

Challenges in representing *Rich Data* and **Annotations**

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Deep Learning Revolution

NLP—No longer only Language

- More accurate computer vision and speech recognition models
- Not just written language, but Multimodal understanding
- Representing data is already challenging
- Representing multi-modal, multi-layered metadata (annotations, in our case) which remains in sync with the data and maintain consistency within and across layers can be quite challenging

Underlying Assumption...

Metadata in the form of annotations—
Linguistic or otherwise—play an important role
in the underlying research

An Example Scenario... **Only Written Text**

Robustness to Tokenization

A Grant is (*finally*) Funded!

- Phase I

- Use existing Treebank-ed text (= use existing **trees** and **tokens**)
- Add a few layers of rich annotation
 - ▶ Word Sense (depends only on tokens)
 - ▶ Named Entities (depends only on tokens)
 - ▶ Propositions (depends on Tree structure)
 - ▶ Coreference (depends on Tree structure)

- Phase II

- It is found to be very important to make minor changes to...
 - ▶ Treebank and PropBank layer guidelines so they more are in sync
 - ▶ A minor change in tokenization to split on some hyphens

*As a result, some tokens
are split into multiple tokens*

New-York-based (single token)



New-York	(first token)
-	(second token)
based	(third token)

Now, Update existing Annotations

Easy, Right?

well, **Not** Necessarily

Factors determining the **Difficulty**

- How the annotation layers are represented?
- How tight is the data coupling between the layers?
- How detailed are the specifications?
 - a. **within** each layer
 - b. **between** the layers

- Depending on the answers to the above questions (and maybe a few more)
 - It might be a nightmarish scenario, or
 - It might be a reasonable task
- **Both** options will very likely require human intervention (annotator)
- The degree of that intervention and the complexity of the task will be determined to a large degree by the above design decisions

This was not a Hypothetical scenario

It happened in the OntoNotes project

- Owing to the design of the underlying representation, it was...
 - a reasonable task
 - ▶ Each layer had a detailed specification
 - ▶ The layers—both *inter-* and *intra-* used a relational data model
 - ▶ The layers were not too tightly coupled

Multiple-Layers in OntoNotes

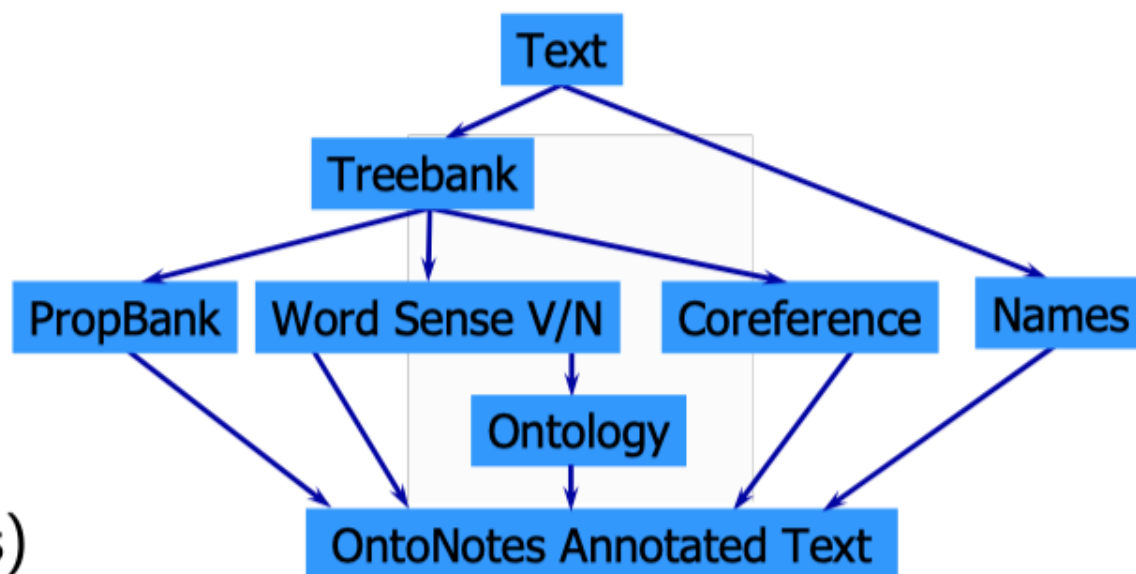
- Multiple layers of annotation

- Syntax
- Propositions
- Word sense
- Coreference
- Names
- Ontology

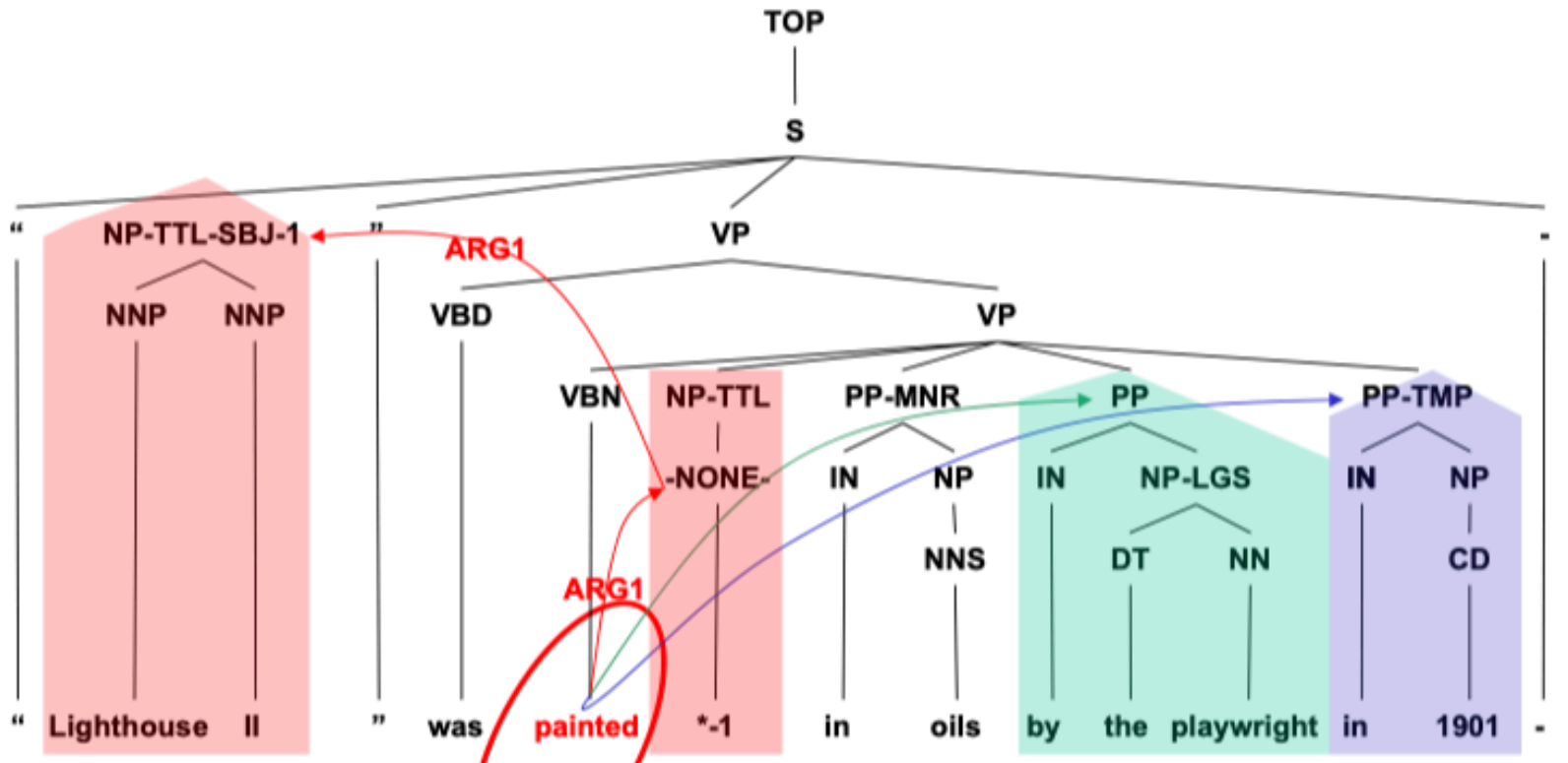
- Multilingual resource

- English (~1M words)
- Chinese (~1M words)
- Arabic (~1M words)

- Parallel Data



Interpreting Tree Pointers



wsj/00/wsj_0037.mrg 67 5 gold set.02 ---- 0:2-ARG0 5:0-rel 6:1-ARG1 10:2-ARGM-TMP
 wsj/00/wsj_0037.mrg 68 5 gold paint.01 ---- 5:0-rel 1:1*6:0-ARG1 8:1-ARG2-in 10:1-ARG0-by 12:1-ARGM-TMP
 wsj/00/wsj_0037.mrg 69 21 gold exchange.01 ---- 17:2-ARG0 21:0-rel 22:1-ARG1 23:1-ARGM-TMP
 wsj/00/wsj_0037.mrg 69 35 gold say.01 ---- 31:1-ARG0 35:0-rel 0:2*37:0-ARG1

Propbank Frames

wsj_0037.mrg 67 5 gold set.02 ----- 0:2-ARG0 5:0-rel 6:1-ARG1 10:2-ARGM-TMP
wsj_0037.mrg 68 5 **paint.01** 5:0-rel 1:1*6:0-**ARG1** 8:1-**ARG2**-in 10:1-**ARG0** 12:1-ARGM-TMP
wsj_0037.mrg 69 21 gold exchange.01 ----- 17:2-ARG0 21:0-rel 22:1-ARG1 23:1-ARGM-TMP
wsj_0037.mrg 69 35 gold say.01 ----- 31:1-ARG0 35:0-rel 0:2*37:0-ARG1

```
<!DOCTYPE frameset SYSTEM "frameset.dtd">
<frameset>
  <predicate lemma="paint">
    <note>
      Frames file for 'paint' based on sentences in wsj and automatic expansion via verbnet.
    </note>
    <roleset id="paint.01" name="put paint on a surface" vncls="25.1">
      <roles>
        <role descr="agent, painter" n="0"> <vnrole vncls="25.1" vntheta="Agent"/></role>
        <role descr="surface" n="1"> <vnrole vncls="25.1" vntheta="Destination"/></role>
        <role descr="explicit mention of paint" n="2"> <vnrole vncls="25.1"
          vntheta="Theme"/> </role>
      </roles>
```

Word Sense Inventories

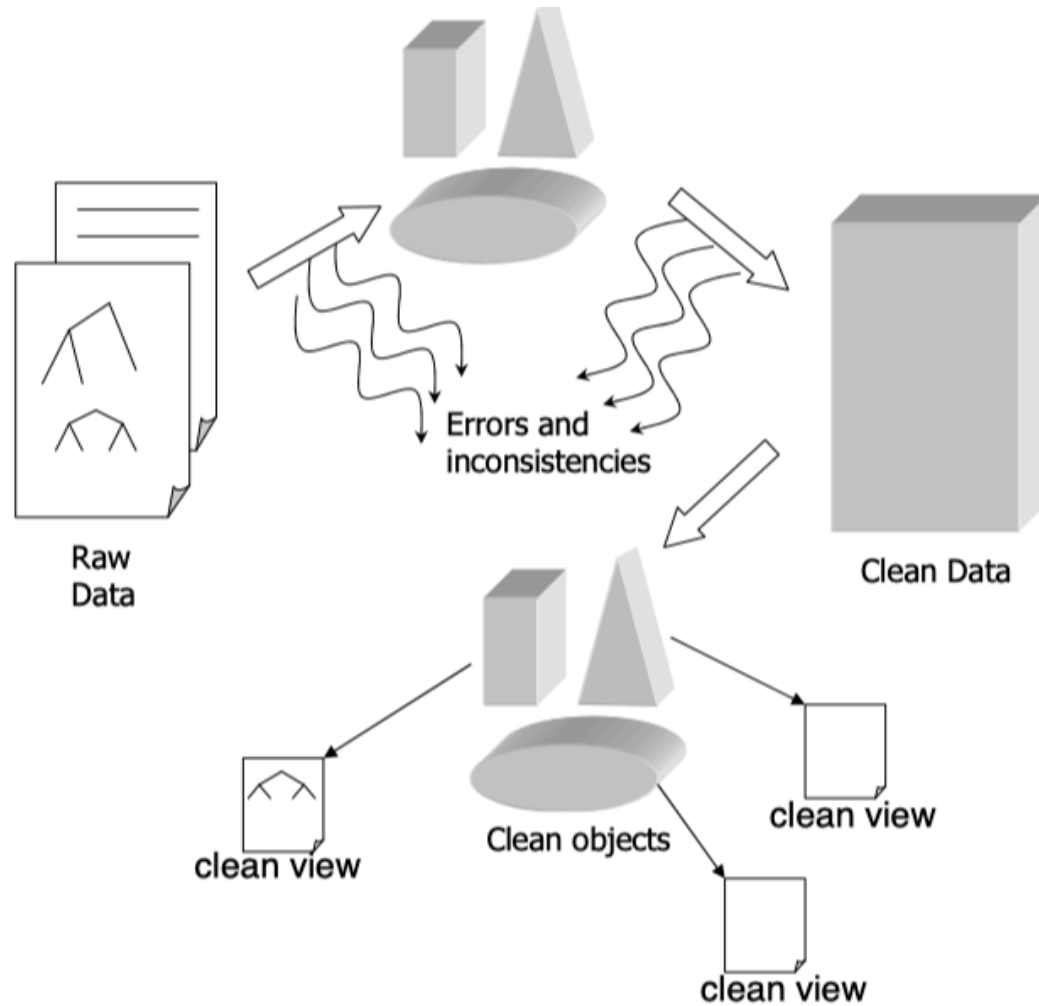
PRO **Judging** from the Americana in Haruki Murakami 's " A Wild Sheep Chase " (Kodansha, 320 pages, \$18.95 *U*) , baby boomers on both sides of the Pacific have a **lot** in common .

wsj/00/wsj_0037.mrg	0 1	judge-v	2
wsj/00/wsj_0037.mrg	0 36	lot-n	1

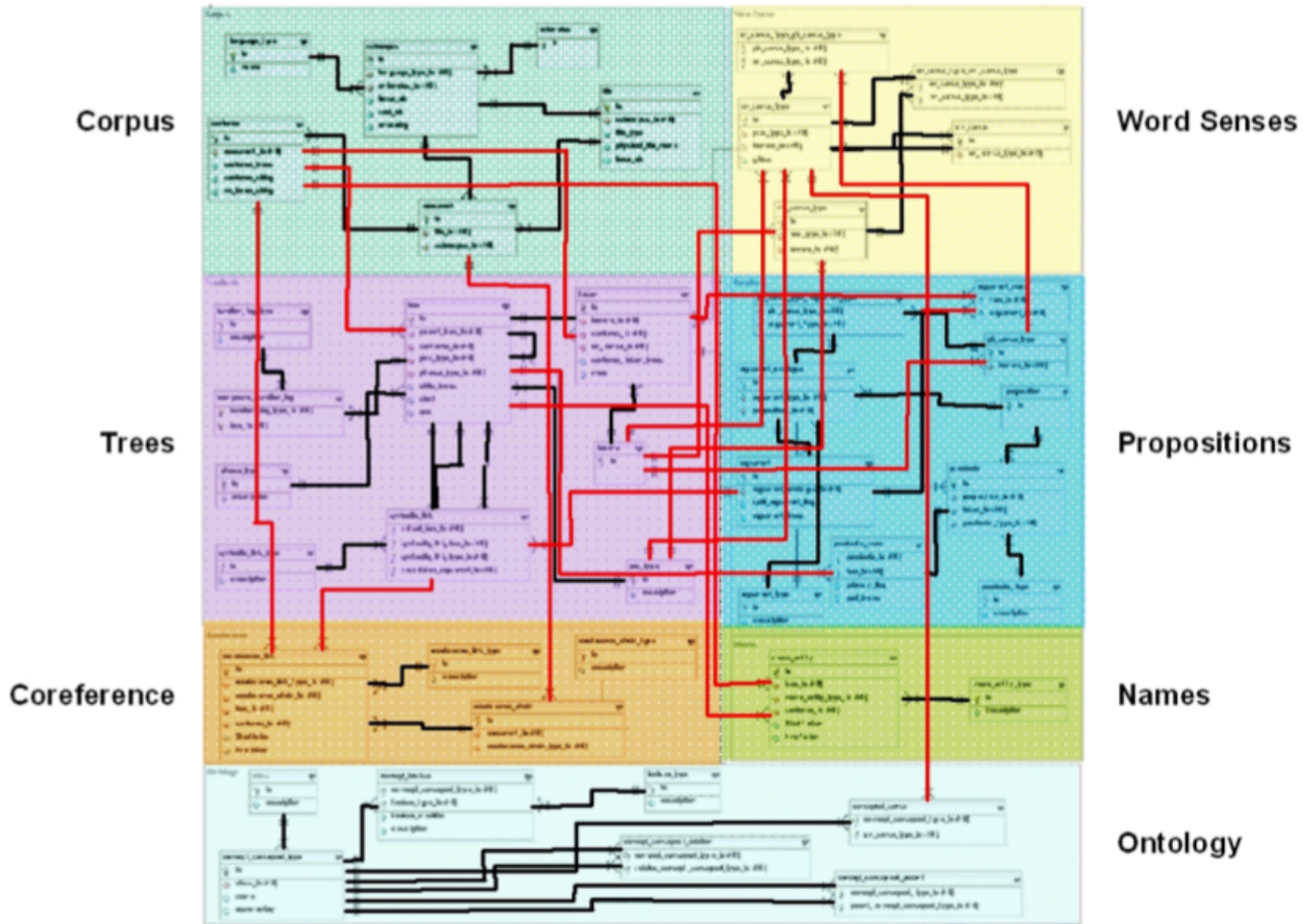
Sense Number

```
<?xml version="1.0" ?>
<!DOCTYPE inventory SYSTEM "inventory.dtd">
<inventory lemma="judge-v">
  <sense group="1" n="1" name="act as an official judge">
    <examples> She was asked to judge the fancy-dress competition. </examples>
    <mappings> <wn version="2.1">1,5</wn> <pb>judge.01</pb> </mappings>
  </sense>
  <sense group="1" n="2" name="form an opinion, or conclusion">
    <examples> They quickly judged him unfit to join the team. </examples>
    <mappings> <wn version="2.1">2,3,4</wn> <pb>judge.01</pb> </mappings>
  </sense>
</inventory>
```

Annotation Lifecycle



Entity-Relationship Diagram



Multiple-Layers in OntoNotes

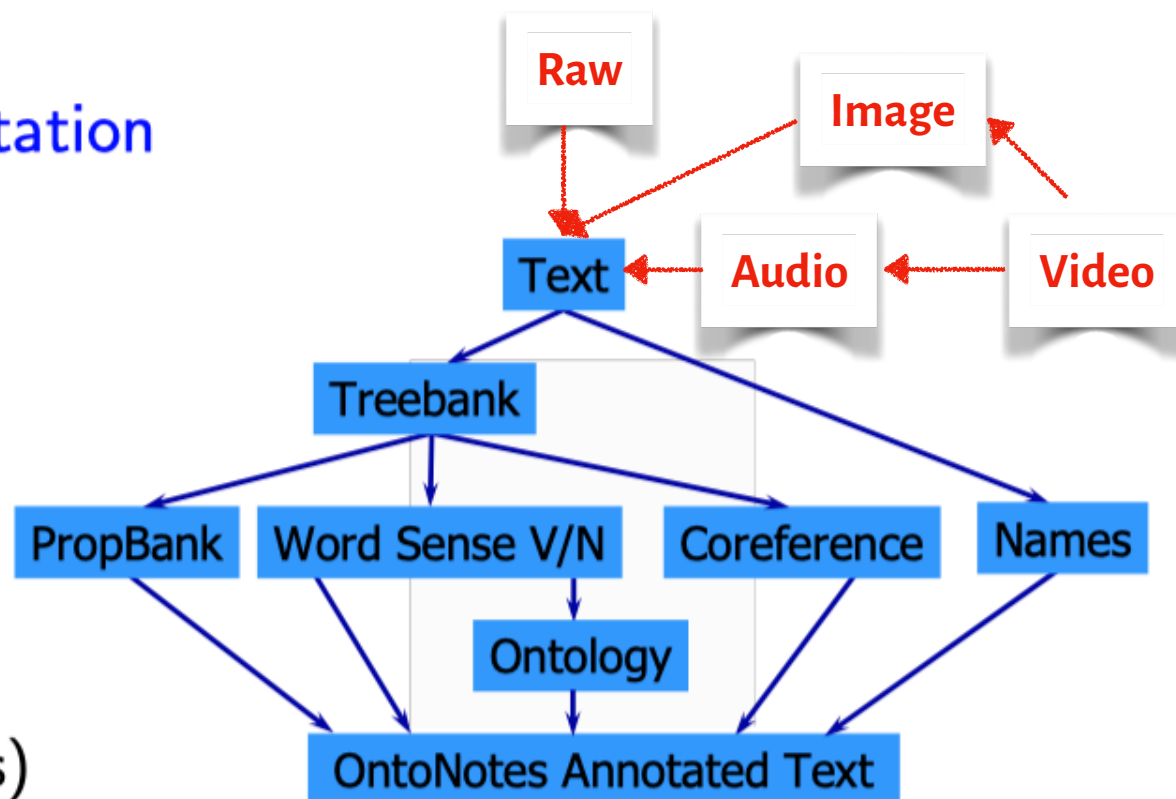
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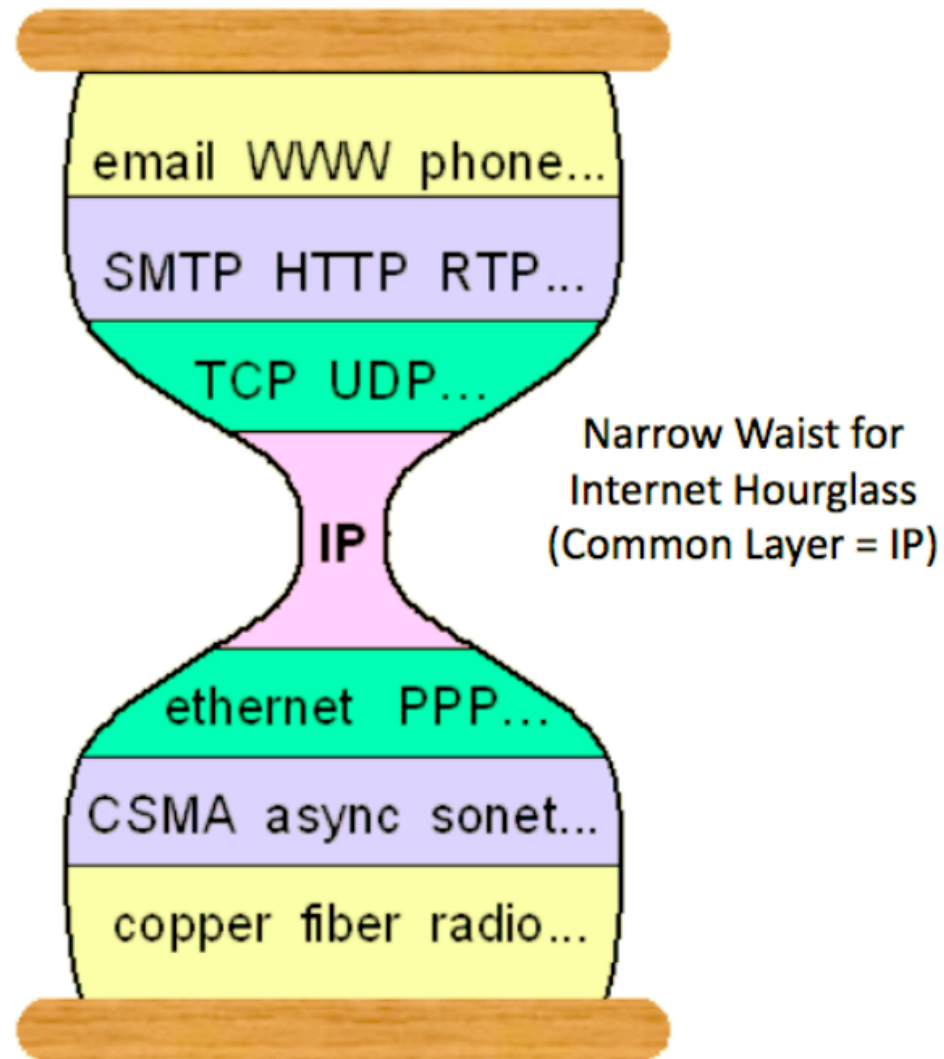
But isn't **SQL** last-century?

But don't take our word for it. Take Google's.



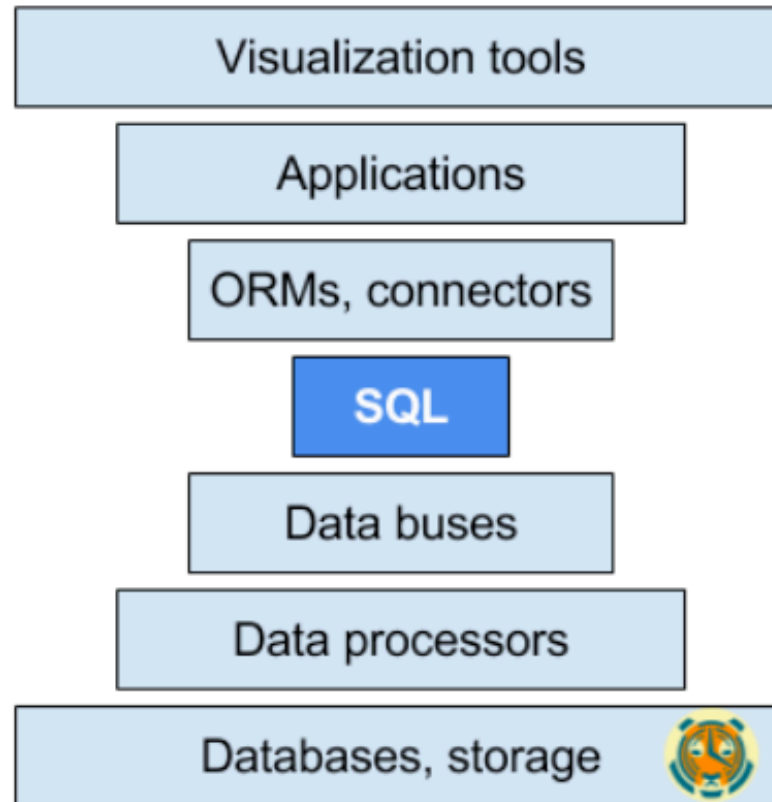
Take a look at Google's second major **Spanner** paper ([Spanner: Becoming a SQL System](#), May 2017), and you'll find that it bolsters our independent findings.

Lessons from the **Internet**



IP as the Networking Universal Interface ([source](#)).

SQL as the **slim**-waist



The Data Universal Interface

Most of the issues are carried over to **Clinical Narrative**
We are adding other modalities involved...

Audio,
Images,
Videos
...

The same task can become **exponentially harder**
or impossible

Annotation as Code

What if...

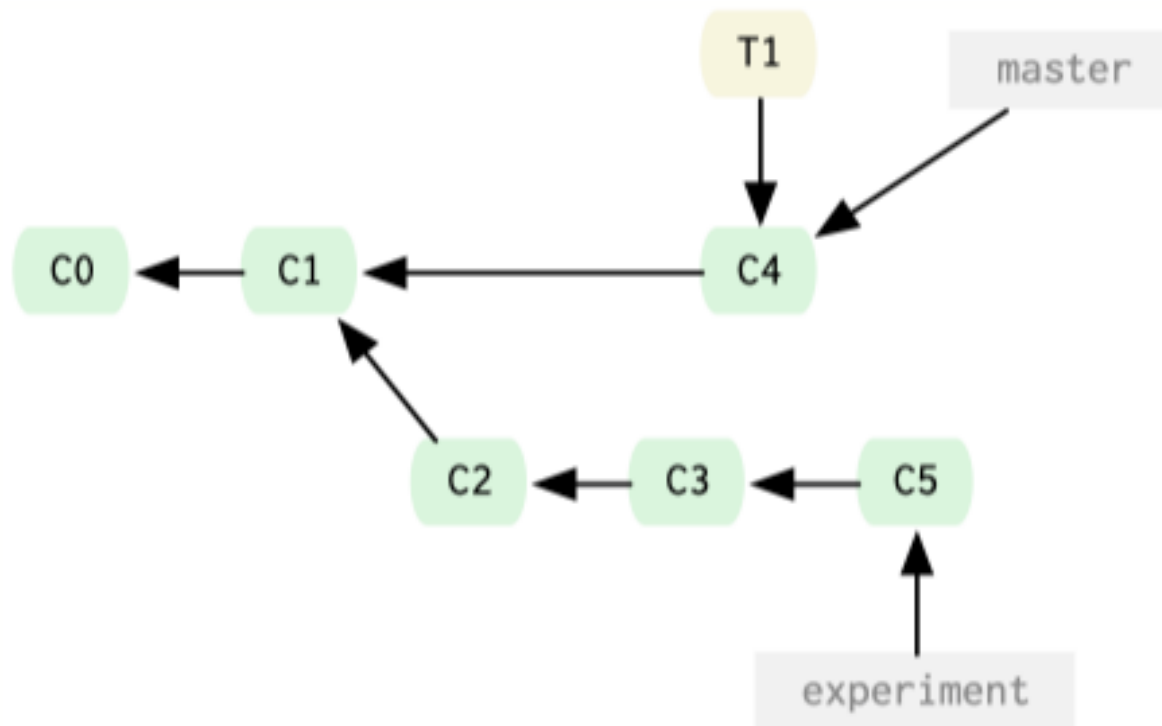
- Annotations are represented as we represent and manage source-code?
- One peculiarity—increased complexity of the semantics for such “language”
- We might benefit from a version control mechanism

Functional Data Structures

Ideas from *git*

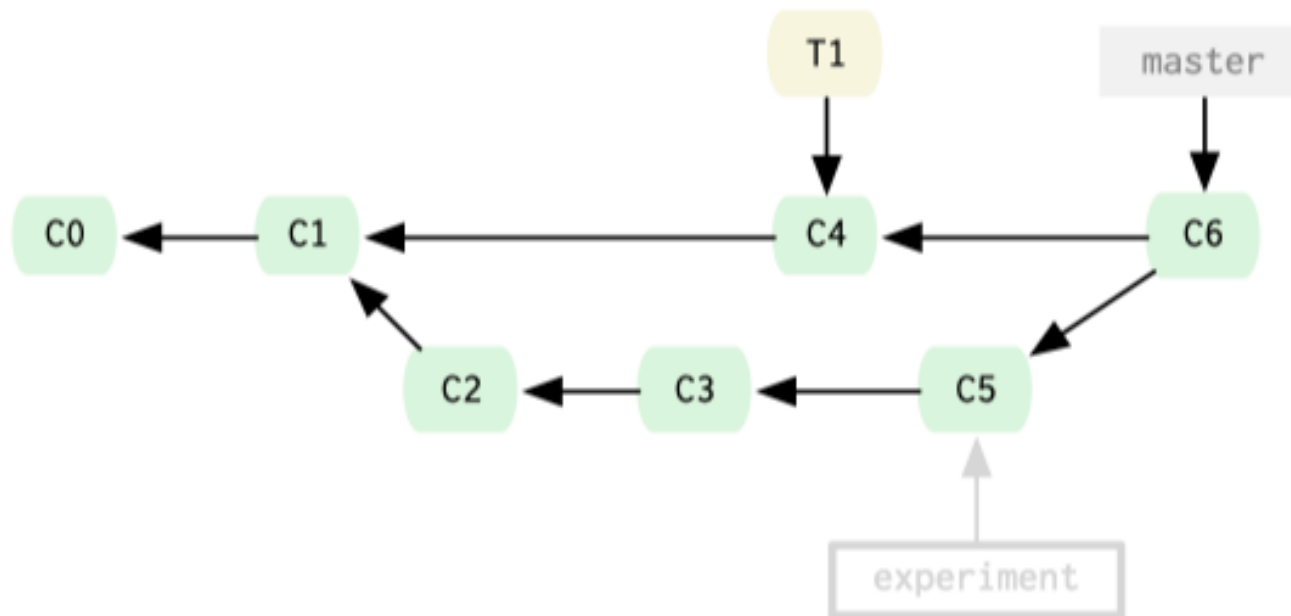
- *Each version* of *each layer* of annotation is an *incremental operation* on top of the *earlier version*.
- Try to maximize *deterministic bi-directional* transformations
- Minimize *lossy uni-directional* transformations
- Track *annotation version* and the *guideline specification* dependencies
- Create new *annotation snapshots*
- Consistency checks using *content hashes*

Past Annotations **Reachable**



Annotation **Guidelines** can be *kept in sync*

Ease of Experimentation

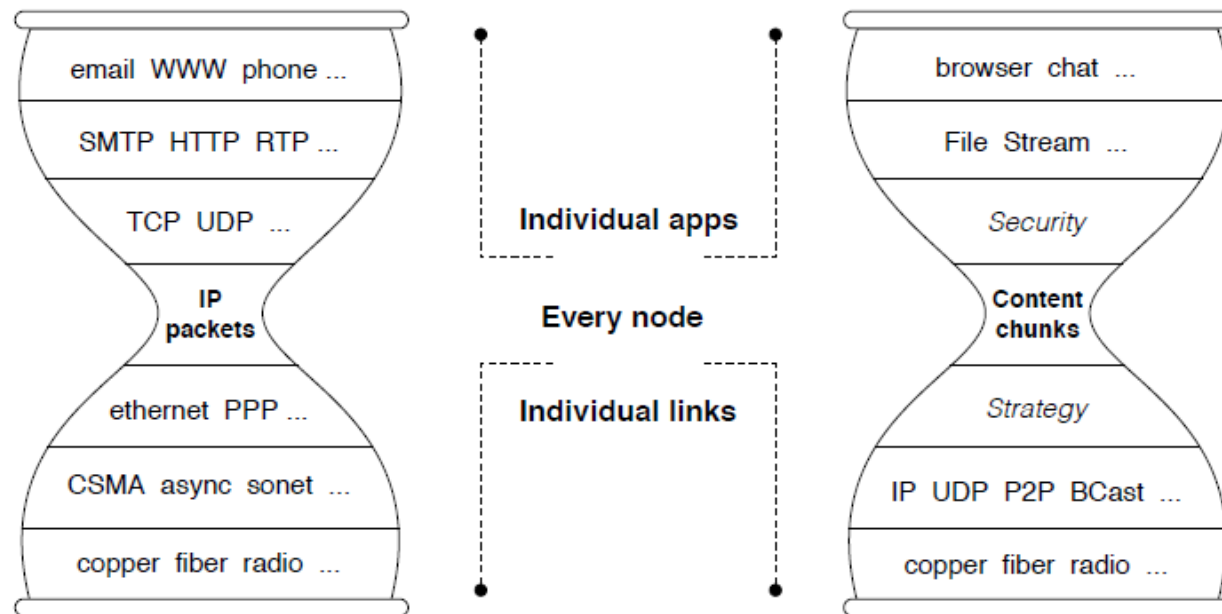


Very important to know exactly what **guidelines** were used for a particular set of annotations

Internet Philosophy

Next Generation of the **Internet**

- **Plain text** files where possible—UTF-8 for serialization or even base64
- **Media Containers**—akin to the next generation of internet that focuses on content—Named Data Networking (NDN) or Content Centric Networking (CCN)



Layer Tags

When a .parse is not the .parse

- File extensions are typically used to determine content, but when it comes to layers of annotation, things can easily get complicated
- Example, a **.parse** might contain one of many *kinds* or *qualities* of parses
 - **Dependency parse**
 - ▶ Universal Dependency v2.0
 - ▶ Custom dependency
 - **Constituency parse**
 - ▶ Gold parse (with traces)
 - ▶ Automatically generated
 - Using Charniak parser, model (A)
 - Using Charniak parser, model (B)
 - Using Berkley parser
 - ...
- Similarly the columns in a **.conll** file might be interpreted differently depending on the year and task involved

When a .parse is not the .parse

- File extensions are typically used to determine content, but when it comes to layers of annotation, things can easily get complicated
- **oohggg_parse** might contain one of many *kinds* or *qualities* of parses
- **cnnnnn_parse** might contain one of many *kinds* or *qualities* of parses
 - **Gold or Automatic**
 - ▶ **o** = Gold parse (using OntoNotes guidelines);
 - ▶ **t** = Gold parse (using original Treebank guidelines)
 - ▶ **c** = Charniak parser (Automatic);
 - ▶ **b** = Berkeley parser (Automatic)
 - **Traces**
 - ▶ **o** = original traces;
 - ▶ **n** = no traces
 - **Hyphens**
 - ▶ **h** = split-at-hyphens;
 - ▶ **n** = not-split at hyphens
 - ...

Cannonical, Compositional Representation

The case of Chinese Characters

Graphical Features—

- Recent work by Wang *et al.*, (2019) has shown that using the radicals in Chinese characters contain semantic information similar to the notion of subwords and suffixes in English and can be used to improve unsupervised learned representations that can improve named entity tagging

Character	Primary Radical
病(illness)	疒(sickness)
癆(tuberculosis)	疒(sickness)
痛(pain)	疒(sickness)
肝(liver)	月(moon)/肉(meat)
胸(chest)	月(moon)/肉(meat)
腦(brain)	月(moon)/肉(meat)

Multiple representations—

- Pinyin representations of Chinese characters also help...

Character	Pinyin
病(illness)	bìng
癆(tuberculosis)	láo
痛(pain)	tòng
肝(liver)	gān
胸(chest)	xiōng
腦(brain)	nǎo

Interoperability Matters

Case of COVID-19

- Shah and Curtis (2020) identify the limitations of current EHR systems
- Difficulties in pooling multiple data sources owing to missing mapping between different medicine nomenclatures
- For a simple query—**Find me patients using Hydroxychloroquine**
 - EHR (A) used **National Drug Code**
 - EHR (A') used **Medi-Span**
- A **Common Data Model** (CDM) would have helped bridge the two variations of the same EHR system and allowed for better and quicker data analysis
- CDM is not automatic and not static, but a better tracking system can be used to manage mapping across multiple versions and nomenclatures.

Learning CDM (Mappings)

- Dong et al. (2020), shows the significance of mapping types of **COVID-19 tests** using **LOINC** codes
 - ~600 Manually mapped codes
 - 99.3% ITA (Cohen's kappa)
 - 98.9% automatic mapping accuracy
- Allowed finer grained analysis of COVID-19 testing data across 8 sites

..

LOINC Code	Total	Percentage	LOINC Long Common Name
Molecular			
94759-8	240	42.25	SARS-CoV-2 (COVID19) RNA [Presence] in Nasopharynx by NAA with probe detection
94500-6	202	35.56	SARS-CoV-2 (COVID19) RNA [Presence] in Respiratory specimen by NAA with probe detection
94309-2	75	13.20	SARS-CoV-2 (COVID19) RNA [Presence] in Unspecified specimen by NAA with probe detection
94502-2	13	2.29	SARS-related coronavirus RNA [Presence] in Respiratory specimen by NAA with probe detection
94660-8	11	1.94	SARS-CoV-2 (COVID19) RNA [Presence] in Serum or Plasma by NAA with probe detection
Antibody			
94563-4	10	1.76	SARS-CoV-2 (COVID19) IgG Ab [Presence] in Serum or Plasma by Immunoassay
94564-2	4	0.70	SARS-CoV-2 (COVID19) IgM Ab [Presence] in Serum or Plasma by Immunoassay