

Full of beans: a study on the alignment of two flowering plants classification systems

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<https://github.com/yiyunyc2/NKOS18>

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I L L I N O I S

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Reasons to align different classifications



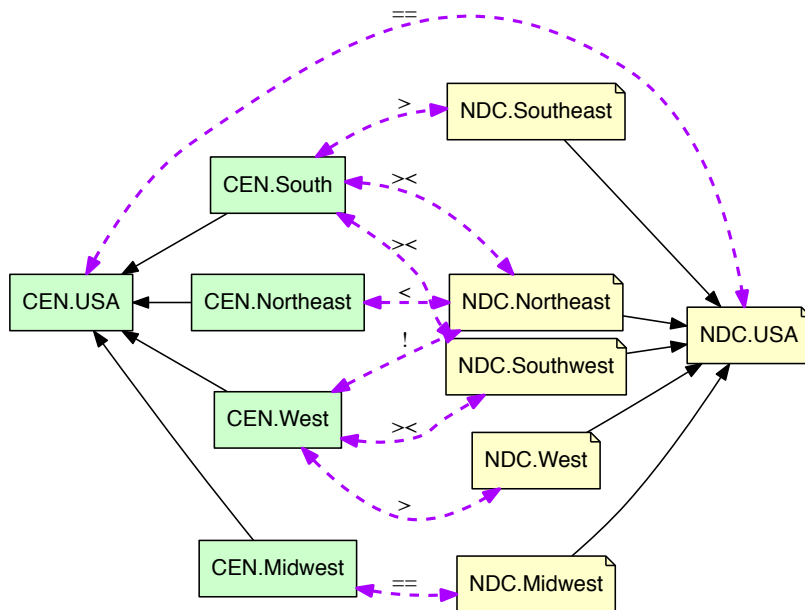
- Difficulty to organize information using a stable unitary classification scheme over time
- In biodiversity domain, It is common for taxonomists to contradict each other's or even their own previous taxonomies
- KOS are dynamic, time-specific, and responsive to both empirical signals and human classification interests

Purpose of this study

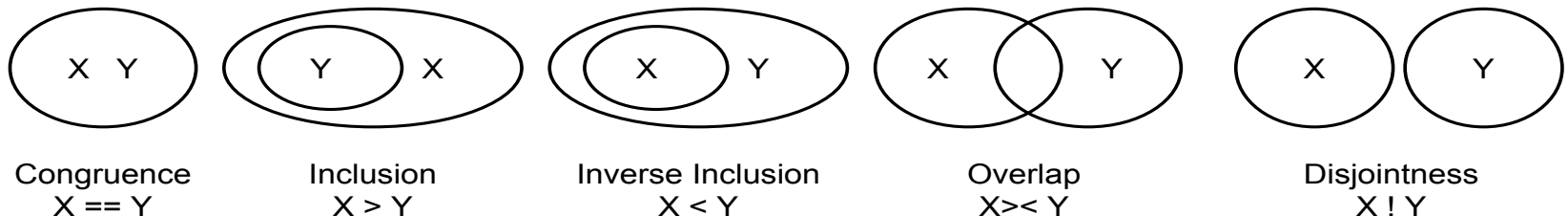


- Demonstrate the feasibility of integrating two classifications that result in numerous possible solutions
 - the computational power that can aid us in aligning KOS which could not have been possible when working with alignments manually
- With the hope that this work will further shed lights on:
 - the possible alignments of the classifications in the information science community
 - bring a novel approach for aligning KOS in the future

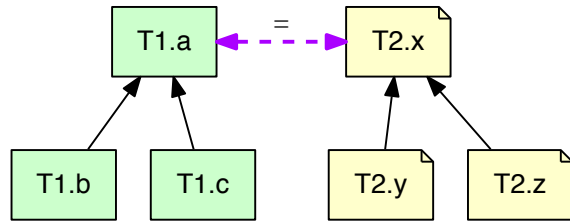
5 ways to relate concepts (regions)



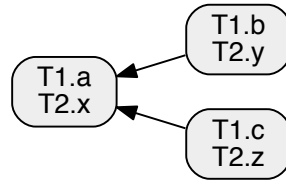
- **Idea:** relate concepts X and Y with *articulations*
- Articulation Language: **Region Connection Calculus (RCC5)**: congruence, inclusion, inverse inclusion, overlap, disjointness



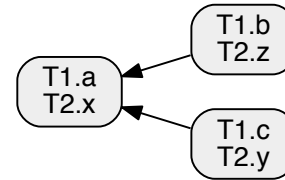
Reasoning about taxonomies



Input Taxonomies



PW1

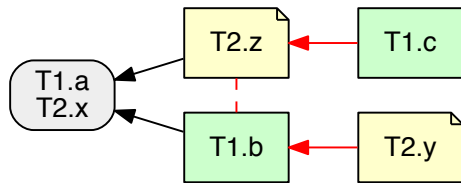


PW2

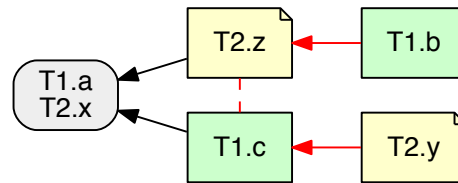
- **Given:**
 - taxonomies T1, T2
 - and relations $T1 \sim T2$ (*articulations*, *alignment*)

- **Find:**
 - merged taxonomy T3

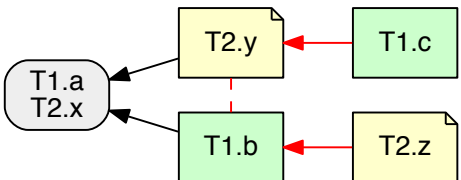
- **Such that:**
 - T1, T2 are **preserved**
 - all pairwise relations are **explicit**



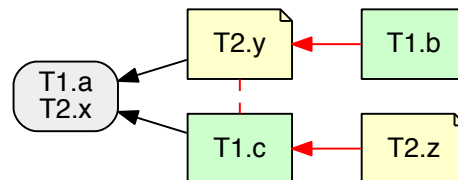
PW3



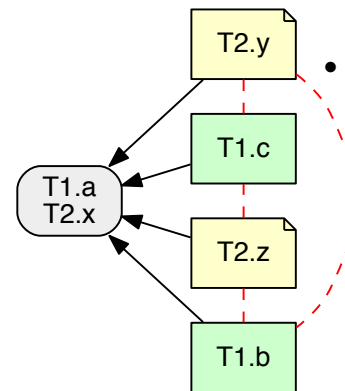
PW5



PW4



PW6



PW7

Two Flowering Plant Classifications



- Cronquist system
 - Arthur Cronquist (1981)
 - classifying plant resemblances based on evolution and *morphological* similarity (similar characteristics)
 - the most fully developed phyletic system of flowering plant classification systems by far
- Angiosperm Phylogeny Group System (APG IV)
 - uses both morphological and *molecular data* to group plants
 - The de facto standard classifications now

Biodiversity Taxonomies

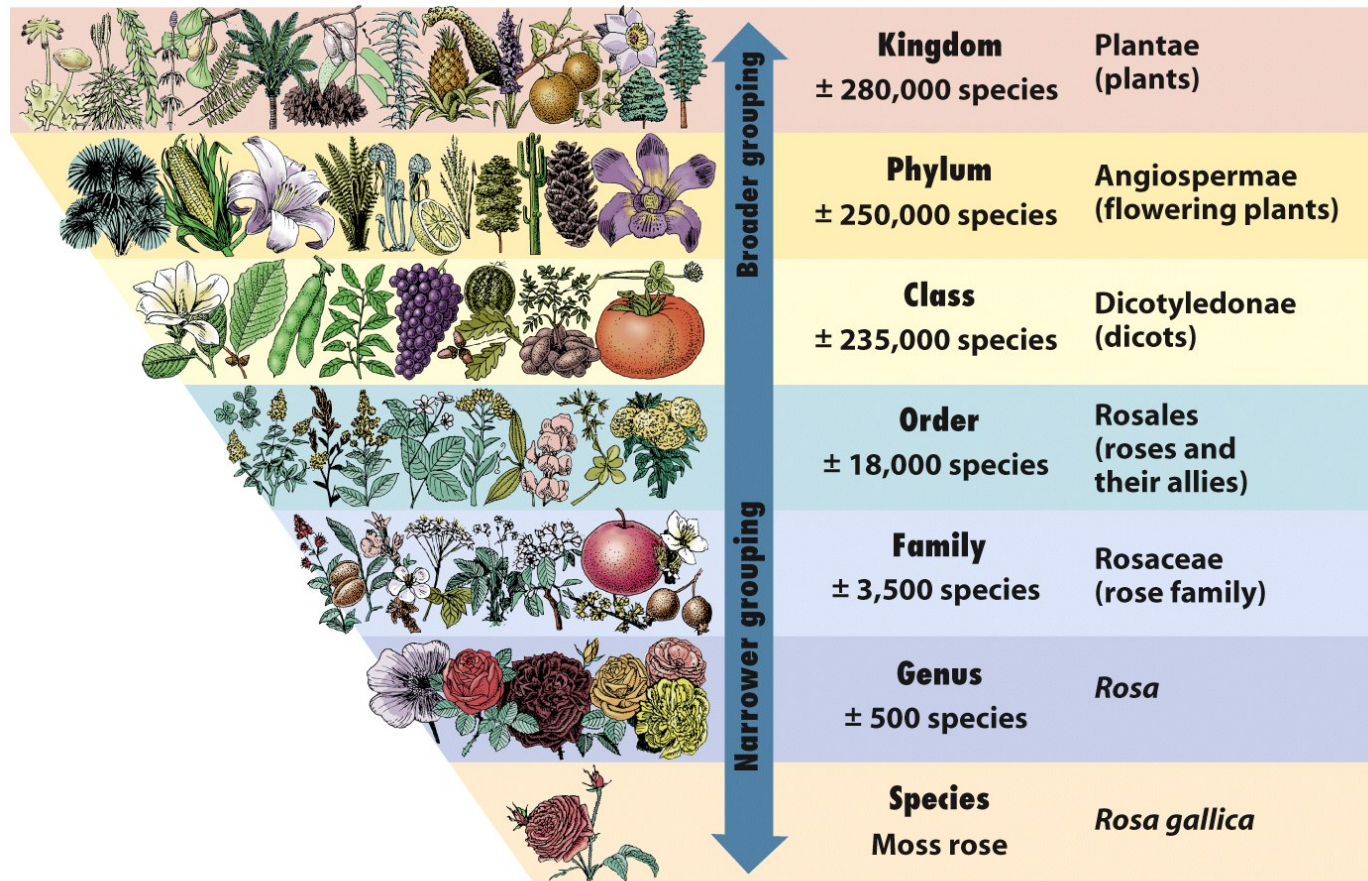


Figure 2-6 Discover Biology 3/e
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What are we aligning in our use case

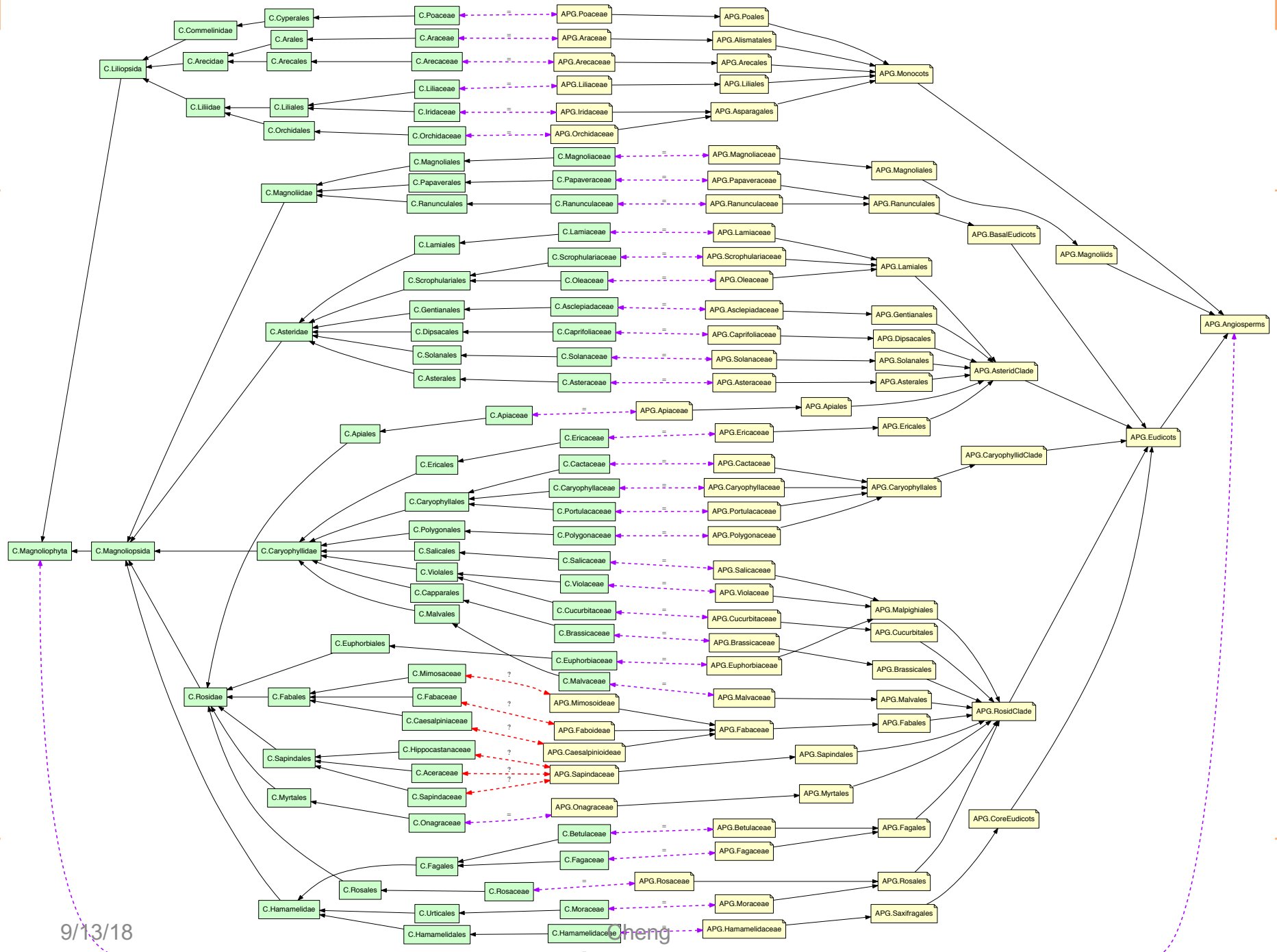


- *Family* level concepts
- Total number of flowering plant families: 416
- Number of families we align
 - 40 most common families/ subfamilies
- Examples: Magnoliaceae, Ranunculaceae, Papaveraceae, Cactaceae, Betulaceae, Fabaceae, Rosaceae

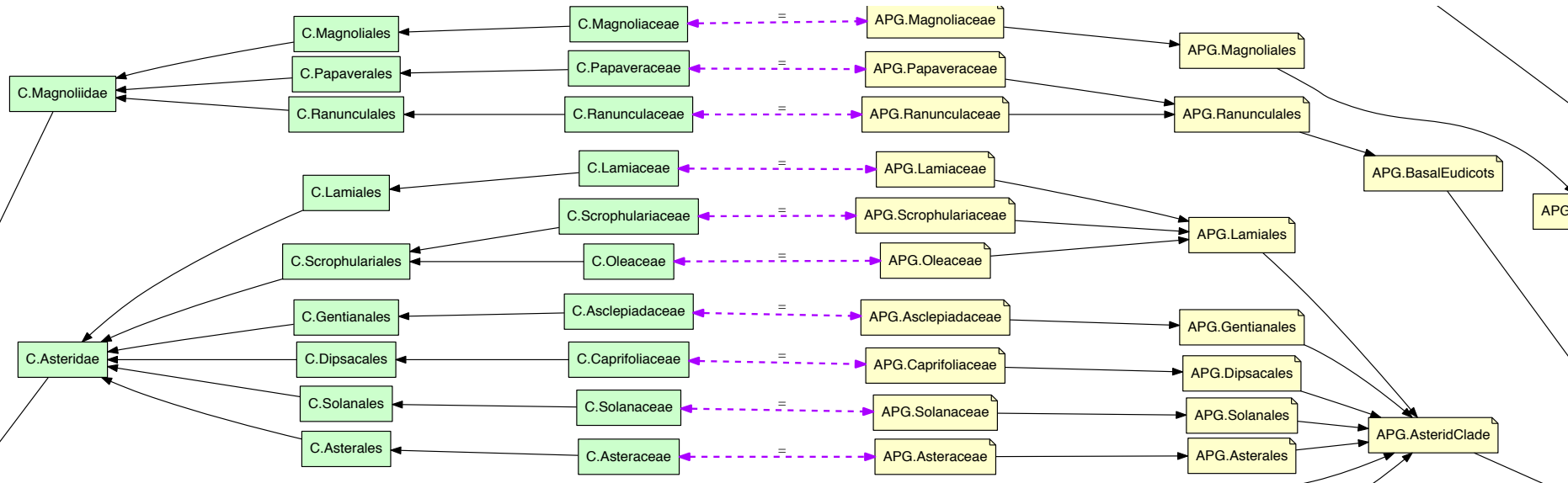
First round of alignment



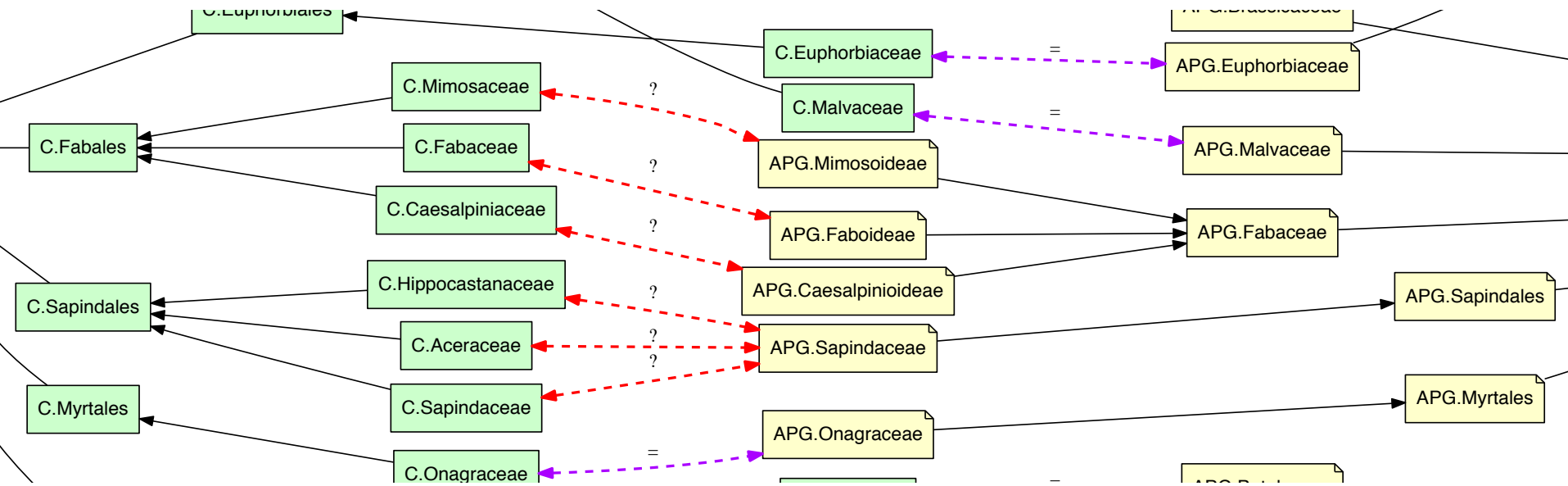
- Goal:
 - Align the 40 families (or the variations of the names) in both classifications
- Rules:
 - if the family in both systems shares the ***exact same name***, we assume (possible incorrectly) that they are ***congruent*** to each other
 - If there are ***similar but different*** names, we will leave the concepts unmapped at first



Zooming in to the input alignments



Six unknown relations



[C.Caesalpin*i*aceae ? APG.Caesalpin*i*oideae]

[C.Mimosa*ce*ae ? APG.Mimoso*i*deae]

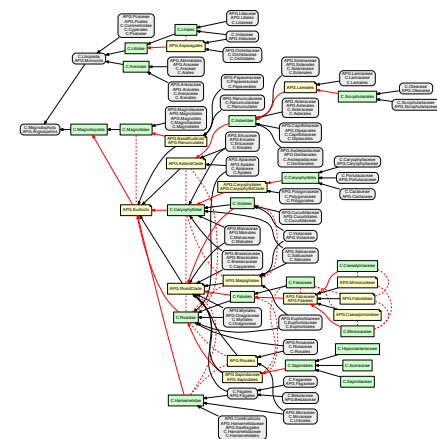
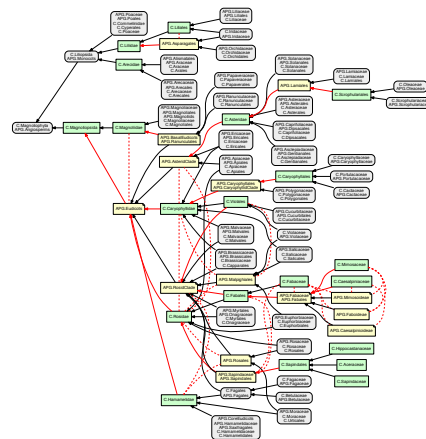
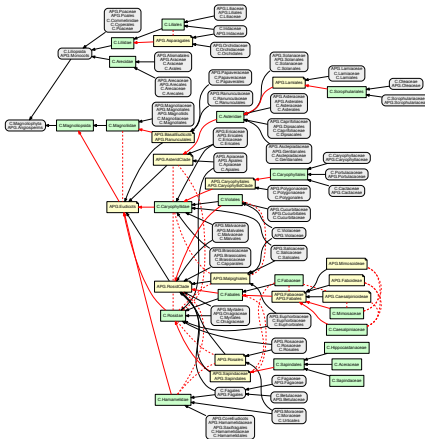
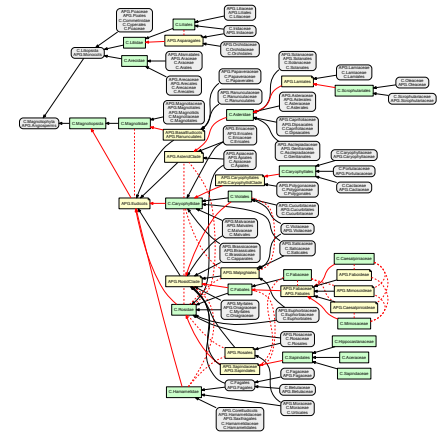
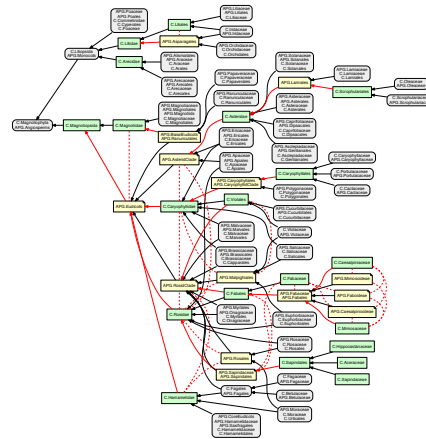
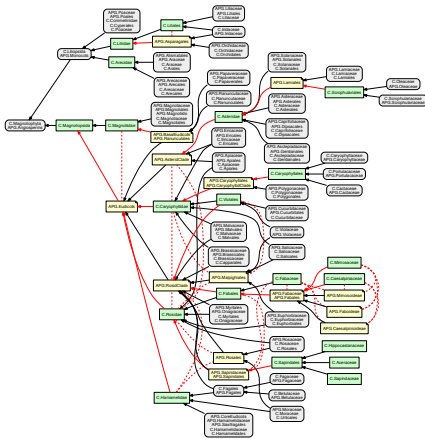
[C.Faba*ce*ae ? APG.Fabo*i*deae]

[C.Aceraceae ? APG.**Sapindaceae**]

[C.Sapindaceae ? APG.**Sapindaceae**]

[C.Hippocastanaceae ? APG.**Sapindaceae**]

Results: 555 Possible Worlds



9/13/18

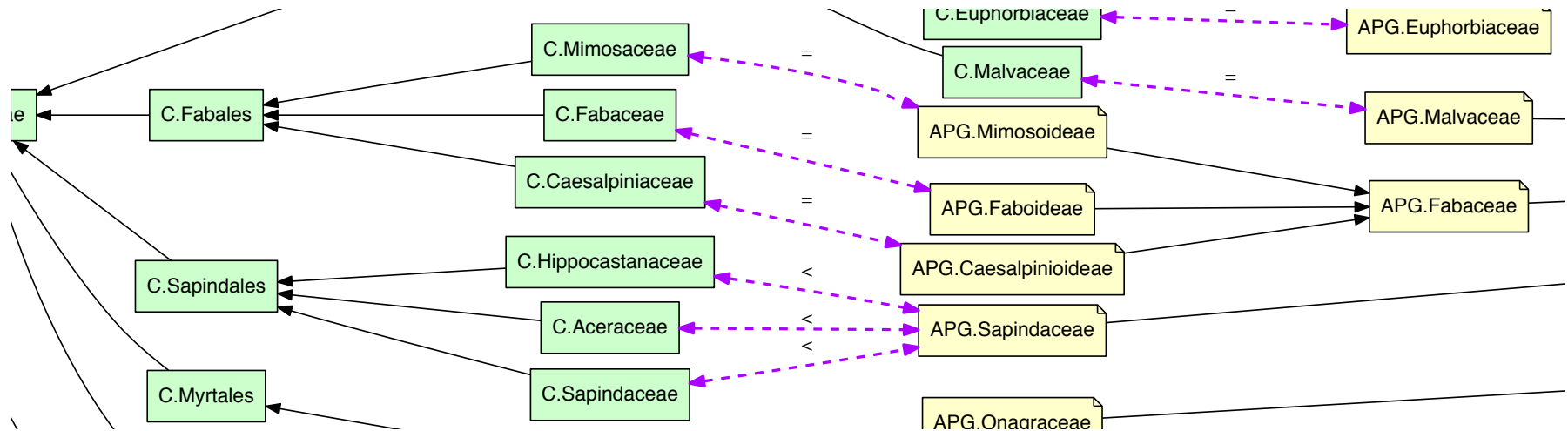


Second round of alignment



- Expert's input
 - To verify our 'congruent' alignments as well as to sort out the underspecified articulations
- Minimum viable product
 - The results from the first stage of alignment help us communicate with the expert and let him/her grasp all possible solutions for the alignment problem

Refining the six unknown relations



[C.Caesalpinaceae {=} APG.Caesalpinioideae]

[C.Mimosaceae {=} APG.Mimosoideae]

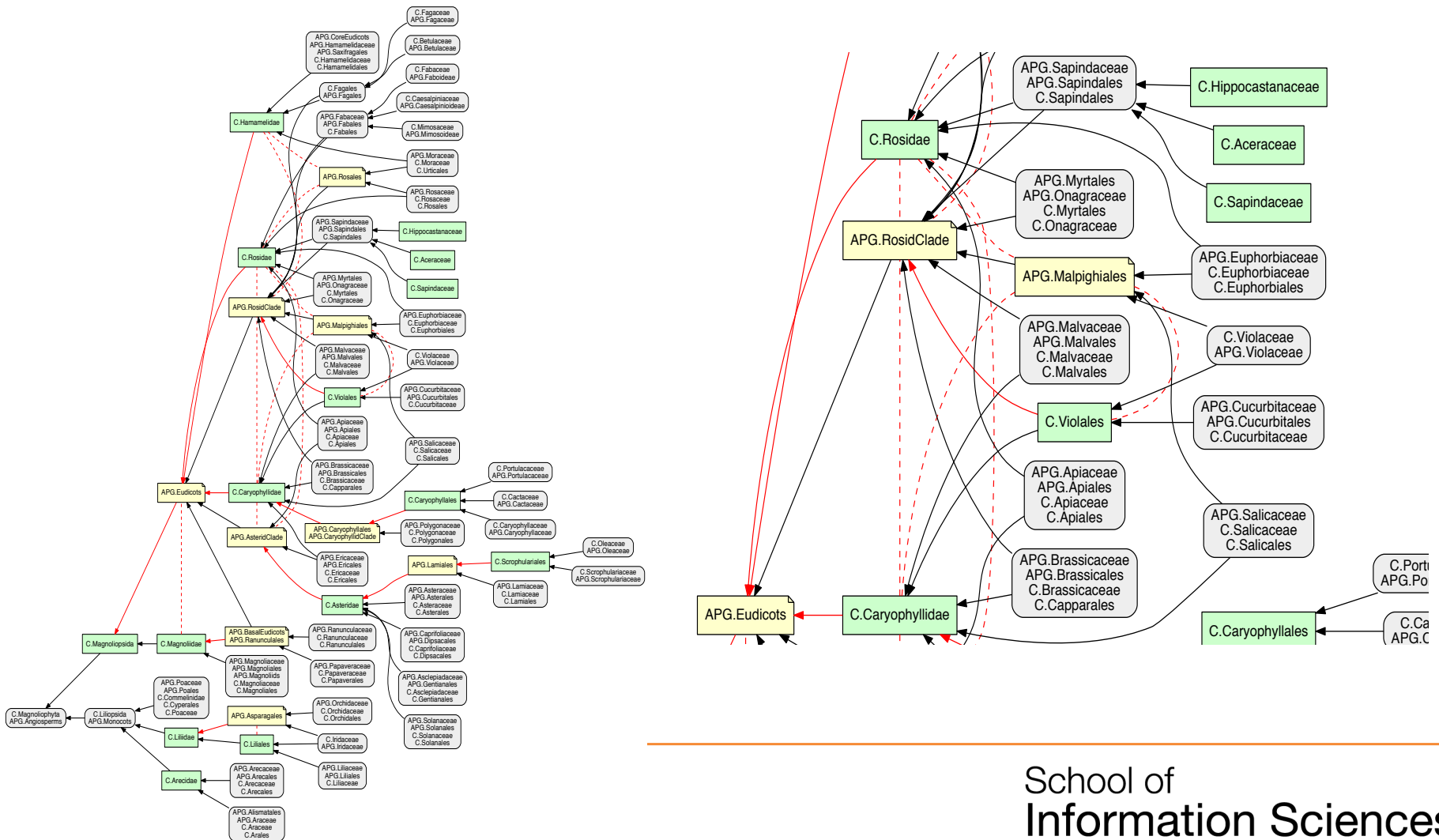
[C.Fabaceae {=} APG.Faboideae]

[C.Aceraceae {<} APG.Sapindaceae]

[C.Sapindaceae {<} APG.Sapindaceae]

[C.Hippocastanaceae {<} APG.Sapindaceae]

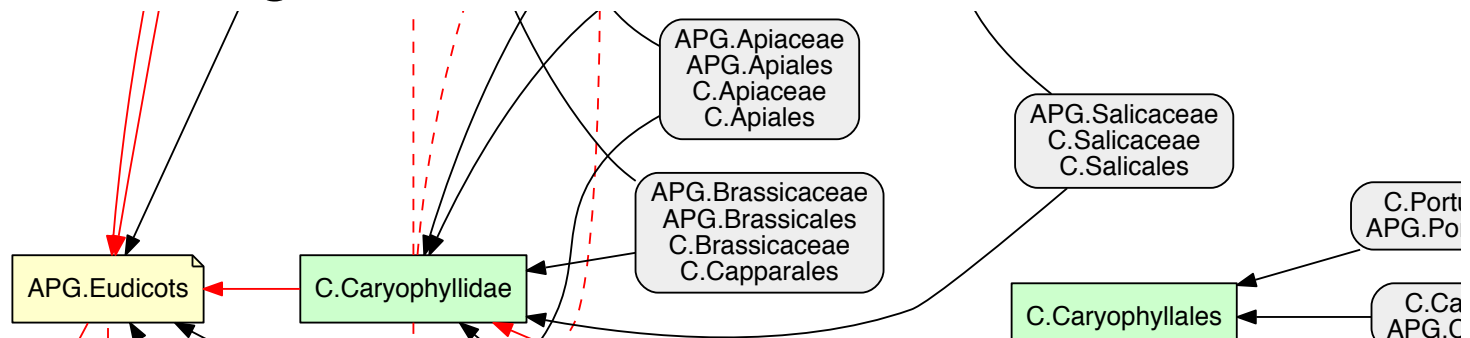
Final result: One Possible World



Limitations



- When a parent node only has one child, the RCC reasoner in Euler/X will collapse the concepts and merge them as the same node
- Our response: we have only chosen some 40 major flower families instead of all 416 families. We could add missing children or artificial children here



Conclusion and discussion



- Why using the logic-based RCC-5, Euler/X approach?
 - classifications can coexist & disambiguating the names among concepts in a merged possible world
 - solving complex alignments for cases where manual efforts would likely fail to yield all 555 different ways to merge and reconcile the two KOS
- Domain experts seems very important..?
 - KOS alignment problems are usually complex-- our logic-based alignment can serve as ***minimum viable product***

Conclusion and discussion



- Why bother aligning the ‘older’ classifications such as the Cronquist system (1981)?
 - the Cronquist system still maintains its esteemed role for its comprehensiveness and preciseness in morphologically classifying the flowering plants
- Some other implications
 - making our classification systems more “full of beans”
 - to enable semantic interoperability, and enrich diversity in classification systems

Thank you!

Github Repo:

<https://github.com/yiyunyc2/NKOS18>

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