

Leveraging your research data using TenWise's KMAP & KMINE

TenWise offers text mining solutions by providing access to its KMAP platform.

Our **KMAP platform** captures validated scientific knowledge about human genes, micro-organisms, metabolites, pathways, phenotypes, diseases, drug compounds and research workflows, in over 200 million biological relations.

"TenWise use intelligent, innovative (bio)informatics techniques, mining data to uncover hidden relationships and networks in disease areas, their expertise providing new insight in pharma research, presented in intuitive, comprehensive reports"

BioAxis Research, customer for research in role of Brown fat in metabolic disorders and obesity

"We use KMAP for several applications both internally and externally. For example, to collect information regarding selected proteins to support our customers or for prioritization of new protein targets."

Olink Proteomics, CRO in the proteomics field

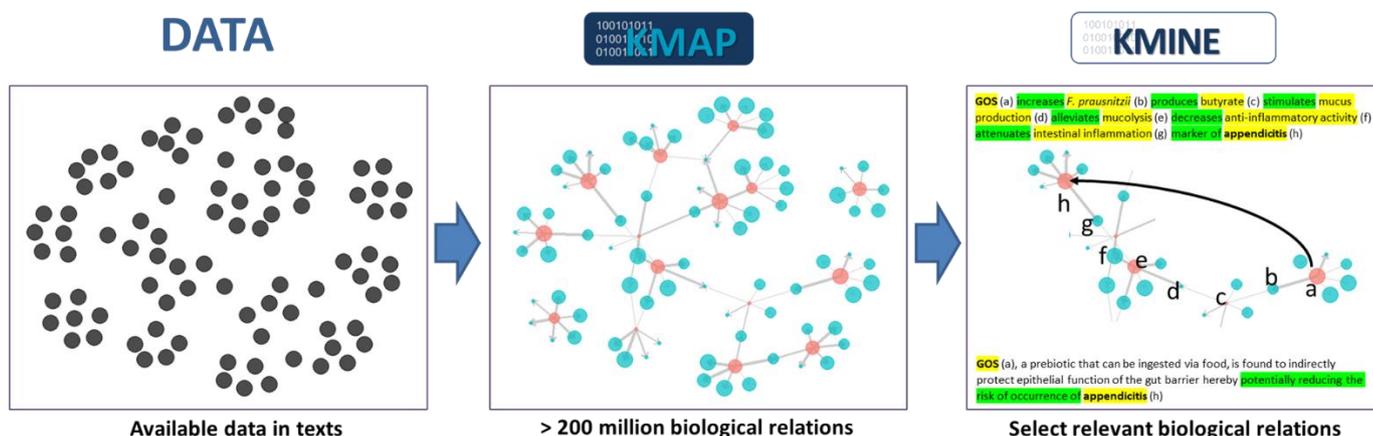
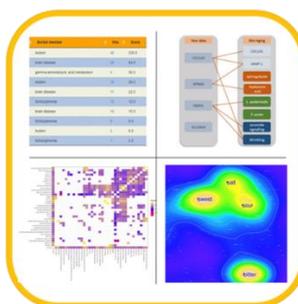


Figure: KMAP captures scientific knowledge in over 200 million biological relations and KMINE uncovers hidden relations

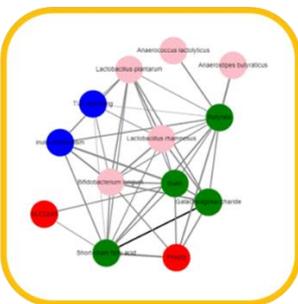
Our **KMINE products** offer access to the KMAP platform. These products are designed specifically for research teams and data engineers to assist in: **✓ discovery** **✓ drug repurposing** **✓ literature scoping** **✓ clinical validation**.



Direct access to KMAP



Web interface for interactive discovery



Dedicated literature reports



Scoping reviews

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Literature Review: Identifying potential new opportunities for microbiome therapies in CNS diseases

We used:



We performed a **KMINE Scoping Review** making use of the TenWise KMAP by applying an expert curated filter with the most important Central Nervous System (CNS) diseases. Subsequently we searched with a selection of microbiome related terms to find those abstracts dealing with the selected CNS disease (eg. anxiety disorder) and the gut microbiome, i.e. the Gut-Brain Axis (GBA). An overview is depicted below that ranks (in order from large to small) the number of abstracts. We performed a full review on abstracts and full texts to retrieve GBA-related mechanism of actions for each disease.

CNS disease	Microbiome terms	Weight factor	Number of abstracts	Score
anxiety disorder	microbiome	1	500	500 link
Alzheimer's disease	microbiome	1	309	309 link
Parkinson's disease	microbiome	1	276	276 link
autism spectrum disorder	microbiome	5	186	930 link
schizophrenia	microbiome	1	141	141 link
dementia	microbiome	0	120	0 link
bipolar disorder	microbiome	1	55	55 link
amyotrophic lateral sclerosis	microbiome	1	48	48 link
epilepsy	microbiome	0	37	0 link
mood disorder	microbiome	1	17	17 link
Huntington's disease	microbiome	0	11	0 link
eating disorder	microbiome	0	10	0 link
acute stress disorder	microbiome	0	7	0 link
Transient ischemic attack	microbiome	0	4	0 link
personality disorder	microbiome	0	3	0 link
vascular dementia	microbiome	0	2	0 link
Lewy body dementia	microbiome	0	1	0 link

Conclusions: Cause-effect relations are still largely unknown for most CNS diseases. Autisms and anxiety disorders are the earliest diseases that have been linked to the gut microbiome. The top 4 diseases showing strong indications for a GBA-related **mechanism of action:** anxiety disorder, Alzheimer's disease, Parkinson's disease and autism spectrum disorder. Potential upcoming areas within GBA (Orange) are: schizophrenia, bipolar disorder, ALS and epilepsy.

Want to know how to access KMAP? Call us or request a demonstration or quote:



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