



**HAL**  
open science

# Understanding odour thresholds: A Quantitative Structure-Property Relationship study of the role of the molecular structure

José Antonio Piornos Martinez, Anne Tromelin

► **To cite this version:**

José Antonio Piornos Martinez, Anne Tromelin. Understanding odour thresholds: A Quantitative Structure-Property Relationship study of the role of the molecular structure. 13. Wartburg symposium on flavour chemistry and biology, Oct 2023, Eisenach, Germany. hal-04228193

**HAL Id: hal-04228193**

**<https://hal.inrae.fr/hal-04228193v1>**

Submitted on 5 Aug 2024

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

# UNDERSTANDING ODOUR THRESHOLDS: A QUANTITATIVE STRUCTURE-PROPERTY RELATIONSHIP (QSPR) STUDY OF THE ROLE OF THE MOLECULAR STRUCTURE

JOSE PIORNOS · ANNE TROMELIN

Centre des Sciences du Goût et l'Alimentation (CSGA), INRAE, CNRS, Université de Bourgogne, Institut Agro, Dijon, France

## THRESHOLDS ARE USEFUL

- ❖ They are commonly used in food science to evaluate the **potency** of an aroma compound, which varies from one compound to another.
- ❖ This characteristic is usually associated with the molecule's **odour detection threshold** (ODT).
- ❖ However, the relationship between a **molecule's structure** and its ODT has not a straightforward answer.

## WE HAD SOME QUESTIONS...

- What is the role of the **chemical structure** on odour detection thresholds?
- What are the most impactful **molecular descriptors** representing these chemical features?

## DATA TREATMENT AND MODELLING

**ODT values in water from the literature**

- 33 research papers
- 407 molecules
- 858 ODT values

**Removal of outliers**

Z-test

**Average ODT for each molecule**

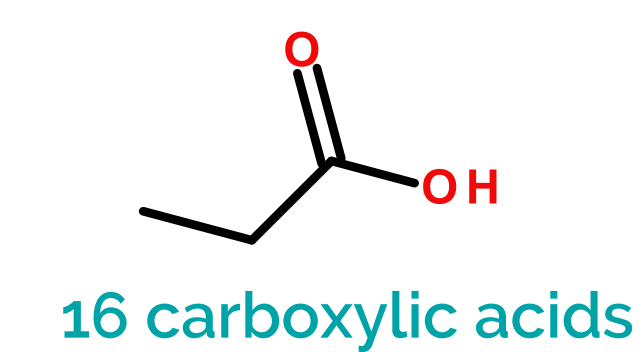
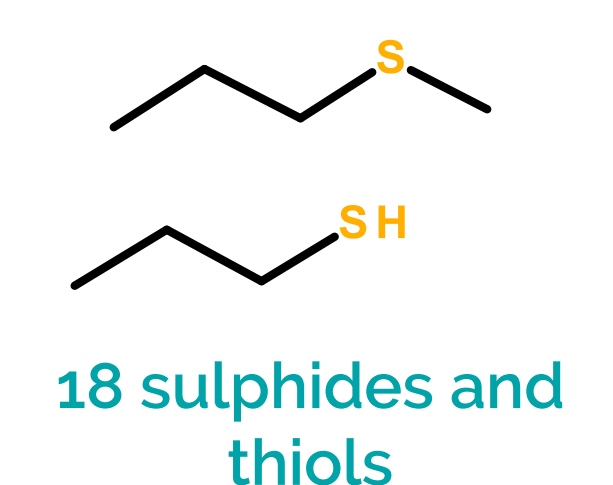
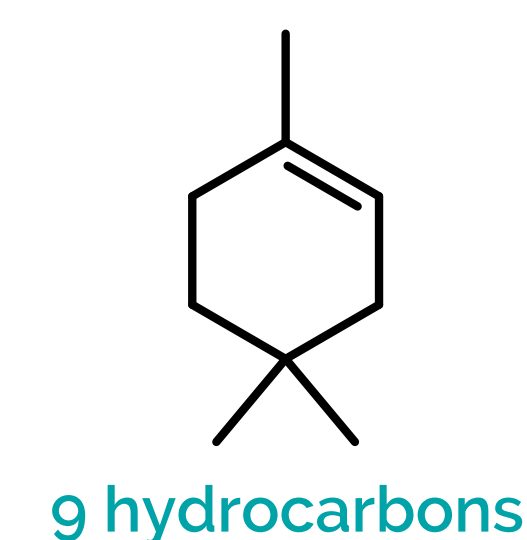
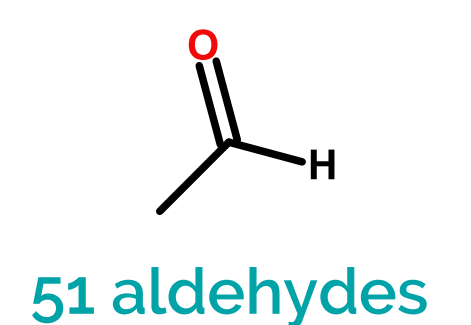
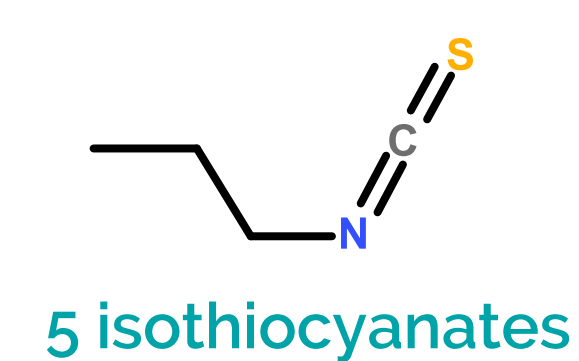
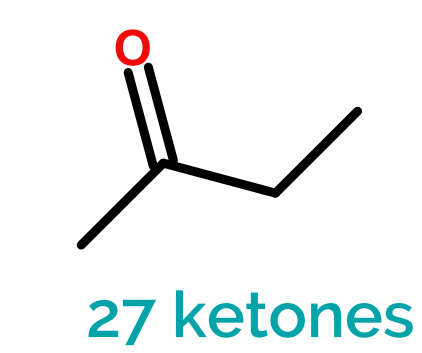
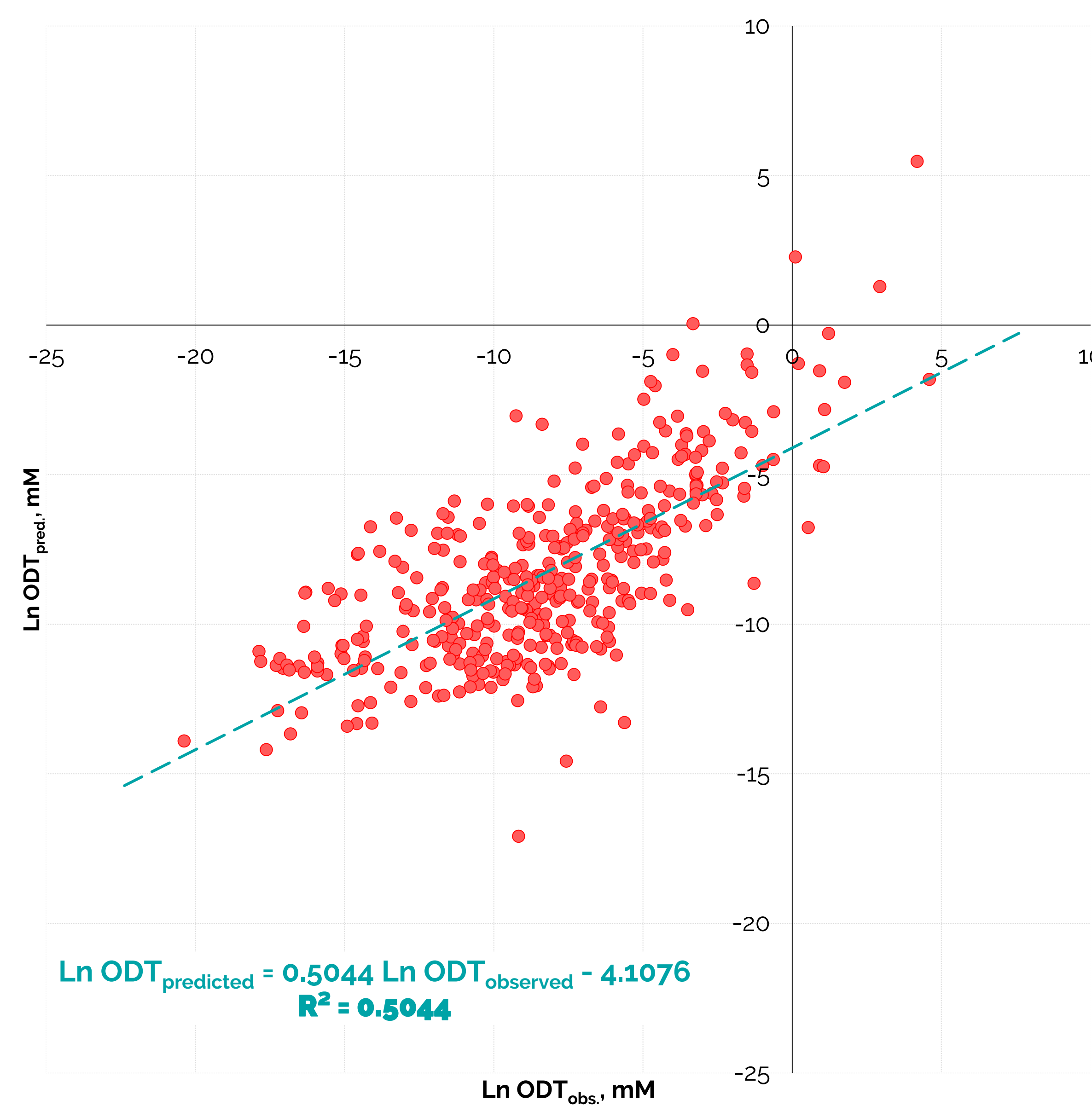
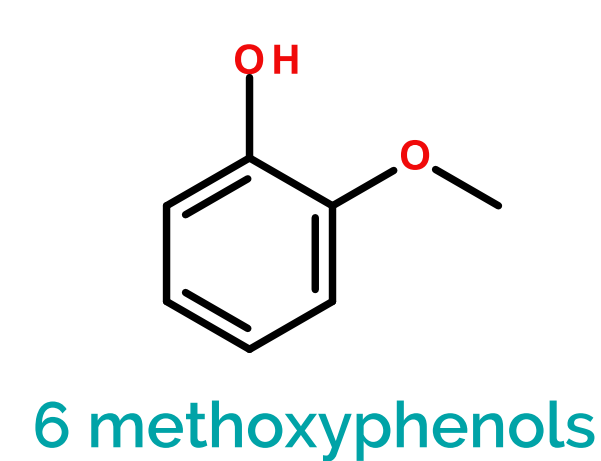
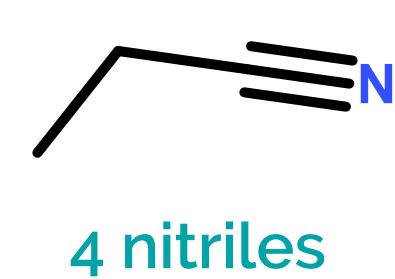
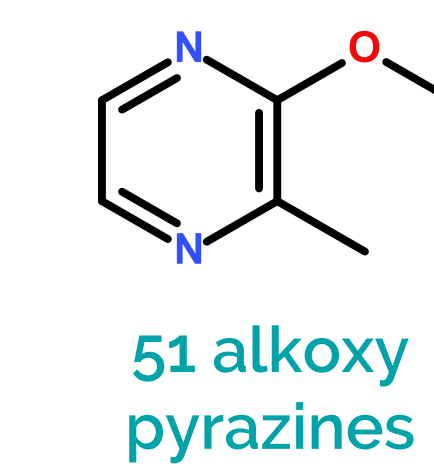
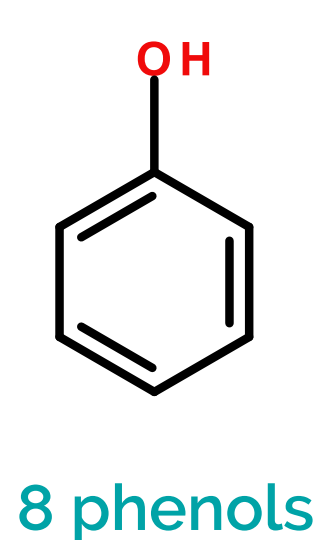
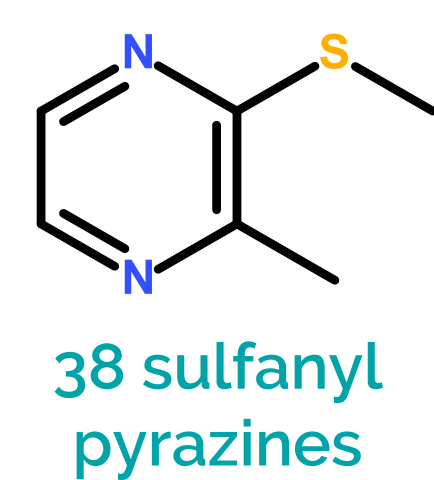
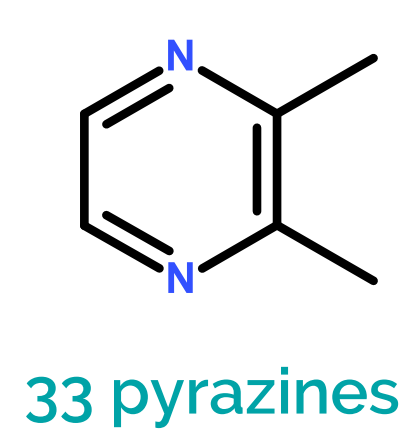
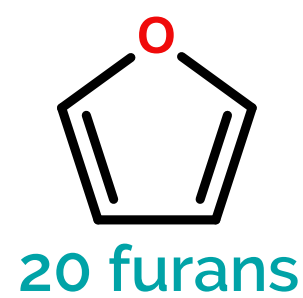
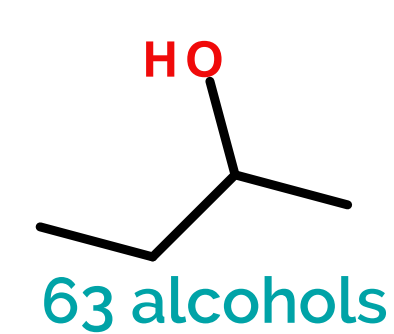
**Calculation of molecular descriptors**

>5,000 descriptors  
Biovia Discovery Studio

### Model development

- Generation of QSPR models by Genetic Functional Analysis (GFA)
- Multiple linear equations for the correlation of ODT values (dependent variable) and molecular descriptors (independent variables)
  - Optimisation by maximisation of R<sup>2</sup>
- A balance between the number of descriptors and the model fit is preferred, i.e. the best fit for the fewer variables.

The **407 aroma** compounds employed belong to a large variety of chemical classes with a large variety of functional groups.



### The proposed 6-variable model:

$$\text{Ln ODT} = 16.56 - 0.93 \cdot \text{ES\_Count\_dsCH} - 2.38 \cdot \text{ES\_Count\_ssO} - 11.23 \cdot \text{Jurs\_RASA} - 0.14 \cdot \text{Jurs\_RNCS} - 0.12 \cdot \text{Molecular\_Weight} + 0.015 \cdot \text{V\_DIST\_equ}$$

## WHAT DO THESE MOLECULAR DESCRIPTORS MEAN?

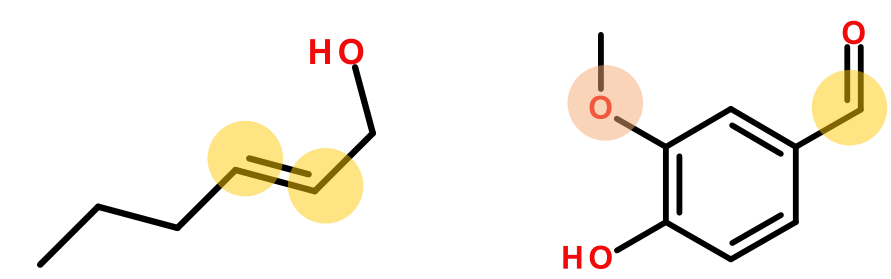
A molecular descriptor is a structural or physicochemical property of a molecule or a part of a molecule.

### ES\_Count\_dsCH

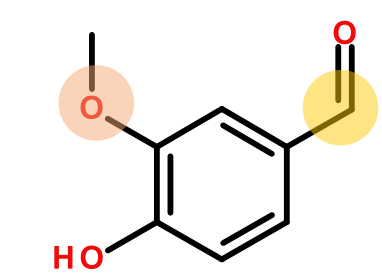
- ❖ Number of =CH- groups in a molecule, i.e. containing a double ('d') and a single ('s') bound (nonaromatic).

### ES\_Count\_ssO

- ❖ Number of O atoms bound by two single ('s') bounds (nonaromatic) to atoms other than H.



ES\_Count\_dsCH = 2  
ES\_Count\_ssO = 0



ES\_Count\_dsCH = 1  
ES\_Count\_ssO = 1

### Jurs\_RASA

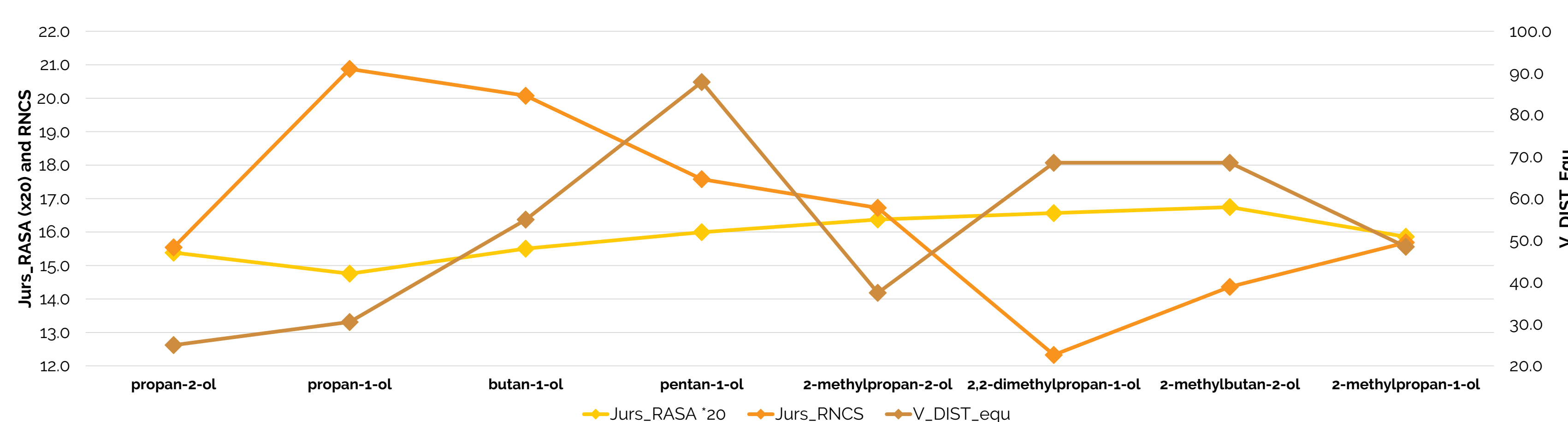
- ❖ Relative Apolar Surface Area. The total hydrophobic surface area (TASA) divided by the total molecular solvent-accessible surface area (SASA).

### Jurs\_RNCS

- ❖ Relative Negative Charged Surface. The solvent-accessible surface area of the most negative atom (SA<sup>-</sup><sub>max</sub>) divided by the relative negative charge (RNCG).

### V\_DIST\_Equ

- ❖ Vertex distance/equality. A descriptor based on a matrix containing information about the distances between the different parts of a molecule.
- ❖ Related to the shape of the backbone of the molecules.



- ❖ The descriptors associated to hydrophobic features (ES\_Count\_dsCH and Jurs\_RASA), to the size of the molecule (**Molecular weight**) and its electronegativity (ES\_Count\_ssO and Jurs\_RNCS) contribute negatively to the ODT values.
- ❖ Only V\_DIST\_Equ, also related to size and shape, contributes positively, counterbalancing the weight of the others.
- ❖ **The more hydrophobic, heavy/big and/or electronegative, the lower the threshold in water.**

Contact details:

José Piornos, PhD

jose-antonio.piornos-martinez@inrae.fr

Research funded by:

TRANSFORM  
INRAE

INRAE