

Toward Community-Driven, Shared Literature Annotation Resources

Jin-Dong Kim
Database Center for Life Science
(DBCLS)

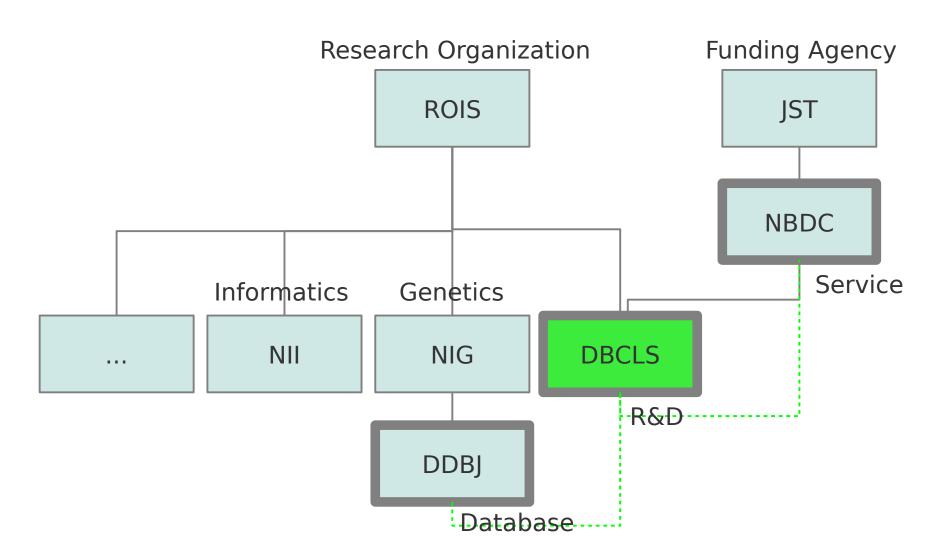


DBCLS

- Database Center for Life Science
 - A government-funded research center
 - For integration of databases of life sciences
 - It annually organizes
 - → BioHackathon series
 - → BLAH series

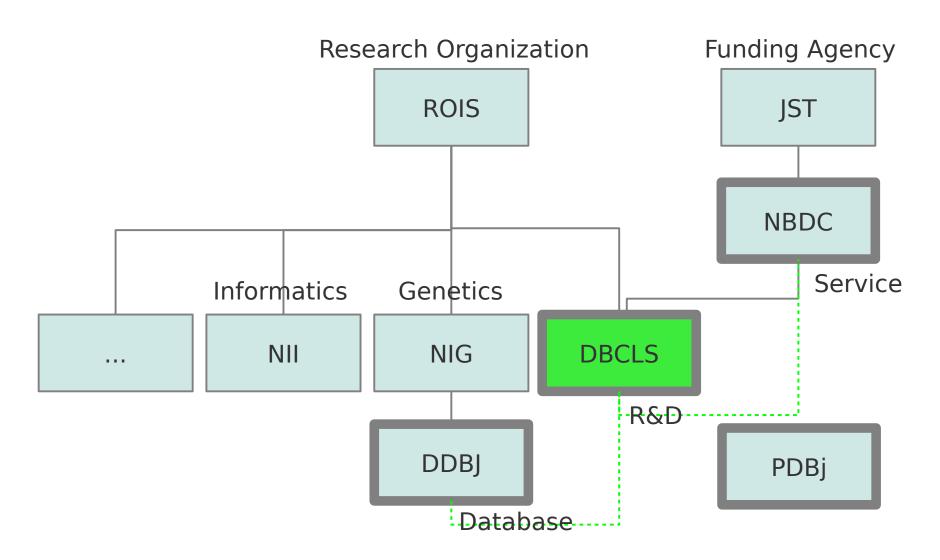


DBCLS



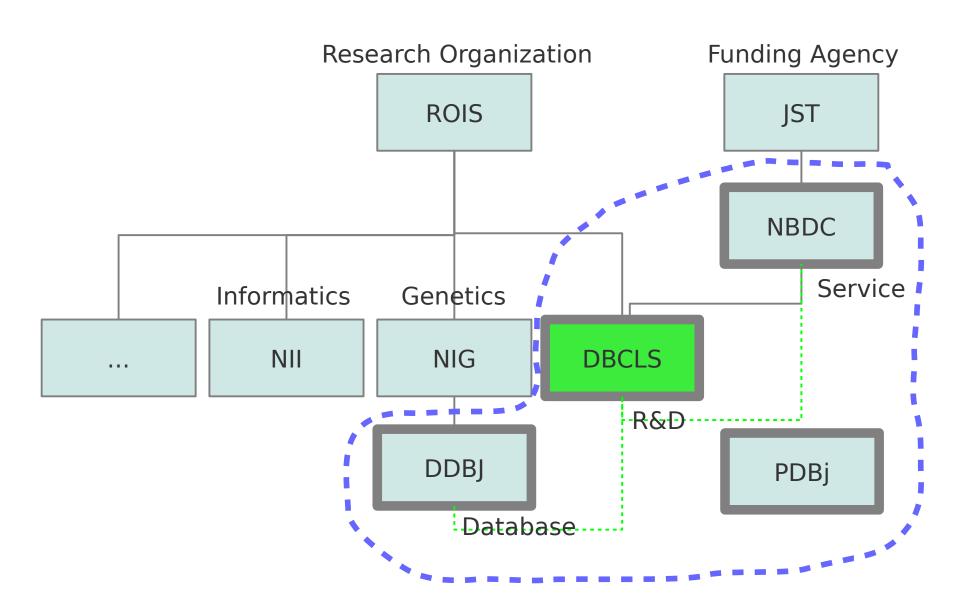


Introduction to DBCLS





Introduction to DBCLS





"Inheritance involves the passing of discrete units of inheritance"

Proceedings of the Natural History Society of Brünn, 1866

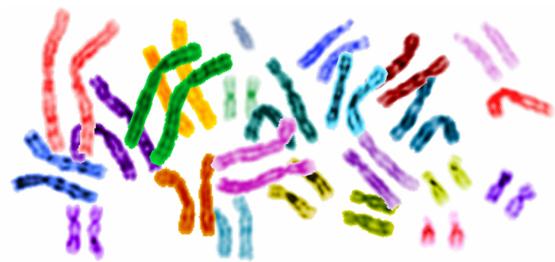


Gregor Mendel



"Inheritance involves the passing of discrete units of inheritance"

Proceedings of the Natural History Society of Brünn, 1866

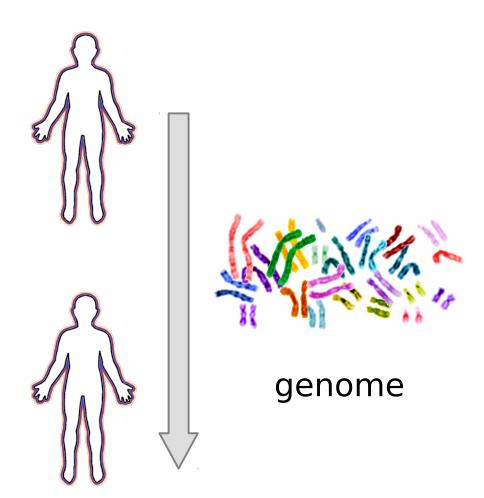




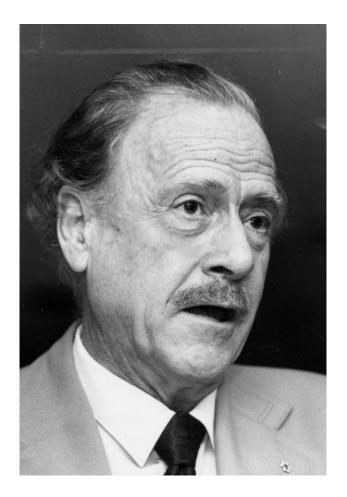
Gregor Mendel



Heredity







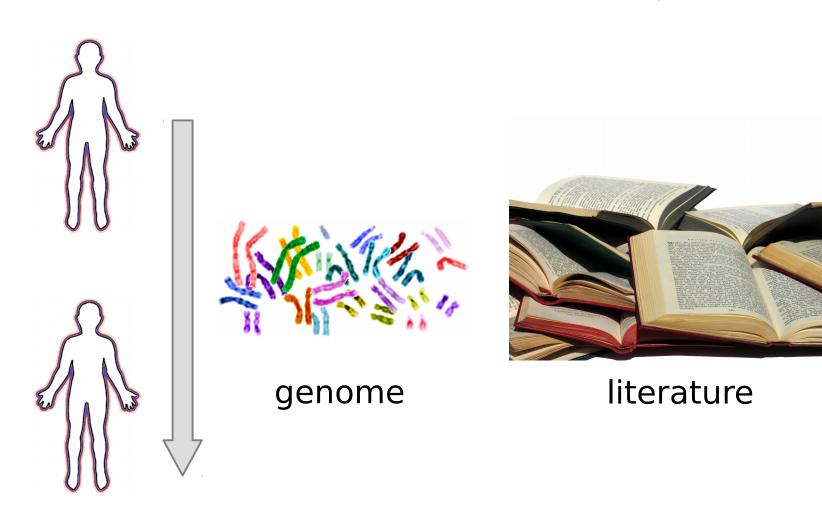
Marshall McLuhan

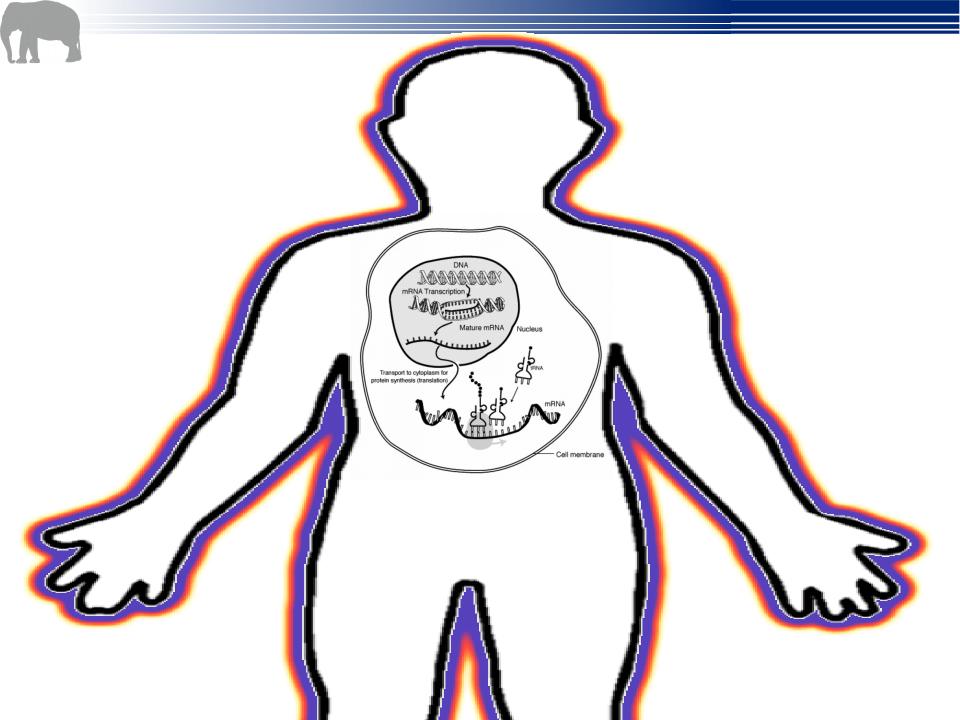
"all media are extensions of our human senses, bodies and minds."

The Medium Is the Massage, 1967



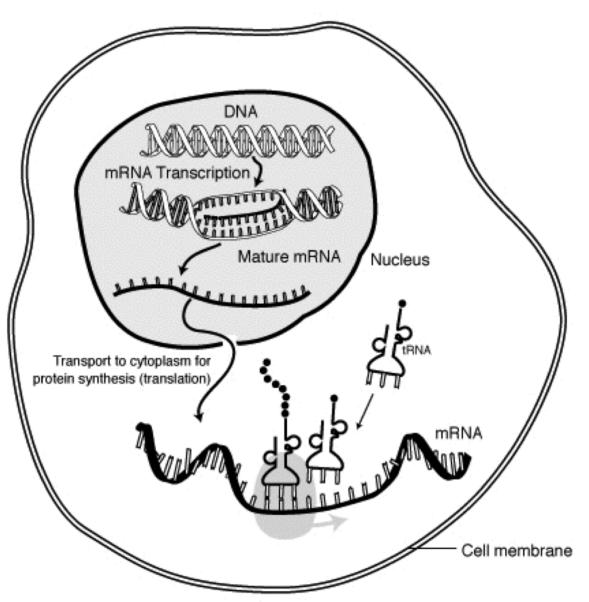
Extension of Heredity







Cell

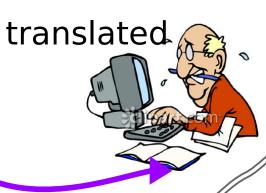




Society



transcribed

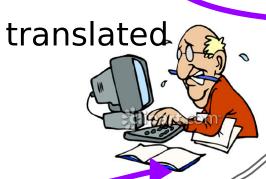


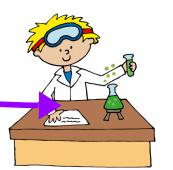


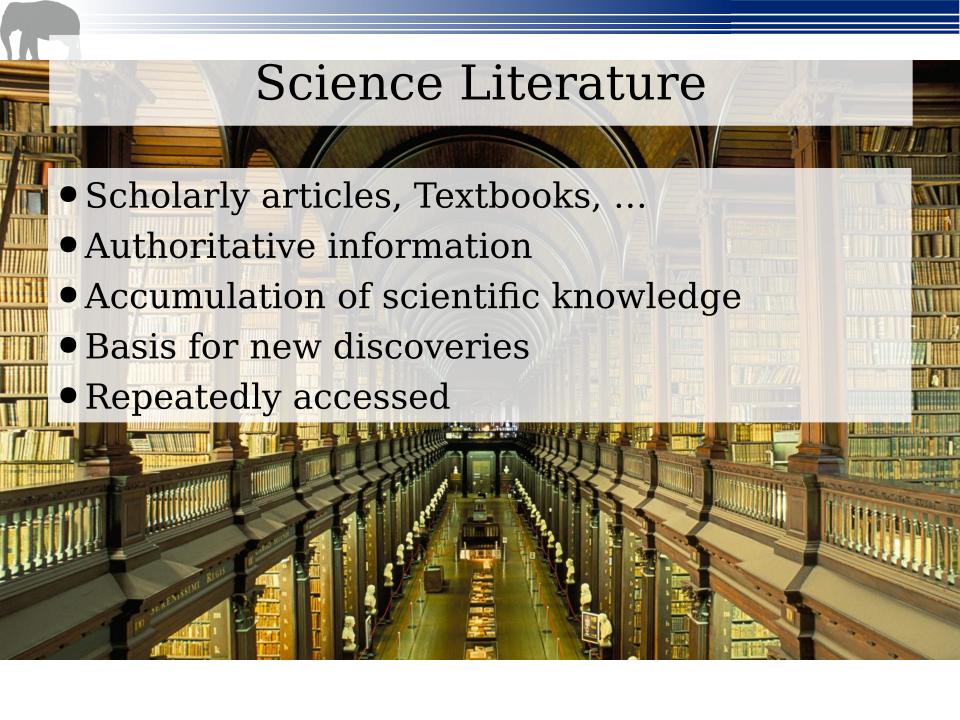
Society

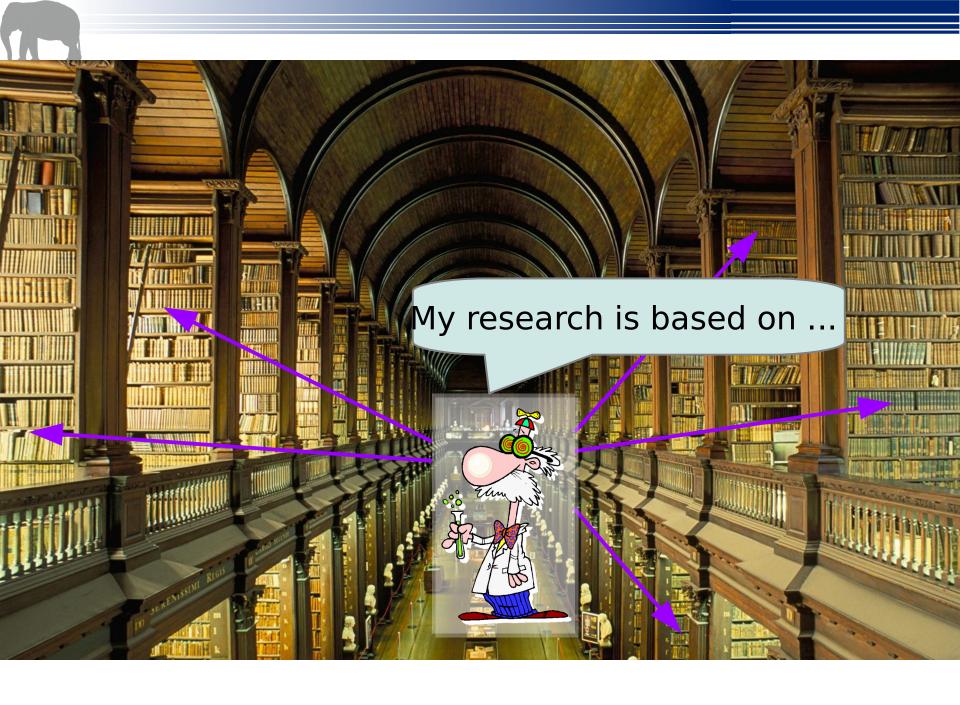


transcribed



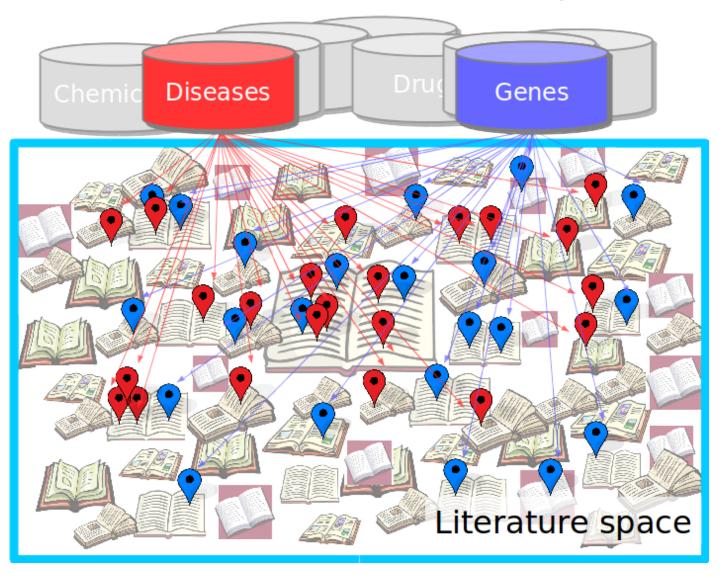






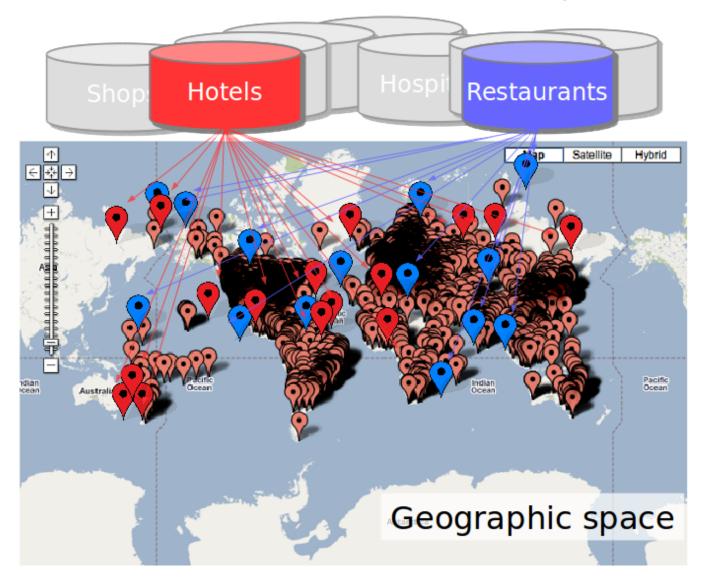


Literature Indexing



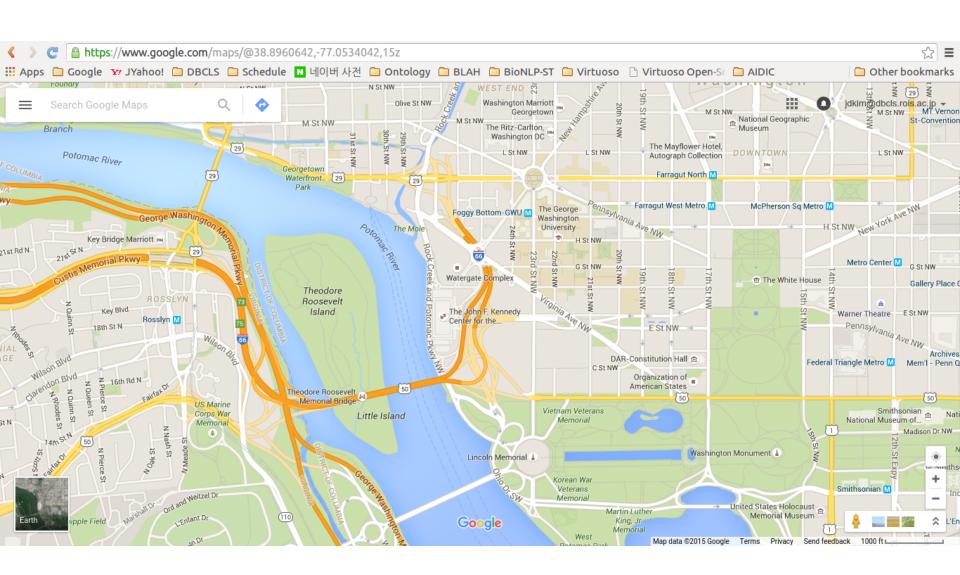


Geospatial Indexing



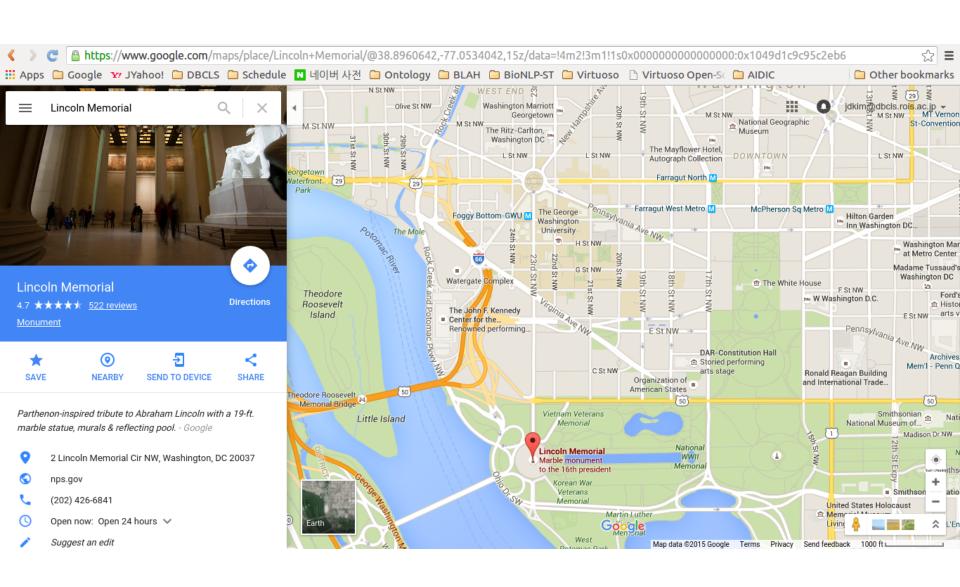


Google Maps (Washington D.C.)



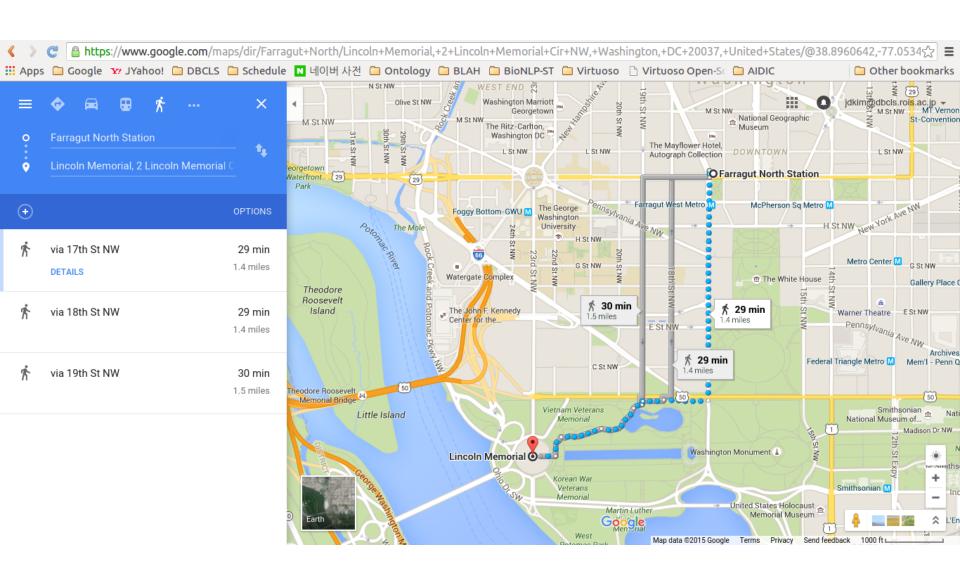


Entities





(Geospatial) Pathways





Entities

Gene_4790

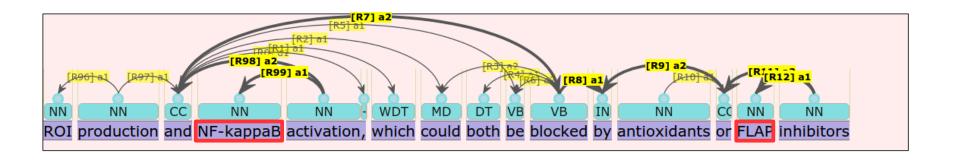
Gene

ROI production and NF-kappaB activation, which could both be blocked by antioxidants or FLAP inhibitors



(Linguistic) Pathways

Gene_4790
ROI production and NF-kappaB activation, which could both be blocked by antioxidants or FLAP inhibitors





Why people use Google Maps?

- Useful
 - Contents
- Easy to use
 - ✓ Interface
 - → To access
 - → To exchange
 - → To create
 - → To reuse

Geospatial annotations



Literature annotation

- Do we have good contents?
 - Many groups are producing annotations.

- Do we have good ways to access them?
 - Ann. resources are scattered and isolated.



Let's link them to each other, and Share them, altogether. *BLAH!*



PubAnnotation

- ✓ Is a repository of literature annotation
- ✓ Is based on a scalable storage system
- Specifically aims at PubMed and PMC
- Solicits contribution of annotations from the community
- Solves various problems for sharing the annotations
 - → Alignment
 - → Global addressing system
 - → REST APIs





REPOSITORY SEARCH NEWS GUIDES ABOUT

Make your annotation public, and more useful!

> top

Documents									
PMC	9,044	keywords Q source ID							
PubMed	8,424,657	keywords Q source ID							
FirstAuthor	8	keywords Q source ID							

Q

News-

- (09 Mar 2016) Recent access problem
- (25 Feb 2016) System problem on 25/02/2016
- (19 Jan 2016) a new project status, Uploading, added.
- (04 Jan 2016) News service begins.

With most annotations

Projects (146)

Name		# Ann.	Updated At	Status
PubmedHPO		12,437,742	2016-01-31	Uploading
DisGeNET		3,117,504	2016-01-28	Beta
NEUROSES		2,151,082	2016-02-24	Beta
CRAFT-treebank		844,123	2015-11-19	Beta
bionlp-st-ge-2016-spacy-par		225,680	2016-05-25	Released
spacy-test		136,597	2016-05-25	Released
FSU-PRGE		59,505	2016-05-17	Released
craft		52,960	2015-10-13	Beta
jnlpba-st-training		51,290	2016-09-09	Released
PennBiolE		23,881	2016-05-17	Released

Recently updated

Name		# Ann.	Updated At	Status
jnlpba-st-training		51,290	2016-09-09	Released
jnlpba-st-test		6,005	2016-09-07	Uploading
bionlp-st-seedev-2016-training		109	2016-09-05	Uploading
bionlp-st-bb3-2016-training		1,292	2016-09-05	Released
bionlp-st-cg-2013-training		10,935	2016-08-22	Released
bionlp-st-pc-2013-traing		7,855	2016-08-22	Released
bionlp-st-id-2011-training		5,609	2016-08-22	Released
bionlp-st-epi-2011-training		7,595	2016-08-22	Released
Ab3P-abbreviations		2,342	2016-07-29	Beta
AnEM_full-texts		689	2016-07-27	Uploading

Challenges for Integration

- Format is not standardized
 - Many proprietary formats
- Texts are changed
 - → PubMed, PMC change texts
 - → Web masters change texts
 - → Annotation projects change texts
 - ✓ For
 - → Cleaning
 - → Convenience for annotation
 - Unicode → ASCII



Challenges for Integration

- Format is not standardized
 - A matter of conversion
- Texts are changed
 - Breaks stand-off annotation
 - → Character offsets become invalid
 - ✓ Solution
 - → Sequence alignment (BLAST!)



PubAnnotation

GATA3-Driven Th2 Responses Inhibit TGF-1Induced FOXP3 Expression and the Formation of Regulatory T Cells Transcription factors act in concert to induce lineage commitment towards Th1, Th2, or T regulatory (Treg) cells, and their counter-regulatory mechanisms were shown to be critical for polarization between Th1 and Th2 phenotypes. FOXP3 is an essential transcription factor for natural, thymus-derived (nTreg) and inducible Treg (iTreg) commitment; however, the mechanisms regulating its expression are as yet unknown. We describe a mechanism controlling iTreg polarization, which is overruled by the Th2 differentiation pathway. We demonstrated that interleukin 4 (IL-4) present at the time of T cell priming inhibits FOXP3. This inhibitory mechanism was also confirmed in Th2 cells and in T cells of transgenic mice overexpressing GATA-3 in T cells, which are shown to be deficient in transforming growth factor (TGF) β -mediated FOXP3 induction. This inhibition is mediated by direct binding of GATA3 to the FOXP3 promoter, which represses its transactivation process. ...

Local Annotation

This inhibitory mechanism was also confirmed in Th2 cells 208-213, Protein transgenic mice over-expressing GATA-3 in T cells, which are single over-deficient in transforming around the (TGF)-beta-mediated FOXP3 induction.



PubAnnotation

GATA3-Driven Th2 Responses Inhibit TGF-1Induced FOXP3 Expression and the Formation of Regulatory T Cells Transcription factors act in concert to induce lineage commitment towards Th1, Th2, or T regulatory (Treg) cells, and their counter-regulatory mechanisms were shown to be critical for polarization between Th1 and Th2 phenotypes. FOXP3 is an essential transcription factor for natural, thymus-derived (nTreg) and inducible Treg (iTreg) commitment; however, the mechanisms regulating its expression are as yet unknown. We describe a mechanism controlling iTreg polarization, which is overruled by the Th2 differentiation pathway. We demonstrated that interleukin 4 (IL-4) present at the time of T cell priming inhibits FOXP3. This inhibitory mechanism was also confirmed in Th2 cells and in T cells of transgenic mice overexpressing GATA-3 in T cells, which are shown to be deficient in transforming growth factor (TGF) β -mediated FOXP3 induction. This inhibition is mediated by direct binding of GATA3 to the FOXP3 promoter, which represses its transactivation process. ...

Local Annotation

Upload & align

This inhibitory mechanism was also confirmed in Th2 cells 208-213, Protein transgenic mice over-expressing GATA-3 in T cells, which are significant to be deficient in transforming around the (TGF)-beta-mediated FOXP3 induction.



PubAnnotation

GATA3-Driven Th2 Responses Inhibit TGF-1Induced FOXP3 Expression and the Formation of Regulatory T Cells Transcription factors act in concert to induce lineage commitment towards Th1, Th2, or T regulatory (Treg) cells, and their counter-regulatory mechanisms were shown to be critical for polarization between Th1 and Th2 phenotypes. FOXP3 is an essential transcription factor for natural, thymus-derived (nTreg) and inducible Treg (iTreg) commitment; however, the mechanisms regulating its expression are as yet unknown. We describe a mechanism controlling iTreg polarization, which is overruled by the Th2 differentiation pathway. We demonstrated 838-833, Protein (IL-4) present at the time of T cell priming inhibits FOXP3. This inhibitory mechan is was all 936-941, Protein 2 cells and in T cells of transgenic mice overexpressing GATA-3 in T cells, which are shown to be deficient in transforming growth factor (TGF) β-mediated FOXP3 induction. This inhibition is mediated by direct binding of GATA3 to the FOXP3 promoter, which represses its transactivation process. ...

Local Annotation

Upload & align

This inhibitory mechanism was also confirmed in Th2 cells 208-213, Protein transgenic mice over-expressing GATA-3 in T cells, which are significant to be deficient in transforming around the (TGF)-beta-mediated FOXP3 induction.

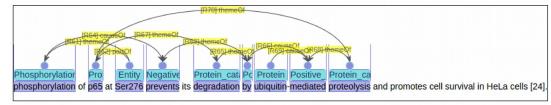


- Jin-Dong Kim, "A generalized LCS algorithm and its application to corpus alignment", Proceedings of the 6th International Joint Conference on Natural Language Processing (IJCNLP), pp.14-18, 2013
 - A definite solution to text variant problem
 - ✓ It can align even full paper articles sourced by two different groups.

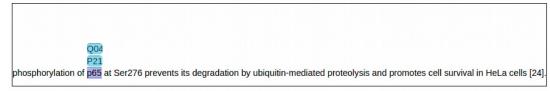


Aligned annotations from different groups

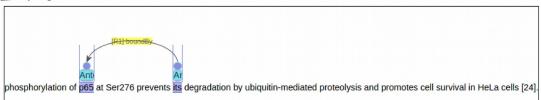
bionlp-st-ge-2016-reference



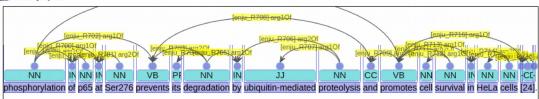
bionlp-st-ge-2016-uniprot



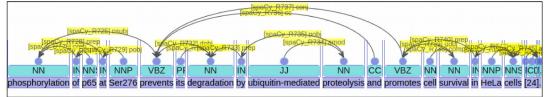
bionlp-st-ge-2016-coref



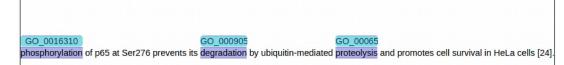
pmc-enju-pas



bionlp-spacy-parsed



GO-BP





- Aligned annotations
 - http://www.pubannotation.org/

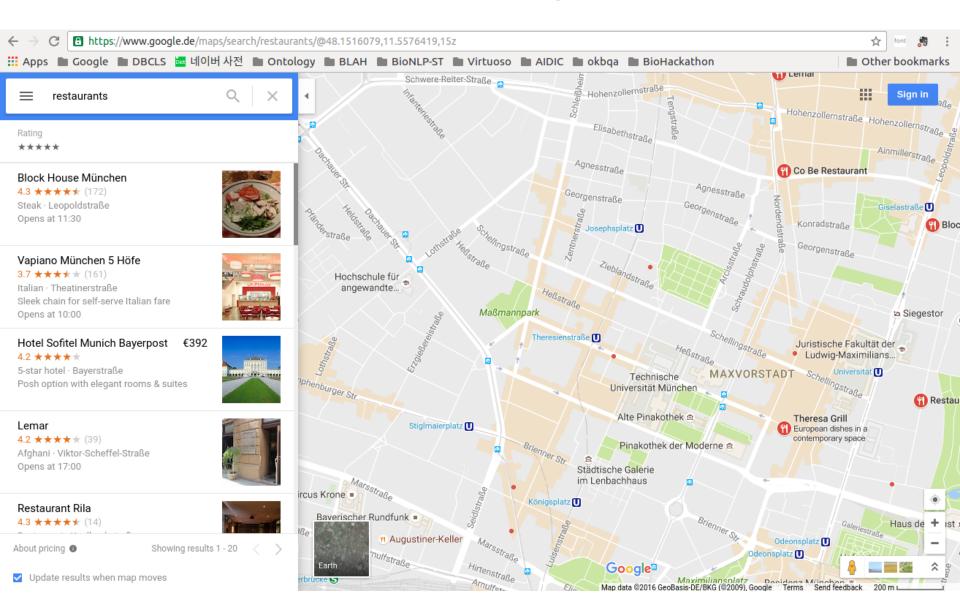


Global addressing system

- Persistently preserve the texts of all the articles from PubMed / PMC(OA)
 - ✓ UTF-8
 - ASCII conversion is provided
- Offset indices are stably maintained



a case of Google Map





A case of PubAnnotation

- Example of URL
 - http://pubannotation.org/docs/sourcedb /PubMed/sourceid/10022882/spans/606-71 0/annotations/visualize
- How to get the URL (Example)
 - http://pubannotation.org/docs/sourced b/PubMed/sourceid/10022882



Global addressing system

- Persistently preserve the texts of all the articles from PubMed / PMC(OA)
 - ✓ UTF-8
 - ASCII conversion is provided
- Offset indices are stably maintained



Exchange

Hi Bill, what is the diagnostic test for MERS infection?

Check this <u>link</u> out.

FYR, I've annotated it using NCIt, OBI, and SNOMEDCT.

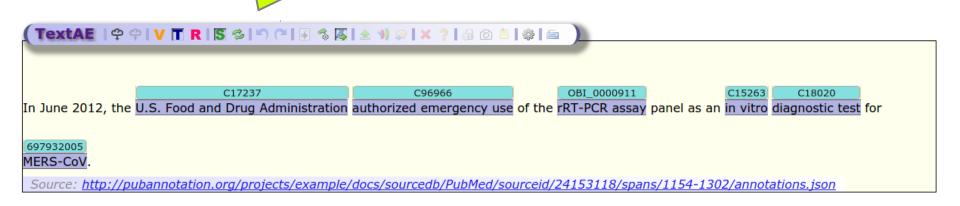


Exchange

Hi Bill, what is the diagnostic test for MERS infection?

Check this <u>link</u> out.

FYR, I've ap ptated it using NCIt, OBI, and SNOMEDCT.





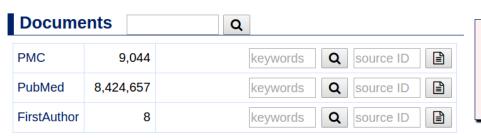
Currently, in PubAnnotation



English 日本語 signup log

Make your annotation public, and more useful!

> ton



Q

News

- (09 Mar 2016) Recent access problem
- (25 Feb 2016) System problem on 25/02/2016
- (19 Jan 2016) a new project status, Uploading, added.
- (04 Jan 2016) News service begins.

Projects (146)

With most annotations

Name		# Ann.	Updated At	Status
PubmedHPO	O ₀	12,437,742	2016-01-31	Uploading
DisGeNET	O ₀	3,117,504	2016-01-28	Beta
NEUROSES	00	2,151,082	2016-02-24	Beta
CRAFT-treebank	9	844,123	2015-11-19	Beta
bionlp-st-ge-2016-spacy-par	O ₀	225,680	2016-05-25	Released
spacy-test	O ₀	136,597	2016-05-25	Released
FSU-PRGE	O ₀	59,505	2016-05-17	Released
craft	9	52,960	2015-10-13	Beta
jnlpba-st-training	9	51,290	2016-09-09	Released
PennBiolE	6	23,881	2016-05-17	Released

Recently updated

Name		# Ann.	Updated At	Status
jnlpba-st-training	6	51,290	2016-09-09	Released
jnlpba-st-test		6,005	2016-09-07	Uploading
bionlp-st-seedev-2016-training		109	2016-09-05	Uploading
bionlp-st-bb3-2016-training		1,292	2016-09-05	Released
bionlp-st-cg-2013-training	6	10,935	2016-08-22	Released
bionlp-st-pc-2013-traing	6	7,855	2016-08-22	Released
bionlp-st-id-2011-training	6	5,609	2016-08-22	Released
bionlp-st-epi-2011-training	6	7,595	2016-08-22	Released
Ab3P-abbreviations	6	2,342	2016-07-29	Beta
AnEM_full-texts	6	689	2016-07-27	Uploading



PubAnnotation

- New version to be released soon
 - Performance improved
 - Interface improved
 - BioC conversion to be supported
 - → Thanks to the NCBI team
 - Bug fixes



TextAE

- To access/edit annotations
 - http://textae.pubannotation.org
 - ✓ Has fully RESTful APIs



PubDictionaries

- To share dictionary resources
 - Current version
 - → http://pubdictionaries.org
 - New version
 - → http://new.pubdictionaries.org

Scientific literature is a central repository of scientific knowledge - every important scientific discovery ha **Shared Annotation Targets** technology of literature annotation thus has played a central role for text mining. While literature annotation still requires enormous effort despite a number of years of concentrated experience, the productivity of literature annotation is recently significantly improved, and there are quite a few groups roducing annotations in large scale. While many groups have released those data sets of literatur nnotation to the public, however, the way of sharing those widely valuable resources still remains at imitive level, e.g., relying on individual exchange of archived files. anwhile, the advancement of internet web technology has enabled much convenient ways of collaborating sharing data. For example, the technology of web 2.0 has enabled crowdsourci ducing generation, and web 3.0 has enabled machine-understandable web of data. annotation in general sense, for example, the Google Map system allows even la ns easily produce geographic annotations and immediately share them by simp ers ending the annotations. Before the era of Google Ma the URL representing producing and sharing geographic annotati What never been a simple work. mal Google Map so convenient may be attributed t sharing of the coordin same system/(latitude and longitud provision dereference URposition to every and annotation

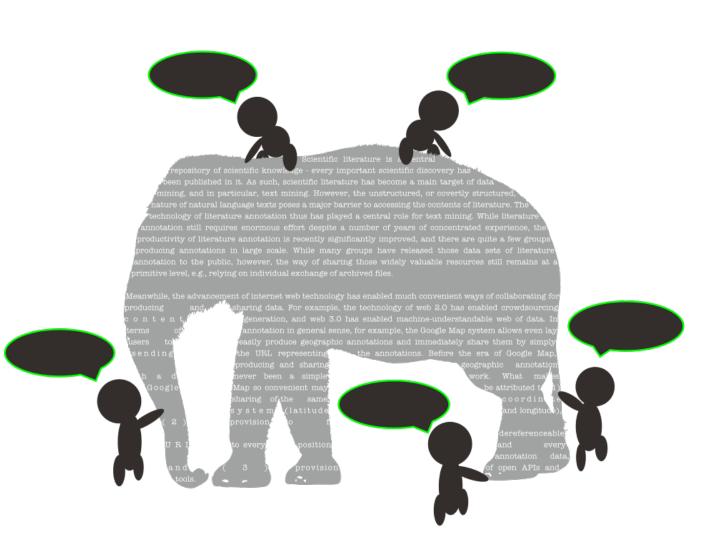


Many literature annotation projects





None of them is complete



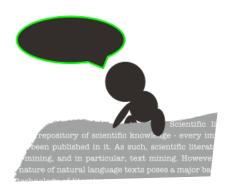


None of them is complete

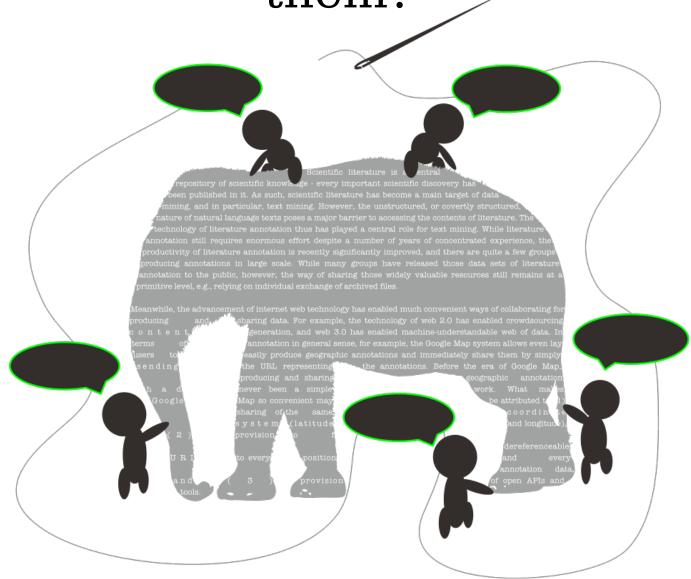




None of them is complete

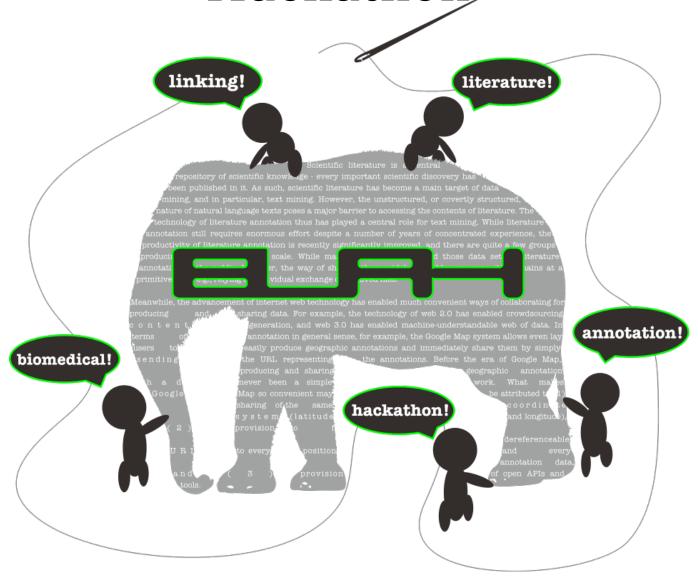


Then, why don't we collect & link them?





Biomedical Linked Annotation Hackathon





BLAH

- Biomedical Linked Annotation Hackathon
 - ✓ BLAH
 - → Feb. 2015, Kashiwa
 - ✓ BLAH2
 - → Nov. 2015, Mishima / Ito
 - **✔** BLAHMUC
 - → Oct. 2016, Munich

BLAH

- Biomedical Linked Annotation Hackathon
 - ✓ BLAH
 - → Feb. 2015, Kashiwa
 - ✓ BLAH2
 - → Nov. 2015, Mishima / Ito
 - ✓ BLAHMUC
 - → Oct. 2016, Munich
 - ✓ BLAH3
 - →Jan. 2017, Tokyo



Thank you! Happy October Blah!