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## Incorporating chirality within the graph kernel framework

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**Abstract.** Molecules being often described using a graph representation, graph kernels provide an interesting framework which allows to combine machine learning and graph theory in order to predict molecule's properties. However, some of these properties are induced both by the covalent bound relationships between atoms and by constraints on the relative positioning of these atoms. Graph kernels based solely on the graph representation of a molecule do not encode the relative positioning of atoms and are consequently unable to predict accurately molecule's properties connected with this relative positioning. In this report, ordered structured object are introduced in order to incorporate spatial constraints within the graph kernel framework. The incorporation of this new features within the graph kernel framework allows to predict accurately chiral information hence overcoming the previous limitation.

Keywords: Graph kernel, Chemoinformatics, Chirality.

#### 1 Introduction

The purpose of Chemoinformatic is to predict properties of molecules, in order to facilitate drug design. Chemoinformatics is based on the similarity principle: two structurally similar molecules should have similar properties.

One common method to predict chemical properties consist to design a vector of descriptors from a molecule and use statistical machine learning algorithms to predict molecule's properties. Such methods [4,3], can use structural information, physical properties or biological activities in order to compute vectors of descriptors. However, such an approach requires to either select a random set of pre defined descriptors (before a variable selection step) or to use an heuristic definition of appropriate descriptors by a chemical expert. In both cases, the transformation of the graph into a finite vector of features induces a loss of information.

Another approach consist to encode a molecule by a graph, and use it to predict properties.

#### Definition 1. Molecular graph

A molecular graph is a labeled graph  $G = (V, E, \mu, \nu)$  representing a molecule. The unlabeled graph (V, E) encodes the structure of the molecule, each node  $v \in V$  encoding an atom and each edge  $e = (v, w) \in E$  a bond between two atoms v and w.  $\mu$  associates to each vertex  $v \in V$  a label  $\mu(v)$  encoding the nature of the atom and  $\nu$  associates to each edge e a type of bond  $\nu(e)$  (single, double, triple or aromatic).

Several methods based on graph theory use this representation to predict properties. One approach consists to search subgraphs with a large difference of frequencies between a set of positive and a set of negative examples [5]. Another approach consists to encode each class of molecules by a graph prototype and to measure the structural similarity between each prototype and an input molecule [6]. However, these methods can not be easily combined with machine learning algorithms. This is not the case of graph kernel methods, which can be coupled to machine learning algorithm provided that the kernel is definite positive. Let  $\mathcal{G}$  be the set of graph. A definite positive kernel is a symmetric function  $k: \mathcal{G} \times \mathcal{G} \to \mathbb{R}$  such that:

$$\sum_{i=1}^{n} \sum_{j=1}^{n} c_i c_j k(G_i, G_j) \ge 0 \text{ where } n > 0, G_1, \dots, G_n \in \mathcal{G}, c_1, \dots, c_n \in \mathbb{R}$$

Such a definite positive kernel corresponds to a scalar product between two vectors  $\psi(G)$  and  $\psi(G')$  in an Hilbert space.

A large family of graph kernel methods, associates a bag of patterns to each graph, and define the kernel value from a measure of similarity between those bags [7–9, 1]. In [7] a graph kernel is defined as a measure of similarity between set of walks extracted from each graph. But those walks are linear features and thus have limited expressiveness. An infinite set of tree patterns is used in [8] to define kernels. However, the similarity between two graphs is based on an implicit enumeration of their common tree patterns which does not allow to readily analyze the influence of a pattern on the prediction. Finally [9] and [1] are based on an explicit enumeration of patterns. In [9], a predefined set of unlabeled subgraphs, called graphlets, is enumerated for each graph and in [1] all subtrees of a labeled graphs up to size 6, called treelets are enumerated. One advantage of [1] is that, unlike [9], the label of the graph are taken into account by the graph similarity measure.

However, some molecules may have a same molecular formula, a same molecular graph but a different relative positioning of their atoms. Such molecules are said to be stereoisomers. Different stereoisomers may be associated to different properties. However, usual graph kernels based on the molecular graph representation are not able to capture any dissimilarity between these molecules. From a more local point of view, an atom or two connected atoms are called stereocenters if a permutation of the positions of two atoms belonging to the union of their neighborhoods produces a different stereoisomer.

In order to get an intuition of stereoisomerism, let us consider an acyclic molecular graph rooted on an atom of carbon with four neighbors, each neighbor

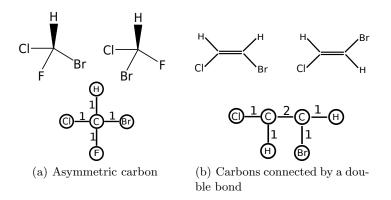


Fig. 1. Two types of stereocenters.

being associated to a different subtree. Such an atom, called an asymmetric carbon, is a stereocenter and has two different spatial configurations of its neighbors encoded by a same molecular graph (Figure 1(a)). Using molecule represented in Figure 1(a), one configuration corresponds to the case where the three atoms (Cl,F,Br) considered from the atom H are encountered in this order when turning counter-clockwise around the central carbon atom. The alternative stereoisomer corresponds to the case where this sequence of atoms is encountered clockwise when considered from the same position. Two carbons, connected by a double bond, can also define stereoisomers (Figure 1(b)). Indeed, on the left side of Figure 1(b) both hydrogen atoms are located on the same side of the double bond while they are located on opposite sides on the stereoisomer represented on the right. In this case both carbon atoms of the double bond correspond to a stereocenter.

To distinguish those configurations, we introduce the two following subsets of the set of vertices V of a molecular graph:

#### **Definition 2. Potential Asymmetric Carbons**

Let us denote  $V_{PAC}$  the subset of V containing all vertices encoding atoms of carbon with four neighbors:

$$V_{PAC} = \{v \in V \mid \mu(v) = C' \text{ and } |V(v)| = 4\}$$

Since being an atom with four neighbors is a necessary condition to define an asymmetric carbon, the set  $V_{PAC}$  contains all vertices which may encode such atoms.

#### Definition 3. Set of double-bonds connecting carbon atoms

The subset of V containing all atoms of carbon which share a double bond with another carbon is noted  $V_{DB}$ :

$$V_{DB} = \left\{ v \in V \mid \exists e(v, w) \in E, \nu(e) = 2, \begin{pmatrix} |V(v)| = |V(w)| = 3 \\ \text{and} \\ \mu(v) = \mu(w) = C' \end{pmatrix} \right\}$$

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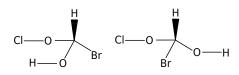


Fig. 2. Asymmetric carbons with identical neighborhood.

An atom of carbon with two double bounds must have a degree equal to two. Hence, each vertex v belonging to  $V_{DB}$  is incident to a single double bound and we denote  $n_{=}(v)$  the other carbon connected by this double bond. Note that  $n_{=}(v) \in V_{DB}$ .

Brown et al. described in [2] a method which includes information related to the spatial configuration of atoms within the tree-pattern kernel [8]. However, this method only considers the direct neighbors of a stereocenter while, as shown by Figure 2, the difference between two subtrees of a stereocenter may not be located on the root of the subtree. In this last case [2] considers as identical two different stereocenters and is thus unable to recover their different properties.

In this paper we propose a method to incorporate the spatial configuration of atoms within a graph kernel based on a subtree enumeration [1]. This method remains valid when the spatial configuration is not encoded in the direct neighborhood of a stereocenter. In Section 2, we define a graph encoding of stereoisomers and we introduce chiral vertices as vertices encoding stereocenters. Next, in Section 3, we restrict our attention to acyclic molecules. Such a restriction allows us to efficiently characterize a chiral vertex by a rooted tree. In Section 4, we define the smallest tree characterizing a chiral vertex and use this information to design a graph kernel between chiral molecules. Finally, we demonstrate the validity of our kernel through experiments in Section 5.

#### 2 Encoding of stereoisomers

#### 2.1 Ordered Structured Object

#### **Definition 4. Structured Objects**

A structured object S is an object to which we can associate an unique labeled graph  $G(S) = (V, E, \mu, \nu)$ .

A structured object can be for example a graph (associated to itself) or a rooted tree (associated to an acyclic graph).

An usual method in chemistry to encode stereoisometry consists to encode a relative order on the neighborhood of each vertex. In order to encode such an information, we introduce the notion of order on structured object.

#### **Definition 5. Ordered Structured Objects**

An ordered structured object  $S = (\hat{S}, ord)$  is a structured object  $\hat{S}$ , associated to a graph  $G(S) = (V, E, \mu, \nu)$ , together with a function ord which maps each vertex v belonging to a subset  $V_{ord}$  of V onto an ordered list of a subset of its neighborhood V(v):

$$ord \begin{cases} V_{ord} \to V^* \\ v \to v_1 \dots v_n \end{cases}$$

where  $\{v_1, \ldots, v_n\} \subset V(v)$  denotes a subset of the neighborhood of v.

We denote |ord(v)| = n the length of the ordered list for any  $v \in V_{ord}$ . We have thus  $0 < |ord(v)| \le |V(v)|$ .

Note that, the notation  $V_{ord}$  which denotes the subset of V for which function ord is defined will be used in the remaining part of this document.

#### Definition 6. Set of Ordered Structured Objects

A set of ordered structured objects  $\mathscr{S}$  is a set  $\mathscr{S} = \{S = (\hat{S}, ord)\}$  from which we can define a set of isomorphism  $\operatorname{Isom}(\hat{S}, \hat{S}') \subset \operatorname{Isom}(G(S), G(S'))$  between any two structured objects  $\hat{S}$  and  $\hat{S}'$ . This set of isomorphism must respects the following properties:

1.  $\forall S_1, S_2 \in \mathscr{S}^2, f \in \operatorname{Isom}(\hat{S}_1, \hat{S}_2) \Leftrightarrow f^{-1} \in \operatorname{Isom}(\hat{S}_2, \hat{S}_1)$ 2.  $\forall S_1, S_2, S_3 \in \mathscr{S}^3, f \in \operatorname{Isom}(\hat{S}_1, \hat{S}_2), g \in \operatorname{Isom}(\hat{S}_2, \hat{S}_3) \Rightarrow g \circ f \in \operatorname{Isom}(\hat{S}_1, \hat{S}_3)$ 3.  $\forall S \in \mathscr{S}, \operatorname{Isom}(\hat{S}, \hat{S})$  is a group. 4.  $\forall (S, S') \in \mathscr{S}^2, \forall f \in \operatorname{Isom}(\hat{S}, \hat{S}')$ 

$$\begin{cases} f(V_{ord}) = V'_{ord} \\ \forall v \in V_{ord}, |ord(v)| = |ord'(f(v))| \end{cases}$$

The first three conditions impose that our restricted set of isomorphim relationships satisfies the usual properties of isomorphisms: the inverse and the composition of two isomorphisms is still an isomorphism (conditions 1 and 2) and considering a set of automorphisms, the identity belongs to our valid set of isomorphisms (condition 3). The last condition imposes that the set of vertices on which the function ord is defined remains stable by an isomorphims. It further imposes that an isomorphism does not modify the number of vertices on which the order relationship is defined for each vertex.

#### Definition 7. Isomorphism between ordered structured objects

Let us consider a set of ordered structured objects  $\mathscr{S}$ . Two ordered structured objects  $S = (\hat{S}, ord)$  and  $S' = (\hat{S}', ord')$  are said to be isomorphic  $S \simeq S'$  iff there is an isomorphism between the structured objects  $\hat{S}$  and  $\hat{S}'$  which is coherent with the order on the subsets of the neighborhoods:

$$\begin{split} S &\simeq_o S' \Leftrightarrow \exists f \in \operatorname{Isom}(\hat{S}, \hat{S}') \ s.t. \\ &\forall v \in V_{ord} \ \text{with} \ ord(v) = v_1 \dots v_n, \ ord'(f(v)) = f(v_1) \dots f(v_n) \end{split}$$

In this case, f is called an ordered isomorphism between S and S', and we denote  $\text{IsomOrd}(S, S') \subset \text{Isom}(\hat{S}, \hat{S}')$  the set of ordered isomorphism between S and S'.

**Proposition 1.** Ordered structured object isomorphism induces an equivalence relationship.

*Proof.* Let us consider a set of ordered structured objects  $\mathscr{S}$ . We have thus to show that the structured object isomorphism relationship is reflexive, symmetric and transitive:

1. The isomorphism between structured object is reflexive.

Let  $S = (S, ord) \in \mathscr{S}$  an ordered structured object associated to a graph  $G(S) = (V, E, \mu, \nu)$ , and let f denotes the identity function on V ( $f(v) = v, \forall v \in V$ ).

Then  $f \in \text{Isom}(\hat{S}, \hat{S})$  (Definition 6, condition 3) and:

$$\forall v \in V_{ord} \subset V, f(v_1) \dots f(v_n) = ord(f(v)) = ord(v) = v_1 \dots v_n$$

where  $\{v_1, \ldots, v_n\} \subset V(v)$  denotes a subset of the neighborhood of v. Therefore,  $S \simeq S$ .

- 2. The isomorphism between ordered structured object is symmetric.
  - Let  $S_a = (\hat{S}_a, ord_a) \in \mathscr{S}$  and  $S_b = (\hat{S}_b, ord_b) \in \mathscr{S}$  two ordered structured object, respectively associated to  $G(S_a) = (V_a, E_a, \mu_a, \nu_a)$  and  $G(S_b) = (V_b, E_b, \mu_b, \nu_b)$ , such that  $S_a \simeq S_b$  and let us further denote by f the ordered isomorphism between  $S_a$  and  $S_b$ .

By definition f is also an isomorphism between  $\hat{S}_a$  and  $\hat{S}_b$ , therefore it exists (Definition 6, condition 1) an isomorphism  $f^{-1}$  between  $\hat{S}_b$  and  $\hat{S}_a$ . Let us consider a vertex  $v_b$  in  $V_{ord_b} \subset V_b$  and  $v_a = f^{-1}(v_b) \in V_{ord_a}$  (Definition 6, condition 4). Since  $S_a \simeq S_b$ , we have:

$$\begin{cases} ord_a(v_a) = v_{a1} \dots v_{an} \text{ and} \\ ord_b(v_b) = ord_b(f(v_a)) = v_{b1} \dots v_{bn} \text{ with } v_{bi} = f(v_{ai}), \forall i \in \{1, \dots, n\} \end{cases}$$

Hence:

 $\begin{cases} ord_b(v_b) = v_{b1} \dots v_{bn} \text{ and} \\ ord_a(v_a) = ord_a(f^{-1}(v_b)) = v_{a1} \dots v_{an} = f^{-1}(v_{b1}) \dots f^{-1}(v_{bn}) \end{cases}$ 

Thus  $S_b \simeq S_a$  and  $f^{-1}$  is an ordered isomorphism between  $S_b$  and  $S_a$ .

3. The isomorphism between ordered structured object is transitive. Let  $S_a = (\hat{S}_a, ord_a) \in \mathscr{S}$ ,  $S_b = (\hat{S}_b, ord_b) \in \mathscr{S}$  and  $S_c = (\hat{S}_c, ord_c) \in \mathscr{S}$ three ordered structured object, respectively associated to  $G(S_a) = (V_a, E_a, \mu_a, \nu_a)$ ,  $G(S_b) = (V_b, E_b, \mu_b, \nu_b)$  and  $G(S_c) = (V_c, E_c, \mu_c, \nu_c)$ , such that  $S_a \simeq S_b$  and  $S_b \simeq S_c$ . We denote by f the ordered isomorphism between  $S_a$  and  $S_b$ , and by g the ordered isomorphism between  $S_b$  and  $S_c$ . As the isomorphism between structured objects is transitive we have  $g \circ f \in$   $\mathrm{Isom}(\hat{S}_a, \hat{S}_c)$  (Definition 6, condition 2). Let us consider a vertex  $v_a$  in  $V_{ord_a} \subset V_a$  with  $v_b = f(v_a) \in V_{ord_b} \subset V_b$  and  $v_c = g(v_b) = g \circ f(v_a) \in V_{ord_c} \subset V_c$ . We have since  $S_a \simeq S_b$  and  $S_b \simeq S_c$ :

$$\begin{cases} ord_a(v_a) = v_{a1} \dots v_{an} \text{ and} \\ ord_b(f(v_a)) = ord_b(v_b) = f(v_{a1}) \dots f(v_{an}) \underset{not.}{=} v_{b1} \dots v_{bn}, \\ ord_c(g(v_b)) = g(v_{b1}) \dots g(v_{bn}) \end{cases}$$

Therefore:

$$ord_c(g(v_b)) = ord_c(g \circ f(v_a)) = g \circ f(v_{a1}) \dots g \circ f(v_{an})$$

Thus  $S_a \simeq S_c$  and  $g \circ f$  is an ordered isomorphism between  $S_a$  and  $S_c$ .

In conclusion, the isomorphism between ordered structured objects is reflexive, symmetric and transitive. It is therefore, an equivalence relationship.  $\Box$ 

#### 2.2 Re-ordering function

A spatial configuration of atoms may be encoded by several equivalent orders. We thus introduce the notion of re-ordering function, which associates to each vertex of an ordered structured object a permutation on a subset of its neighborhood.

#### Definition 8. Re-ordering functions

Let us consider a set of ordered structured objects  $\mathscr{S}$ . A re-ordering function  $\sigma_S$  on an ordered structured object  $S = (\hat{S}, ord)$ , associated to a graph  $G(S) = (V, E, \mu, \nu)$ , associates to each vertex  $v \in V_{ord}$  a permutation  $\varphi_v$  on  $\{1, \ldots, |ord(v)|\}$ .

$$\sigma_S \begin{cases} V_{ord} \to \mathcal{P} \\ v \to \varphi_v \in \Pi_{|ord(v)|} \end{cases}$$

where  $\Pi_n$  is the group of permutations of n elements and  $\mathcal{P}$  is the union of  $\Pi_n$  for all  $n \in \mathbb{N}$ .

Application of a re-ordering function on an ordered structured object provides a new ordered structured object defined as follows:

#### Definition 9. Re-ordered structured objects

Let us consider a set of ordered structured objects  $\mathscr{S}$ . Let  $S = (\hat{S}, ord)$  denotes an ordered structured object,  $\sigma_S(S) = (\hat{S}, ord_{\sigma_S})$  is defined as the ordered structured object obtained after applying the re-ordering function  $\sigma_S$  on the order of the object:

$$\forall v \in V_{ord} \ s.t. \left( \begin{array}{c} ord(v) = v_1, \dots, v_n \\ and \\ \sigma_S(v) = \varphi_v, \end{array} \right) \ ord_{\sigma_S}(v) = v_{\varphi_v(1)}, \dots, v_{\varphi_v(n)}$$

Note that  $\hat{S} = \sigma_S(\tilde{S})$ . In other words, a re-ordering of an ordered structured object does not change the associated structured object. Re-ordering operations being defined as functions, these functions may be combined using composition operations:

#### Definition 10. Composition of re-ordering functions

Let us consider a set of ordered structured objects  $\mathscr{S}$ . Let  $\sigma_S$  and  $\sigma'_S$  denote two re-ordering functions on an ordered structured object  $S = (\hat{S}, ord)$ . The composition of  $\sigma_S$  and  $\sigma'_S$  is a re-ordering function denoted by  $\sigma_S \circ \sigma'_S$  and defined as follows:

$$\sigma_S \circ \sigma'_S \begin{pmatrix} V_{ord} \to \mathcal{P} \\ v \to \sigma_S(v) \circ \sigma'_S(v) \in \Pi_{|ord(v)|} \end{pmatrix}$$

where  $\Pi_n$  is the group of permutations of n elements and  $\mathcal{P}$  is the union of  $\Pi_n$  for all  $n \in \mathbb{N}$ .

The identity for the composition is the re-ordering function  $Id_S$  such that  $\forall v \in V_{ord}, Id_S(v) = Id_{|ord(v)|}$  where  $Id_n$  is the identity permutation on  $\Pi_n$ .

#### 2.3 Structured object having equivalent order

Re-ordering functions previously defined may apply any re-ordering on a structured object hence removing the notion of order on these objects. In order to obtain a useful notion of re-ordering, we have to define more precisely which properties should satisfies a valid family of re-ordering functions.

#### Definition 11. Valid re-orderings

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Let us consider a set of ordered structured objects  $\mathscr{S}$ . For each  $S = (\hat{S}, ord) \in \mathscr{S}$ , let us denote by  $\Sigma_S$  a set of re-ordering functions on S. A valid family of re-ordering functions is a set  $\Sigma = \{\Sigma_S, S \in \mathscr{S}\}$  which satisfies the two following properties :

- For any  $S \in \mathscr{S}, \Sigma_S$  is a group for the composition.
- For any two ordered structured objects  $S = (\hat{S}, ord)$  and  $S' = (\hat{S}', ord')$ whose associated un-order structured objects are isomorphic by a function f, any re-ordering function  $\sigma \in \Sigma_S$  is equal, up to the isomorphism  $f^{-1}$ , to a re-ordering function of  $\Sigma_{S'}$ .

$$\forall f \in \operatorname{Isom}(\hat{S}, \hat{S}'), \\ \forall \sigma \in \Sigma_S \ \right) \sigma \circ f^{-1} \in \Sigma_{S'}.$$

The first constraint of Definition 11 states that the set of re-ordering functions of an ordered structured objects may be combined freely using composition operations. The second constraint, involves that two ordered structured objects with isomorphic un-ordered structured objects should have, up to the isomorphism function, equivalent set of re-ordering functions. Note that this last constraint is equivalent to the following equation:

$$\forall f \in \operatorname{Isom}(\hat{S}, \hat{S}'), \\ \forall \sigma \in \varSigma_S \ ) \exists \sigma' \in \varSigma_{S'} \mid \sigma' \circ f = \sigma$$

**Proposition 2.** Let us consider a set of ordered structured objects  $\mathscr{S}$ , and a valid family of re-ordering functions  $\Sigma$ . The group of re-ordering functions  $\Sigma_{\sigma(S)}$  of any re-ordered structured object  $\sigma(S)$  is equal to the group  $\Sigma_S$  of re-ordering functions of S:

$$\begin{array}{l} \forall S \in \mathscr{S}, \\ \forall \sigma \in \varSigma_S, \end{array} \right) \varSigma_{\sigma(S)} = \varSigma_S$$

*Proof.* Let us consider  $\sigma \in \Sigma_S$ . Since S and  $\sigma(S)$  only differ by the order defined on each vertices, the identity function Id is a valid isomorphism between the unorder structured objects associated to  $\sigma(S)$  and S. Then, using Definition 11, for any  $\sigma' \in \Sigma_{\sigma(S)}$ , it exists a re-ordering function  $\sigma'' \in \Sigma_S$  such that:

$$\forall v \in V_{ord}, \ \sigma'(v) = \sigma''(Id(v)) = \sigma''(v)$$

We have thus  $\sigma' = \sigma''$  and thus  $\Sigma_{\sigma(S)} \subset \Sigma_S$ . The reverse inclusion is shown in the same way.

#### Definition 12. Equivalent orders

Let us consider a set of ordered structured objects  $\mathscr{S}$  and two of its ordered structured objects  $S_a = (\hat{S}_a, ord_a) \in \mathscr{S}$  and  $S_b = (\hat{S}_b, ord_b) \in \mathscr{S}$ . These structured objects are said to be equivalent  $S_a \simeq S_b$  according to a valid family of re-ordering functions  $\Sigma$  if:

$$\exists \sigma \in \Sigma_{S_a} \in \Sigma, \ \sigma(S_a) \simeq S_b \tag{1}$$

In other word, we consider that two ordered structured objects are equivalent if, up to a valid re-ordering  $\sigma$  we can establish an ordered structured object isomorphism f between them. In that case the ordered isomorphism f is called an equivalent ordered isomorphism through  $\sigma$  between  $S_a$  and  $S_b$  and we denote IsomEqOrd<sub> $\sigma$ </sub>( $S_a, S_b$ ) the set of equivalent ordered isomorphism through  $\sigma$  between  $S_a$  and  $S_b$ . We further denote by IsomEqOrd( $S_a, S_b$ ) the union of all IsomEqOrd<sub> $\sigma$ </sub>( $S_a, S_b$ ) for all  $\sigma \in \Sigma_{S_a}$ .

$$\operatorname{IsomEqOrd}(S_a, S_b) = \bigcup_{\sigma \in \Sigma_{S_a}} \operatorname{IsomEqOrd}_{\sigma}(S_a, S_b)$$

We will now prove that the equivalence order relationship is, as suggested by its name, an equivalence relationship.

**Proposition 3.** Let  $\mathscr{S}$  be a set of ordered structured objects and  $\Sigma$  denotes a valid family of re-ordering functions. The equivalent order relationship based on this family is reflexive.

$$\forall S \in \mathscr{S}, \ S \underset{\Sigma}{\simeq} S$$

*Proof.* Let  $\Sigma$  denotes a valid family of re-ordering functions and  $S = (\hat{S}, ord)$  an ordered structured object.

By Definition 11,  $\Sigma_S$  is a group and therefore  $Id_S \in \Sigma_S$ . We have by definition of  $Id_S$  (Definition 10):

$$\forall v \in V_{ord}, ord_{Id_S}(v) = ord(v).$$

We have by Definition 7,  $Id_S(S) \simeq S$  with  $Id_S \in \Sigma_S$ .

Thus  $S \simeq S$ .

**Lemma 1.** Let  $\mathscr{S}$  be a set of ordered structured objects and  $\Sigma$  a valid family of re-ordering functions. Let us consider two ordered structured objects  $S_a = (\hat{S}_a, ord_a) \in \mathscr{S}$  and  $S_b = (\hat{S}_b, ord_b) \in \mathscr{S}$  such that  $S_a \simeq S_b$ . Let  $\sigma_a \in \Sigma_{S_a}$  and  $\sigma_b \in \Sigma_{S_b}$  two re-ordering functions such that:

$$\forall v \in V_{ord_a}, \, \sigma_a(v) = \sigma_b(f(v)),$$

where f is an ordered isomorphism between  $S_a$  and  $S_b$ . Then we have  $\sigma_a(S_a) \simeq \sigma_b(S_b)$ .

*Proof.* Let us consider  $f \in \text{IsomOrd}(S_a, S_b)$ , and a vertex  $v \in V_{ord_a}$  with u = f(v). We have by Definition 7:

$$\forall v \in V_{ord_a} \begin{cases} ord_a(v) = v_1 \dots v_n \text{ and} \\ ord_b(u) = u_1 \dots u_n, \text{ with } u_i = f(v_i), \forall i \in \{1, \dots, n\} \end{cases}$$

Let us further denote by  $\varphi_v$  the permutation defined on vertex v both by  $\sigma_a$ and  $\sigma_b$ :  $\varphi_v = \sigma_a(v) = \sigma_b(f(v))$ .

Given the re-ordered structured objects  $\sigma_a(S_a) = (\hat{S}_a, ord_{\sigma_a})$  and  $\sigma_b(S_b) = (\hat{S}_b, ord_{\sigma_b})$  we have by Definition 9:

$$\begin{cases} ord_{\sigma_a}(v) = v_{\varphi_v(1)} \dots v_{\varphi_v(n)} \text{ and} \\ ord_{\sigma_b}(u) = u_{\varphi_v(1)} \dots u_{\varphi_v(n)}. \end{cases}$$

Since sequences  $v_1 \dots v_n$  and  $u_1 \dots u_n$  satisfy  $u_i = f(v_i)$  and since a same permutation  $\varphi_v$  is applied on both sequences we have:

$$\forall i \in \{1, \dots, n\} \ u_{\varphi_v(i)} = f(v_{\varphi_v(i)})$$

The isomorphism f maps thus the order encoded by  $\sigma_a(S_a)$  around each vertex of  $S_a$  onto the order defined by  $\sigma_b(S_b)$  on the corresponding vertex of  $S_b$ .

Moreover, since f is an isomorphism between ordered structured objects, it also corresponds to an isomorphism between un-ordered structured objects and we have by Definition 7,  $\sigma_a(S_a) \simeq \sigma_b(S_b)$ .

**Proposition 4.** Let  $\mathscr{S}$  be a set of ordered structured objects and  $\Sigma$  a valid family of re-ordering functions. The equivalent order relationships based on this family is symmetric:

$$\forall (S_a, S_b) \in \mathscr{S}^2, \ S_a \underset{\overline{\Sigma}}{\simeq} S_b \Leftrightarrow S_b \underset{\overline{\Sigma}}{\simeq} S_a.$$

*Proof.* Let us consider a set of ordered structured objects  $\mathscr{S}$ ,  $\Sigma$  a valid family of re-ordering functions and two ordered structured objects  $S_a = (\hat{S}_a, ord_a) \in \mathscr{S}$  and  $S_b = (\hat{S}_b, ord_b) \in \mathscr{S}$  such that  $S_a \simeq S_b$  We have by Definition 12:

$$\exists \sigma \in \Sigma_{S_a} s. t. \sigma(S_a) \simeq S_b.$$

As  $\Sigma_{S_a}$  is a group (Definition 11), it exists a re-ordering function  $\sigma^{-1} \in \Sigma_{S_a}$ such that  $\sigma^{-1}(\sigma(S_a)) = S_a$ .

Let us denote by f the ordered isomorphism between  $\sigma(S_a)$  and  $S_b$ . By Definition 11, since  $\sigma(S_a) \simeq S_b$ , the re-ordering function  $\sigma^{-1} \in \Sigma_{S_a}$  should be equivalent to some re-ordering function  $(\sigma^{-1})'$  in  $\Sigma_{S_b}$ . In other words:

$$\exists (\sigma^{-1})' \in \Sigma_{S_b} \text{ such that } \forall v \in V_{ord_a}, \sigma^{-1}(v) = (\sigma^{-1})'(f(v)).$$

We have thus by Lemma 1:  $\sigma^{-1}(\sigma(S_a)) \simeq (\sigma^{-1})'(S_b)$ . Therefore  $S_a \simeq (\sigma^{-1})'(S_b)$ , and by symmetry of the ordered isomorphism  $(\sigma^{-1})'(S_b) \simeq S_a$ . So by Definition 12,  $S_b \simeq S_a$ . Both cases being symmetric, the reverse implication is proved in the same way.

**Proposition 5.** Let  $\mathscr{S}$  a set of ordered structured objects and  $\Sigma$  a valid family of re-ordering functions. The equivalent order relationship based on this family is transitive:

$$\forall (S_a, S_b, S_c) \in \mathscr{S}^3, \ \left( \begin{matrix} S_a \underset{\Sigma}{\simeq} S_b \text{ and} \\ S_b \underset{\Sigma}{\simeq} S_c \end{matrix} \right) \Rightarrow S_a \underset{\Sigma}{\simeq} S_c$$

*Proof.* Let us consider a set of ordered structured objects  $\mathscr{S}$ ,  $\Sigma$  a valid family of re-ordering functions and three ordered structured objects  $S_a = (\hat{S}_a, ord_a) \in \mathscr{S}$ ,  $S_b = (\hat{S}_b, ord_b) \in \mathscr{S}$  and  $S_c = (\hat{S}_c, ord_c) \in \mathscr{S}$  such that  $S_a \underset{\Sigma}{\simeq} S_b$  and  $S_b \underset{\Sigma}{\simeq} S_c$ .

Using Definition 12, it exists two re-ordering functions  $\sigma_a \in \Sigma_{S_a}$  and  $\sigma'_b \in \Sigma_{S_b}$  such that  $\sigma_a(S_a) \simeq S_b$  and  $\sigma'_b(S_b) \simeq S_c$ .

Let us denote the isomorphism between un-ordered structured objects  $S_b$  and  $\widehat{\sigma_a(S_a)}$  by  $f_{ba}$ . Since  $S_b$  and  $\sigma(S_a)$  are isomorph and since  $\sigma'_b \in \Sigma_{S_b}$ , it must exists (by Definition 11) a re-ordering function  $\sigma'_a \in \Sigma_{\sigma_a(S_a)}$  such that:

$$\forall v \in V_{ord_b}, \ \sigma'_b(v) = \sigma'_a(f_{ba}(v)).$$

Then by Lemma 1, we have  $\sigma'_b(S_b) \simeq_o \sigma'_a(\sigma_a(S_a))$ . Since the ordered isomorphism is an equivalence relationship (Proposition 1) and  $\sigma'_b(S_b) \simeq_o S_c$ , we have  $\sigma'_a(\sigma_a(S_a)) \simeq_o S_c$ .

Since  $\Sigma_{S_a}$  is a group (Definition 11) and since  $\sigma_a \in \Sigma_{S_a}$  and  $\sigma'_a \in \Sigma_{\sigma_a(S_a)} = \Sigma_{S_a}$  (Proposition 2) we have:  $\sigma'_a \circ \sigma_a \in \Sigma_{S_a}$ . So by Definition 12,  $S_a \simeq S_c$ .

**Theorem 1.** Let  $\mathscr{S}$  be a set of ordered structured objects and  $\Sigma$  a valid family of re-ordering functions. The equivalent order relationship based on this family is an equivalence relationship.

*Proof.* The equivalent order relationship is reflexive (Proposition 3), symmetric (Proposition 4) and transitive (Proposition 5).  $\Box$ 

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#### 2.4 Ordered graphs

We now apply the definition of ordered structured objects to labeled graphs, in order to define ordered graphs.

#### Definition 13. Set of Ordered Graphs

An ordered graph  $S = (G = (V, E, \mu, \nu), ord)$  is an ordered structured object with G(S) = G and a function ord which maps each vertex of  $V_{ord} \subset V$  to an ordered list of its neighbors:

$$ord \begin{cases} V_{ord} \to V^* \\ v \to v_1 \dots v_n \end{cases}$$

where  $V(v) = \{v_1, \ldots, v_n\}$  denotes the neighborhood of v.

The set of ordered graph is denoted  $\mathcal{OG}$ . The set of isomorphism defined between two un-ordered graphs (Definition 6) is the usual set of isomorphism between labeled graphs.

As the set of ordered graphs is a set of structured objects, we can define, for ordered graphs, re-ordering functions (Definition 8). The definition of orders and re-ordering functions depends on the application at end. Nevertheless, if re-orderings fulfill the definition of a valid family of re-ordering functions (Definition 11) we can define the equivalence relationship between ordered graphs defined by Definition 12 (Theorem 1).

Let us now define a chiral vertex, which encodes a stereocenter when ordered graphs represents molecules.

#### Definition 14. Chiral vertices

Let  $\Sigma$  be a valid family of re-ordering functions on  $\mathcal{OG}$ . Let  $G = (V, E, \mu, \nu, ord)$  be an ordered graph. A vertex  $v \in V_{ord} \subset V$  of degree n is called a chiral vertex iff:

$$\forall (i,j) \in \{1,\ldots,n\}^2$$
 with  $i \neq j, \nexists f \in \text{IsomEqOrd}(G,\tau_{i,j}(G))$  with  $f(v) = v$ .

where  $\tau_{i,j}$  is a re-ordering function equals to the identity on all vertices except v for which it permutes the vertices of index i and j in ord(v).

In other words, a vertex is chiral if any permutation of its neighbors produces an ordered graph with a non-equivalent order.

## 2.5 Ordered graphs and re-ordering functions encoding of a molecule

We now restrict our attention on molecular graphs (Definition 1) and let us define from them molecular ordered graphs.

The molecular ordered graph of a molecule is defined by first defining its molecular graph (Definition 1)  $G = (V, E, \mu, \nu)$  which encodes relationships between atoms together with the type of atom and bond respectively associated to each vertex and each edge.

#### Definition 15. Molecular ordered graph

A molecular ordered graph is a couple S = (G, ord) where G corresponds to a molecular graph. The function ord is defined on a set  $V_{ord}$  defined as:

$$V_{ord} = V_{PAC} \cup V_{DB}$$

where  $V_{PAC}$  and  $V_{DB}$  denote respectively the set of potential asymmetric carbons (Definition 2) and the set of carbons of degree 3 connected by a double bound (Definition 3).

The function ord is defined as follows for each vertex  $v \in V_{ord}$ :

- If  $v \in V_{PAC}$ :

We set randomly one of its neighbor  $v_1$  at the first position. The three other neighbors of v are ordered such that if we look at v from  $v_1$ , the three remaining neighbors are ordered clockwise (Section 1). One of the three orders (defined up to circular permutations) fulfilling this condition is chosen randomly (Figure 3(a)).

- If  $v \in V_{DB}$ :

Let us consider  $w = n_{=}(v)$  and the two neighborhoods  $V(v) = \{w, a, b\}$  and  $V(w) = \{v, c, d\}$ . The order on the neighborhood of v is set as ord(v) = w, a, b and the order on w's neighborhood is set as ord(w) = v, c, d, whereby a, b, c, d are traversed clockwise when turning around the double bond for a given plane embedding (Figure 3(b)).

We denotes  $\mathcal{OM}$  the set of ordered molecular graphs.

#### Definition 16. Set of molecular re-ordering functions

We define for each molecular ordered graph S, a set of re-ordering function  $\Sigma_S^M$ .  $\Sigma_S^M$  contains all the re-ordering functions  $\sigma$  such that:

- For each v in  $V_{PAC} \sigma(v)$  is an even permutation:

$$\forall v \in V_{PAC}, \ \epsilon(\sigma(v)) = 1.$$

- For each v in  $V_{DB}$ ,  $\sigma(v)$  and  $\sigma(n_{=}(v))$  have the same parity:

 $\forall v \in V_{DB}, \ \epsilon(\sigma(v)) = \epsilon(\sigma(w)) \text{ with } w = n_{=}(v).$ 

where  $\epsilon$  denotes the signature of a permutation.

**Proposition 6.** For any molecular graph S,  $\Sigma_S^M$  is a group for the composition.

*Proof.* We have to show that  $\Sigma_S^M$  admits an identity element, is closed under composition and admits for each re-ordering function an inverse element.

1.  $\Sigma_S^M$  admits an identity element  $Id_S$ .

Let us consider the re-ordering function  $Id_S$  such that:

$$\forall v \in V_{ord}, Id_S(v) = Id_{|V(v)|}$$

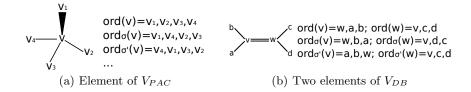


Fig. 3. Example of elements of  $V_{PAC}$  and of  $V_{DB}$  with their ordered list (top) and the ordered lists obtained using two permutations  $\sigma \in \Sigma_S^M$  and  $\sigma' \in \Sigma_S^M$ 

where  $Id_n$  is the identity permutation on  $\Pi_n$ . Since the identity is an even permutation we have:

$$\begin{cases} \forall v \in V_{PAC} \ \epsilon(Id_S(v)) = 1\\ \forall v \in V_{DB} \ \epsilon(Id_S(v)) = \epsilon(Id_S(n_{=}(v)) = 1) \end{cases}$$

Thus by definition of  $\Sigma_S^M$ ,  $Id_S \in \Sigma_S^M$ . 2.  $\forall (\sigma, \sigma') \in (\Sigma_S^M)^2$ ,  $\sigma \circ \sigma' \in \Sigma_S^M$ . Let  $\sigma$  and  $\sigma'$  denote two re-ordering functions of  $\Sigma_S^M$ .

- If  $v \in V_{PAC}$ :
  - Since  $\epsilon$  is a morphism between  $\Pi_{[ord(v)]}$  and  $(\{-1,1\},\times)$  we have:

$$\forall v \in V_{PAC}, \ \epsilon(\sigma(v) \circ \sigma'(v)) = \epsilon(\sigma(v))\epsilon(\sigma'(v)) = 1.1 = 1$$

- If  $v \in V_{DB}$ :

Let us consider  $w = n_{=}(v)$ . Since  $\sigma(v)$  and  $\sigma(w)$  on one hand and  $\sigma'(v)$ and  $\sigma'(w)$  on the other hand have a same signature, we have:

$$\forall v \in V_{DB}, \, \epsilon(\sigma(v) \circ \sigma'(v)) = \epsilon(\sigma(v))\epsilon(\sigma'(v)) \\ = \epsilon(\sigma(w))\epsilon(\sigma'(w)) \\ = \epsilon(\sigma(w) \circ \sigma'(w))$$

Permutations  $\sigma(v) \circ \sigma'(v)$  and  $\sigma(w) \circ \sigma'(w)$  have thus a same parity.

Thus by definition of  $\Sigma_S^M$ ,  $\sigma \circ \sigma' \in \Sigma_S^M$ . 3.  $\forall \sigma \in \Sigma_S^M$ ,  $\exists \sigma^{-1} \in \Sigma_S^M$  such that  $\sigma \circ \sigma^{-1} = Id_S$ , where  $Id_S$  is the identity element of  $\Sigma_S^M$ .

Let us consider  $\sigma^{-1}$  such that:

$$\forall v \in V_{ord}, \ \sigma^{-1}(v) = (\sigma(v))^{-1}.$$

We have by Definition 10,  $\sigma \circ \sigma^{-1} = Id_S$ . We have to prove that, for each  $\sigma \in \Sigma_S^{\widetilde{M}}$ ,  $\sigma^{-1} \in \Sigma_S^M$ .

- If  $v \in V_{PAC}$ :

$$\epsilon(\sigma^{-1}(v)) = \epsilon(\sigma(v)^{-1}) = \epsilon(\sigma(v)) = 1$$

Thus  $\sigma^{-1}(v)$  is even.

- If  $v \in V_{DB}$ :

Let us consider  $w = n_{=}(v)$ . Since  $\sigma(v)$  and  $\sigma(w)$  have a same parity and:

$$\begin{cases} \epsilon(\sigma^{-1}(v)) = \epsilon(\sigma(v)) \\ \epsilon(\sigma^{-1}(w)) = \epsilon(\sigma(w)), \end{cases}$$

 $\sigma^{-1}(v)$  and  $\sigma^{-1}(w)$  have also a same parity. Thus by definition of  $\Sigma_S^M$ ,  $\sigma^{-1} \in \Sigma_S^M$ .

As the composition of functions is associative and  $\Sigma_S^M$  admits an identity element for the composition, is closed under composition, and admits for each re-ordering function an inverse element we can conclude that  $\Sigma_S^M$  is a group for the composition.

**Proposition 7.** For any two molecular ordered graphs S and S' whose associated un-order graphs are isomorphic by a function f, and for any re-ordering function  $\sigma \in \Sigma_S^M$ , we have  $\sigma \circ f^{-1} \in \Sigma_{S'}^M$ :

$$\begin{array}{l} \forall f \in \operatorname{Isom}(\hat{S}, \hat{S}'), \\ \forall \sigma \in \varSigma_{S}^{M} \end{array} \right) \sigma' = \sigma \circ f^{-1} \in \varSigma_{S'}^{M}.$$

*Proof.* Let us consider two molecular ordered graphs S and S' whose associated un-order graphs G and G' are isomorphic by a function f. We denote by  $f^{-1}$ the isomorphism between G' and G and define the re-ordering function  $\sigma'$  as  $\sigma \circ f^{-1}$ .

Let us prove that  $\sigma' \in \Sigma_{S'}^M$ .

- If  $v' \in V'_{PAC}$ : We have by definition of  $\sigma'$ ,  $\sigma'(v') = \sigma(v)$  with  $v = f^{-1}(v')$ . As  $v' \in V'_{PAC}$  we have ,  $\mu(v') = C'$  and |V(v')| = 4. Since  $f^{-1}$  is an isormorphism between G' and G we have:

$$\begin{cases} \mu(v) = \mu(f^{-1}(v')) = \mu(v') = {}^{?}C' \\ |V(v)| = |V(f^{-1}(v'))| = |V(v')| = 4 \end{cases}$$

Thus by definition of  $V_{PAC}$ ,  $v \in V_{PAC}$ . Moreover, since  $\sigma'(v') = \sigma(v)$ ,  $\sigma(v)$ has the same even parity than  $\sigma'(v')$ .

If  $v' \in V'_{DB}$ : Given  $w' = n_{=}(v')$ , we have by definition of  $\sigma'$ :

$$\begin{cases} \sigma'(v') = \sigma(v) & \text{with } v = f^{-1}(v') \\ \sigma'(w') = \sigma(w) & \text{with } w = f^{-1}(w') \end{cases}$$

As v' and  $w' \in V'_{DB}$  we have  $\mu(v') = \mu(w') = C'$  and |V(v')| = |V(w')| = 3. Since  $f^{-1}$  is an isormorphism between G' and G we have:

$$\begin{cases} \mu(v) = \mu(f^{-1}(v')) = \mu(v') = C' \text{ and} \\ \mu(w) = \mu(f^{-1}(w')) = \mu(w') = C' \\ |V(v)| = |V(f^{-1}(v'))| = |V(v')| = 3 \text{ and} \\ |V(w)| = |V(f^{-1}(w'))| = |V(w')| = 3 \end{cases}$$

As  $v' \in V'_{DB}$ , it exists an edge e'(v', w') connecting v' and w' with a label  $\nu(e') = 2$ . Such an edge is preserved by the isomorphism  $f^{-1}$  between G' and G and it thus exists an edge e(v, w) in G between  $v = f^{-1}(v')$  and  $w = f^{-1}(w')$  with  $\nu(e) = 2$ .

We have thus by definition of  $V_{DB}$ ,  $\{v, w\} \subset V_{DB}$  and since  $\sigma \in \Sigma_S^M$  we have by definition of  $\Sigma_S^M$ :

$$\epsilon(\sigma(v)) = \epsilon(\sigma(w))$$

Therefore, by definition of  $\sigma'$ :

$$\epsilon(\sigma'(v')) = \epsilon(\sigma(v)) = \epsilon(\sigma(w)) = \epsilon(\sigma'(w'))$$

Permutations  $\sigma'(v')$  and  $\sigma'(w')$  have thus the same parity.

Thus  $\sigma' = \sigma \circ f^{-1} \in \Sigma_{G'}^M$ .

**Theorem 2.** The set of molecular re-ordering functions  $\Sigma^M = \{\Sigma_S^M, S \in \mathcal{OM}\}\$  is a valid family of re-ordering functions. Therefore the equivalent order relationship based on this family is an equivalence relationship.

*Proof.* The set of molecular re-ordering functions  $\Sigma^M$  is a valid family of reordering functions by Proposition 6 and 7. Thus by Theorem 1, the equivalent order relationship based on this family is an equivalence relationship.

 $\square$ 

**Remark 1.** Let S = (G, ord) with  $G = (V, E, \mu, \nu)$  denote a molecular ordered graph. We have, by construction of re ordering functions, the following property:

If we select for each vertex  $v \in V_{ord}$  a neighbor  $n_v$ , we can always find a re-ordering function  $\sigma$  of  $\Sigma_S^M$  such that the ordered list of each vertex v of  $V_{ord}$  in  $\sigma(S)$  starts by its selected neighbor  $n_v$ .

Given our encoding of the relative positioning of atoms by orders defined in this section, we encode the spatial configuration of atoms within the neighborhood of each of its vertex. Our equivalence relationship between molecular ordered graphs allows to check if two molecules have a same spatial configuration.

Stereocenters are defined as chiral vertice (Definition 14). Indeed, a vertex is chiral if any permutation of its neighbors produces an ordered graph with a nonequivalent order, called a different stereoisomer within the chemistry framework.

#### 3 Isomorphism between labeled partially ordered tree

Let us now restrict our attention to acyclic graphs in order to obtain an efficient way to determine if two molecular graphs have equivalent orders.

Given a rooted tree, the father of each node v is denoted by  $p_v$ . The tree itself is denoted by  $\hat{T} = (r, G)$  where r denotes the root of the tree and  $G = (V, E, \mu, \nu)$ the acyclic graph associated to  $\hat{T}$ .

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#### Definition 17. Ordered rooted tree

An ordered rooted tree  $T = (\hat{T}, ord)$  with  $\hat{T} = (r, G)$  is an ordered structured object. Its associated acyclic labeled graph is  $G = (V, E, \mu, \nu)$ . The function ord maps each internal vertex to an ordered list of its children:

$$ord \begin{cases} V_{ord} \to V^* \\ v \mapsto v_1 \dots v_n \end{cases}$$

where  $\{v_1, \ldots, v_n\}$  denotes the children of v.

We denote by  $\mathcal{OT}$  the set of ordered trees.

Note that the function ord is defined on all internal vertices of the tree we have thus:

$$\forall T = (\hat{T}, ord) \in \mathcal{OT}, \text{ with } \begin{pmatrix} \hat{T} = (r, G), \\ G = (V, E, \mu, \nu) \end{pmatrix} V_{ord} = V - Leaf(T)$$

where Leaf(T) denotes the set of leaves of T.

Moreover, the order relationship of each vertex of an ordered rooted tree is defined on all its children, i.e. all its neighbors but its parent. We have thus:

$$\forall T = (\hat{T}, ord) \in \mathcal{OT}, \text{ with } \hat{T} = (r, G) \begin{cases} |ord(r)| = |V(v)| \\ \forall v \in V_{ord}, v \neq r \ |ord(v)| = |V(v)| - 1 \end{cases}$$

An isomorphism between rooted trees may be considered as an isomorphism between graphs which maps the roots of both trees one on the other. Hence the set of ismorphisms  $\text{Isom}(\hat{T}, \hat{T}')$  between two rooted tree may be considered as a subset of the isomorphisms between the associated acyclic graphs:  $\text{Isom}(\hat{G}, \hat{G}')$ (Definition 6).

Given these isomorphisms we define for  $\mathcal{OT}$  ordered isomorphisms between ordered rooted trees (Definition 7). Such isomorphisms preserve both the structure of both trees and their orderings. The order defined on each vertex of the trees belonging to  $\mathcal{OT}$  depends of the considered application. The valid family of re-ordering functions  $\Sigma$  (Definition 11), which may be defined on  $\mathcal{OT}$  also depends of this application. Given both orders and a valid family of re-ordering functions we can build an equivalence relationship (Definition 12 and Theorem 1) between ordered trees encoding the fact that up to re-orderings two rooted trees are structurally similar and have a same order.

Following [10], we associate to each ordered rooted tree T, an unique depthfirst string encoding DFSE(T). This string is based on the sequence of node and edge labels obtained by traversing the tree in a depth-first order and uses respectively \$ and # to represent backtracks and the end of the string encoding.

#### Definition 18. Depth-First String Encoding

The depth first string encoding of an ordered rooted tree  $T = (\hat{T}, ord)$  denoted by DFSE(T) is defined as the sequence of node and edge labels encountered when traversing T using a depth-first method based on the order defined by function ord. Each backtrack during this traversal is encoded by the symbol '\$' while the end of the string is encoded by the symbol '#'.

**Remark 2.** As shown by [10](Lemma 2.2), two isomorphic ordered trees have the same depth-first string encoding and conversely.

$$T_1 \simeq T_2 \Leftrightarrow \text{DFSE}(T_1) = \text{DFSE}(T_2)$$

Definition 19. Depth-first canonical string and Depth-first canonical form of a tree

Let us consider an ordered tree T and a valid family of re-ordering functions  $\Sigma$ :

- The depth-first canonical string  $DFCS_{\Sigma}(T)$  of T is the minimal depth-first string encoding among all possible ordered trees  $\sigma(T)$  obtained by applying  $\sigma \in \Sigma_T \in \Sigma$  on T:

$$DFCS_{\Sigma}(T) = \min_{\sigma \in \Sigma_T} DFSE(\sigma(T))$$

- The depth-first canonical form  $\text{DFCF}_{\Sigma}(T)$  of T according to  $\Sigma$  is the ordered tree  $\sigma(T)$  whose depth-first string encoding is minimal (and thus equals to  $\text{DFCS}_{\Sigma}(T)$ ):

$$DFSE(DFCF_{\Sigma}(T)) = DFCS_{\Sigma}(T)$$

The depth-first canonical form is unique up to ordered isomorphism (Remark 2).

**Proposition 8.** Two ordered rooted trees  $T_a = (\hat{T}_a, ord_a)$  and  $T_b = (\hat{T}_b, ord_b)$  have equivalent orders according to a valid family of re-ordering functions  $\Sigma$  iff their depth-first canonical string according to  $\Sigma$  are equals:

$$T_a \simeq T_b \Leftrightarrow \mathrm{DFCS}_{\Sigma}(T_a) = \mathrm{DFCS}_{\Sigma}(T_b)$$

*Proof.* Let us first prove that  $DFCS_{\Sigma}(T_a) = DFCS_{\Sigma}(T_b) \Rightarrow T_a \simeq T_b$  and then the reverse implication.

1. We suppose that  $DFCS_{\Sigma}(T_a) = DFCS_{\Sigma}(T_b)$ . Let us denote  $\sigma_a \in \Sigma_{T_a}$  the re-ordering function such that  $\sigma_a(T_a) = DFCF_{\Sigma}(T_a)$ and  $\sigma'_b \in \Sigma_{T_b}$  the one such that  $\sigma'_b(T_b) = DFCF_{\Sigma}(T_b)$ . By definition  $DFCS_{\Sigma}(T_a) = DFSE(\sigma_a(T_a))$  and  $DFCS_{\Sigma}(T_b) = DFSE(\sigma'_b(T_b))$ . Since  $DFCS_{\Sigma}(T_a) = DFCS_{\Sigma}(T_b)$  we have  $DFSE(\sigma_a(T_a)) = DFSE(\sigma'_b(T_b))$ , and thus  $\sigma_a(T_a) \simeq \sigma'_b(T_b)$  (Remark 2).

As  $\Sigma_{T_a}$  is a group (Definition 11), it exists a re-ordering function  $\sigma_a^{-1} \in \Sigma_{T_a}$ such that  $\sigma_a^{-1}(\sigma_a(T_a)) = T_a$ .

Let us denote by f the ordered isomorphism between  $\sigma_a(T_a)$  and  $\sigma'_b(T_b)$ . By Definition 11, since  $\sigma_a(T_a) \simeq \sigma'_b(T_b)$ , the re-ordering function  $\sigma_a^{-1} \in \Sigma_{T_a}$ should be equivalent to some re-ordering function  $\sigma_b^{-1}$  in  $\Sigma_{T_b}$ . In other words:

$$\exists \sigma_b^{-1} \in \Sigma_{T_b} \text{ such that } \forall v \in V_a, \sigma_a^{-1}(v) = \sigma_b^{-1}(f(v)).$$

We have thus by Lemma 1:  $\sigma_a^{-1}(\sigma_a(T_a)) \simeq \sigma_b^{-1}(\sigma_b'(T_b))$ . Therefore  $T_a \simeq \sigma_b^{-1}(\sigma_b'(T_b))$ . As  $\Sigma_{T_b}$  is a group,  $\sigma_b^{-1} \circ \sigma_b' \in \Sigma_{T_b}$ , so  $T_a \simeq T_b$ .

2. We suppose that  $T_a \simeq T_b$ .

We denote  $\sigma'_b \in \Sigma_{T_b}$ , the re-ordering function such that  $\sigma'_b(T_b) = \text{DFCF}_{\Sigma}(T_b)$ . As  $T_a \simeq T_b$ ,  $\exists \sigma \in \Sigma_{T_a}$  such that  $\sigma(T_a) \simeq T_b$ . Let us denote by f the ordered isomorphism between  $T_b$  and  $\sigma(T_a)$ .

isomorphism between  $T_b$  and  $\sigma(T_a)$ . By Definition 11, since  $T_b \simeq \sigma(T_a)$ , the re-ordering function  $\sigma'_b \in \Sigma_{T_b}$  should be equivalent to some re-ordering function  $\sigma'_a$  in  $\Sigma_{T_a}$ . In other words:

$$\exists \sigma'_a \in \Sigma_{T_a} \text{ such that } \forall v \in V_b, \sigma'_b(v) = \sigma'_a(f(v)).$$

We have thus by Lemma 1:  $\sigma'_a(\sigma(T_a)) \simeq \sigma'_b(T_b)$ . Therefore

$$DFSE(\sigma'_a(\sigma(T_a))) = DFSE(\sigma'_b(T_b)) = DFCS_{\Sigma}(T_b)$$

Since  $\Sigma_{T_a}$  is a group, we have  $\sigma'_a \circ \sigma \in \Sigma_{T_a}$ . Therefore:

$$\operatorname{DFCS}_{\Sigma}(T_b) = \operatorname{DFSE}(\sigma'_a(\sigma(T_a))) \ge \operatorname{DFCS}_{\Sigma}(T_a).$$

Both cases being symmetric the reverse inequality is shown by considering  $\sigma'_a \in \Sigma_{T_a}$  such that  $\sigma'_a(T_a) = DFCF_{\Sigma}(T_a)$ . Therefore:

$$\mathrm{DFCS}_{\Sigma}(T_a) = \mathrm{DFCS}_{\Sigma}(T_b).$$

By 1 and 2 we have proven that

$$T_a \simeq T_b \Leftrightarrow \mathrm{DFCS}_{\Sigma}(T_a) = \mathrm{DFCS}_{\Sigma}(T_b).$$

Let  $\Sigma$  be a valid family of re-ordering functions. An ordered tree T can have two vertices connected to a same parent and whose associated subtrees are equivalent according to  $\Sigma$ . Any path from a leaf of T passing through one of these two vertices is equivalent to another path passing through the other vertex. From a more global point of view, a permutation exchanging these two subtrees on the depth-first canonical form of T would lead to an isomorphic ordered tree. We thus consider that these two vertices are equivalent.

#### Definition 20. Equivalent ordered sub-tree

Let us consider an ordered rooted tree T and a valid family of re-ordering functions  $\Sigma$ .

– Two child of a same parent whose associated rooted sub-trees are equivalent according to  $\Sigma$  are said to be equivalent:

$$v_i \sim v_j \Leftrightarrow \exists (v, \sigma) \in V \times \Sigma_T \, s.t. \begin{cases} p_{v_i} = p_{v_j} = v \text{ and} \\ (\sigma(v))(i) = j \text{ and } \mathrm{DFCF}_{\Sigma}(\sigma(T)) \simeq \\ 0 \end{cases} \text{DFCF}_{\Sigma}(T)$$
(2)

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- The representative of each class is defined as the vertex with the minimal index within the ordered list of children of its parent:

$$\forall i \in \{1, \dots, n\} rep(v_i) = min\{j \mid v_j \sim v_i\}.$$
(3)

The representative of a class is properly defined since two equivalent nodes must have a same parent.

# 3.1 Ordered trees and re-ordering functions encoding of an acyclic molecule

To define a molecular ordered tree  $T = (\hat{T}, ord_T)$ , from an acyclic molecular ordered graph  $G = (\hat{G}, ord_G)$ , we have to define a root and for each vertex an order on its children. By definition, the function  $ord_T$  of T is defined on (Definition 17):

$$V_{ord}^T = V - Leaf(T)$$

where Leaf(T) denotes the set of leaves of T.

On the other end, the set of vertices of an ordered molecular graph  $G = (\hat{G}, ord_G)$  on which the function  $ord_G$  is defined (Definition 15) is equal to:

$$V_{ord}^M = V_{PAC} \cup V_{DB}$$

Since a molecular ordered graph does not provide an order for all vertices, while all vertices of an ordered tree but its leaves are ordered, the definition of an order on a molecular tree imposes to fix a priori the order on the child of some vertices if this order is not provided by the molecular ordered graph.

#### Definition 21. Molecular ordered tree

A molecular ordered tree  $T = (\hat{T}, ord_T)$  with  $\hat{T} = (r, G_T)$  is defined from a molecular ordered graph  $G = (\hat{G}, ord_G)$  with  $\hat{G} = (V, E, \mu, \nu)$  by setting  $G_T = \hat{G}$ ,  $r \in V$  and by defining  $ord_T$  from  $ord_G$  as follows:

Given the root r, let us consider a re-ordering function of  $\Sigma_G^M$  such that (Remark 1):

$$\forall v \in V_{ord}^M - \{r\}, \ ord_{\sigma(G)}(v) = p_v.v_1....v_n$$

If  $V \neq \{r\}$ , the function  $ord_T$  on r is set equals to:

$$ord_T(r) = \begin{cases} ord_{\sigma(G)}(r) & \text{if } r \in V_{ord}^M \\ random(child(r)) & \text{otherwise} \end{cases}$$

where random(child(r)) denotes a random ordering of the child of r (child(r)). For all other vertices:

$$- \text{ If } v \in V_{ord}^T - V_{ord}^M - \{r\}$$

$$ord_T(v) = random(child(v))$$

 $- \text{ If } v \in V_{ord}^T \cap V_{ord}^M - \{r\}$ 

We have  $ord_{\sigma(G)}(v) = p_v . v_1 ... . v_n$  where  $v_1 ... . v_n$  corresponds to an ordering of child(v). We set thus  $ord_T(v)$  as:

$$ord_T(v) = v_1 \dots v_n$$

#### Definition 22. Set of molecular re-ordering functions for tree

The set of re-ordering functions  $\Sigma_T^M$  is defined by :

- If  $v \in V_{ord}^T V_{ord}^M$ , Permutation  $\sigma(v)$  can be any permutation. Therefore the order on those vertex have no influence, it only allows us to determine the depth-first string encoding and the depth-first canonical string of a molecular ordered tree.
- If  $v \in V_{PAC}$ ,  $\sigma(v)$  is an even permutation:

$$\epsilon(\sigma(v)) = 1$$

- If  $v \in V_{DB}$ ,

Permutations  $\sigma(v)$  and  $\sigma(n_{=}(v))$  have a same parity :

$$\epsilon(\sigma(v)) = \epsilon(\sigma(w))$$
 with  $w = n_{=}(v)$ 

Given a unique code associated to an ordered rooted tree, the chirality of a vertex may be efficiently tested if one can transpose Definition 14 to ordered rooted trees:

**Proposition 9.** Let  $T = (\hat{T}, ord_T)$  with  $\hat{T} = (r, G)$  be an ordered rooted tree encoding an acyclic molecule, and  $\Sigma_T^M$  the set of molecular re-ordering functions for T. r is a chiral vertex if:

$$\forall (i,j) \in \{1,\ldots,|V(r)|\}^2 \text{ with } i \neq j, T \not\in \tau_{i,j}(T)$$

where  $\tau_{i,j}$  is a re-ordering function equals to the identity on any vertex but r where it permutes children of index i and j in the ordered list of r.

*Proof.* Using acyclic molecular graphs, an equivalent ordered isomorphism between ordered rooted trees corresponds to an equivalent ordered isomorphism between ordered graphs with an additional constraint on the mapping of both roots. If we can find an isomorphism between T and  $\tau_{i,j}(T)$  such an isomorphism f satisfies f(r) = r and also corresponds to an isomorphism between ordered graphs. Conditions of Definition 14 are thus violated and r is not chiral. The reverse implication may be demonstrated using the same type of reasoning.

#### 4 From a global to a local characterization of chirality

Proposition 9 allows us to determine if a vertex induces a chiral property for a molecule. Such a proposition concerning the whole molecule induces a global characterization of chirality. However, such a proposition does not allow to characterize the minimal subgraph of a molecule which induces the chiral property of a vertex. Using acyclic graphs, such a minimal subgraph corresponds to the smallest ordered sub-tree, rooted on a chiral vertex v which allows to characterize the chirality of v using Proposition 9.

#### 4.1 Minimal chiral subtree of an asymmetric carbon

Let v be a chiral vertex representing an asymmetric carbon  $(v \in V_{PAC})$ . We denote its neighbors  $v_1, \ldots, v_4$ . We consider the ordered tree T rooted on v and described in Section 3 and the family of re-ordering functions for molecular tree  $\Sigma_T^M$ . We note  $T_1, \ldots, T_4$  the subtrees of T rooted on the children of v. For any  $i \in \{1, 2, 3, 4\}$  we denote  $T_i^j$  the subtree of  $T_i$  composed of all nodes with a depth lower than j. According to Proposition 9, the chirality of v may be characterized from its subtrees  $T_i^j$  iff all pairs of subtrees are not equivalent. Indeed, in such a case no transposition of two subtrees  $T_i^j$  and  $T_k^{j'}$  can induce a rooted tree with equivalent order. Therefore for each  $i \in \{1, 2, 3, 4\}$ , we define the minimal subtree associated to  $v_i$  as  $T_i^{j^*(i)}$  with:

$$j^*(i) = \min\{j \mid \forall k \in \{1, \dots, 4\} - \{i\}, \ T_i^j \not\simeq T_k^j\}$$

For example in Figure 4, the root of  $T_1$  is a Chloro atom while the root of each other  $T_i$  is an oxygen atom, thus the subtree  $T_1^1$  reduced to the Chloro atom has a sufficient depth to be distinguished from all other subtrees  $T_i^k$ ,  $i \neq 1$  and we have  $j^*(1) = 1$ . The minimal chiral subtree of v is the subtree of T rooted on v, where v has for children  $T_1^{j^*(1)}, \ldots, T_4^{j^*(4)}$ . The asymmetric carbon is then represented by the depth-first canonical string of this tree according to  $\Sigma_T^M$ .

To find  $j^*(i)$ , we increase j for each  $T_i^j$  until  $T_i^j \not\simeq T_k^j$  for each  $k \in \{1, \ldots, 4\}$ ,

 $k \neq i$ . At each iteration we compute  $\operatorname{DFCS}_{\Sigma^M}(T_i^j)$  for each  $i \in \{1, \ldots, 4\}$ . Therefore the calculus of the minimal chiral subtree of v is performed in  $\mathcal{O}((\max_i |T_i^{j^*(i)}|)^2)$  which is bounded by  $\mathcal{O}(|V|^2)$ .

#### 4.2 Minimal chiral subtree of double bond

Let  $v_a$  be one carbon of a double bond,  $v_a \in V_{DB}$ . We denote  $n_=(v_a) = v_b$  and  $e = (v_a, v_b)$  the double bond between them. Let us denote by  $v_a^1$  and  $v_a^2$  the two remaining neighbors of  $v_a$ . Considering the ordered tree T rooted on  $v_a$ ,  $v_a$  is chiral only if the subtrees rooted on the children of  $v_a$  do not have equivalent orders (Proposition 9). This implies that the two subtrees rooted on  $v_a^1$  and  $v_a^2$  do not have equivalent orders. This last necessary condition is however not

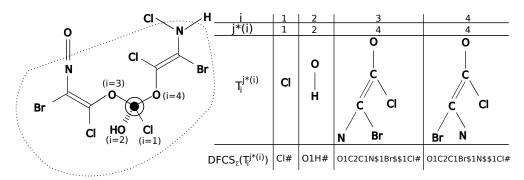


Fig. 4. Left: An asymmetric carbon () with its minimal chiral subtree (surrounded by a dotted line). Right: minimal subtrees rooted on its children.

sufficient. Indeed if the subtrees rooted on the remaining neighbors  $v_b^1$  and  $v_b^2$  of  $v_b$  have equivalent orders, then one can apply a re-ordering function  $\sigma \in \Sigma_T^M$ on T which simultaneously permutes the subtrees rooted on  $v_a^1$  and  $v_a^2$  and the subtrees rooted on  $v_b^1$  and  $v_b^2$  (by definition of  $V_{DB}$  and  $\Sigma^M$ ). The resulting rooted tree  $\sigma(T)$  has an equivalent order to T (Definition 12) but also to  $\tau(T)$ , where  $\tau$  permutes only vertices  $v_a^1$  and  $v_a^2$  in the ordered list of children of  $v_a$ . In such a case,  $v_a$  is non chiral (Proposition 9). Therefore, the non chirality of  $v_b$  induces the non chirality of  $v_a$  and conversely.

Hence  $v_a$  and  $v_b$  are chirals, only if the two following conditions are satisfied:

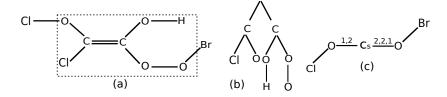
- subtrees rooted on  $v_a^1$  and  $v_a^2$  do not have equivalent orders and subtrees rooted on  $v_b^1$  and  $v_b^2$  also do not have equivalent orders.

In order to encode this constraint, we define as in Section 4.1 the minimal subtrees rooted on  $v_a^1$   $(T_a^1)$  and  $v_a^2$   $(T_a^2)$  with non-equivalent orders together with the minimal subtrees rooted on  $v_b^1$   $(T_b^1)$  and  $v_b^2(T_b^2)$  with non-equivalent orders. We denote by  $T_a$  and  $T_b$  the two ordered rooted trees rooted on  $v_a$  and  $v_b$ . The subtrees of these two roots being respectively  $(T_a^1, T_a^2)$  and  $(T_b^1, T_b^2)$ .

The tree encoding the chirality of the double bond is then defined as an ordered rooted tree, whose root corresponds to a virtual vertex (not corresponding to any atom) connected to the two subtrees  $T_a$  and  $T_b$ . As in Section 4.1, the computation of the minimal chiral subtree is bounded by  $\mathcal{O}(|V|^2)$ . Figure 5a represents a double bond between two carbon atoms with its minimal chiral subtree (Figure 5b).

#### **Graph Contraction** 4.3

Using results in Section 4.1 and 4.2, each stereocenter may be associated to a minimal chiral subtree and a depth-first canonical string according to  $\Sigma^M$  representing it (Section 3). However, properties of a molecule are both determined by its set of minimal chiral subtrees and by relationships between these trees and



**Fig. 5.** A double bond (a), its minimal chiral subtree (b) and its contraction (c).

the remaining part of the molecule. In order to obtain a local characterization of such relationships, we propose to contract the minimal chiral subtree of each stereocenters.

Let us consider a stereocenter s and its minimal chiral subtree  $T = (\hat{T}, ord_T)$ , with  $\hat{T} = (r, G_T), G_T = (V_T, E_T, \mu, \nu)$  associated to a depth-first canonical string according to  $\Sigma^M$ , we denote this string  $c_s = \text{DFCS}_{\Sigma^M}(T)$ . We define for this tree a set of connection vertices:

$$V_{\text{con}} = \{ v \in \text{Leaf}(T) \mid d(v) > 1 \}$$

and a set of edges to contract:

$$K_T = E_T - E_{\text{con}}$$
 with  $E_{\text{con}} = \{(v, p_v) \in V_{\text{con}} \times V_T\}$ .

The contraction of  $K_T$  creates a new graph  $G_s = (V_s, E_s)$ , with a contracted node  $n_s$  labeled by  $c_s$  and  $V_s = V - (V_T - V_{\text{con}}) \cup \{n_s\}; E_s = E - K_T$  (Figure 5c).

Each edge of  $E_{\text{con}}$  connects an element l of  $V_{\text{con}}$  to  $n_s$  in  $G_s$ . The label of  $e = (n_s, l)$  has to encode the position of l in the minimal chiral subtree. We thus consider the path connecting r to l in the minimal chiral subtree:

$$CP(l) = v_1, ..., v_n$$
 with  $v_1 = r$  and  $v_n = l$ 

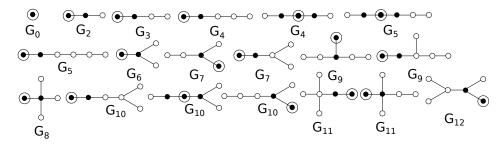
Let us denote  $i_j$  the index of  $v_j$  in the ordered list of children of  $p_{v_j}$ . The sequence  $i_2 \ldots i_n$  defines a unique path in the chiral subtree associated to  $n_s$ . Such a sequence may thus be considered as a proper label of edge e. However as mentioned in Section 3, some paths may pass through equivalent subtrees and should thus be considered as equivalent. In order to encode such an equivalence relationship we define the label of e as:

$$\nu(e) = \bigotimes_{i=2}^{n} rep(v_i)$$

where rep is defined by Equation 3, Definition 20 and  $\bigcirc$  denotes the concatenation operator.

#### 4.4 StereoTreelet

For each stereocenter s we have a graph  $G_s$ . The stereotreelets of  $G_s$  are defined as all subtrees of  $G_s$  whose size is lower than 6 and which include  $n_s$ . Since each



**Fig. 6.** The set of stereotreelet with  $n_s(\bigcirc)$ , elements of  $V_{con}(\bullet)$ , elements of  $V-V_{con}(\circ)$ 

neighbors v of  $n_s$  corresponds to a leaf of the minimal chiral tree of s, the edge  $(v, n_s)$  is already encoded within the code  $c_s$  of  $n_s$ . Consequently, we impose that each neighbor v of  $n_s$  in a stereotreelet must have at least another neighbor (different of  $n_s$ ). This constraint induces the set of stereotreelets represented in Fig. 6. The set of stereotreelet  $\mathcal{T}(G)$  of G is defined as the union of stereotreelets of each  $G_s$ .

When all stereotreelets of G have been enumerated, we compute its spectrum s(G) which corresponds to a vector representing the treelet distribution. Each component of this vector is equal to the frequency of a given stereotreelet t:  $s(G) = (f_t(G))_{t \in \mathcal{T}(G)}$  with  $f_t(G) = |(t \subseteq G)|$ . The kernel between two graphs G and G' is defined as a sum of kernels between the different number of treelets common to both graphs:

$$k(G,G') = \sum_{t \in \mathcal{T}(G) \cap \mathcal{T}(G')} K(f_t(G), f_t(G')).$$

#### 5 Experiments

We have tested our method on a dataset of acyclic chiral molecules [11] related to a regression problem. This dataset is composed of 90 molecules together with their optical rotations. In practice, we only select 35 molecules, since almost all molecules have only one stereocenter, and for 55 molecules this stereocenter is unique in the dataset. Such molecules correspond to a property represented only once in the dataset which can thus not be accurately predicted. The property to predict, the optical rotation, is connected with chirality and has a standard deviation of 38.25 for the 35 selected molecules.

For our experiment we use a leave-one-out cross-validation on the dataset to predict the optical rotation of each molecule. The predicted rotations are computed by using both kernel ridge regression and the weighted mean of known values using the similarity measure provided by the kernel

$$\hat{y} = \frac{\sum\limits_{i} k(G_i, G) \times y_i}{\sum\limits_{i} k(G_i, G)}$$

Method	Kernel Ridge		0 0		Gram's matrix
	Average Error	RMSE	Average Error	RMSE	computations (s)
Random Kernel	31.7	39.5	32.0	39.3	0.03
KMean [12]	31.0	38.7	32.3	39.6	153.84
Treelet Kernel [1]	26.0	33.9	28.9	37.4	0.49
Stereotreelet Kernel	21.0	25.6	11.6	16.3	0.13

 Table 1. Optical rotation prediction for the acyclic chiral dataset.

We present in Table 1 the average errors, Root Mean Squared Errors (RMSE) and computation times of the Gram matrix for our method and the ones of [12, 1] which do not take into account chirality. Results obtained by using a random Gram matrix are also shown.

Weighted mean provides much better results for our kernel since on this dataset each molecule has a non null similarity with a reduced number of molecules (less than 10). Such a reduced number of data do not allow kernel ridge regression to perform reliable prediction. Other methods provide similar results than those obtained using a random Gram matrix. These results are also comparable with the variance of the dataset. Such a result means that the similarity measures provided by alternative kernels are not correlated with the property to predict. This last point may be explained by the fact that optical rotation is connected to chirality which is not encoded by these kernels.

#### 6 Conclusion

In this report we proposed a new model which allows to encode the relative positioning of vertices within a graph. Such a model is quite flexible since it does not require any coordinate information and may be defined on only some vertices of a graph. We also introduced the notion of Minimal chiral subtree which, in the acyclic case, corresponds to the minimal subgraphs which allows to explain the chiral property of a vertex.

We applied this model to encode the chiral information of molecules and vertices. Based on the minimal chiral subtree of each vertex we defined a graph kernel between chiral molecules. Our experiments show promising results and our future work will consist to create larger datasets and to extend our method to graphs including cycles.

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