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# JULIE Lab & Med Uni Graz @ TREC 2019 Precision Medicine Track

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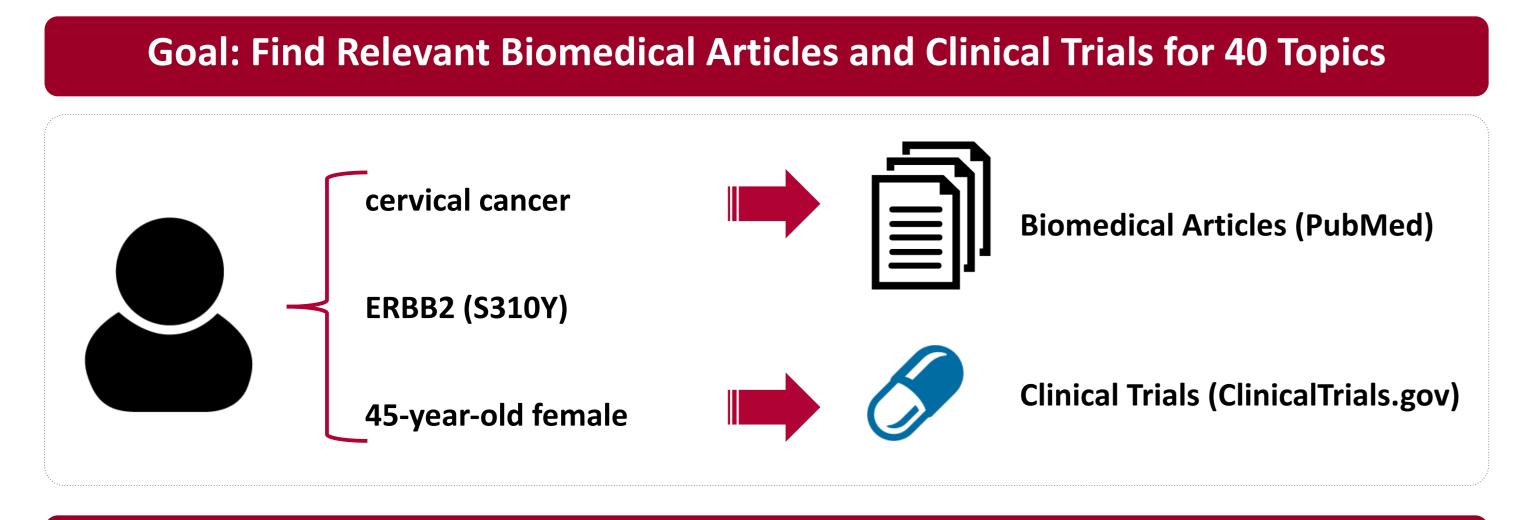
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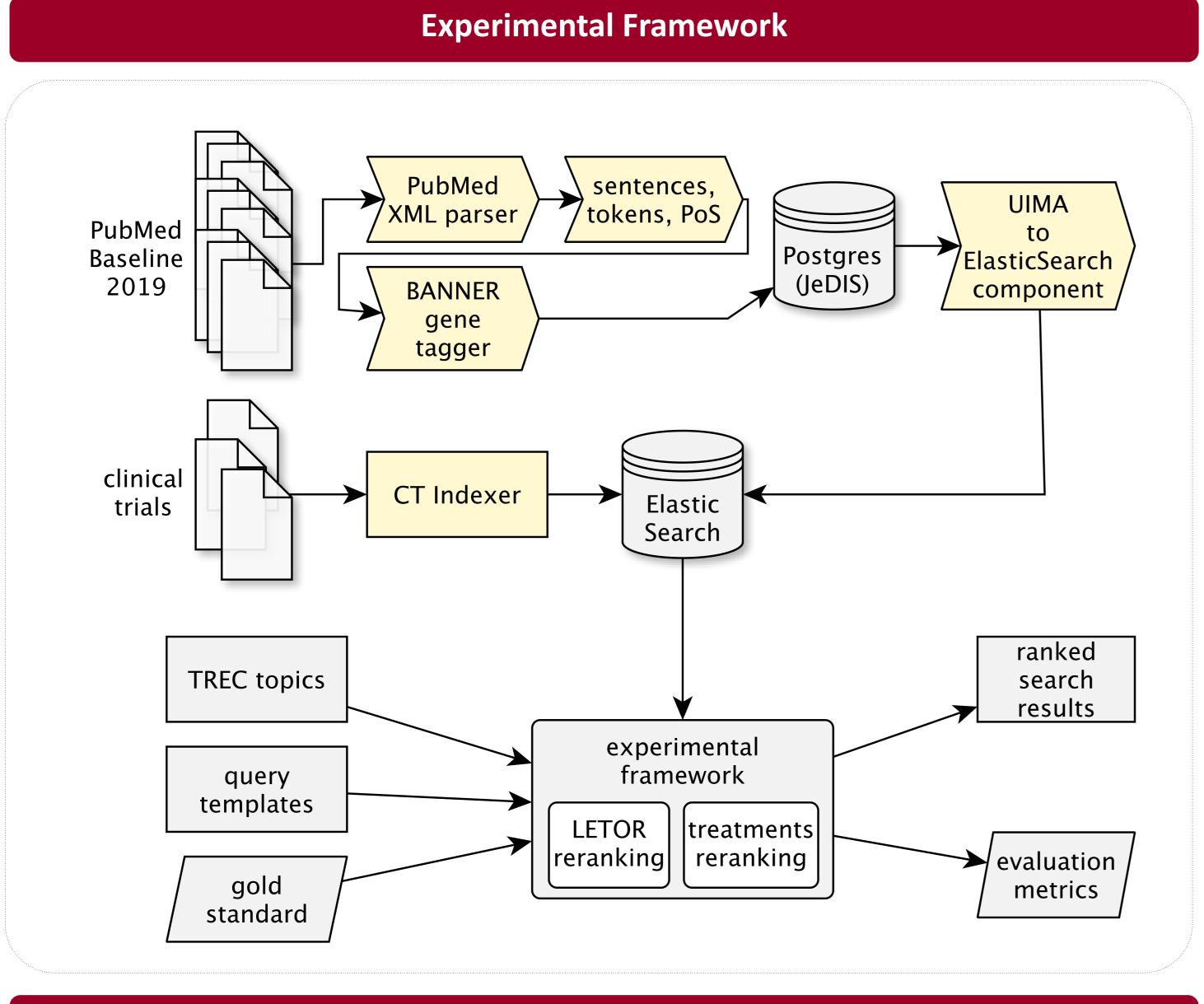
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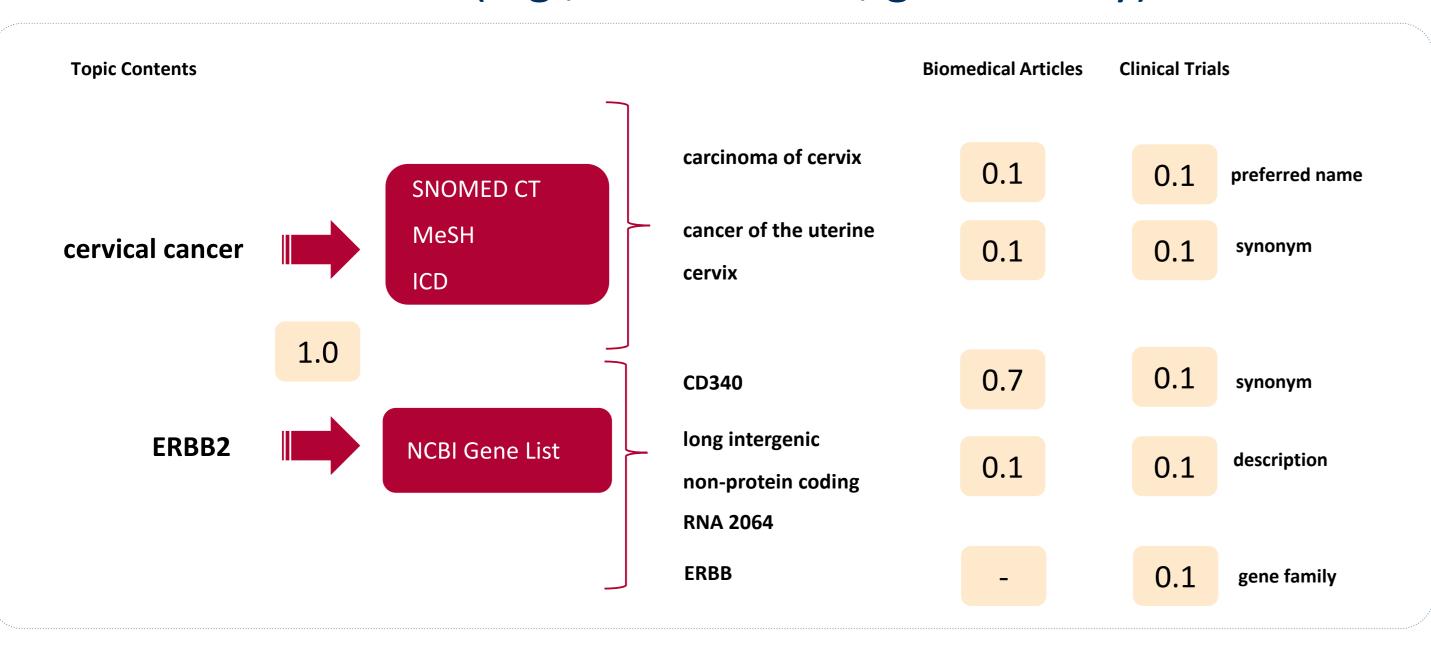






#### **Baseline Retrieval Strategies**

- Removal of domain stop words from query (e.g. cancer, cell)
- Hypernym and synonym weighting (using dis\_max queries)
- Positive & negative keyword relevance signals
- Hand-crafted rules (e.g., solid tumors, gene family)



#### **2019** Reference Standard: **2** Annotators

- 405 scientific abstracts (Cohen's kappa: 75.22%)
- 403 clinical trials (Cohen's kappa: 80.90%)

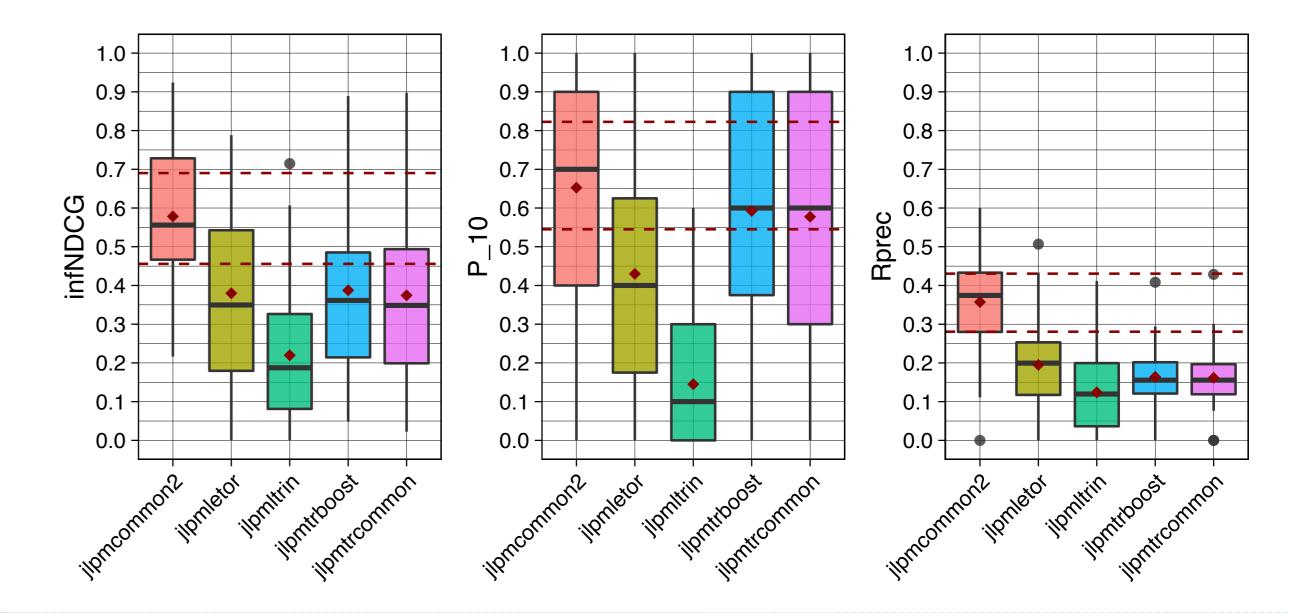
#### **Treatment Reranking**

- 1. Semantic type filtering
- 2. Word removal: handcrafted blacklist of 230 words
- 3. Removal of treatments mentioned in higher positions
- 4. Treatment ranking by frequency
- 5. Removal of documents without treatments

#### **Reranking: Learning to Rank (LETOR)** vords occurring ir top 500 TF-IDF bag-of-words FastText document Ranking embedding BM25 scores - whole topic RankLib disease LambdaMART - gene keywords n. disease match topic gene match and document gene variant match

**Results: Biomedical Articles** 

Strategy	jlpmcommon2	jlpmletor	jlpmltrin	jlpmtrboost	jlpmtrcommon
Baseline strategies	Υ	Υ	Υ	Υ	Υ
Filtered treatment match	Υ	N	N	Υ	N
LETOR training data	-	2017/18	2019	-	-
Treatments	-	-	-	11,128	9,443
Mean Top-10 Recall	-	-	-	0.2857	0.2698
Mean Top-10 F1	-	-	-	0.3118	0.3019
Mean Top-25 Recall	-	-	-	0.4603	0.4469
Mean Top-25 F1	_	-	-	0.3793	0.3716
infNDCG	0.5783	0.3798	0.2198	0.3876	0.3745
P@10	0.6525	0.4300	0.1450	0.5925	0.5775
R-Prec	0.3572	0.1948	0.1241	0.1639	0.1615



## Results: Clinical Trials

Strategy	jlctgenes	jlctletor	jlctltrin	jlctphrase	jlctprec
Baseline strategies	Υ	Υ	Υ	Υ	Υ
Match all	Υ	Υ	Υ	Υ	N
Match extracted genes	Υ	N	N	N	N
LETOR training data	-	2017/18	2019	-	-
infNDCG	0.6451	0.3731	0.4492	0.6244	0.6245
P@10	0.5474	0.2500	0.2921	0.5474	0.5447
R-Prec	0.4814	0.2066	0.2898	0.4799	0.4820
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