

Development of An Efficient Algorithm to Enumerate the Number of Constitutional Isomers of Alkyne Series

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ABSTRACT

Computerized Enumeration of alkane's isomers has long been a topic of interest to the researchers. The double bond carbon content and triple bond carbon content isomers were mostly neglected for many decades by the researchers and claimed in the literature that there is no simple algorithm for these problems. In this paper an efficient and simple algorithm has been proposed to count the number of constitutional isomers of alkynes series. By using this recursive algorithm the resulting computation of number of isomers for alkynes with any number of carbon contents [limited to system specifications] are easily enumerated. Further the time complexity of the proposed algorithm has also been investigated. The algorithm has been implemented using object oriented programming language-Java.

Keywords

Carbon, Isomer, Counting, alkynes, enumeration, time complexity

1. INTRODUCTION

Before outlining the background and scope of the problem that we have consider, it will be appropriate to review briefly the basic facts about chemical compounds. For our purpose we can regard a molecule of a chemical compound as an assemblage of atoms in which some atoms are linked to other by "valency bonds". These may be single, double and triple. In this paper we shall be concerned only about hydrocarbons. A hydrocarbon is any chemical compound containing only carbon and hydrogen. The carbon atom has valency four and the hydrogen one. Hydrocarbons can be classified into acyclic and cyclic types. In this paper we have discussed only acyclic type. The acyclic hydrocarbons are characterized by a branched tree structure, and can be separated into three most categories [3].

- Alkanes: Contain single bonds, and have the general formula C_nH_{2n+2} .
- Alkenes Contain double bonds, and have the general formula C_nH_{2n} .
- Alkynes contain a triple bond, and have the general formula C_nH_{2n-2} .

Isomers are any set of chemical compounds that have the same chemical formula, but a different arrangement of the atoms. For example pentanes have the three isomers.

2. RELATED WORK

In the past a Simple algorithm for counting constitutional isomers of alkanes, single C-atom isotopically labelled alkanes, and monosubstituted alkanes was reported[4] and the algorithm was implemented in a high level programming language-Turbo Pascal. Mathematically recount of alkenes with geometric isomerism due to a double bond was taken into consideration[5]. Further the mathematical relationship between the number of isomers of alkenes and alkynes was established from the enumeration of isomers of alkenes and alkynes[2]. Computerised isomer Enumeration of the Alkane Series was presented[1] and an algorithm to enumerate the isomers of eighteen carbon atoms alkane series was implemented in the Object Oriented programming language- C++.

3. THE ALGORITHM

3.1 Process

The main goal of this algorithm is to enumerate the isomers of any number of carbons of alkynes series in an accurate and efficient manner The proosed algorithm is using recursion and has been developed using an object oriented programing language-Java. A custom data structure has been used in this program.

3.2 Method

The algorithm is based on the formation of alkyl biradicals [8]. These alkyl biradicals can be imagined to be formed by a hypothetical chemical reaction in which an alkane molecule loses two protons and becomes an alkyl biradical, i.e. is with two unpaired electrons. Interestingly, not all possible alkane isomers are required to form all possible alkyl biradicals as alkane molecule has many different sites where the two protons can be lost. When a single alkyl biradical combines with another alkyl biradical, the four electrons will be paired up and double bond will be formed [2].

3.3 Procedure

ALKYNES_ISOMER (n , $Count=0$, Ptr , $nofcarbon$) $n = number$ of isomeric alcohols, $nofcarbon=no$ of carbon of alkynes, $count$ is number of isomer of alkenes

1. Initialize array{ $n1, n2, n3, \dots, ni$ }
2. Set $Ptr=0, b=0, Count=0$
3. Repeat step 5 and 6 until $ptr \leq n$,
4. if $nofcarbon$ is even go to step 5 else go to step 6
5. Repeat until $n < (nofcarbon/2)$
 - I. $Count = Count + array[n] * array[nofcarbon - (n+1)]$

- II. if $(n \neq \text{nofcarbon}/2)$ go to step (a) else go to step (III).
 - a) Break
 - b) Return Count
 - c) Exit
- III. $\text{Ptr} = \text{Ptr} + 1$
- 6. Repeat step (I) to (c) until $n < (\text{nofcarbon} - n)$
 - I. if $(n \neq (\text{nofcarbon} - 1)/2)$ go to step (II) else go to step (III)
 - II. $\text{Count} = b + (\text{array}[n] * (\text{array}[n] + 1)) / 2$
 - a) break
 - b) Return Count and exit
 - III. $b = b + \text{array}[n] * \text{array}[\text{nofcarbon} - (n - 1)]$
 - IV. $\text{Ptr} = \text{Ptr} + 1$
 - V. Return Count
 - VI. Exit

4. RESULT AND DISCUSSION

4.1 Outputs

The enumeration of isomer is not only of particular interest in theoretical chemistry, but also expanding field of computational chemistry.

The procedure described in section III(C) will enumerate the isomers of any member of the alkynes family exhaustively and irredundantly. In this research works using the said procedure isomers of alkynes series with carbon content from 1 to 47 have been calculated without any error. Isomers of higher carbon content could not be calculated because of system limitation.

Table 1 Alkynes Isomes

| Number of carbon | Number of constitutional Isomer of alkynes series |
|------------------|---|
| 1 | 1 |
| 2 | 1 |
| 3 | 2 |
| 4 | 3 |
| 5 | 7 |
| 6 | 14 |
| 7 | 32 |
| 8 | 72 |
| 9 | 171 |
| 10 | 405 |
| 11 | 989 |
| 12 | 2426 |
| 13 | 6045 |
| 14 | 15167 |
| 15 | 38422 |

| | |
|----|---------------------------|
| 16 | 97925 |
| 17 | 251275 |
| 18 | 648061 |
| 19 | 1679869 |
| 20 | 4372872 |
| 21 | 11428365 |
| 22 | 29972078 |
| 23 | 78859809 |
| 24 | 208094977 |
| 25 | 550603722 |
| 26 | 1460457242 |
| 27 | 3882682803 |
| 28 | 10344102122 |
| 29 | 27612603765 |
| 30 | 73 844 151 259 |
| 31 | 197 818 389 539 |
| 32 | 530 775 701 520 |
| 33 | 1 426 284 383 289 |
| 34 | 3 838 066 701350 |
| 35 | 10 341 758 769 406 |
| 36 | 27 900 947 721 908 |
| 37 | 75 362 644 825 968 |
| 38 | 203 787 850 635 992 |
| 39 | 551 645 375 673 949 |
| 40 | 1 494 781 478 155 753 |
| 41 | 4 054 242 571 711 886 |
| 42 | 11 006 161 817 116 528 |
| 43 | 29 904 564 722 290 758 |
| 44 | 81 319 947 893 937 569 |
| 45 | 221 308 699 013 145 314 |
| 45 | 602 735 147 429 051 222 |
| 47 | 1 642 733 167 881 428 721 |

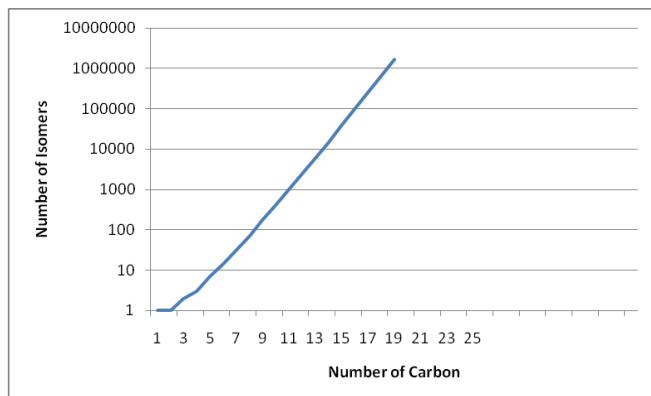


Figure 1

Figure 1 show that, after the carbon content eight, the graph takes on a linear outward show, which indicates that the number of isomers increases logarithmically with the carbon content. Before eight however, there does not appear a clear cut pattern or relationship between numbers of carbon contents with its isomers however beyond carbon content eight there is a linear relationship which could be used to estimate the number of isomers for higher carbon contents.

4.2 PERFORMANCE OF THALGORITHM

4.2.1 Accuracy

The developed algorithm has produced the number of isomer exhaustively irredundantly up to carbon contents 47 without any error as compare to the published results results[2] hence the proposed algorithm is accurate .

4.2.2 Time Complexity & Efficiency:

The algorithm denotes a significant improvement over previous algorithms developed earlier in respect of computation time. The time complexity of this algorithm is by far its strongest point. As established in the figure 2, the amount of computational time required varies approximation linearly with the number of carbons. This is notable because in past algorithms, computational time have increased logarithmically or greater, with the increasing number of isomers. The time complexity is dependent on the creativity of a programmer and system configuration.

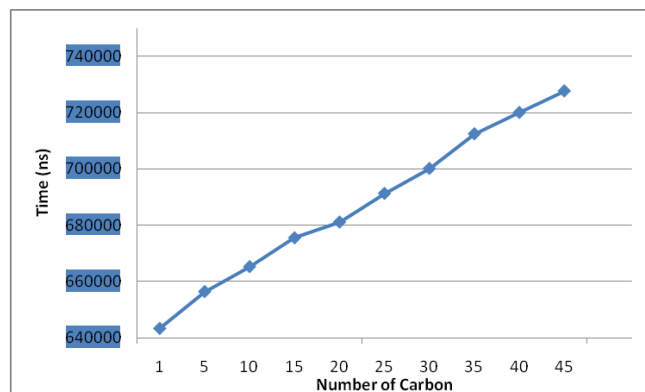


Figure 2

5. CONCLUSIONS AND FUTURE SCOPE

The developed algorithm has been successful in accomplishing the main goals of the current research work. However isomers of higher carbon contents could not be calculated because of system limitations.

The algorithm can be modified to enumerate the isomers of alkenes series with great accuracy

6. REFERENCES

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