
POTENZIALE DER ALGORITHMISCHEN NUTZUNG MEDIZINISCHEN WISSENS FÜR DIE VERBESSERUNG DER VERSORGUNG



Prof. Dr. Martin Hofmann–Apitius

Leiter der Abteilung Bioinformatik

Fraunhofer Institut für Algorithmen und Wissenschaftliches Rechnen (SCAI)

Fraunhofer Sankt Augustin & Universität Bonn

- Largest research centre for informatics and applied mathematics in Germany
- Approx. 700 employees, about 500 scientists, about 200 students and trainees
- Bonn-Aachen International Center for Information Technology (B-IT)
- International curriculum on Life Science Informatics (LSI)



Wissenschaftliche Problemlösungs-Kompetenz



Development (DEV)

Semantics (SEM)

AI & Data Science (AI-DAS)



Dr. Marc Jacobs



Dr. Alpha Tom Kodamullil

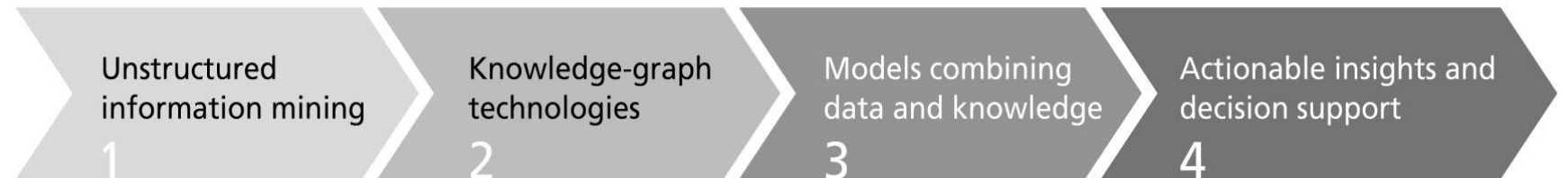


Prof. Dr. Holger Fröhlich

INFORMATIONSEXTRAKTION

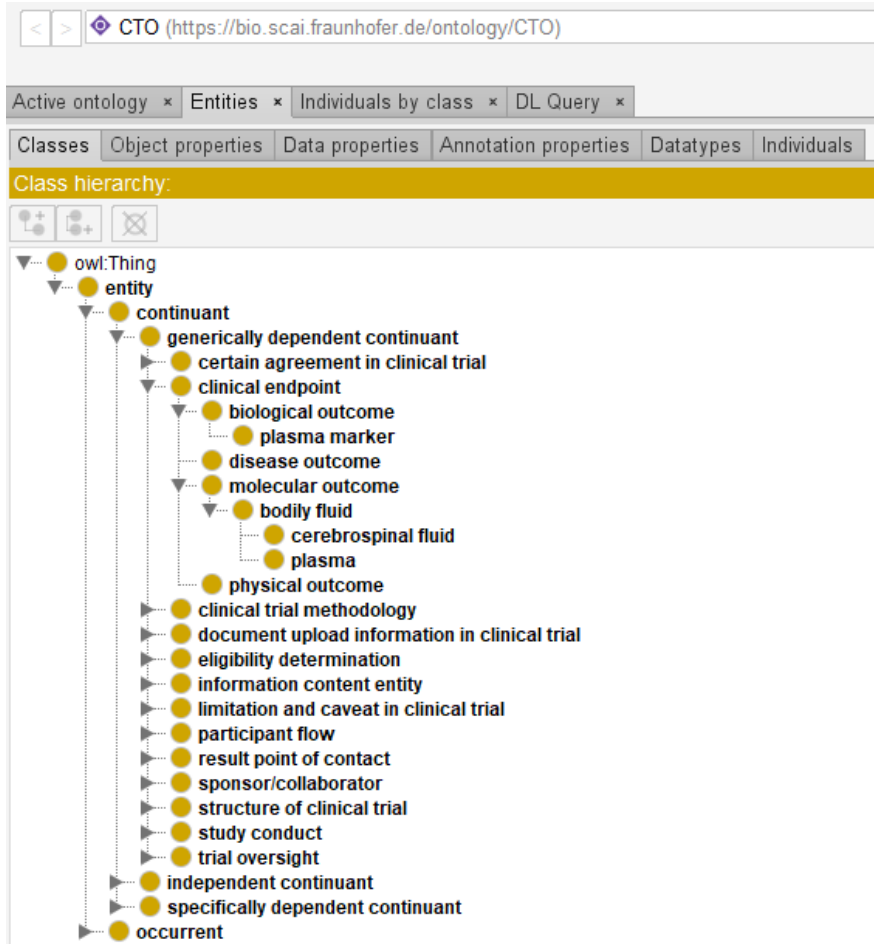
From **shared semantics** to **computable knowledge**

We are here



Formalisierte Semantik für Studiendaten: CTO

Clinical Trial Ontology (CTO)



CTO: a Community-Based Clinical Trial Ontology and its Applications in PubChemRDF and SCAIView

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Download: <http://purl.obolibrary.org/obo/cto.owl>

Ontologie – Kooperation von SCAI, FDA, NCBI and U. Michigan

COMPUTEBARES WISSEN: VON UNSTRUKTURIERTEN TEXTEN ZU ALGORITHMISCH NUTZBAREN WISSENSGRAPHEN

From **information extraction** to **re-usable knowledge graphs**

we are here



EXTRAKTION VON KAUSALEN UND KORRELATIVEN ZUSAMMENHÄNGEN AUS DER WISSENSCHAFTLICHEN LITERATUR

OpenBEL: Capturing of Knowledge and “Encoding” of Data

“Phosphorylation of **glycogen synthase kinase 3beta** at **Threonine, 668** **increases** the **degradation** of **Amyloid precursor protein**.”

p (HGNC:GSK3B, pmod (P,T,668)) -> deg (p (HGNC:APP))

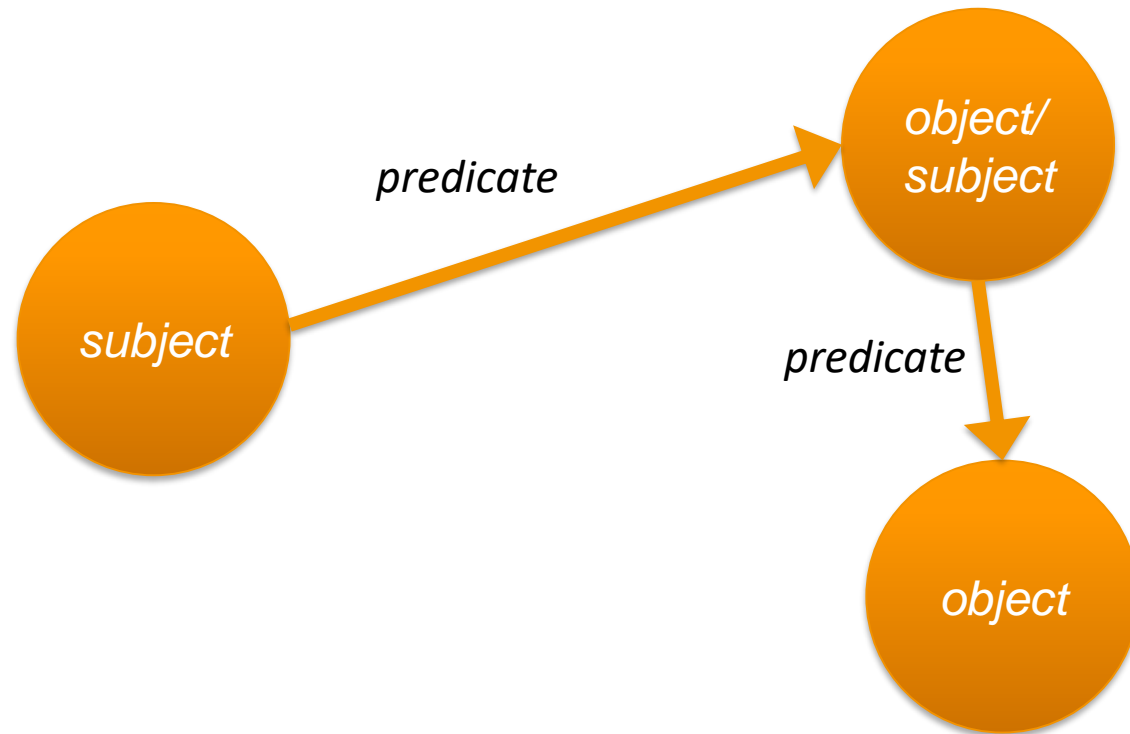
BEL Functions

Namespace Identifiers

Entity Definitions

TRIPLES AGGREGIEREN ZU GRAPHEN

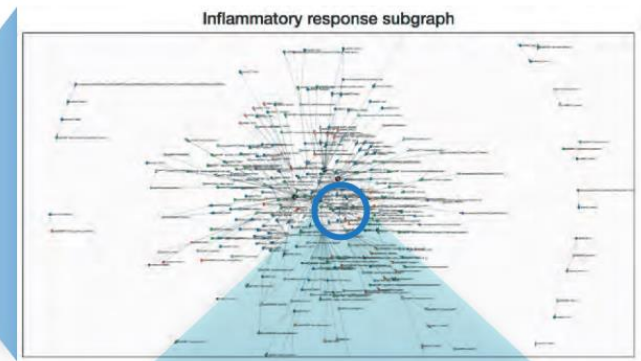
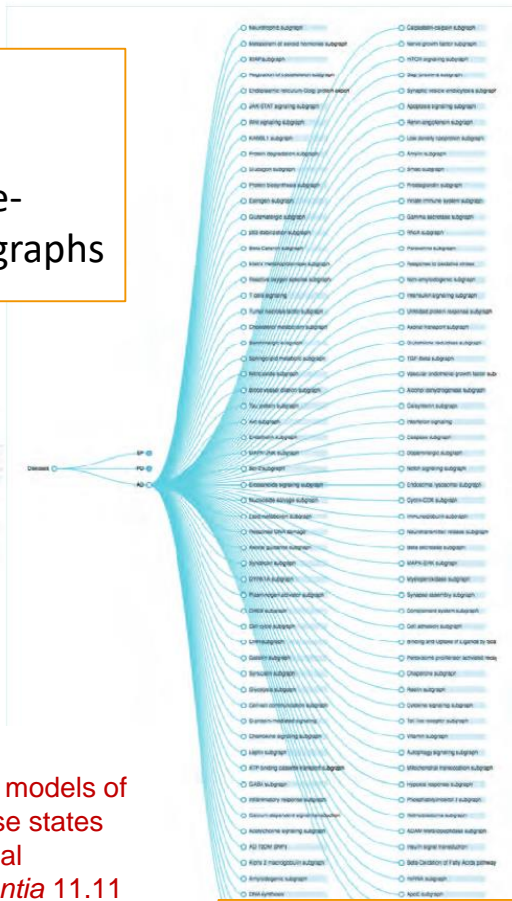
Das **OBJEKT** eines Triples kann das **SUBJEKT** des nächsten Triples sein



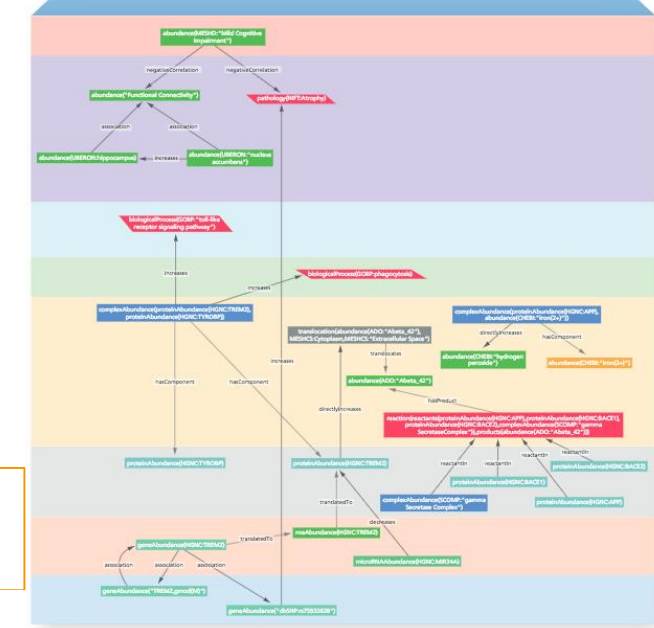
- Subject-predicate-object “triples”
- Object of one triple can be subject of another
- Putting them together makes arbitrarily large knowledge graphs
- Reasoning over causal relationships becomes a graph traversal

Multimodal Mechanism Models in Neurodegeneration Research

Text Mining extracts cause-and-effect relationships, disease-specific triples form graphs



Neuroinflammation mechanistic subgraph representing multiple biological levels



Clinical level
(e.g., cognition tests, imaging)

Biological processes

Molecular level (e.g., proteins, chemicals)

Genetic level (e.g., genes, epigenetics, variations)

Classifying each relation in the network into the mechanism(s) they participate

Kodamullil, Alpha Tom, et al. "Computable cause-and-effect models of healthy and Alzheimer's disease states and their mechanistic differential analysis." *Alzheimer's & Dementia* 11.11 (2015): 1329-1339.

Domingo-Fernández, Daniel, et al.

"Multimodal mechanistic signatures for neurodegenerative diseases (NeuroMMSig): a web server for mechanism enrichment." *Bioinformatics* 33.22 (2017): 3679-3681.

Cause-and-Effect Models & Applications

■ Identifying early mechanisms in AD aetiology

- Kodamullil AT, Younesi E, Naz M, Bagewadi S, Hofmann-Apitius M. *Alzheimers Dement.* 2015
- Kodamullil AT, Zekri F, Sood M, Hengerer B, Canard L, McHale D, Hofmann-Apitius M. *Nat Rev Drug Discov.* 2017

■ Functional impact of genetic and epigenetic variants

- Karki R, Kodamullil AT, ... Hofmann-Apitius M. *BMC Bioinformatics.* 2019
- Naz M, Kodamullil AT, Hofmann-Apitius M. *Brief Bioinformatics.* 2016

■ Analysis of comorbidity

- Khanam Irin A, Kodamullil AT, Gündel M, Hofmann-Apitius M. *J Immunol Res.* 2015

■ Repurposing of Drugs

- Karki R, Kodamullil AT, Hofmann-Apitius M. *J Alzheimers Dis.* 2017
- Karki R,... Kodamullil AT, Hofmann-Apitius M. *J Alzheimers Dis.* 2020

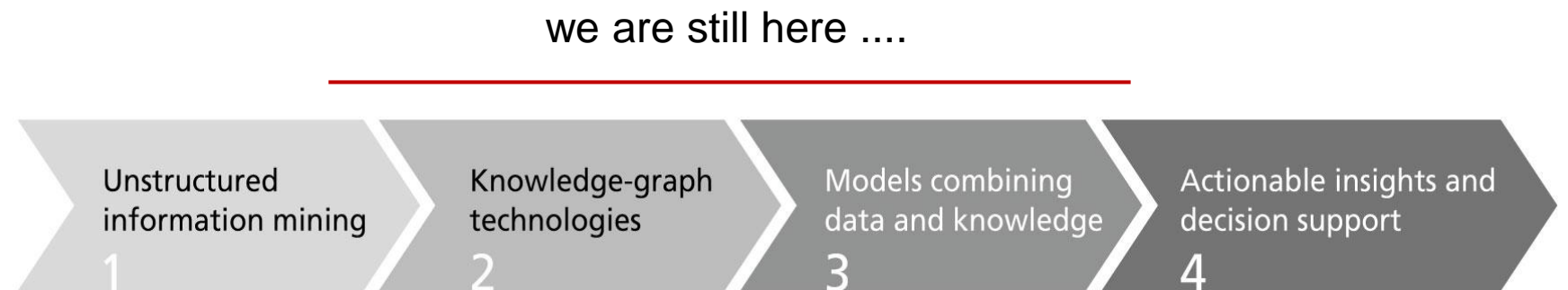
■ Comparitive analysis of pathophysiology mechanisms mouse/human

- Emon MA, Kodamullil AT, Karki R, Younesi E, Hofmann-Apitius M. *J Alzheimers Dis.* 2017
- Kodamullil AT, Ivappan A, Karki R, Madan S, Younesi E, Hofmann-Apitius M. *J Alzheimers Dis.* 2017

- Alzheimer Disease
- Parkinson Disease
- Epilepsies
- Amyotrophic lateral sclerosis
- Type 2 diabetes
- Schizophrenia
- Bipolar Disorder
- Traumatic Brain Injury
- PTSD
- [COVID-19]

ALGORITHMEN FÜR KNOWLEDGE GRAPHS

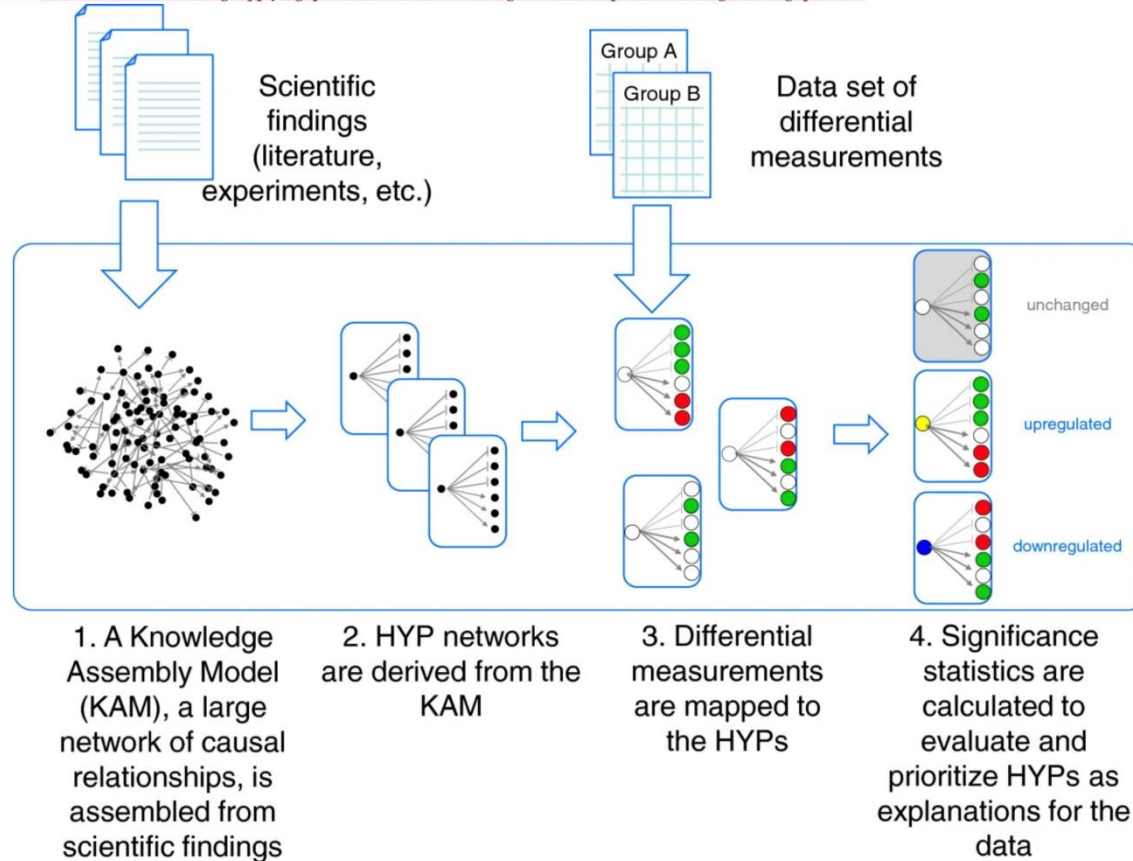
From **reverse causal reasoning** to **guilty targets**



REVERSE CAUSAL REASONING:

Figure 1

From: [Reverse causal reasoning: applying qualitative causal knowledge to the interpretation of high-throughput data](#)

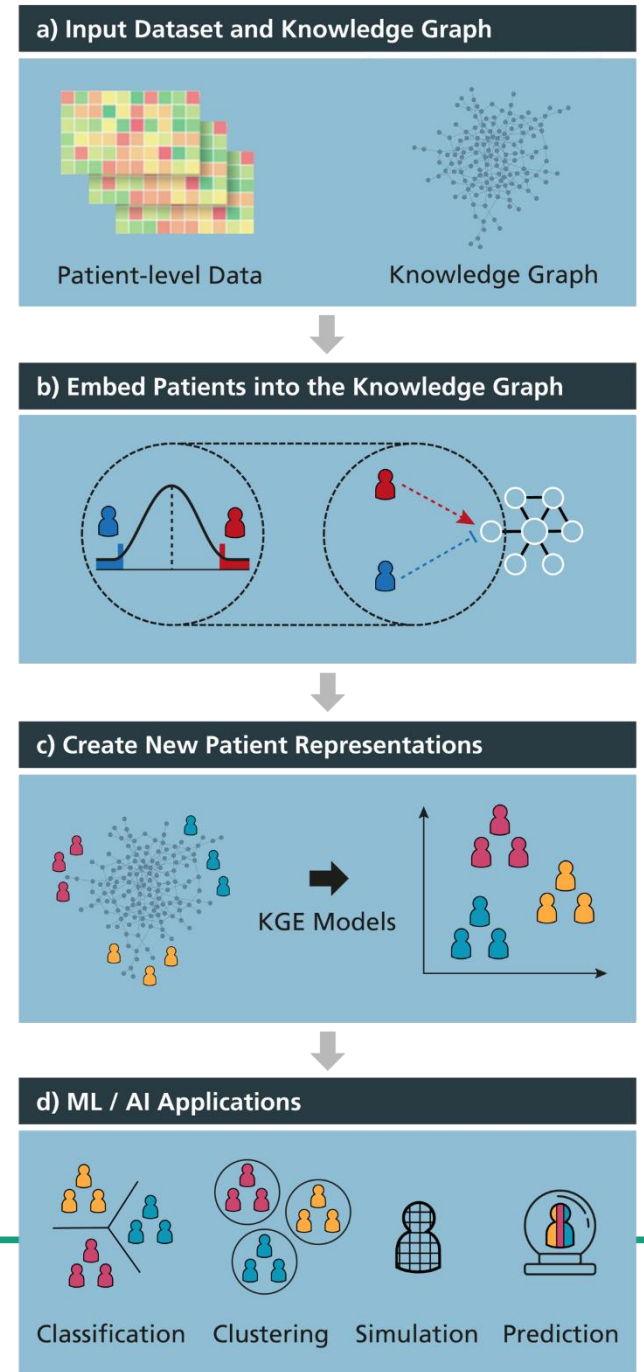


- Wissensgraph „kodiert“ kausale und korrelative Beziehungen
- Subgraphen repräsentieren Hypothesen zu Mechanismen
- Subgraphen werden auf die Übereinstimmung mit Daten getestet
- Die Signifikanz der Übereinstimmung zwischen Subgraphen und Daten wird berechnet
- Die am besten zu den Daten passenden Subgraphen werden selektiert

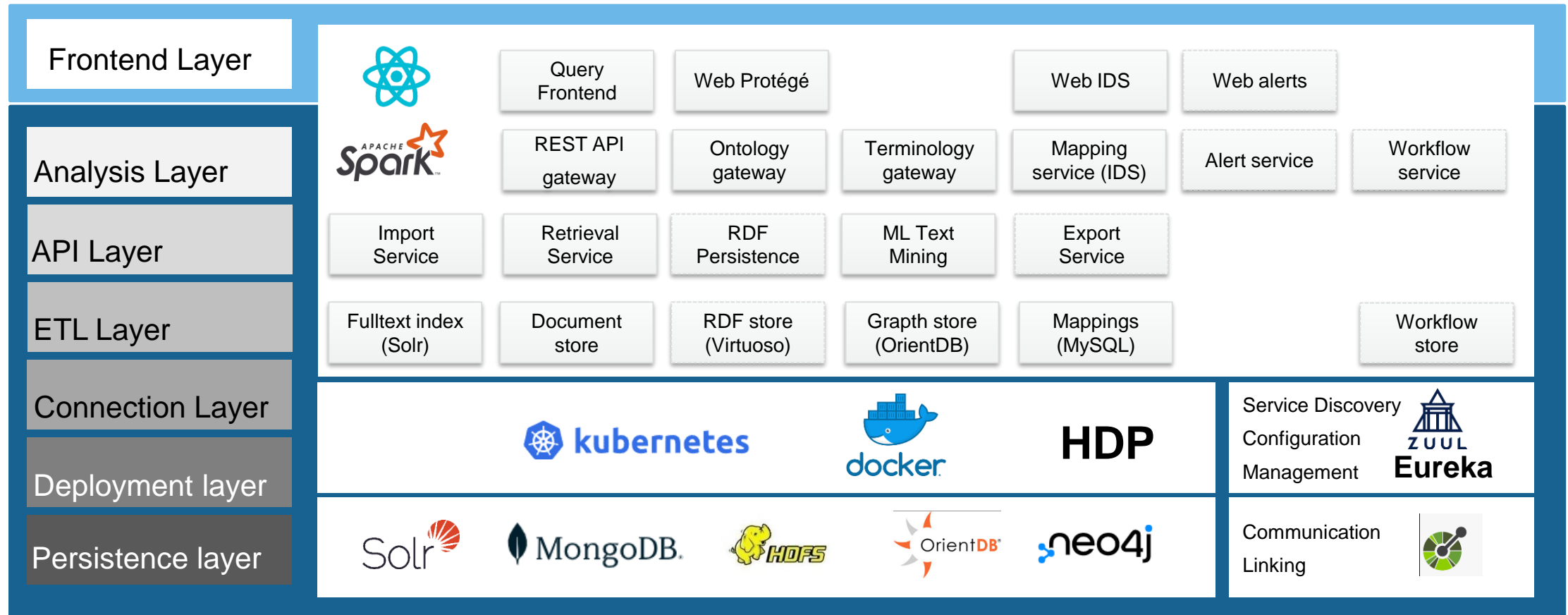
Patient Embeddings in Knowledge Graphs

Schematic illustration of the CLEP framework.

- (a) CLEP requires two inputs: (i) a **patient-level dataset** such as multi-omics, and (ii) a **KG comprising relations between features** measured in the previously mentioned dataset.
- (b) Using one of the proposed methods, CLEP incorporates patients into the KG by connecting them to their most distinctive features in the dataset.
- (c) KGEMs are then used to generate new patient representations based on both data- and knowledge-driven features.
- (d) These patient representations can subsequently be used for several downstream tasks, such as patient classification and stratification.

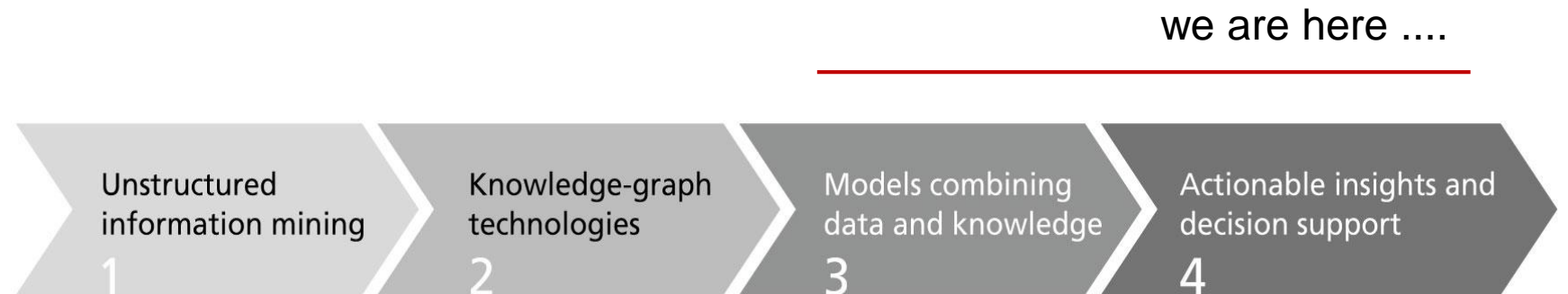


Aus dem Maschinenraum: Knowledge Graph Service Infrastruktur



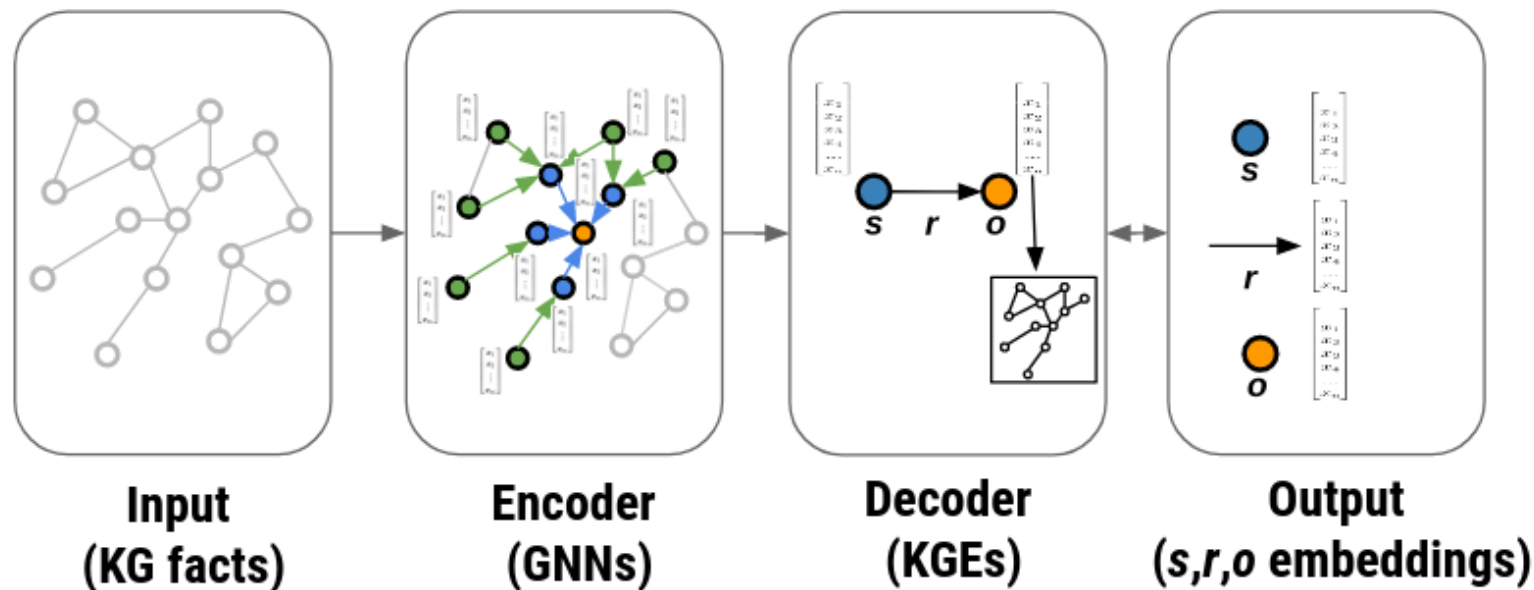
NUTZUNG VON WISSENSGRAPHEN: KG-EMBEDDINGS

From **Understanding Pathophysiology Mechanisms** to the **Identification of Patient Subgroups**



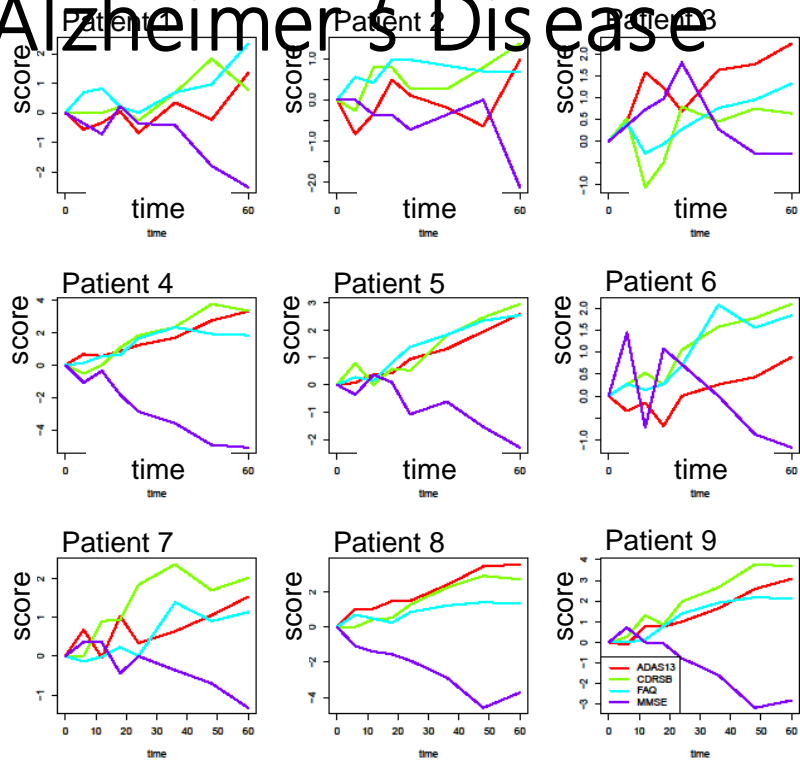
KNOWLEDGE GRAPH EMBEDDINGS

Reduktion der Dimensionalität von komplexen Graphen

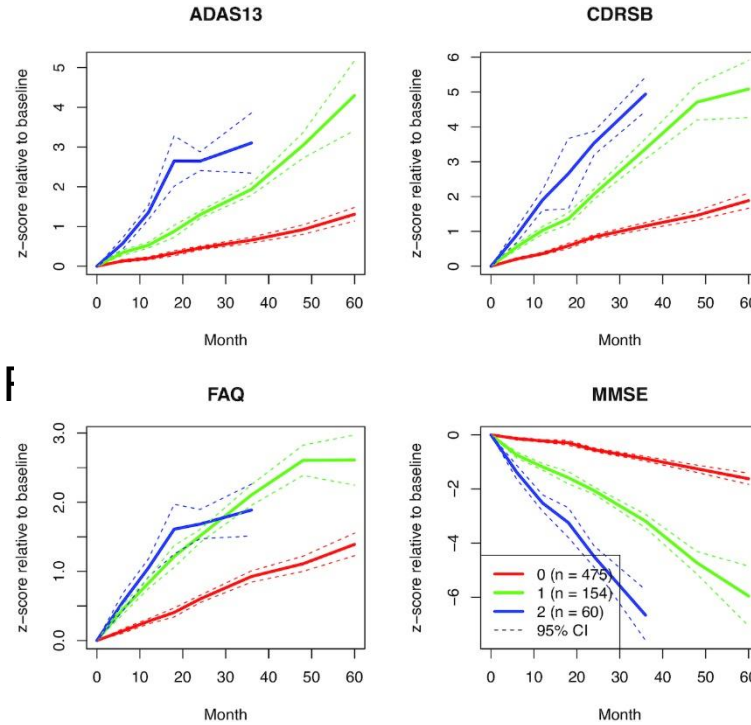


- Wissensgraph wird in einen Vektor umgewandelt
- Die gängigen Algorithmen des „Data Mining“ können sehr gut mit Vektorräumen umgehen.
- Hierdurch wird er „leichter und besser nutzbar“.

Stratification of Disease Trajectories in Parkinson's and Alzheimer's Disease



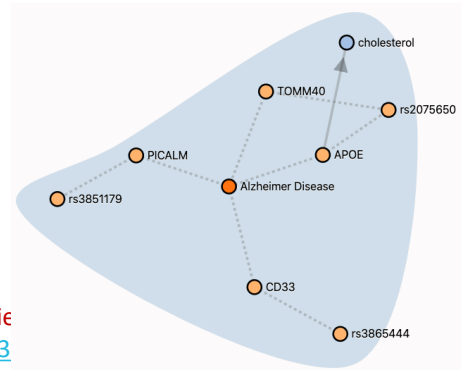
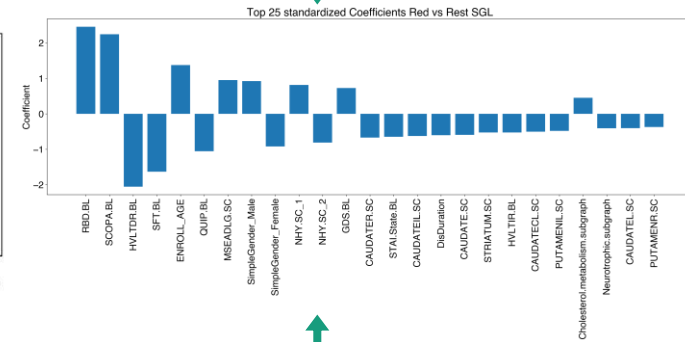
VADEI



Baseline

- Biomarkers
- Clinical
- Genetics
- Imaging

AI



Predictive molecular mechanisms

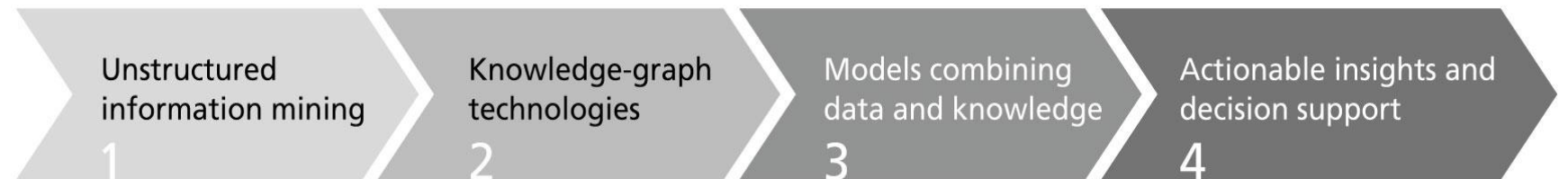
Impact: Clinical trial

Johann de Jong, Mohammad Asif Emon, Ping Wu, Reagon Karki, Meemansa Sood, Patrice Godard, Ashar Ahmad, Henri Vrooman, Martin Hofmann-Apitius, Holger Fröhlich, Deep learning for clustering of multivariate clinical patient trajectories with missing values, *GigaScience*, Volume 8, Issue 11, November 2019, giz134, <https://doi.org/10.1093/gigascience/giz134>

PERSPEKTIVEN FÜR DIE ZUKUNFT: GUIDELINES, EVIDENZEN AUS DER LITERATUR, REAL-WORLD DATA

From **Clinical Guidelines** to **Graph-based Analytics & Decision-Support**

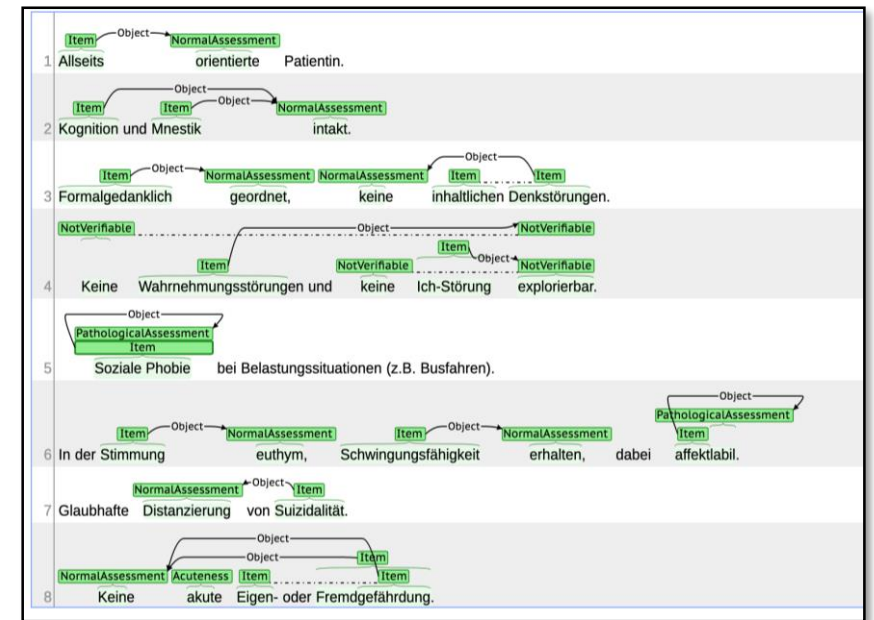
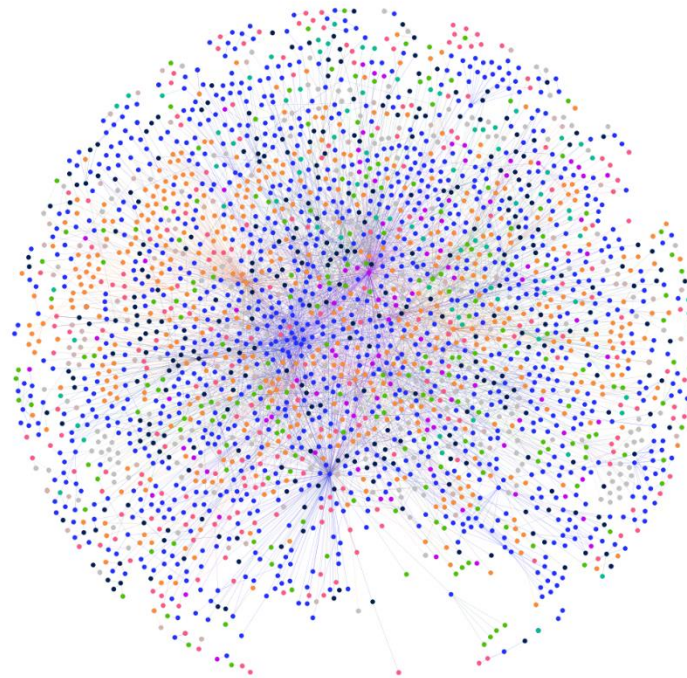
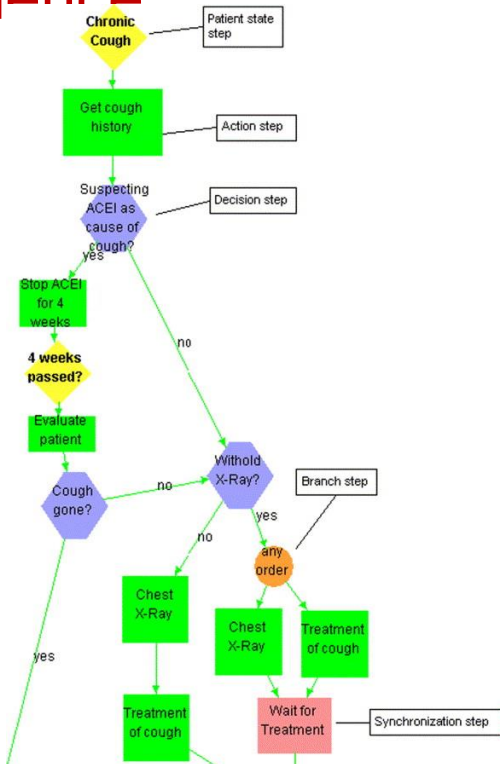
we are here



EVIDENZEN AUS DREI WELTEN

Clinical Guidelines – Scientific Studies & Publications – Real-World-Evidence

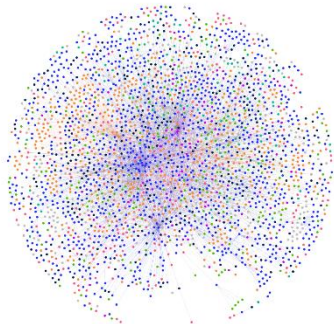
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KNOWLEDGE GRAPH EMBEDDINGS FÜR DIE VERSORGUNG (S-FORSCHUNG)

Informationsextraktion, Wissensgraphen und AI-Verfahren für die Versorgungsforschung

Literatur-Wissensgraph



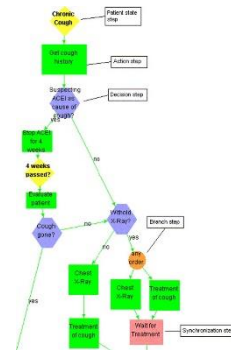
Aktualisierung



Erweiterungen

(co-Morbiditäten,
Nebenbedingungen,
neue Erkenntnisse /
Evidenzen, neue
Subgruppen)

Leitlinien-Graph



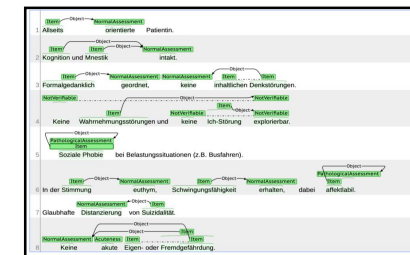
Validierung



Anpassung

(Konkordanz
zwischen Leitlinie
und realen
Behandlungspfaden)

Real-World Evidenzen



DIE WICHTIGEN BOTSCHAFTEN DIESES VORTRAGS

- WissensGraphen sind super geeignet, **um Wissen algorithmisch zugänglich** zu machen
- **Text-2-Graph Verfahren** sind inzwischen sehr weit entwickelt und werden ständig besser
- Patienten-Daten und WissensGraphen können gemeinsam genutzt werden, **um Patienten-Strata zu identifizieren**
- **Leitlinien** können als Graph-Modelle repräsentiert werden
- Es gibt **kein prinzipielles Hindernis** für das “Maschinelle Erlernen“ spezifischer, aktueller Leitlinien-Graphen
- Wie bei allen AI-Verfahren ist die **Verfügbarkeit von Daten das Haupthindernis**
- Die große Herausforderung für eine durch AI erheblich verbesserte Versorgungsforschung ist die Verfügbarkeit **großer Mengen an medizinischen Routine(Versorgungs-)daten**.

THE TEAM @ FRAUNHOFER SCAI

My team with currently 18 nationalities / 3 groups / 12 scientists / 13 PhD students



Information Extraction

Marc Jacobs
Jürgen Klein
Tim Adams
Negin Babaiha
Sumit Madan
Johannes Darms

Knowledge Graphs

Alpha Tom Kodamullil
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Stephan Gebel
Christian Ebeling
Bruce Schultz
Daniel Domingo-Fernandez
Sepehr Golritz-Khatami
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Data Science & AI

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Tamara Raschka