

http://sabio.h-its.org/chemHits/

# NORMALIZATION AND MATCHING OF CHEMICAL COMPOUND NAMES

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Select Database:

Perform Acid-Ba

Match List



#### Background

We have developed ChemHits (http://sabio.h-its.org/chemHits/), an application which detects and matches synonymic names of chemical compounds and thereby facilitates the bundling of corresponding data referring to the same compound, but described with different names. The tool that we have developed is based on natural language processing (NLP) methods and applies rules to systematically normalize chemical compound names. Subsequently, matching of synonymous names is achieved by comparison of the normalized name forms. The tool is capable of normalizing a given name of a chemical compound and matching it against names in (bio-)chemical databases, like SABIO-RK, PubChem, ChEBI or KEGG, even when there is no exact name-to-name-match.

#### **Chemical Compound Name Matching** ChemHits chemical compound name normalization and matching Short Documentation Match Single Compound Name ame of Compound: 2-aminoprop-2-enoic acid elect Database: Chebi200711TEST Groovy Results in Chebi200711TEST\_Groovy for 2-aminoprop-2-enoic acid erform Acid-Base Equalisation? 🗵 T Groovy) Name (Chebi200711TEST Gro arch for Normalised Substrings 2-aminoacrylic acid 🙆 ChemHitsApp Match List of Compour File Help

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Options: Perform Acid-Bas Results: ID (pubchem020211TEST) 123991	Name (pubchem020211TEST)	Normalised Dubschings: Normalised Name 2-aminoacrylate		Export Re Input Name 2-aminoprop-2-enoic acid

The tool is capable of normalizing a given name of a chemical compound and matching it against names in (bio-)chemical databases, like ChEBI (http://www.ebi.ac.uk/chebi/), KEGG COMPOUND (http://www.genome.jp/kegg/compound/), SABIO-RK (http://sabio.h-its.org/), or PubChem (http://pubchem.ncbi.nlm.nih.gov), even when there is no exact name-to-name-match. The tool is also able to match a complete list of compound names against these databases which makes it useful for the automatic cross-annotation of chemical data in databases. We offer a web-based service with pre-normalized reference lists (upper screenshot) and a tool that can be installed and run locally with any given reference list (lower screenshot).

# The Future: From Name to Structure

**Combining Name Matching and Semantic Analysis** 

Alternative name:

Alternative name:

Alternative name:

## **Terminology of Chemical Compounds**

Synonymous notations of chemical compounds

Trivial name and systematic chemical description

Valproic acid

= 2-Propylpentanoic acid

Different parts of the molecule could be considered as lead structure

**Acetylphenol** 

= Phenzylacetate

Abberrant order of the substituents of a lead structure (prefixes)

- 2-Amino-6-methyl-4-pyrimidol = 6-Methyl-2-amino-4-pyrimidol
- Description of substituents as prefix (like amino-) or suffix (like –amine)
  - 2-Aminopropane
- = Propan-<mark>2-amine</mark>
- 2-Methylpropan-2-ol = 2-Hydroxy-2-methyl-propane

**Different nomenclature systems (e.g. abberrant order of the morphems)** 

2-Amino-6-methyl-4-pyrimidol = 2-Amino-6-methylpyrimidin-4-ol

A chemical compound can have many different names - trivial, as well as systematic names. Hence, the identification of a chemical compound solely based on its name requires comprehensive chemical knowledge and often extensive searches in chemical databases. However, this identification is crucial for the integration of biochemical data e.g. for the setup of biochemical models based on published data. As many publications exclusively describe a chemical compound by its name the matching of these diverging notations can be tedious.



The tool that we have developed applies transformation rules to systematically normalize the notation of chemical compound names [1]. Subsequently, matching of synonymous names is achieved by comparison of the normalized name forms. The normalization rules include, among others, reordering of substituent descriptions in the name and replacement of synonymous name constituents (e.g. equivalent trivial names). Matching of conjugated acid-base pairs is optional for biochemicals.



After normalization, synonymous notations could potentially be matched to the corresponding systematic name as defined by the International Union of Pure and Applied Chemistry (IUPAC). When combined with our approach to construct chemical structures from systematic names [2, 3], notations could be translated into a chemical structure (SMILES) and classified by functional groups [4], resulting in the unambiguous identification of these compounds.

### References

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- 2) Henriette Engelken: "A System for Semantic Analysis of Chemical Compound Names", in Proceedings of the ACL-IJCNLP 2009 Student Research Workshop, Singapore, August 2-7, 2009, pages 36-44. Association for Computational Linguistics.
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- 4) Wittig U, Weidemann A, Kania R, Peiss C, Rojas I: "Classification of chemical compounds to support complex queries in a pathway database", Comparative and Functional Genomics, 5: 156-162 (2004)









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