 (3.8% of tokens)



1. saliency(term w) = frequency(w) \* [sum\_t p(t|w) \* log(p(t|w)/p(t))] for topics t; see Chuang et. al (2012)  
 2. relevance(term w l topic t) =  $\lambda * p(w|t) + (1 - \lambda) * p(w|t)/p(w)$ ; see Sievert & Shirley (2014)

Topic model:

Topic models are trained on selected searches from the Pubmed corpus. Visualization based on the Python port of the LDAvis package.

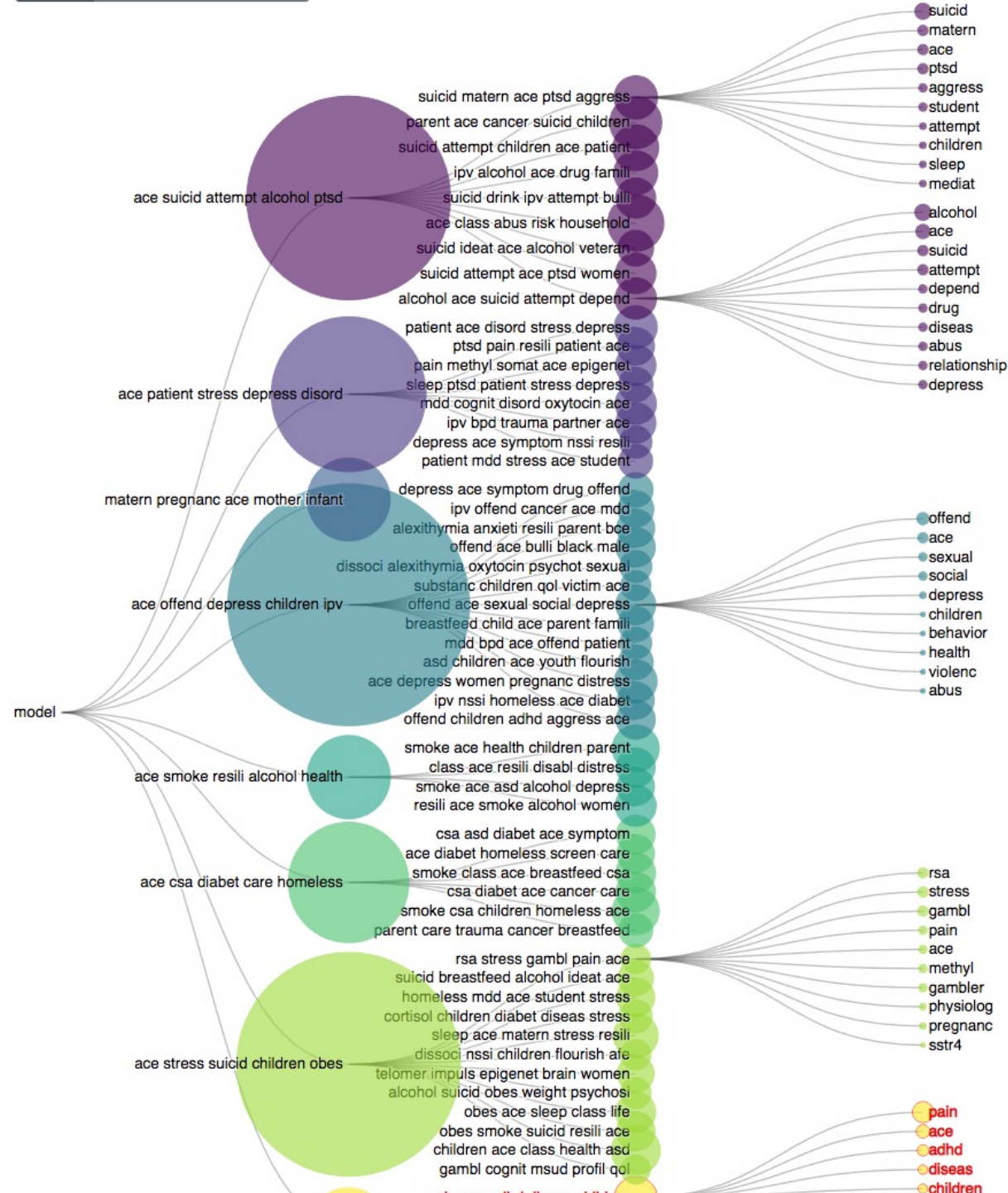
| Select a topic to see the related documents.                            |   |
|---|---|
| Topic 6   |   |
| [19723248] Phylogenetic origin of Phyllobium with a further implicati   | 0.64  |
| [23071663] Genetic diversity and population structure of cucumber (Cucu | 0.54  |
| [16622792] Biodiversity of Streptomyces of high-mountainous ecosystems  | 0.39  |
| [26467618] Phylogeography of Phytophagous Weevils and Plant Species in  | 0.35  |
| [23962409] Phylogeography sheds light on the central-marginal hypothesi | 0.34  |
| [24065181] Contemporary pollen-mediated gene immigration reflects the h | 0.31  |
| [17191876] Biologically active ibogan and vallesamine derivatives from  | 0.30  |
| [18206283] Reduced nitrogen has a greater effect than oxidised nitrogen | 0.28  |
| [27886271] Speciation and genetic diversity in Centaurea subsect. Phalo | 0.28  |
| [22081412] Establishing the phylogenetic origin, history, and age of th | 0.26  |
| [22546007] Phylogeographic analysis reveals significant spatial genetic | 0.26  |
| [24498103] Genetic differentiation and genetic diversity of Castanopsis | 0.26  |
| [17638329] Topoisomerase-II-inhibitory principles from the stems of Spa | 0.25  |
| [20397228] Prenylated benzophenone peroxide derivatives from Hypericum  | 0.25  |
| [16876446] Linking patterns and processes of species diversification in | 0.24  |
| [17509846] Diversity of viruses in Cryphonectria parasitica and C. nits | 0.24  |
| [18807258] Ectomycorrhizal characterization of an American chestnut (Ca | 0.24  |
| [20854478] Forest refugia revisited: nSSRs and cpDNA sequences support  | 0.24  |
| [23469278] Effect of degradation intensity on grassland ecosystem servi | 0.24  |
| [23478944] Post-Boreotropical dispersals explain the pantropical disjun | 0.24  |
| [27974324] The evolutionary history of Eugenia sect. Phyllocalyx (Myrta | 0.24  |
| [25072783] Temperate pine barrens and tropical rain forests are both ri | 0.23  |
| [22182994] A phylogeny of Delphinieae (Ranunculaceae) shows that Aconit | 0.22  |
| [23418542] Phylogeographic evidence for a link of species divergence of | 0.22  |
| [24028582] Large-scale pattern of genetic differentiation within Africa | 0.22  |
| [22730022] Range expansion of a selfing polyploid plant despite widespr | 0.21  |
| [23629053] Diversification of plant species in arid Northwest China: sp | 0.21  |
| [23560070] Tertiary origin and pleistocene diversification of dragon bl | 0.20  |
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| Mesh  | Africa [D000349]<br>Bayes Theorem [D001499]<br>Cluster Analysis [D016000]<br>DNA, Chloroplast [D018742]<br>genetics [Q000235]<br>DNA, Plant [D018744]<br>genetics [Q000235]<br>Evolution, Molecular [D019143]<br>Gene Flow [D051456]<br>Gene Pool [D005788] |
| Headings  | Genetics, Population [D005828]<br>Microsatellite Repeats [D018895]<br>Models, Genetic [D008957]<br>Moraceae [D029586]<br>genetics [Q000235]<br>Pollen [D011058]   |

enter a search term

| 1570              | 1580              | 1590                | 1600                 | 1610               | 1620              | 1630              | 1640                  | 1650                   | 1660                   | 1670                   | 1680                   |
|-------------------|-------------------|---------------------|----------------------|--------------------|-------------------|-------------------|-----------------------|------------------------|------------------------|------------------------|------------------------|
| turks<br>100%     | turks<br>100%     | moors<br>100%       | turks<br>100%        | turks<br>100%      | moors<br>100%     | turks<br>100%     | saracens<br>64%, 68%  | moors<br>100%, 69%     | moors<br>100%, 76%     | moors<br>100%, 69%     | turks<br>71%, 100%     |
| turkes<br>74%     | turkes<br>80%     | turks<br>100%       | moors<br>100%        | moors<br>100%      | turks<br>100%     | moors<br>100%     | parthians<br>59%, 59% | turks<br>69%, 100%     | turks<br>75%, 100%     | turks<br>69%, 100%     | moors<br>100%, 71%     |
| saracens<br>70%   | saracens<br>71%   | turkes<br>78%       | turkes<br>91%        | turkes<br>81%      | turkes<br>79%     | turkes<br>79%     | moors<br>100%         | saracens<br>73%, 75%   | tartars<br>69%, 74%    | tartars<br>70%, 71%    | spaniards<br>72%, 81%  |
| assyrians<br>65%  | persians<br>69%   | tartars<br>68%      | spaniards<br>69%     | saracens<br>70%    | saracens<br>71%   | barbarians<br>72% | turks<br>100%         | tartars<br>69%, 77%    | spaniards<br>66%, 72%  | saracens<br>67%, 71%   | saracens<br>74%, 72%   |
| hunnes<br>65%     | pagans<br>68%     | hungarians<br>67%   | hungarians<br>67%    | moores<br>68%      | moores<br>68%     | saracens<br>71%   | turkes<br>80%         | spaniards<br>67%, 70%  | saracens<br>66%, 68%   | persians<br>66%, 69%   | tartars<br>70%, 75%    |
| persians<br>64%   | barbarians<br>66% | tartarians<br>67%   | romans<br>66%        | pagans<br>67%      | tartarians<br>65% | tartars<br>68%    | pagans<br>75%         | tartarians<br>68%, 67% | turkes<br>63%, 67%     | indians<br>64%, 68%    | indians<br>71%, 67%    |
| vandales<br>64%   | armenians<br>65%  | saracens<br>66%     | portugals<br>66%     | mahumetans<br>66%  | arabians<br>62%   | moores<br>68%     | vandals<br>70%        | scythians<br>66%, 66%  | sarazens<br>65%, 64%   | sarazens<br>64%, 66%   | barbarians<br>66%, 71% |
| saracenes<br>63%  | assyrians<br>64%  | spaniards<br>65%    | barbarians<br>66%    | tartars<br>64%     | parthians<br>61%  | scythians<br>67%  | gothes<br>67%         | turkes<br>83%          | portugals<br>68%, 61%  | arabians<br>63%, 64%   | persians<br>65%, 72%   |
| tartarians<br>61% | egyptians<br>62%  | dalmatians<br>64%   | tartars<br>65%       | arabians<br>64%    | vandales<br>59%   | persians<br>64%   | infidels<br>67%       | moores<br>78%          | muscovites<br>62%, 64% | tartarians<br>61%, 65% | germans<br>68%, 68%    |
| arabians<br>61%   | arabians<br>61%   | nomades<br>64%      | sarazins<br>65%      | barbarians<br>64%  | portugals<br>59%  | romans<br>63%     | tartars<br>65%        | sarmatians<br>72%      | hungarians<br>70%      | armenians<br>67%       | venetians<br>73%       |
| spaniards<br>60%  | infidels<br>61%   | persians<br>63%     | moores<br>65%        | infidels<br>64%    | visigothes<br>59% | tartarians<br>62% | persians<br>63%       | pagans<br>70%          | moores<br>67%          | mahometans<br>67%      | imperialists<br>68%    |
| gothes<br>60%     | gothes<br>61%     | gaules<br>62%       | sicilians<br>63%     | armenians<br>63%   | pagans<br>58%     | danes<br>61%      | moores<br>63%         | huns<br>69%            | carthaginians<br>66%   | scythians<br>66%       | hungarians<br>67%      |
| medes<br>59%      | scythians<br>60%  | moores<br>61%       | britans<br>62%       | mahometans<br>63%  | spaniards<br>58%  | indians<br>61%    | arabians<br>63%       | mahometans<br>69%      | venetians<br>65%       | mahumetans<br>66%      | romans<br>67%          |
| normans<br>59%    | romans<br>59%     | scythians<br>61%    | sarasins<br>62%      | persians<br>63%    | persians<br>58%   | venetians<br>61%  | carthaginians<br>63%  | infidels<br>69%        | portuguez<br>65%       | barbarians<br>65%      | arabians<br>67%        |
| moores<br>59%     | sarasins<br>59%   | lombardes<br>61%    | persians<br>61%      | grecians<br>61%    | infidells<br>58%  | hungarians<br>61% | danes<br>61%          | vandals<br>68%         | indians<br>65%         | moscovites<br>65%      | scythians<br>66%       |
|                   |                   | parthians<br>60%    | carthaginians<br>61% | aethiopians<br>61% | indians<br>58%    | polonians<br>61%  | heathens<br>61%       | mahumetans<br>68%      | castilians<br>64%      | infidels<br>64%        | muscovites<br>66%      |
|                   |                   | gothes<br>60%       | sirians<br>52%       | pomerania<br>60%   | alani<br>57%      | getes<br>57%      | grecians<br>61%       | sarazens<br>68%        | persians<br>64%        | moores<br>63%          | islanders<br>65%       |
|                   |                   | sacrata<br>60%      | mahumetists<br>52%   | epire<br>56%       | tartars<br>58%    | spartan<br>54%    | tartarians<br>60%     | goths<br>67%           | germans<br>63%         | goths<br>63%           | cossacks<br>65%        |
|                   |                   | florentia<br>60%    | perspicua<br>51%     | benin<br>57%       | goths<br>57%      | patricians<br>54% | scythians<br>59%      | persians<br>67%        | pyrates<br>63%         | portugals<br>62%       | numidians<br>64%       |
|                   |                   | aider<br>60%        | longi<br>50%         | aria<br>57%        | infidels<br>57%   | scythians<br>53%  | turkish<br>59%        | turk<br>67%            | mamalukes<br>63%       | spaniards<br>61%       | swedes<br>64%          |
|                   |                   | celtiberians<br>60% | bramas<br>50%        | abassia<br>57%     | herules<br>57%    | libyan<br>53%     | goths<br>59%          | barbary<br>66%         | arabians<br>63%        | chineses<br>60%        | gauls<br>64%           |

| topic ↓↑ | over time | top words   | proportion of corpus    |
|----------|-----------|---|-------------------------|
| Topic 1  |           | increas given breath decreas secret apn cbg patient chang rate prn retract desatur morn rr thick                            | 7.4%                    |
| Topic 2  |           | secret patient suction note cloudi thick will frequent continu desatur progress requir see window amount monitor stabl thin | 27.7%                   |
| Topic 3  |           | secret stabl cloudi suction set thin will small vent current moder no desatur toler thick                                   | 54.2%                   |
| Topic 4  |           | vital suction axillari resp secret temp will desatur puls bp file hc height tempsrc weight                                  | 4.3%                    |
| Topic 5  |           | trach patient today mom rt secret care cuff rn back dad suction bag educ leak   | 6.3%                    |
| topic    | over time | top words   | proportion of corpus ↓↑ |
| Topic 10 |           | stabl no patient desatur current set will remain vent note toler ventil issu overnight night                                | 23.3%                   |
| Topic 5  |           | desatur stabl patient will suction current vent one overnight recov set increas breath agit remain                          | 15.7%                   |
| Topic 6  |           | secret cloudi suction small thin desatur stabl will vent set clear no toler well moder thick                                | 15.4%                   |
| Topic 9  |           | patient desatur see note progress increas expect bag recov suction wean back requir rn bradi satur ventil will              | 9.4%                    |
| Topic 8  |           | vital axillari stabl resp puls bp file temp tempsrc weight hc height night will desatur                                     | 8.6%                    |
| Topic 7  |           | window no mom today desatur min trach dad hour well patient place toler parent room time                                    | 7.9%                    |
| Topic 2  |           | desatur increas night patient breath suction rr time retract apn given decreas frequent no awar morn                        | 6.8%                    |
| Topic 4  |           | note monitor sx frequent desatur cloudi vent distress suction time close continu thin will requir                           | 5.7%                    |
| Topic 3  |           | cuff desatur leak trach volum low no chang ml high breath increas alarm place post time                                     | 4.4%                    |

| Title  | Score              |
|--|--------------------|
| Associations between adverse events in childhood and chronic widespread pain in adulthood: are they  | 0.995311439037323  |
| Adverse Childhood Experiences and Psychological Distress in Juvenile Offenders: The Protective Influ | 0.9952369332313538 |
| Dying to be famous: retrospective cohort study of rock and pop star mortality and its association wi | 0.9951603412628174 |
| Impact of early personal-history characteristics on the Pace of Aging: implications for clinical tri | 0.9951075911521912 |
| The relationship between childhood adversity and food insecurity: 'It's like a bird nesting in your  | 0.9950807690620422 |
| Enhanced startle reflexivity during presentation of visual nurture cues in young adults who experien | 0.9949988722801208 |
| The evidence base for routine enquiry into adverse childhood experiences: A scoping review.          | 0.9949987530708313 |
| On Becoming Trauma-Informed: Role of the Adverse Childhood Experiences Survey in Tertiary Child and  | 0.99470454454422   |
| Screening for Toxic Stress Risk Factors at Well-Child Visits: The Addressing Social Key Questions fo | 0.99470454454422   |



| cluster | #para, | #  | terms                             |
|---------|--------|----|-----------------------------------|
| 3       | 2106,0 | 13 | ace offend depress children ipv   |
| 6       | 1873,0 | 12 | ace stress suicid children obes   |
| 0       | 2309,0 | 9  | ace suicid attempt alcohol ptsd   |
| 1       | 1258,0 | 8  | ace patient stress depress disord |
| 5       | 1159,0 | 6  | ace csa diabet care homeless      |
| 2       | 744,0  | 4  | matern pregnanc ace mother infant |
| 4       | 1082,0 | 4  | ace smoke resili alcohol health   |
| 7       | 778,0  | 4  | pain ace disabl adhd diseas       |

# The transgenerational transmission of maternal adverse childhood experiences (ACEs): Insights from placental aging and infant autonomic nervous system reactivity.

<PubmedArticle>OBJECTIVE: To test alterations in placental cellular aging as one pathway by which maternal early adversity influences physiologic development in her offspring. METHODS: Maternal report of her adverse childhood experiences (ACE) was obtained prenatally along with measures of prenatal stress and demographic information. Placentas (N = 67) were collected at birth and telomere length (TL) was measured in four separate fetally-derived placental tissues: amnion, chorion, villus, and umbilical cord. At four months of age, infants completed the still-face paradigm (SFP)



