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# The BEL Information Extraction Workflow (BELIEF): Updates and Evaluation

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(part of Philip Morris International group of companies)***

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# Overview

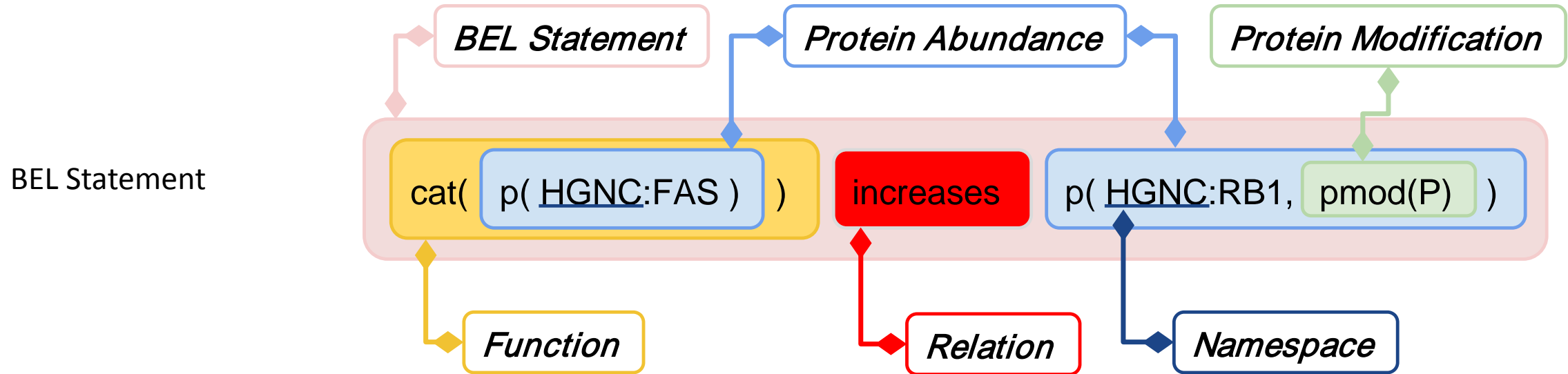
- Biological Expression Language (BEL)
- The Text Mining Pipeline (BELIEF Pipeline)
- Performance and Limitations of Text Mining
- The Curation Interface (BELIEF Dashboard)
- Performance and Limitations of the Curation Interface
- Summary



Openbel.org

# Biological Expression Language (BEL)

## BEL Nanopub



Citation

SET Citation = {“PubMed”, “Regulation of Rb and E2F by signal transduction cascades: divergent effects of JNK1 and p38 kinases.”, “EMBO J. 1999 Mar 15; 18(6):1559-70.”, “10075927”}

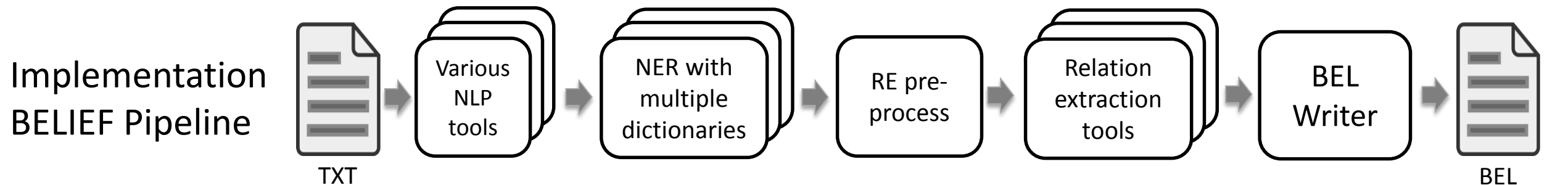
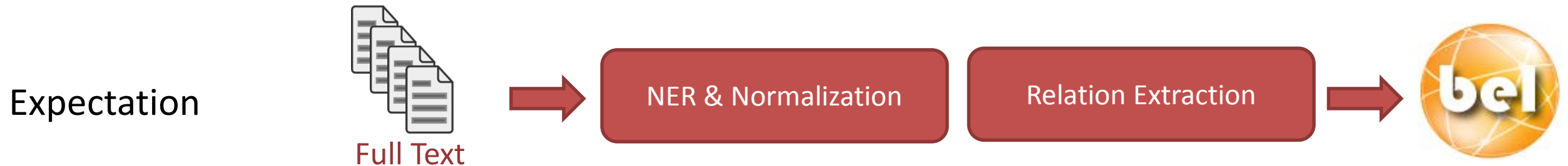
Support

SET Evidence =  
“Fas stimulation of Jurkat cells is known to induce p38 kinase and we find a pronounced increase in Rb phosphorylation within 30 min of Fas stimulation”

Experiment Context

SET Ti ssue = “jurkat cells”

# Expectation from Text Mining BELIEF Pipeline





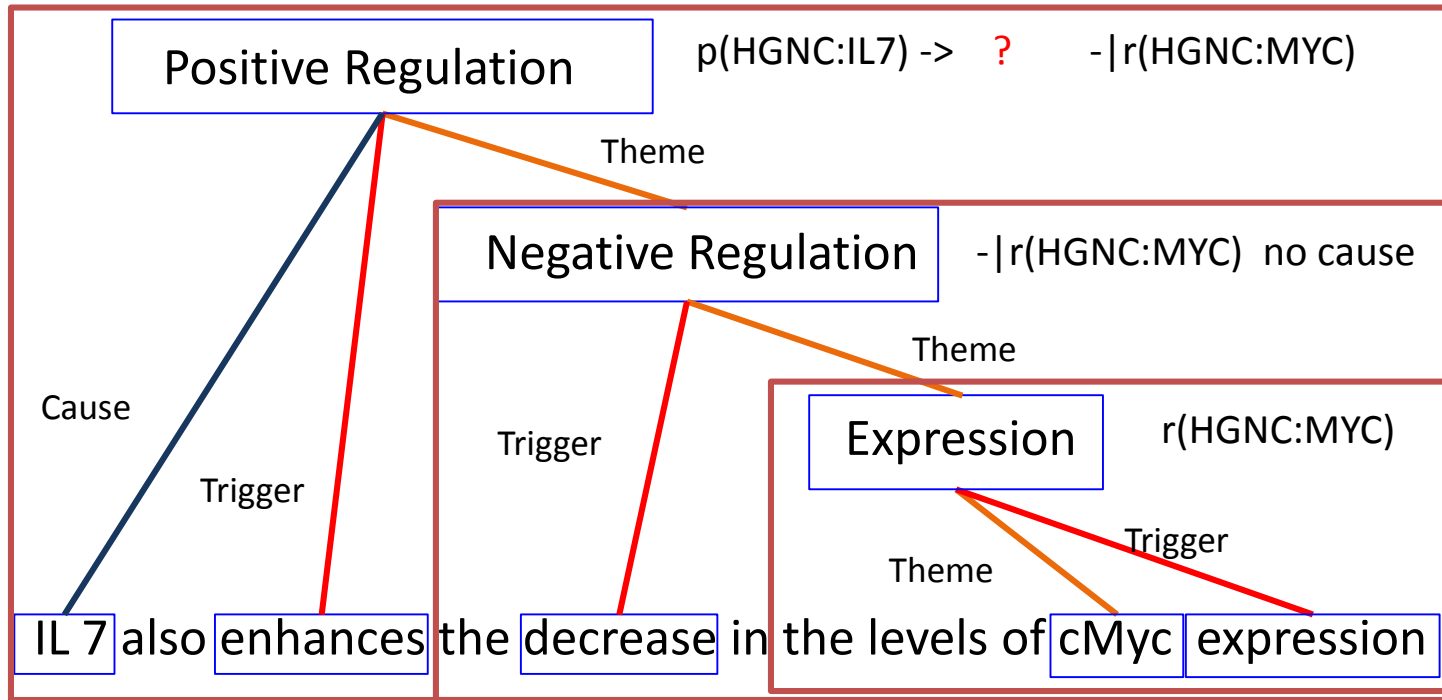
# BELIEF Pipeline - Named Entity Recognition

Entity class	Resources	BEL namespace
Human genes/proteins	EntrezGene/Uniprot	HGNC
Mouse genes/proteins	EntrezGene/Uniprot	MGI
Rat genes/proteins	EntrezGene/Uniprot	RGD
Protein family names	OpenBEL	SFAM
Protein complex names	OpenBEL	SCOMP
Protein complex names	Gene Ontology	GOCC
Biological processes	Gene Ontology	GOBP
Chemical names	OpenBEL	SCHEM
Chemical names	ChEBI	CHEBI
Chemical names	ChEMBL	CHEMBL
Disease names	MeSH	MESHD
Anatomical names	MeSH	MeSHAnatomy
Cell lines	Cell Line Ontology	CellLine
Cell structures	MeSH	CellStructure

Dictionary	Recall rate application adapted
Genes/Protein: (HGNC)	93 %
Chemical compounds: ChEBI	66 %
Chemical compounds: SCHEM	75 %
Chemical compounds: ChEBI + SCHEM+ ChEMBL	91 %
Selventa-human-complex	46 %
GO-Complex	64 %
Selventa-human-complex + Complex	82 %
Selventa-human-families	77 %

# BELIEF Pipeline - Relation Extraction

- The BioNLP shared tasks delivers a very detailed annotation for relationship extraction similar to the information needed for BEL (TEES2.1):



- Simpler binary classification (LibLINEAR):

IL 7 also enhances the decrease in the levels of cMyc ....

IL7 - cMyc Relation: Yes

$p(\text{HGNC:IL7}) - - p(\text{HGNC:MYC})$

Classifies if a relation between 2 entities is existing but gives no information about the direction or type

# BELIEF Pipeline - Limitations

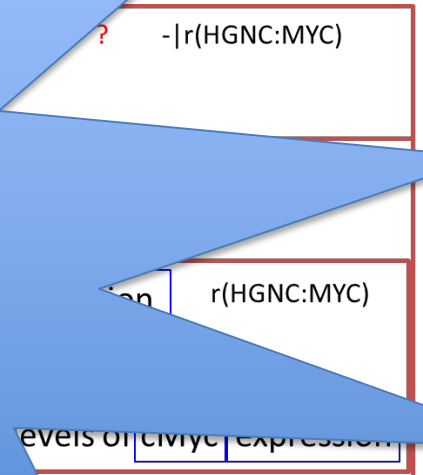
- NLP (Sentence Detection ~6% error)
- NLP (Tokenization ~6% error)
- NER (Different Classes)
- Relation Extraction (Multi)

We need manual curation!

Class	Precision	Recall
Term	81.34	72.6
Function	51.16	33.1
Relationship	67.37	41.68
Statement	59.15	20.79

Manual statement:

p(HGNC:IL7) -| r(HGNC:MYC)



statement

IL7) -> p(Plac) -| r(HGNC:MYC)

# BELIEF Dashboard - Curation Interface

Detected concepts:

- Mouse over highlights text
- Provides a fast overview of all entities in the evidence

[Return to document list](#)

SIRT1 knockdown caused a slight yet highly significant decrease in LXRalpha expression in both fed and fasted **livers**.

However, SIRT1 knockdown decreased expression of PGC-1beta only the fasted state.

Evidence: 1/2

Delete evidence

Detected concepts

**SIRT1**

MGI:Sirt1

HGNC:SIRT1

RGD:Sirt1

**LXRalpha**

MGI:Nr1h3

RGD:Nr1h3

**livers**

MeSHAnatomy:Liver

Browse namespaces

type e.g. HGNC:CDK1

Pubmed Information

Pubmed Id: 17646659

Update

PMID: 17646659

Title: Fasting-dependent glucose and lipid metabolic response through hepatic sirtuin 1.

Journal: Proceedings of the National Academy of Sciences of the United States of America; Vol. 104; Iss. 31

Authors: Joseph T Rodgers, Pere Puigserver

Published: 2007-07-31 00:00:00 CEST

Create table evidence

BEL statement

Context annotation

Export

Statements with "very low" confidence:

1 (id:44394): p(FIXME) -| (p(HGNC:SIRT1) -| r(MGI:Nr1h3))

(id:90567):

MeSHAnatomy

Liver

CellLine

Enter annotation value

Enter new statement

Add/edit/delete/export

BEL namespace search

Document PubMed metadata



# BELIEF Dashboard - Other features

BELIEF Dashboard Home BEL-Documents Upload BEL-Document Projects Documents

## Create Project

Name

Enter name

Description

Enter description

Create

BELIEF Dashboard Home BEL-Documents Upload BEL-Document Projects Documents

List

Create

## Document Lists

Search:

Id	Pubmed Id
1	2234234
2	12165281
Id	Pubmed Id

Showing 1 to 2 of 2 entries

Show 10 entries

Search:

Id	Statement	EvidenceId	Evidence	Export	Curate	Delete
14580	p(HGNC:PARK7, sub(L,166,P)) -> path(MESHD:"Parkinsonian Disorders")	1	In the present study, we investigated the effects of edaravone on the neurotoxicity in PD-induced isoforms of DJ-1 containing the mutation L166P.	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
14581	p(HGNC:PARK7) -- p(MESHD:"Parkinsonian Disorders")	1	In the present study, we investigated the effects of edaravone on the neurotoxicity in PD-induced isoforms of DJ-1 containing the mutation L166P.	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
14582	a(CHEBI:edaravone) -> p(HGNC:SLC18A2)	2	Interestingly, our result also demonstrated that edaravone was able to up-regulate VMAT2 expression in N2a cells in a dose-dependent manner.	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
14583	a(CHEBI:edaravone) -  bp(MESHPP:"Oxidative Stress")	2	Interestingly, our result also demonstrated that edaravone was able to up-regulate VMAT2 expression in N2a cells in a dose-dependent manner.	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
14584	a(CHEBI:edaravone) -  path(MESHD:"Parkinsonian Disorders")	3	Our findings enhance the understanding of the neuro-protective effects of edaravone in cell models and suggest that edaravone offers significant protection in a PD-related in vitro model.	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
14586	a(CHEBI:edaravone) -  bp(MESHPP:Apoptosis)	2	Interestingly, our result also demonstrated that edaravone was able to up-regulate VMAT2 expression in N2a cells in a dose-dependent manner.	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>

Id Statement EvidenceId Evidence Export Curate Delete

Showing 1 to 6 of 6 entries

Previous

1

Next

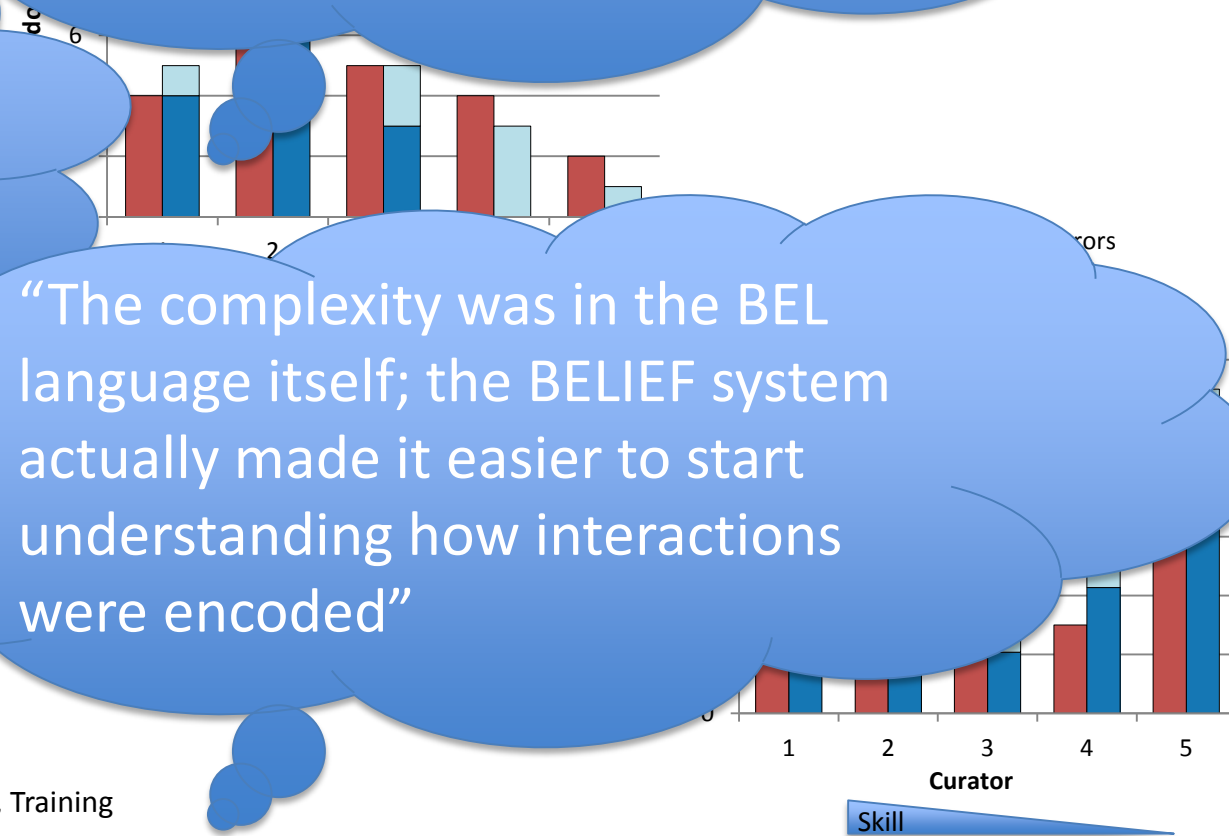
# BELIEF Dashboard - Performance Impression

“The system is very easy to learn for a user who is already familiar with BEL.”

“In particular, the preselected protein identifiers were immensely useful (which I only found out when I tried to find them by hand).”

“The complexity was in the BEL language itself; the BELIEF system actually made it easier to start understanding how interactions were encoded”

Protein		
RNA abundance		
Complex abundance	3	
Biological process	10	14



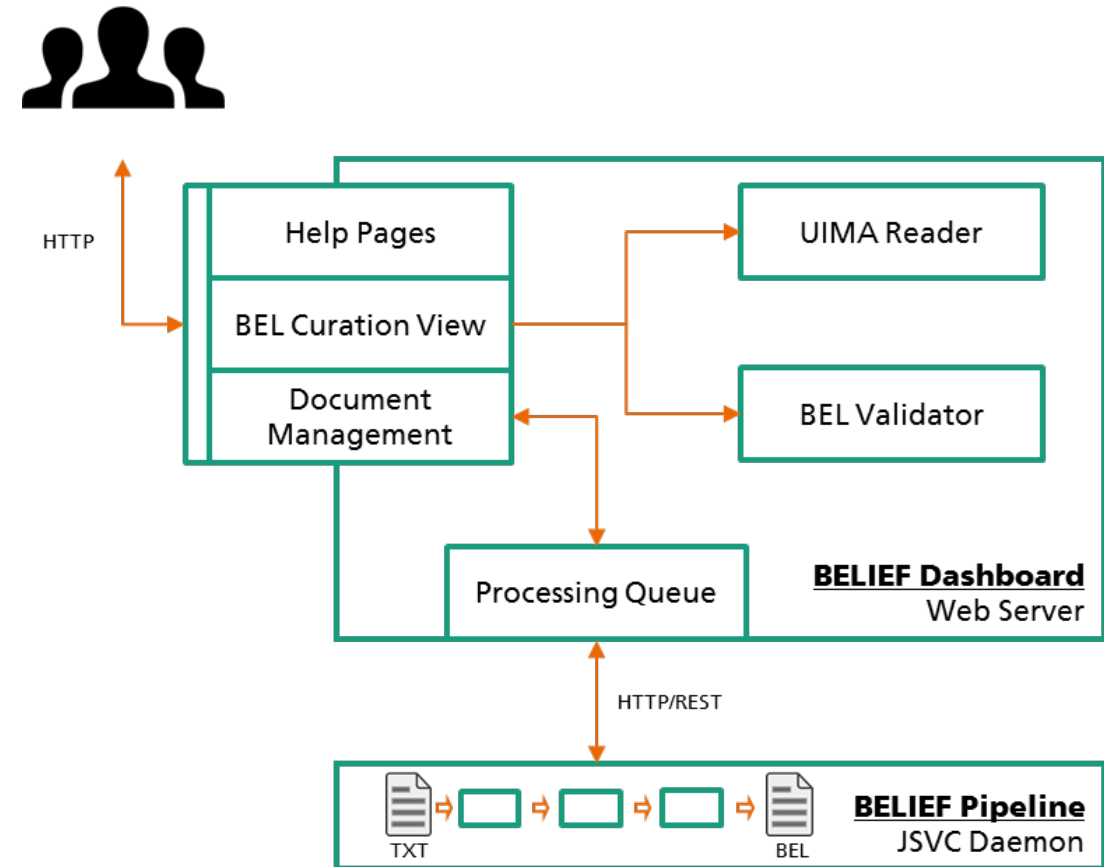
# Summary

BELIEF features a text mining pipeline and a complementing curation interface with the goal to support domain experts in different stages of knowledge acquisition and network model creation

The performance of the pipeline improved and has an acceptable level

Independent tester have given BELIEF a good systems usability score and appreciate the curation interface

The overall impression of all untrained testers was that BELIEF speeds up and further simplifies the creation of BEL statements



# Summary cont'd

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## *The new and impactful features are:*

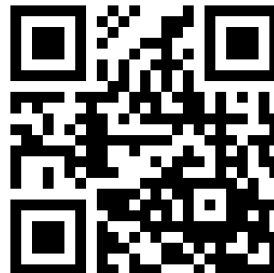
- Single point of entry including document and task management
- Reduced BEL coding effort due to full and partial BEL statement generation and validation on modifications
- Automatic citation from the Pubmed ID
- Two curation views to facilitate curation (evidence and statement centric view)
- Possibility to use custom dictionaries and re-running the text mining pipeline with these
- Show adjacent sentences to support curation

## *Outlook:*

Additional research is foreseen to improve a series of features from workflow management to user guides.

## *Accessibility:*

<http://www.scaiview.com/belief/>



# Acknowledgement



Fraunhofer SCAI

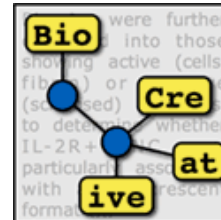
**Sumit Madan**  
**Philipp Senger**  
**Juliane Fluck**



**William Hayes**  
**Natalie Catlett**

[www.OpenBEL.org](http://www.OpenBEL.org)

**BioCreativeV**  
**IAT task**  
**BEL task**



**BELIEF Testers**

**Justyna Szostak**  
**Marja Talikka**  
**Julia Hoeng**  
**Filipe Bonjour**  
**Manuel Peitsch**

PMI R&D

