# Binary Substructure Descriptors for Organic Compounds

## K. VARMUZA\*, W. DEMUTH, H. SCSIBRANY

Vienna University of Technology Institute of Chemical Engineering



Laboratory for ChemoMetrics

\* Corresponding and presenting author

kvarmuza@email.tuwien.ac.at www.lcm.tuwien.ac.at Getreidemarkt 9/166-2 A-1060 Vienna. Austria

Acknowledgment Austrian Science Fund, project P14792-CHE M. Karlovits, A. Kerber, R. Laue, S. Stein, R. Neudert

Poster Presentation: Mathematics, Chemistry & Computer Sciences - MATH/CHEM/COMP 2004 21 - 26 June 2004, Dubrovnik, Croatia

# Introduction / Overview

A set of 1365 substructures has been defined for the representation of organic compounds by binary vectors.

- Substructure encoding is evident to chemists and easily interpretable.
- Substructure encoding is capable to cover the great diversity of chemical structures.

**Software SubMat** has been developed for an easy and flexible generation of binary substructure descriptors [1,2].

Only 2-dimensional (connectivity) data of chemical structures are considered.

### **Applications** are reported for

- characterization of structural diversity [2],
- search for similar structures [2],
- test of spectra similarity measures [3,4],
- cluster analysis of structures [1,2].
- [1] K. Varmuza, H. Scsibrany: J. Chem. Inf. Comput. Sci. 40 (2000) 308-313.
- [2] H. Scsibrany, M. Karlovits, W. Demuth, F. Müller, K. Varmuza: Chemom. Intell. Lab. Syst. 67 (2003) 95-108.
- [3] K. Varmuza, M. Karlovits, W. Demuth: Anal. Chim. Acta **490** (2003) 313-324.
- [4] W. Demuth, M. Karlovits, K. Varmuza: Anal. Chim. Acta (2004) in print.

# Software SubMat

SubMat calculates binary substructure descriptors for an input file with molecular structures, and an input file with substructures (all in Molfile format).

SubMat runs under MS Windows operating systems.



#### Example

- n = 1000 molecular structures, and p = 200 substructures 1 s computation time (Pentium IV, 2.6 GHz)
- 5  $\mu$ s per descriptor value

Demo version and User Guide free at www.lcm.tuwien.ac.at (Software)

Authors: H. Scsibrany and K. Varmuza

# Substructures

#### Groups

1	Elements (single atoms)		46
2	Two-atom substructures		78
3	Single rings (not aromatic)		404
4	Condensed rings (not aromatic)		130
5	Aromatic rings		97
6	Other rings		39
7	Trees (chains and branches)		418
8	Functional groups		153
		sum	1365

Bonds: single, double, triple, aromatic, any type. Pseudo elements: A (hetero atom), Q (any atom, except H).

#### **Examples**

#### Group 3: single rings, not aromatic

	₽°	N N	$\bigcap^{\circ}$	$\bigcirc$
IR:	0.69%	IR: 1.61%	IR: 0.59%	IR: 0.02%
MS:	1.15%	MS: 0.20%	MS: 2.32%	MS: 1.40%

#### Group 5: aromatic rings



Frequency of compounds containing the substructure are given for two spectroscopic databases. IR, 13,484 compounds; MS, 106,955 compounds.

# **Applications**

### **Characterization of structural diversity**

Frequency distributions of **Tanimoto indices** (t, a measure for structural similarity) from 10<sup>6</sup> randomly selected structure pairs from two spectroscopic databases. IR, 13,484 compounds; MS, 106,955 compounds.



### Search for similar structures

The query structure has been searched in two spectroscopic databases. IR, 13,484 compounds; MS, 106,955 compounds. Hits 1 - 3 are shown, including Tanimoto indices, t.



# **Applications**

### Spectral versus structural similarity

Three methods (F862B, M200, M1000) for mass spectra similarity searches have been compared with 10,000 "test-unknowns". Method F862 yields hitlists with best chemical structure information. Database: 106,955 comps.



### **Cluster analysis of structures**

A spectral similarity search (IR) for the "test-unknown" **3-amino-benzylalcohol** gave 25 compounds. PCA with 18 binary substructure descriptors (selected by maximum variance) shows potential substance classes for the test-unknown. PC1,2: 36%, 28% of total variance, resp.

