

Molecular descriptors based on automorphism data

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Basic definitions / Overview

Chemical structures are represented by mathematical **graphs** [1,2].
 Graph vertices: atoms; graph edges: bonds; hydrogen-depleted;
 connected coloured graphs; no atomic charge info; no 3D info.

Topologically (constitutionally) **equivalent atoms (bonds)** have identical neighborhoods in terms of connectivity as described by the graph - considering the whole graph (molecular structure).

The **automorphism group** of a graph describes all mappings of the graph onto itself - preserving the connectivity (not cutting bonds). It contains data about **topologically equivalent atoms (bonds)**.

Asymmetric structures: only a single (trivial) mapping exists;
 highly symmetric structures: several (many) mappings.

Size of the automorphism group: number of possible mappings of a graph onto itself; it is a **symmetry measure** for the graph.

Orbits (in graph theory)

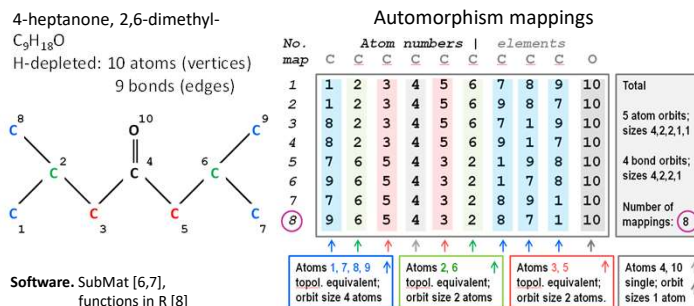
Atom (vertex) orbit: set of topologically equivalent atoms

Bond (edge) orbit: set of topologically equivalent bonds

Molecular descriptors [3] can be derived from the size of the automorphism group and from the frequencies of the different orbit sizes [4]. They are evaluated here together with other descriptors [5] for multivariate QSPR models: quantitative structure property relationships for the prediction of melting points.

Continues a **similar study with exhaustive sets of alkanes** [6].

Automorphism with a demo structure



Examples of molecular descriptors based on automorphism data

- * **Number of orbits**; separately for atom types (C, N, O) and bond types (single, double, triple, aromatic).
- * **Size of automorphism group** (number of mappings); absolute, logarithm, normalized by number of atoms, bonds.
- * Number of asymmetric C-atoms.
- * Based on **frequencies of orbits with sizes 1, 2, 3, ...**; e. g., entropy, symmetry index [3,9], root of orbit polynomial [4]; separately for atom and bond types.

Application example [QSPR model for melting point]

Data

Origin: Tetko et al., 2014 [10]; Bradley J.C., 2014 [11]

Selected:

$n = 1161$ chemical compounds (C_{5-30} , H, N, O only),

y, melting point 20 – 300 °C; $sd(y) = 61$

X, $m = 478$ molecular descriptors;

$m_1 = 428$ (Dragon) [5]; $m_2 = 50$ (Automorphism);

autoscaled.

X_{SEL} , $m = 21$, **selected stepwise** with BIC criterion [12];

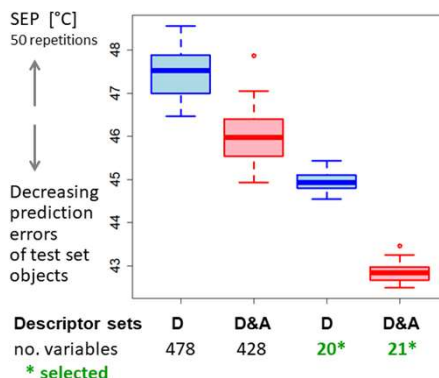
$m_{1,SEL} = 15$ (Dragon); $m_{2,SEL} = 6$ (Automorphism)

Multivariate models

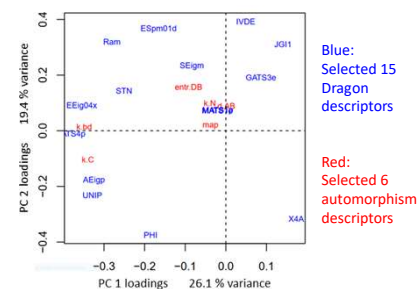
Linear regression with **PLS**; strategy **repeated double cross validation (rdCV)** separates optimization of model complexity (no. of PLS components) and estimation of prediction performance [13, 14].

Performance criterion: SEP = standard deviation of prediction errors for test set objects (boxplot for 50 repetitions).

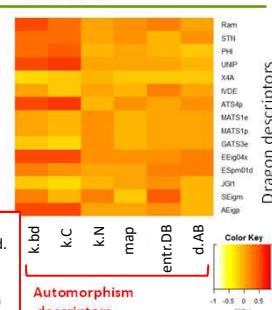
Comparison of descriptors sets without/with automorphism descriptors



PCA loading plot of selected descriptors



Heat map for correlation coefficients between selected Dragon and automorphism descriptors



Tentative conclusion. Perhaps a useful complement to other descriptors. More tests required.

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