



A COMPUTATIONAL MODELLING STUDY OF CHLORINE
DIOXIDE
AND ITS ROLE IN WATER PURIFICATION CYCLES

by

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Submitted in partial fulfilment for the degree of

Doctor of Philosophy of Science (PhD)

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(2023)

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DECLARATION

I, *Natasha Misheer* declares that this dissertation, which I hereby submit for the degree of Doctor of Philosophy of Science (PhD), in Chemistry at the University of Pretoria, is my own work and has not previously been submitted by me for a degree at this or any other tertiary institution.

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Date: 16 November 2023

Published output of this work:

Submitted to: *Heliyon Journal of Science (Chemistry)*

“Semi-empirical supported, *Ab Initio* derived Thermodynamic properties for ClO_2 and its sub- and extended- species, applied in water treatment cycles”, by: Natasha Misheer, Jan A Pretorius and Patrick Ndungu

DEDICATION

I dedicate this achievement to my parents Rabichund and Bindu Misheer and my late brother Nitesh Misheer. A special mention to my mum Bindu Misheer, my heartfelt gratitude to you mum for your support. I appreciated your efforts to sometimes “burn the midnight oil with me”. My parents unwavering belief in my abilities helped me to push through the innumerable difficulties and achieve what seemed like an unattainable feat.

ACKNOWLEDGEMENTS

- I would like to thank my parents for their constant encouragement and motivation. You both encouraged me to persevere against all odds.
- My heartfelt gratitude to my brother Ajay Misheer. for your steadfast support and lending a sympathetic ear to my issues.
- My sincere gratitude to my nieces Kajal and Keasha Misheer for their motivation to achieve this accomplishment. The mere fact that you both look to me for inspiration to achieve success in your own lives impelled me to refuse to accept defeat.
- I would like to acknowledge my supervisor Dr Jannie Pretorius
- My heartfelt gratitude to Prof Vinesh Maharaj, Prof Patrick Ndungu and Prof Barend Erasmus. Your support motivated me to endure tremendous difficulties and to achieve my ultimate goal.
- I take this opportunity to convey my sincerest appreciation to Prof Patrick Ndungu and Prof Vinesh Maharaj for your guidance.
- I would like to thank my friends for supporting me throughout the duration of my studies. Your encouragement helped me succeed at a task that seemed impossible at times.
- I thank God for always giving me the strength to achieve my goals. Nothing is ever possible without your presence.

ABSTRACT

This study involves the computational investigation into the perceived reactivity of a group of sixty (60) sub- and extended- chlorine oxide species with the general formulae of Cl_xO_y applied to water purification. In order to achieve the ultimate objective of elucidating their chemical role in aqueous media, acting as oxidative agents, an intensive computational approach has been followed to determine their thermochemical properties. The extended species of ClO_2 display a complex sequence of bonding character with an appreciable charge dissipation (extracted as partial charges), which complicates the effective selection of basis sets and electronic structural optimization, during *Ab Initio* analyses. Besides a single molecular computational analysis, an alternative grand canonical ensemble approach was introduced, applying *Gibbs* ensemble *Monte Carlo* simulations, supported by revised force field parameters to derive at optimum model sizes. In this context, this approach proved highly efficient, resulting in consistent thermochemical properties for all species, through optimum selection of Hamiltonians and appropriate basis sets, during quantum chemical analyses.

Excellent correlations with published Heat of Formation energies were obtained for almost all the ensemble derived species. A few energy discrepancies identified during *Ab Initio* (*VASP* and *Gaussian* software) calculations will need to be investigated more thoroughly in a further study. Chemical structure geometries were typically maintained for all models and self-consistency reached in all the quantum chemical refinement cycles.

Some of these species are presented as radical and ionic entities, which complicates their quantum atom potential representations. This observation specifically applies to species exhibiting variable spin conditions. This spin variability can further promote spin contamination, through extensive polarization contributions. Charged species were unfortunately not exposed to empirical ensemble simulations (reasons given) and had to be considered as single molecular entities.

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LIST OF ABBREVIATIONS

ΔH_f	Heat of Formation
ΔG_f	Free Energy of Formation
Å	Angstroms
AM1	Austin Model 1 (MOPAC)
ATcT	Active Thermochemical Tables
BFGS	Broyden-Fletcher-Goldfarb-Shanno
B3LYP	Becke's three-parameter nonlocal exchange functional with the correlation functional of Lee, Yang, and Parr
CCSD(T)	Coupled Cluster Single-Double Excitation (with Triples)
CG	Conjugate Gradient (<i>VASP</i>) minimization method
Cl ⁻	Chloride
ClO ₂ ⁻	Chlorite
ClO ₂	Chlorine dioxide
ClO ₃ ⁻	Chlorate
cvff	Consistent-Valence force field
DBPs	Disinfection byproducts
DFT	Density Functional Theory
DOS	Density of states (<i>VASP</i>)
DPD	N, N-diethyl-p-phenylenediamine
\bar{e}	electron
EF	Eigenvector Following
FACT	The Facility for the Analysis of Chemical Thermodynamics (<i>FactSage software</i>)
FFT	Fast Fourier Transform
GAC	Granular activated carbon
GEMC	Gibbs ensemble Monte Carlo (<i>GIBBS</i>)
GGA	Generalized Gradient Approximation
HAAs	Haloacetic acids
HANs	Haloacetonitriles
HClO ₂	Chlorous acid
HF	Hartree Fock theory
HLC	Higher-level correction
KClO ₃	Potassium chlorate
LBFGS	Limited memory Broyden-Fletcher-Goldfarb-Shanno
MBJLDA	Modified Becke Johnson Local Density Approximation (<i>VASP</i>)
MCL	Maximum contaminant level (MCL)

MedeA	Materials Exploration and Design Analysis (Software from Materials Design Inc.)
MNDO	Modified Neglect of Diatomic Overlap (<i>MOPAC</i>)
MINDO	Modified Intermediate Neglect of Diatomic Overlap (<i>MOPAC</i>)
MNDOD	Modified Neglect of Diatomic Overlap with d Atomic Orbitals (<i>MOPAC</i>)
MO	Molecular orbital
MOPAC	Molecular Orbital Package
MRDL	Maximum residual disinfectant limit
MS0	Made Simple 0 (<i>Gaussian</i>)
MS1	Made Simple 1 (<i>Gaussian</i>)
MS2	Made Simple 2 (<i>Gaussian</i>)
mw	Molecular weight
NDC	Severe nondynamical correlation
NDDO	Neglect of diatomic differential overlap
NLLSQ	Non-Linear Least Squares gradient minimisation method
NOM	Natural Organic Matter
NTP	Normal Temperature and Pressure
pcff	Polymer Consistent force field
PM6	Parametric Method 6 (<i>MOPAC</i>)
PM3	Parametric Method 3 (<i>MOPAC</i>)
PM7	Parametric Method 7 (<i>MOPAC</i>)
QCISD(T)	Quadratic configuration interaction single-double excitation (with triples)
revTPSS	Revised Tao-Perdew-Staroverov-Scuseria
RHF	Restricted Hartee Fock
UHF	Unrestricted Hartee Fock
RM1	Recife model 1 (<i>MOPAC</i>)
ROHF	Restricted Open Shell Hartree Fock
SCAN	Strongly Constrained and Appropriately Normed (<i>Gaussian</i>)
THMs	Trihalomethanes
TOX	Total organic halogen
TPPS	Tao-Perdew-Staroverov-Scuseria
TS	Transition energies (<i>VASP</i>)
VASP	Vienna Ab Initio Simulation Package

CHAPTER-1: INTRODUCTION

1.1 Problem Statement

Natural Organic Matter (NOM) removal from water has become increasingly difficult, due to the wide variety of organic compounds found in an aquatic environment [1]. NOM has the ability to disrupt or interfere in many processes in the water treatment cycle [2]. It is well known that the enhanced coagulation process which comprises flocculation and coagulation, alters the chemical/physical nature of dissolved and suspended solids, allowing for removal in a clarifier. However, coagulation can only efficiently remove hydrophobic and high molecular fractions of NOM [3]. The remaining NOM species persists in the aquatic environment and interfere with the rest of the process treatment steps.

Application and the online chemical production of ClO_2 in the presence of a water treatment plant, will demand strict compliance to *Occupation Health and Safety* conditions and can be considered to be the main inhibiting factor, in favor of purging chlorine from pressurized cylinders.

1.2 Objective of this Study

The objective of the investigation focuses on a computational construction of ClO_2 extended- and sub- species, to elucidate their thermochemical reaction schemes, in an attempt to identify the main species persisting in an aqueous medium.

The ultimate objective is to demonstrate the exceptional behaviour of ClO_2 as a water pre-treatment agent, but also to derive at optimum conditions for these species to survive sufficiently long, to act as pathogen controlling agent, a topic of a further study.

This study aims to understand if the added oxidant (ClO_2), can alter the remaining NOM species' chemical properties, such that they can further aid the enhanced coagulation process resulting in higher organic removal.

1.3 Background and Study Step

The study involves a selection of a significant group of chemically extended- and sub-species of ClO_2 , including a series of hydrogen derived HCl_xO_y species. These chemical species are mostly unstable and therefore difficult to sustain an environment to measure their properties through spectroscopic or other chemical instrumental techniques [4]. An extended sub-set of species (some as radical and ionic in nature) can be constructed from their common denominating radical compound ClO_2 . This 'synthetic' step is applied, mainly due to their short life span and extreme kinetics in aqueous based chemical reaction cycles [5].

- a) An electronic and dynamic structural optimization step was first undertaken to derive effective model geometries, validated against published Heats of Formation. Both *Ab Initio* (*VASP*) and

semi-empirical (MOPAC) quantum chemistry software systems have been used for this step and acted as prelude to Monte Carlo Simulations at the *Grand Canonical* ensemble model scale

- b) A *Grand Canonical* ensemble (GEMC) step was introduced to formulate larger (bulk) models, in an attempt to correlate associated Heats of Formations against single molecular properties. As a result of the restriction of force field definitions, only neutral and radical species were exposed to Monte Carlo, *Grand Canonical* ensemble simulations, to derive at their respective optimum model sizes (based on their optimum bulk model Internal Energies U_i).
- c) An extensive set of empirical force field parameters for the radical species of ClO_2 have had to be derived, in order to perform the GEMC empirical simulations. The *pcff+* set of force field parameters were employed and adjusted for this study. The materials based *Medea/GIBBS* software was employed for the GEMC simulations.
- d) These extended optimum GEMC model presentations were then exposed to semi-empirical quantum chemistry vibrational analysis (*MOPAC-2016* software) to derive an active set of thermochemical properties.
- e) Single species were exposed to the *Ab Initio* derivation of internal electronic energies (including the zero-point energy) for all species, using the *Gaussian-16* and *MOPAC-2016* software, to derive at a complementary set of thermochemical properties.
- f) A *Density Functional* (DFT) approach was then followed, applying the *VASP* software system in periodic environments, for all single species (initially to derive at optimum geometries, but also to obtain thermochemical properties)
- g) The final phase of this study involved a species-based thermodynamic analysis, applying the PC-based *FactSage* software program, to predict the most relevant chemical reaction phenomena in an aqueous medium.

1.4 Limitation of the study

The investigation into the effects that the ClO_2 species (oxidant) have on NOM species characteristics, are extremely vast and requires multiple studies which are time consuming. The simulation of the interaction of various NOM species with the known persistent ClO_2 species, could not be examined in this study. Furthermore, the ability of ClO_2 to act as a pathogen controlling agent must also be undertaken as further investigations.

1.5 References

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CHAPTER-2: LITERATURE REVIEW

2.1 Background

This chapter offers a detailed summary of the extensive chemical applications, chemical and physical properties and vast array of wider phenomena (e.g. medical applications, paper and pulp, pathogen control) already recorded for ClO_2 and its extended and sub species. This also demonstrates the complexity contained in such a “small” chemical compound, having electronic properties of a radical, reduced or oxidized into ions as well. It is furthermore important to realise the related complexity, to undertake electronic level computations on these species. Their geometries and stereo-chemical presentations are generally known to be insufficiently ‘described’ by the quantum chemical software systems of today.

2.2 Historic Statement

Chlorine dioxide (ClO_2) was discovered in 1814 by Sir Humphrey Davy whose likeness is illustrated in Figure-2.1 [1]. He produced ClO_2 in the gaseous phase by reacting sulphuric acid (H_2SO_4) with potassium chlorate (KClO_3) [2]. Later he replaced H_2SO_4 with hypochlorous acid (HOCl) [1]. Equation (1) highlighted below has recently gained popularity as the typical method used to yield huge quantities of ClO_2 [2]. Sodium chlorate (NaClO_3) replaced potassium chlorate in the reaction [3]:

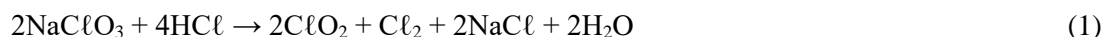


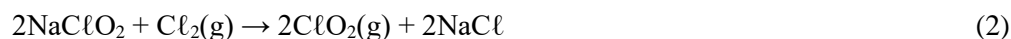
Figure-2.1: Sir Humphrey Davy [4]

The ClO_2 molecule contains an odd number of valence electrons, and is classified as a paramagnetic radical in the triplet spin state. L.O. Brockway [5] presented a three-electron bond electronic structure for ClO_2 . Chemist Linus Pauling carried out investigations which led to the discovery of two resonance structures encompassing a double bond on one side and a single bond which comprised of a $3\bar{e}$ bond on the other [6]. It is an extremely energetic and volatile molecule considering its size and is the only substance that can still persist as a monomeric free radical even in diluted aqueous solutions. [7].

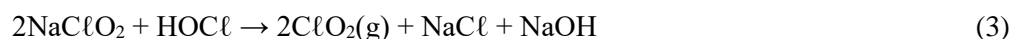
2.3 Properties, Applications and Chemical Character

2.3.1 ClO_2 generation

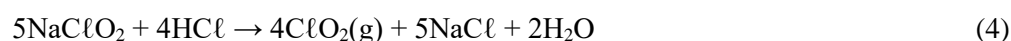
Three methods of ClO_2 generation have been identified in open literature. It can be generated by reacting sodium chlorite with reacting with gaseous chlorine.



Secondly, sodium chlorite can be reacted with hypochlorous acid (HOCl) to produce ClO_2 .

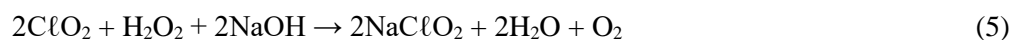


And the reaction of sodium chlorite with hydrochloric acid (HCl) also produces ClO_2 .



Reactions (2), (3), and (4) demonstrate how generation steps can differ [8], [9]. The most efficient conversion of sodium chlorite to ClO_2 (80%) when using hydrochloric acid is achieved through reaction (4) above. This method is currently widely used in industry in most water process applications [10].

Chlorite (NaClO_2) is a stable precursor of ClO_2 that is safely transported and utilised to generate ClO_2 as per the process reflected in Equation (4) [11]. This stable precursor is produced commercially by reacting ClO_2 with the reducing agent hydrogen peroxide [12].



2.3.2 ClO_2 Physico-chemical properties

The gas phase of ClO_2 has the following thermochemical properties [13] at Normal Temperature and Pressure (NTP):

1. Enthalpy of formation: 102.5 kJ/mol [13]
2. Gibbs energy of formation: 120.5 kJ/mol [13]
3. Entropy: 256.8 J/mol K [13]
4. Heat capacity: 42.0 J/mol K [13]
5. The dipole moment of ClO_2 is 1.69 Debye units [14], [15]

The reactivity of ClO_2 and its physical chemical properties have been studied extensively. Some key physical-chemical parameters are listed below:

1. ClO_2 is reduced to chlorite (ClO_2^-) for it to react predominantly as a highly selective oxidant [16], [8], [17]. The equivalent electron transfer reactions are similar to those occurring when singlet oxygen acts as an oxidant [18], [19].
2. The chlorine-oxygen bonds have been described as double bond comprising an angle of about 117.5° [14], [20] and a chlorine oxygen bond length of 1.47 Å.

3. ClO_2 has a maximum absorption response at 359 nm and a molar absorptivity of 1250 litres/mol
4. cm [14], [20].
5. When ClO_2 comes in contact with aqueous pollutants, it typically reduces to chlorite ion ($\text{ClO}_2 + \text{NOM} \rightarrow \text{Products} + \text{ClO}_2^-$) [21], [22].
6. A significant difference between ClO_2 and chlorine is the ability of ClO_2 to exist as a dissolved true gas in solution and has superior solubility in water, ten times greater than the solubility of chlorine in water [7]. This ability supports ClO_2 in maintaining its biocidal effectiveness over a wide pH range [23], [24]. Hence, the use of ClO_2 is favoured for the operation of cooling systems that have poor pH control.
7. ClO_2 is an oxidizing agent, not a chlorinating agent [7].
8. Its application in water treatment, containing high organic content, has the benefit of hardly forming trihalomethanes (THMs) [15].
9. It has a “lower oxidation strength than ozone but is more powerful than chlorine” [25].

2.3.3 ClO_2 Structure

According to Pauling’s theory, the molecular geometry (shape) of ClO_2 is considered angular whilst the electronic shape is tetrahedral due to the presence of five valence electrons [26]. The five valence shell π -electrons are oriented in the $(1b_1)^2(1a_2)^2(2b)^1$ molecular orbital (MO) configuration [27], [28]. The removal of the 3d-atomic orbitals from the chlorine atom results in the modification of $(1b_1)^2(1a_2)^2(2b)^1$ configuration whereby the five electrons become distributed across three overlapping $p-\pi$ atomic orbitals (AOs) [27], [28].

2.3.4 ClO_2 as a Disinfectant

ClO_2 is being applied progressively to control microbiological growth in various industries. ClO_2 has a stronger oxidizing power, hence small doses produce effective pathogen control. This property makes ClO_2 an economical dosing option which also highlights its safety aspects in terms of storage and deployment. In terms of microbiological control or disinfection, ClO_2 finds use in the following:

- Pulp and paper industries for paper pulp bleaching [29], [30]
- Fruit and vegetable process industries where it is used to clean and preserve fruit and vegetables, extending the shelf life of the produce [31].
- Various canning plants for meat and poultry add ClO_2 as a process aid to reduce the microbial plate counts [32].

ClO_2 is utilised as a primary or secondary disinfectant in water treatment, for the production of potable water. The key factors that favour its use include [33]:

- Taste and odor control

- THM/halo-acetic acids (HAA) reduction
- Iron (Fe) and manganese (Mn) control
- Colour removal
- Sulphide and phenol degradation.

2.4 Microbiological Properties of ClO_2

ClO_2 has a more efficient antimicrobial activity than chlorine [34]. ClO_2 has been documented to have the ability of eliminating *Cryptosporidium*, *Giardia* and *Legionella sp.* effectively [35], [36]. It is very efficient in killing bacteria and especially successful in deactivating viruses [37]. Organic components in bacterial cells interact with ClO_2 which results in the inhibition and destruction of cellular processes. ClO_2 penetrates the cell walls of bacteria and inactivates amino acids and the RNA present in the cell [38]. ClO_2 adsorbs into the surface area of bacterial cells and concentrates around the bacterial cell, where it can act effectively due to the physical adsorption process. ClO_2 hence, infiltrates the bacterial cell thereby destroying enzymes [39], [40], [41].

An example of the action ClO_2 on bacterial cells is demonstrated by the reaction of ClO_2 on *E. coli*. ClO_2 interferes with the external membrane of *E. coli* by promoting substantial leakage of K^+ ions out of the specimen via inhibition of enzyme β -D-galactosidase [42]. This point again suggests that ClO_2 acts by targeting the cell wall of micro-organisms.

Virus elimination differs from bacterial elimination, by directly reacting with water soluble peptone which in turns prevents the formation of amino acids that is required for protein synthesis. Hence, protein formation is prevented which leads to death of the viruses. [43]. It inactivates microorganisms by directly oxidizing tyrosine-, methionyl- or cysteine-containing proteins that disrupt important structural functions in the enzyme metabolic processes [44], [43].

2.4.1 Microbial action of ClO_2 in alkaline medium

In the presence of an alkaline solutions, ClO_2 disproportionate to chlorite (ClO_2^-) and chlorate (ClO_3^-) via the following reaction [24]:



This reaction is catalysed by hydrogen (H^+) ions. The half-life of aqueous mixtures of ClO_2 decreases with increasing pH.

Since the “organic matter” is oxidised by undissociated chlorine dioxide as the ClO_2 ion [45], it is possible that ClO_3^- is the most reactive species that is responsible for the inactivation of the poliovirus in an alkaline environment. Alvarez *et al.*, [46], [47], studied the mechanism of inactivation of the polio virus by ClO_2 and reported that addition of ClO_2 in an alkaline medium, resulted in the separation of the RNA from the capsids. ClO_2 penetrates into HeLa cells of the virus and initiates the protein uncoating process [46], [47]. The inactivation process takes place at pH 6.0 where ClO_2

penetrates the protein coat and changes the pH thereby altering the viral RNA reducing its ability of producing templates for RNA replication [46], [47].

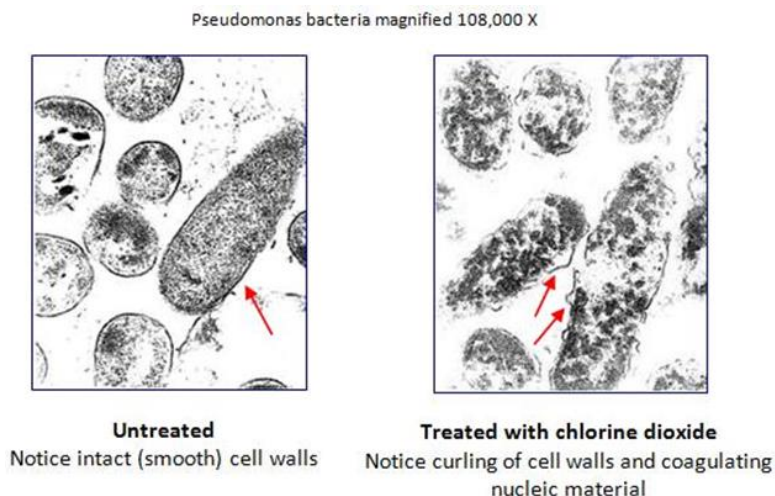


Figure- 2.2: Impact of ClO_2 on micro-organism cell depletion [33]

High pH also increases the susceptibility of the virus to ClO_2 attack [46], [47]. The reaction is not time dependent nor concentration dependent [38]. In contrast to non-oxidising disinfectants, ClO_2 kills micro-organisms regardless of it being in an active or passive state. Therefore, a lower concentration of ClO_2 is required to effectively kill micro-organisms, which cannot build up any resistance against ClO_2 .

ClO_2 has a higher ‘oxidative capacity’ than all disinfectants [48]. The capacity of chemicals denotes the number of electrons one molecule can accept from its surrounding molecules. In the case ClO_2 , this means that it can gain five electrons from microbial species per molecule, making it a superior biocide to alternative oxidisers, which typically are only able to gain two electrons. The oxidative capacity of ClO_2 compared to other disinfectants is presented in Table 2.1.

Table-2.1: Chlorine Dioxide Oxidation Capacity compared with other Disinfectants [48]

Oxidant	ClO_2	H_2O_2	NaClO_2	KMnO_4	Cl_2	NaClO
Capacity Oxidation	263%	209%	157%	111%	100%	93%

- Oxidation capacities tabulated in table 2.1 indicate that ClO_2 has the highest oxidation capacity compared to other disinfectants.
- This oxidizing capacity substantiates why ClO_2 is considered to be the most effective disinfectant agent.

- ClO_2 is highly soluble in water and, unlike ozone, does not react with the extracellular polysaccharides of biofilms, hence it can rapidly eliminate biofilms [49], [50].
- ClO_2 penetrates into biofilms swiftly, to destroy the microbes present inside the film and oxidises the polysaccharide matrix that retains the bio-film structure [51], [52]. As a result, the biofilm dissociates into pieces that remain steady. An acidic environment forms as the biofilm begins to grow again, and the chlorite ions are transformed into ClO_2 in this environment. Hence, the remaining biofilm is eliminated [50]. Many researchers have described exceptional results whilst using ClO_2 as a biocide [49], [53], [54], [50].

2.5 Radicals and Other Reactive by-products of ClO_2

There are various chlorine oxide species that are either classified as radicals or highly reactive species i.e., ClO , ClO_2 , ClO_3 , ClO_4 , HOCl , HClO_2 , Cl_2O , Cl_2O_3 , Cl_2O_4 and Cl_2O_6 . The mono-chlorine oxides are highly reactive *radicals* which are challenging to detect since they are fairly short-lived species [55]. The di-chlorine oxides are highly unstable [55].

ClO_2 undergoes a decomposition reaction in aqueous medium that produces chlorite and chlorate ions:



A solution of ClO_2 in water will degrade upon exposure to light, through a complicated reaction mechanism. Chloride (Cl^-), chlorite (ClO_2^-) and chlorate (ClO_3^-) are the products that form, from the disproportionation reaction of ClO_2 . Other radical formation products observed are: HClO_2 , ClO_2^- , HOCl , and Cl_2 [56]. “Approximately 70% of applied ClO_2 forms chlorite, while about 10% forms chlorate [57]”. The free radical is found in gaseous form above 11 to 12°C [58], [8], [20].

2.6 Decomposition and Dissociation of ClO_2

Chlorate (ClO_3^-) and chlorite (ClO_2^-) concentrations are both present in the pH range between 1 and 4. Chlorite (ClO_2^-) is the main species above pH 4. Although chlorite is also a byproduct of a complicated mechanism involving the decomposition of ClO_2 in water, it is predominantly produced during NOM oxidation [59], [52]. The reactions that follow (equations (8) – (9)), describe what potentially can happen when ClO_2 dissociates spontaneously:

First, ClO_2 abstracts an electron via the reduction process resulting in chlorite:



The chlorite ion is reduced to form a chloride ion:



These reactions suggest that ClO_2 accepts five electrons and is reduced to chloride (Cl^-). The chlorine atom does not change until stable chloride is formed. Hence formation of chlorinated substances is prevented. Chlorine gas reactions on the other hand are driven by addition and substitution reactions whereby, chlorine atoms incorporate into organic material. The chlorous acid equilibrium reaction produces a fairly low pKa of 1.8 for the chlorite ion (ClO_2^-). However, base ion pair equilibrium reaction pH for hypochlorous acid/hypochlorite has been recorded to fall near pH 7. "This suggests that the chlorite ion (ClO_2^-) exists as the dominant species in drinking water and in the human body [12], [60].

2.7 Decomposition of Alkaline NaClO_2 Solutions

Since NaClO_2 is used to generate ClO_2 (**refer to section 2.3.1**), its decomposition products in aqueous medium are also emphasized below. ClO_2 does not result from the decomposition reaction of NaClO_2 in hot, highly alkaline solution; instead, ClO_3^- is the main by-product of the breakdown reaction [61].



2.7.1 Decomposition of NaClO_2 in Neutral Solutions

Neutral solutions of sodium chlorite are stable. The solution will decompose slowly upon exposure to light and heat [62], [61]. A chlorite ion solution becomes less stable as the pH decreases. The disintegration of a neutral sodium chlorite mixture does not produce ClO_2 , only chlorite (ClO_2^-) and chlorate (ClO_3^-) ions are formed [63].

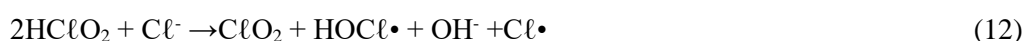
2.7.2 Dissociation of chlorous acid (HClO_2)

Chlorous acid is the least stable of all the chlorine oxyacids and is also occasionally used in the generation of ClO_2 [11]. Its behaviour in aqueous medium is highlighted in equations (11-14). In relation to chlorous acid, the decomposition reaction of chlorous acid is second order.

Bohmlander [64] found that the initial reaction is second order with a delayed transition to a first order reaction [64]. He then proposed the dissociation reaction for chlorous acid. The initial reaction follows the path between pH 0.5 to pH 2 (below):



And:



Followed by:



After 10 minutes, the reaction is no longer rate limiting and is written as:



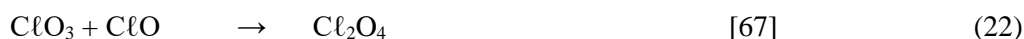
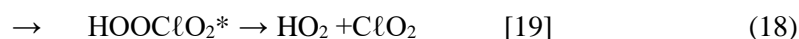
In water, ClO_2 forms a solid polyhydrate [65]. Above about 18.2 °C, the hydrates lose some of their solubility and transform back into their unhydrated form [66]. ClO_2 hydrolyzes relatively slowly, although light and heat speed up the decomposition process. ClO_2 decomposes at 0 and 60°C according to:



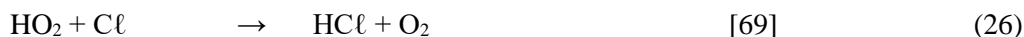
The reaction is accelerated by the hydrogen and chloride ion. There is some evidence that ClO_2 breaks down into chlorine and oxygen [11].

Various reaction schemes have been published for chlorine oxide molecule. “It is commonly assumed that terminal oxygens form dative bonds with chlorine whereas bridging-oxygen-chlorine bonds are covalent [55]. Most studies have focused on photo excitation processes in the stratosphere and not in liquid medium. These reaction schemes are nevertheless represented below to give an idea of possible species formation.

With: M= Molecule (**non-reactive molecule**)



Francisco and Sanders [68] confirmed the existence of a $\text{ClO} \cdot \text{H}_2\text{O}$ radical complex that enhances the formation of a dimer as seen in Equation 24.

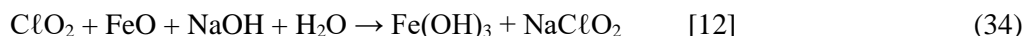


The oxidation of chloride ions in aqueous solutions is known to produce the $\bullet\text{Cl}_2^-$ radical as an intermediate [73], [74].

2.8 Chlorine Dioxide Reaction/Reactivity Properties

ClO_2 will likely react with any inorganic species present in water being treated for potable use. Based on the following selected example reactions (reaction 34 - 38), ClO_2 may likely form insoluble or aqueous species that may be removed by flocculation processes, although beyond the scope of this thesis, these are areas of study that should be looked into when utilising ClO_2 for potable water treatment.

2.8.1 Oxidation of iron



FeO is responsible for influencing the aesthetic property of water as it is capable of imparting undesirable colour to potable water [75]. It can be removed during the flocculation (clarification) process.

2.8.2 Oxidation of manganese



The formation of MnO_2 is noted for its use in water softening process, hence its presence can assist the water purification process substantially [76].

2.8.3 Oxidation of sodium sulphide



Ingestion of potable water with excessive levels of Na_2SO_4 can cause intestinal discomfort leading to diarrhea [77].

2.8.4 Oxidation of nitrogen oxide pollutant



Nitrogen oxide is considered to be flue gas pollutant. ClO_2 can be used to remove this pollutant from flue gas.

2.8.5 Aluminum, magnesium, zinc & cadmium (M) react with ClO_2

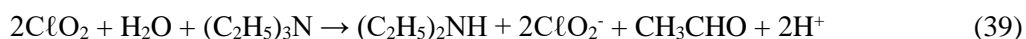


ClO_2 bleaches aluminum, magnesium, zinc & cadmium by oxidizing them

2.8.6 Organic Reactions with ClO_2

When reacting with organic compounds in water, ClO_2 has converted such compounds to aldehydes, carboxylic acids, ketones and quinones [12]. Hydrocarbons chain lengths that exceed eight carbons are easily oxidised by ClO_2 [39].

ClO_2 oxidizes triethylamine in aqueous solution with the production of diethylamine, acetaldehyde, hydrogen ion, and chlorite ion [12].



ClO_2 changes large aromatic and long aliphatic chain organic compounds into tiny and hydrophilic organics, largely destroying the aromatic bond structure of NOM [78].

The interactions between various organic compounds and ClO_2 in water has been tabulated in Table 2.2.

Table-2.2: Interactions of organic compounds with ClO_2 in water

Reactants	Products/ Description of reaction	Reference
$\text{H}_2\text{O} + \text{ClO}_2 + \text{olefins}$	aldehydes, epoxides, chlorohydrins, dichloro-derivatives and chloro-unsaturated ketones	[12], [79]
$\text{H}_2\text{O} + \text{ClO}_2 + \text{ethylenic double bonds}$	ketones, epoxides, alcohols	[12]
$\text{H}_2\text{O} + \text{ClO}_2 + \text{toluene}$	CH_3 , CH_2Cl , CH_2OH	[12]
$\text{H}_2\text{O} + \text{ClO}_2 + \text{anthracene at } 45^\circ\text{C}$	anthraquinone and 1,4-dichloroanthracene	[12]
$\text{H}_2\text{O} + \text{ClO}_2 + \text{phenanthrene}$	diphenic acid and 9-chlorophenanthrene	[12]
$\text{H}_2\text{O} + \text{ClO}_2 + 3,4\text{-benzopyrene}$	quinones, traces of chlorinated benzopyrene (no longer considered carcinogenic)	[12]
$\text{H}_2\text{O} + \text{ClO}_2 + \text{aldehydes}$	carboxylic acids	[12]
$\text{H}_2\text{O} + \text{ClO}_2 + \text{ketones}$	alcohols	[12]
$\text{H}_2\text{O} + \text{ClO}_2 + \text{aliphatic amines: tertiary}$	rupture of CN bond, no N-oxides formed	[12], [79]
$\text{H}_2\text{O} + \text{excess ClO}_2 + \text{phenol}$	Para-benzoquinone and 2-chlorobenzoquinone	[12], [79]
$\text{H}_2\text{O} + \text{ClO}_2 + \text{phenol}$	maleic acid and oxalic acid	[12], [79]
$\text{H}_2\text{O} + \text{ClO}_2 + \text{thiophenols}$	sulfonic acids	[12]
$\text{H}_2\text{O} + \text{ClO}_2 + \text{tocopherol}$	demethylated derivatives	[12]
$\text{H}_2\text{O} + \text{ClO}_2 + \text{anhydrides}$	no reaction but catalyses hydrolysis reactions	[12]
$\text{H}_2\text{O} + \text{ClO}_2 + \text{amino acids: glycine, leucine, serine, alanine, phenylamine, valine, hydroxyproline, phenylaminoacetic acid, aspartic, glutamic acids}$	little, or no reaction	[12]
$\text{H}_2\text{O} + \text{ClO}_2 + \text{amino acids containing sulphur}$	reactive	[12]

Table 2.2 continued

Reactants	Products/ Description of reaction	Reference
H ₂ O + ClO ₂ + methionine sulfoxide	sulfone	[12]
H ₂ O + ClO ₂ + aromatic amino acids	reactive	[12]
H ₂ O + ClO ₂ + tyrosine	dopaquinone, dopachrome	[12], [22]
H ₂ O + ClO ₂ + tryptophan	idoxyl, isatine, indigo red and trace chlorinated products	[12], [22]
H ₂ O + ClO ₂ + thiamine	slow reaction	[12]
H ₂ O + ClO ₂ + keratin	hydrosoluble acids	[12]
H ₂ O + ClO ₂ + carbohydrates CHO and CH ₂ OH	carboxylic functions	[12]
H ₂ O + ClO ₂ + pectic acid	mucic acid, tartaric acid and galacturonic acid	[12]
H ₂ O + ClO ₂ + chlorophyll and plant dyes	colour removed	[12]
H ₂ O + ClO ₂ + latex and vinyl enamels	delays polymerisation	[12]
H ₂ O + ClO ₂ + biacetyl	acetic acid, carbon dioxide	[12]
H ₂ O + ClO ₂ + 2,3-butaneodiol	acetic acid and carbon dioxide	[12]
H ₂ O + ClO ₂ + cyclohexene	aldehydes, carboxylic acids, epoxides, alcohols, halides, dienes and ketones	[12]
H ₂ O + ClO ₂ + cyanides	oxidised	[12]
H ₂ O + ClO ₂ + nitrites	oxidised	[12]
H ₂ O + ClO ₂ + sulphides	oxidised	[12]
H ₂ O + ClO ₂ + unsaturated fatty acids and their esters	typically undergo double bond oxidation	[12]
H ₂ O + ClO ₂ + phenols	quinones, malonic acid and oxalic acid	[80], [81], [82]
H ₂ O + ClO ₂ + lignins	quinones and carboxylic acids	[29], [83]
H ₂ O + ClO ₂ + humic acid	quinones and carboxylic acids	[29], [83]
H ₂ O + ClO ₂ + methoxylic groups (complex organic groups)	aliphatic hydroxylic groups	[84], [85]
H ₂ O + ClO ₂ + methoxylic groups	aldehydes, ketones and esters (oxidises aliphatic complexes between the p-electron system of the olefin fragments)	[83], [84], [85]

2.8.7 *Non-Reactive Species*

ClO_2 does not react with benzene, naphthalene, ethanol, maleic acid, crotonic acid, hippuric acid, cinnamic acid, betaine, creatine, alanine, phenylalanine, valine, leucine, asparaginic acid, asparagine, glutamic acid, serine, hydroxyproline and taurine [12], [85]. No reaction takes place when ClO_2 is in contact with aliphatically combined NH_2 groups, amido and imido compounds, HO groups in acids and alcohols [12], [83]. ClO_2 does not react with free CO_2H groups or CO_2H groups that have been esterified in a medium of mono and polybasic acids, nitrile groups, CH_2 groups that are present in a homologous series, ring structures such as C_6H_6 , C_{10}H_8 , cyclohexane, and the salts of $\text{C}_5\text{H}_5\text{N}$, quinoline and piperidine [12].

Under normal water treatment conditions, the majority of the aliphatic and aromatic hydrocarbons will not react with ClO_2 . The exception to this rule only relates to the hydrocarbons containing specific reactive groups. These reactive groups include phenol type compounds, secondary and tertiary amines, organic sulphides and certain hydrocarbon polycyclic aromatics such as benzopyrene, anthracene and benzoanthracene [84].

2.9 Water Treatment Processing

In the mid to late 1970s, numerous studies linked higher cancer mortality rates to the chlorination of potable water [60]. Trihalomethane (THM) concentrations, specifically chloroform, were found to be associated with an increase in cancer mortality [86], [87]. The formation of chloroform was a direct consequence of the reaction of chlorine and naturally occurring organics. As a result of this research, the USEPA established a maximum THM containment level for drinking water of 0.1 ppm.

ClO_2 is becoming more widely used in the water treatment industry. The first reported water treatment plant to incorporate ClO_2 was in Niagara Falls in 1944 [88]. In 1986, approximately 200-300 USA potable water treatment process plants introduced ClO_2 into their treatment cycles, whilst thousands of ClO_2 applications were recorded in Europe [51], [22].

In an aqueous medium, the species aids processes that drive coagulation and the removal of turbidity [89], [90] and prevents the formation of THMs [91].

The Poznan Water Treatment and Sewage Co in Poland uses ClO_2 as a pre-oxidation step before granular activated carbon (GAC) filtration [92]. This plant has shown that even low doses of $0.2 \text{ mg ClO}_2 \text{ L}^{-1}$ can have a substantial impact on the extent to which NOM is absorbed, which in turn improves the capacity of GAC filters [92]. The molecular weight (mw) of NOM showed very slight variations. The larger molecular weight compounds were found to be more inclined to react with ClO_2 [93].

2.9.1 Advantages and Disadvantages of the Use of ClO_2 in Water Treatment

There are several advantages associated with the use of ClO_2 in water. A few main advantages are listed below:

- Giardia, Cryptosporidium, and other viruses are more effectively inactivated by ClO_2 than by chlorine and chloramines [94]. Also see section 2.4.
- ClO_2 oxidises iron, sulphides and manganese. [Eq. 34 -38]
- ClO_2 may enhance the clarification process.
- ClO_2 is easy to generate [Refer to section 2.3.1]
- The pH of the system has no impact on the biocide capabilities of ClO_2 [94].
- ClO_2 provides residuals for long-term disinfection because it does not hydrolyze when it enters water, but instead persists as a dissolved gas in solution and can thus act more efficiently over time [95]. ClO_2 is also more soluble than chlorine gas in water [11].

Refer to section 2.5

Disadvantages

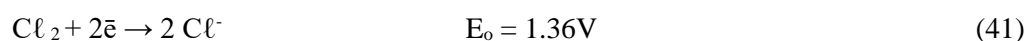
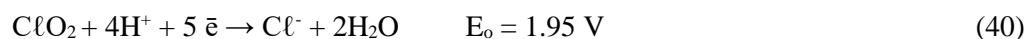
- The ClO_2 process forms the specific by-products chlorite (ClO_2^-) and chlorate (ClO_3^-). (These by-products are both highly toxic compounds and require Occupational Health Care during scaled processes) [60].
- The costs of sampling and laboratory testing for chlorite and chlorate are exorbitant [51], [12].
- Rental of generators and cost of the sodium chlorite are excessive [51].
- Safety precautions are required whilst measuring chlorine dioxide gas, since it is explosive [51], [12]
- ClO_2 decomposes in sunlight, so it must be produced in close proximity to the application [94].

2.9.2 Toxicity of ClO_2

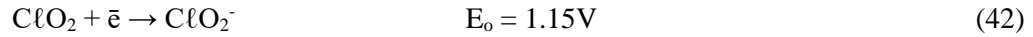
Chlorite (ClO_2^-) causes hemolytic stress in animals when consumed in food at low levels of 50 ppm. [23]. The drinking water standard for total ClO_2 , chlorite (ClO_2^-) and chlorate (ClO_3^-) was therefore limited to 1.0 ppm, taking into account individuals prone to oxidative stress [7], [96], [23].

2.9.3 ClO_2 Residuals

Monitoring of ClO_2 residuals in a water treatment plant environment can be problematic. The Palin DPD method [97] is used to measure ClO_2 residuals. This test usually gives a higher residual reading for ClO_2 than for chlorine. According to these results, one would expect chlorine to have a higher oxidizing potential than ClO_2 . This, however, is not the case with respect to reactions (40) and (41).



Hence ClO_2 is the stronger oxidising agent. To clarify the discrepancy in findings acquired using Palin's DPD method and reactions (42) and (43), one must make the assumption that ClO_2 is reduced as follows:

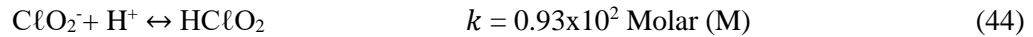


ClO_2 has two distinct oxidising capacities [98], [10].

When there are adequate hydronium ions present, ClO_2 proceeds to reduce to chlorite (equation 39), which in turn reduces to chloride (equation 44).

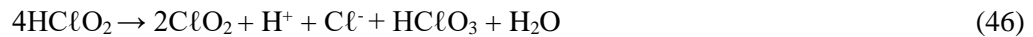
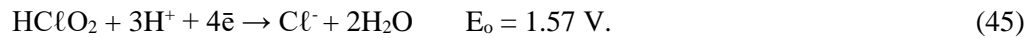


The conversion of chlorite to chloride only occurs at low pH as described in reaction (4). Chlorite undergoes protolysis to produce chlorous acid at pH 6–8 (reaction 44) [99].



The equilibrium concentrations of chlorous acid and chlorite at pH 7 are 1:100,000 [98].

Chlorite is only partially protolyzed. If chlorous acid prevails, it might be disproportionate to products as shown in reaction (41).



The absence of chlorate (ClO_3^-) following the addition of ClO_2 [98] is possibly due to analytical challenges in detecting trace levels of chlorate in close proximity of other chlorine containing species. However, if chlorous acid is present in water with a pH of 6 to 8, a disproportionation reaction is perhaps most likely to occur (reaction 46). As a result, the situation above suggests that Palin's DPD approach produces chlorite residuals rather than ClO_2 residuals.

The following can be postulated regarding the reactivity of ClO_2 in water with oxidizable organic matter at pH 6–8, according to Myhrstad, and Samdal [98], Nawrocki et al, [100]:

1. A reduction of ClO_2 to chlorite (reaction 38)
2. Protolysis to a very limited extent of chlorite to chlorous acid (reaction 40)
3. Disproportionation of chlorous acid in ClO_2 and chloride (reaction 42)

The above would apply to most South African raw water treatment processes. As a result, ClO_2 degrades in two stages:

1. The conversion of ClO_2 to chlorite

2. Chlorite ion reduction occurs when the reductant produces a low redox potential and exists in the presence of excess hydrogen ions [101], [12].

2.10 Primary Photo Processes of Electronically Excited OClO in Water Solution

The symmetric **OClO** and the asymmetric **ClOO** are the two isomeric forms of ClO₂. The ClOO isomer is extremely reactive, thermodynamically more stable than OClO by around 3 kcal/mol [102], [55], [103] and challenging to examine spectroscopically.

OClO is kinetically stable at ambient temperature and does not react thermally with water over several days [104], [105]. Excitation of the ClO₂ molecule usually occupies three reactive states namely: ²B₂, ¹A₁ and ²A₁. The two competitive pathways depicted in reactions 47 & 48, represents the photochemistry of excited-states of ClO₂:



There has been limited research on the rate at which OClO decays via isomerization in aqueous medium and the potential impact of solvent on its reactivity [106], [107]. Given the dynamic nature of the solvent cage, a branching ratio is anticipated to exist between the two pathways. Both the liquid's static and dynamic qualities affect the branching ratio. Since the charge distribution of the reactive ²B₂ and ²A₁ states differs from one another and from the ²A₂ state with Franck-Condon population, this interaction will be very sensitive to the liquid's dielectric properties [102], [108].

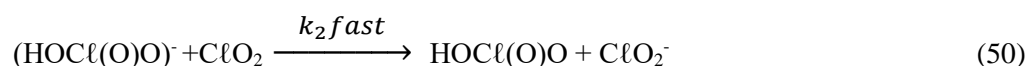
The photochemistry of OClO yields the following chemical species in the absence of a solvent reaction: ClO, O, Cl, O₂ and ClOO. When photochemistry of OClO in water takes place at ambient temperature, excitation of the ²A₂ - ²B₁ absorption band at 355 nm occurs. 90% of the OClO molecule undergoes dissociation and 10% undergoes isomerisation [109], [110]. Competitive bond dissociation creates ClO + O and isomerization creates ClOO from the excited OClO molecule. The isomerized product that thermally dissociates into Cl and O₂ has a first order reaction rate constant of 6.7 x 10⁹ s⁻¹ at ambient temperature [106]·[20].

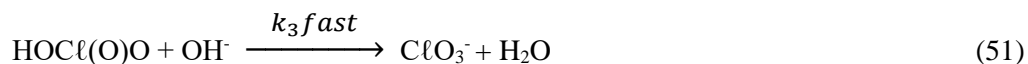
2.11 Decomposition and Mechanism of ClO₂ in Basic Medium

ClO₂ decomposes rather slowly in neutral aqueous solutions [80], whereas the presence of a base increases the rate of decay [66], [111]. The disproportionation reaction is presented in reaction (6).

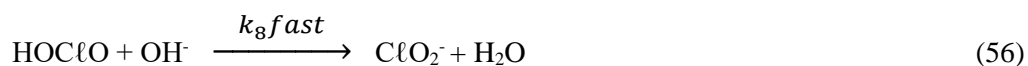
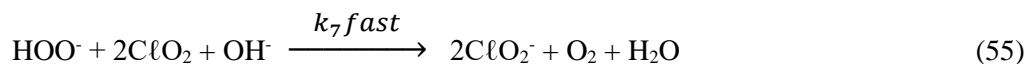
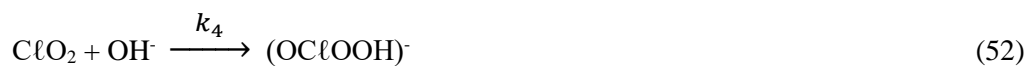
Decomposition pathways [112]:

- Pathway 1: First – order in [ClO₂]: products: [ClO₂⁻], [ClO₃⁻]

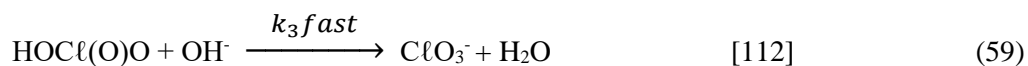
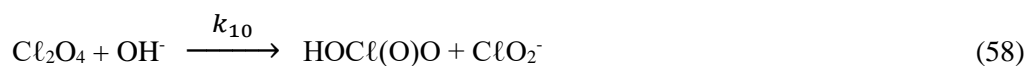




- Pathway 2: First-order in $[\text{ClO}_2]$: products: ClO_2^- and O_2



- Pathway 3: Second – order in $[\text{ClO}]$: products: $[\text{ClO}_2^-] = [\text{ClO}_3^-]$



Reaction Mechanism

$(\text{HOCl(O)O})^-$ is produced by pathway 1 when OH^- adducts with the Cl in ClO_2 . Similar adducts have been postulated for $\text{ClO}_2 + \text{HO}_2^-$ reactions [113]. The reaction between Cl and OH is weak but considerable, according to an ab initio computation involving the $(\text{HOCl(O)O})^-$ adduct [114]. ClO_2^- and HOClO_2 are produced in the next subsequently rapid electron-transfer step (k_2), which also rapidly produces OH^- (k_3) to give ClO_3^- . Equimolar ClO_2^- and ClO_3^- are formed in a first order reaction involving $[\text{ClO}_2]$ and $[\text{OH}^-]$.

An electron transfer that is facilitated by the presence of base occurs during the k_2 step of the reaction [112]. It is assumed that pathway 2 involves OH^- combining with a ClO_2 oxygen atom to produce an adduct to produce the reactive intermediate $(\text{OClOOH})^-$ (k_4). A weak link exists between an OOH segment and an OCl segment, the latter of which has a net charge of -0.553 (Figure 2.3).

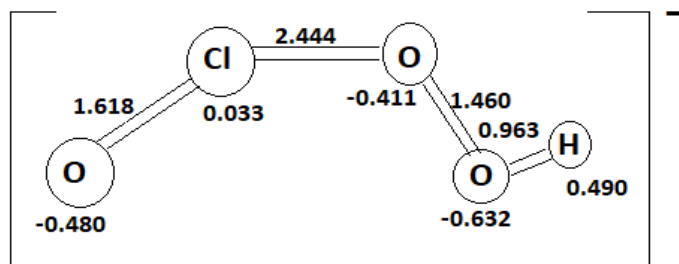


Figure-2.3: Equilibrium geometries (bond distances in Å) and atomic charges for the steady-state species (OClOOH). [112]

This adduct can swiftly transfer electrons to a second ClO_2 (k_5) to produce ClO_2^- and OClOOH. HOClO and HOO^- (k_6) are formed when the latter species reacts favourably with OH. The interaction of HOO^- and ClO_2 results in ClO_2^- and O_2 , wherein k_7 denotes a succession of steps [115]. The stoichiometry for the pathway 2 reaction is presented in reaction (56). Further substantiation of processes using a base-assisted electron transfer are illustrated by the reactions in steps k_4 and k_5 . The formation of $(\text{OClOH})^-$ is a potential intermediate, according to *Ab Initio* calculations (Figure-2.4).

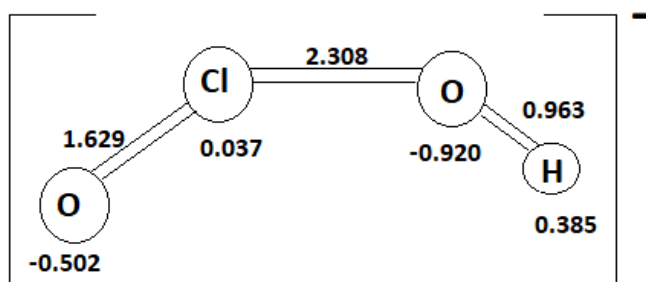


Figure-2.4: Equilibrium geometries (bond distances in Å) and atomic charges for the steady state species $(\text{OClOH})^-$. [112]

A second-order reaction in ClO_2 depicted in pathway 3 is shown to proceed through an intermediate Cl_2O_4 which is in pre-equilibrium with two ClO_2 molecules (k_9/k_{-9}). The Cl_2O_4 intermediate is similar to the BrO_2^- , ClO_2 [114] and Br_2O_4 intermediates hypothesised by Halperin and Taube [116]. Another instance of a base-assisted electron transfer occurs in step k_{10} when Cl_2O_4 reacts with OH^- , this time between two weakly linked ClO_2 molecules.

2.12 Chlorine Oxide Species (This study)

Sixty (60) chlorine oxide species have been identified for the purpose of this study. Heat of Formation values available in open literature were measured using various UV spectroscopic and quantum techniques i.e., W8 theory, G2, G3 and G4 theory and various *Ab Initio* methods (**Refer to Chapter 4 of this manuscript**).

The chemical formula, structure, name and published Heat of Formation (ΔH_f kcal/mol at 298 K) of the sixty (60) chlorine oxide species are tabulated in Table 2.3. Active Thermodynamic Tables

(ATcT) [117] offer a comprehensive reference for many of these chlorine oxides. For the purposes of this study, a new structural naming convention has been adopted, to accent particular atoms, in highlighting their bonding sequence and conformation/configuration within the structure. The chlorine oxide species presented in Table-2.3 will be examined in this study.

Table-2.3: Structure, formulae and names of selected chlorine species

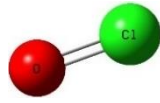
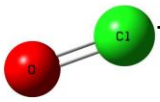
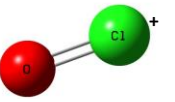
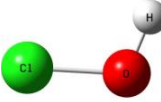
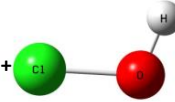
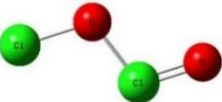
Formula	Chemical Name	Structure	ΔH_f (kcal mol ⁻¹) at 298K	Reference (ΔH_f)
ClO (g)	Chloro-oxidanyl		24.31	[117]
[ClO] ⁻ (g)	Hypochlorite		-28.26	[117]
[ClO] ⁺ (g)	Oxochloronium		274.84	[117]
HOCl (g)	Hypochlorous acid		-18.35	[117]
[HOCl] ⁺ (g)	Hypochlorous acid cation		238.01	[117]
ClOClO (g)	Chlorine chlorite		39.77	[117]

Table-2.3 continued

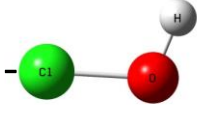
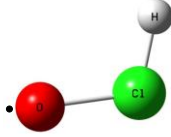
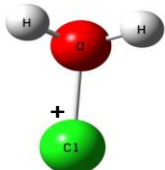
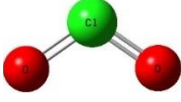
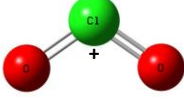
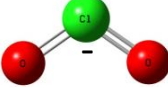
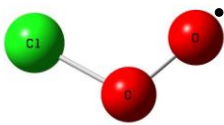
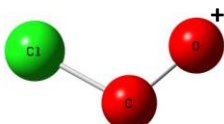
Formula	Chemical Name	Structure	ΔH_f (kcal mol ⁻¹) at 298K	Reference (ΔH_f)
[HOCl] ⁻ (g)	Hypochlorous acid anion		47.41	[117]
HClO (g)	Chlorosyl hydride		33.24	[117]
[ClOH ₂] ⁺ (g)	Aquachlorine cation		194.66	[117]
OClO (g)	Chlorine dioxide		24.36	[117]
[OClO] ⁺ (g)	Chloryl ion		262.13	[117]
[OClO] ⁻ (g)	Chlorite		-25.74	[117]
ClOO (g)	Chlorodioxidanyl		24.55	[117]
[ClOO] ⁺ (g)	Chlorodioxidenium		286.71	[117]

Table-2.3 continued

Formula	Chemical Name	Structure	ΔH_f (kcal mol ⁻¹) at 298K	Reference (ΔH_f)
[ClOO] ⁻ (g)	Peroxyhypochlorite		-59.28	[117]
HOClO (g)	Chlorous acid		4.94	[117]
[HOClO] ⁺ (g)	Chlorous acid cation		236.81	[117]
[HOClO] ⁻ (g)	Chlorous acid anion		-37.33	[117]
HOOCℓ (g)	Peroxyhypochlorous acid		-0.31	[117]
[HOOCℓ] ⁺ (g)	Peroxyhypochlorous acid cation		244.86	[117]
[HOOCℓ] ⁻ (g)	Peroxyhypochlorous acid anion		-8.12	[117]
HClOO (g)	Chloryl hydride		191.20	[117]

Table-2.3 continued

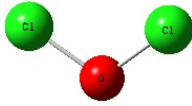
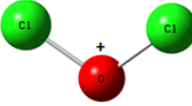
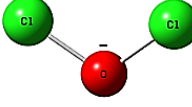
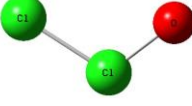
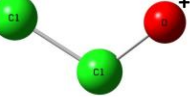
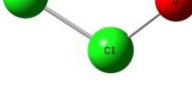

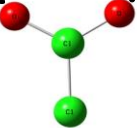
Formula	Chemical Name	Structure	ΔH_f (kcal mol ⁻¹) at 298K	Reference (ΔH_f)
ClOCl (g)	Chloro hypochlorite		18.63	[117]
[ClOCl] ⁺ (g)	mu-Oxodichlorine cation		269.31	[117]
[ClOCl] ⁻ (g)	mu-Oxodichlorate anion		-35.30	[117]
ClClO (g)	Chlorosyl chloride		31.83	[117]
[ClClO] ⁺ (g)	Chlorosyl chloride cation		270.26	[117]
[ClClO] ⁻ (g)	Chlorosyl chloride anion		-44.14	[117]
ClOOCl (g)	Chlorooxy hypochlorite		31.38	[117]
ClClO ₂ (g)	Chloryl chloride		29.18	[117]

Table-2.3 continued

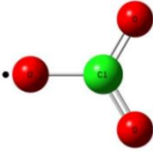
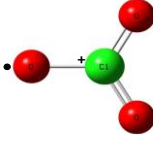
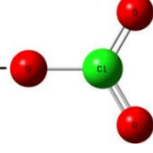
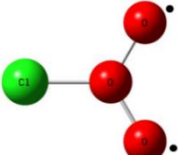
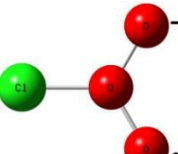
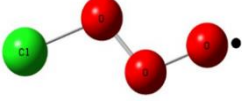
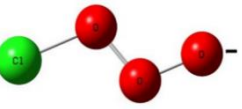
Formula	Chemical Name	Structure	ΔH_f (kcal mol ⁻¹) at 298K	Reference (ΔH_f)
ClO ₃ (g)	Perchloryl		44.00	[117]
[ClO ₃] ⁺ (g)	Perchloryl cation		297.67	[117]
[ClO ₃] ⁻ (g)	Chlorate		-50.86	[117]
Cl(O)O ₂ (g)	2-chloro ozone		63.36	[117]
[Cl(O)O ₂] ⁻ (g)	2-chloro ozone anion		-25.22	[117]
ClOOO (g)	1-Chloro ozone		53.32	[117]
[ClOOO] ⁻ (g)	1-Chloro ozone anion		-26.62	[117]

Table-2.3 continued

Formula	Chemical Name	Structure	ΔH_f (kcal mol ⁻¹) at 298K	Reference (ΔH_f)
OCℓOO (g)	Chlorine oxide peroxide		54.34	[117]
[OCℓOO] ⁻ (g)	Peroxychlorite		-8.19	[117]
HOℓO ₂ (g)	Chloric acid		-1.19	[117]
HℓO ₃ (g)	Chlorine hydride oxide		37.11	[117]
HOOCℓ (g)	Hypochloro-peroxoous acid. Hydroxy ester		10.10	[117]
HOOCℓO (g)	Peroxychlorous acid		21.55	[117]
ℓO ₄ (g)	Perchloryloxy		57.60	[117]
[ℓO ₄] ⁻ (g)	Perchlorate		-64.99	[117]

Table-2.3 continued

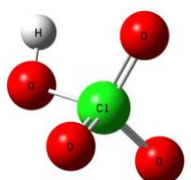
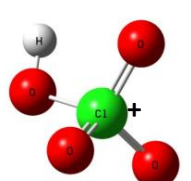
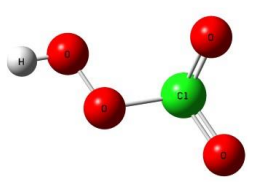
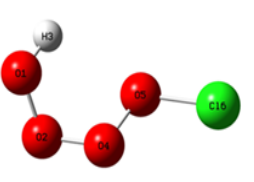
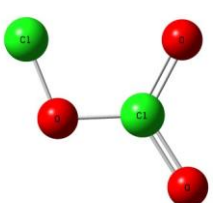
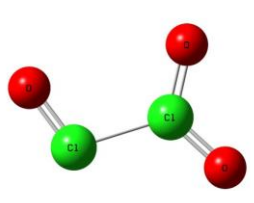
Formula	Chemical Name	Structure	ΔH_f (kcal mol ⁻¹) at 298K	Reference (ΔH_f)
HOClO ₃ (g)	Perchloric acid		0.43	[117]
[HOClO ₃] ⁺ (g)	Perchloric acid cation		282.55	[117]
HOOCLO ₂ (g)	Peroxychloric acid		16.56	[117]
HOOOOCl (g)			22.7	[119]
ClOClO ₂ (g)	Dichlorine trioxide		37.40	[117]
OClClO ₂ (g)	Chlorineoxide		46.2	[119]

Table-2.3 continued

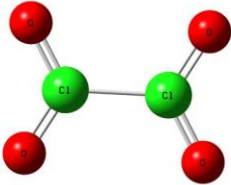
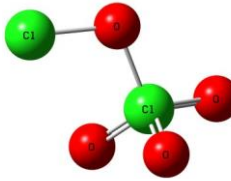
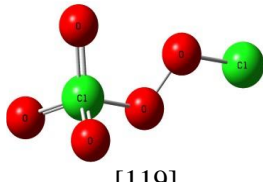
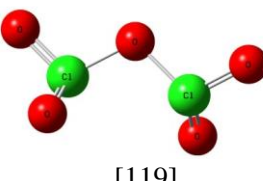
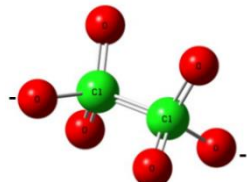
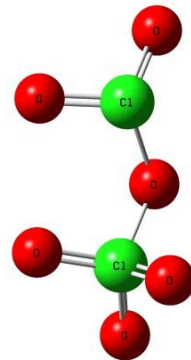
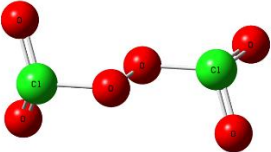
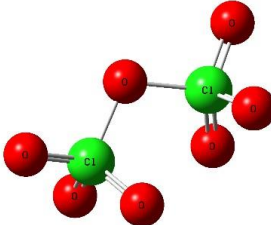
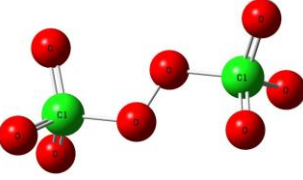
Formula	Chemical Name	Structure	ΔH_f (kcal mol ⁻¹) at 298K	Reference (ΔH_f)
ClO_2ClO_2 (g)		 [72]	Not reported	Not analysed
ClOClO_3 (g)	Chlorine perchlorate and (Chlorooxy) chlorane trioxide		37.40	[118]
ClOOClO_3 (g)	Dichlorine Pentoxide	 [119]	69	[118]
$\text{ClO}_2\text{-O-ClO}_2$ (g)	Chloryl chlorate	 [119]	Not reported	Not reported
$[\text{ClO}_3\text{ClO}_3]^{-2}$ (g)	Oxido-[oxido(dioxo)- lambda7- chloranylidene]-dioxo- lambda7-chlorane	 [119]	Not reported	Not reported
$\text{ClO}_2\text{-O-ClO}_3$ (g)	Chloryl perchlorate		72.4	[118]

Table-2.3 continued

Formula	Chemical Name	Structure	ΔH_f (kcal mol ⁻¹) at 298K	Reference (ΔH_f)
ClO ₂ -O-O- ClO ₂ (g)	Chloryloxy chlorate	 [119]	Not reported	Not reported
ClO ₃ -O-ClO ₃ (g)	Chlorine heptoxide	 [119]	86.2	[118]
ClO ₃ -O-O- ClO ₃ (g)	Perchloryloxy perchlorate	 [55]	111.48*	[55]

*Heat of formation (ΔH_f ClO₃-O-O-ClO₃) calculated from energies published by Beltran et al. (1999) [55].

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CHAPTER 3: CHEMICAL APPROACH AND COMPUTATIONAL STRATEGY

3.1 Introduction

Some of the chlorine dioxide (ClO_2) species investigated, are classified as ionic and radicals and are involved in extensive intra- and inter-molecular interactions in aqueous medium. Although Heats of Formation for most of the species were available from the open literature, it was important to construct chemical structures and derive their optimum geometries, validated against reported energies to be applied in all subsequent calculations. It was therefore important to elucidate their thermochemical properties, applying an empirical supported first principal computational approach.

The study involved a few technical stages:

- a) Force field parameter preparation
- b) Model preparation
- c) Model optimization
- d) GCMC Canonical ensemble simulations (radical and neutral species only)
- e) Ab Initio and semi-empirical derivation of Heats of Formation
- f) Species based thermodynamics to extract reaction schemes.

Simulations performed on extended (Grand ensemble) model representations, have been undertaken to determine the optimum potential energy model stages. This was followed by semi-empirical and *Ab Initio* quantum chemical optimizations (*MOPAC* and *VASP* software) to derive thermochemical properties.

This was done to verify the authenticity and achievable electronic structures of the compounds, against published energy figures and furthermore correlate their ensemble chemical properties against single molecular figures, as proof of a technique to determine organic compound characteristics.

Model preparation Stage

A group of chlorine oxide species listed towards the end of Chapter 2, was selected for this study. The group comprised of neutral, radical and ionic species (+ve and -ve) with some extended compounds, hosting a higher oxygen content, including a number of hydrogen derivatives of ClO_2 .

- a) Periodic models were constructed using the MedeA-3.3.1 software, followed by a *VASP* (DFT) refinement cycle (conditions listed), used in all further software simulations
- b) Molecular models were prepared with the *GaussView-6* model preparation interface of the *Gaussian-16* software suite

3.1.1 *Initial Stage*

The empirical *pcff+* [8] force field, lacked some parameters for a number of the chlorine oxide species, in support of *Grand Canonical* ensemble simulations and several force field parameters were “borrowed” and derived from the *cvff* and *pcff* force field repositories to complement the *pcff+* set.

All adjusted *pcff+* force field entries are presented in Tables- (5.1–5.6).

3.1.2 *GEMC Simulation stage*

For all neutral and radical samples, *GIBBS* GEMC (GCMC) ensemble model simulations were undertaken, to portray realistic and effective inter-molecular interactions that take place between species.

3.1.3 *Semi-empirical Stage*

The GEMC models extracted at the minima Internal Energy (U) of the simulated ensemble and beyond the minimum Saddle Points for each of the neutral species (considered to be their optimum model sizes) were subjected to *MOPAC* (semi-empirical quantum) analyses, to determine their ΔH_f (reported as for single species). The ΔH_f of the simulated optimum ensemble models, was computed applying all seven Hamiltonians of the *MOPAC-16* software.

3.1.4 *Ab Initio simulation Stage*

Ensemble models of structures from the *GIBBS Grand Canonical* ensemble outcome, were transferred to the *VASP* and *MOPAC* software, for a final ‘bulk’ structural optimization at the electronic level. Heats of Formation (ΔH_f) derived from these simulations were verified against some of the available, literature referenced energies, as validation of their geometries and stereochemistry. Charged ClO₂ species, not possible to be exposed to force field simulations, were optimized through *Gaussian-16* calculations, applying B3LYP theory and the cc-pv5z basis set, to determine their Heats of Formation as single molecular entities.

All these species have computationally been treated in the gas phase, for the thermodynamic derivations to be reported as single species (molecules) in a vacuum environment. It remains to compensate for coulomb and dipole-dipole interactions, van der Waals dispersion and inter-species hydrogen bonding in an attempt to present a true representation of their energies of formation.

3.1.5 *Final Stage*

Finally, the *FactSage-7.3* species-based thermodynamic software was used to determine the prevalent species of chlorine dioxide in an aqueous medium. A consolidated set of chemical reaction schemes were derived to demonstrate the major reactions involved and highlights specific product species, considered to be harmful during scaled water purification processes.

3.2 Computational Software Programs

The software programs and related techniques that were used to refine and optimize model structures and determine thermochemical properties, are listed below:

- *MedeA (GIBBS-9.7.4)* 1998-2017 -- *GIBBS* ensemble Monte Carlo simulation. This statistical-mechanical method predicts the equilibrium conditions of interaction at a molecular level [1], confined to periodic simulation space.
- *MedeA (VASP-6.2.1)* --The Vienna *Ab Initio* Simulation Package (*VASP*) was used to conduct DFT electronic structure refinement, for to derive at thermochemical properties [2]
- *MedeA (VASP-5.4)* –*Mulliken* molecular partial charge distributions
- *MedeA (MOPAC-2016, v.17.048)* – Semi-empirical quantum chemical software system which incorporates molecular orbital theory, combined with force field structural support. The system was used to calculate thermochemical properties of species and Bader (partial) charges [3]
- GaussView-6 -- A software modelling interface, applied to prepare Gaussian models for numeric analysis [4]
- *Gaussian-16, Rev.C.01* -- Computational software to conduct *Ab Initio* single molecular calculations to predict Thermochemical properties and structure refinement [5]
- *FactSage-7.3* -- Thermochemical software package to predict water-based species interactions [6] and extract chemical reaction schemes.

3.3 GIBBS-9.7.4 (GIBBS ensemble Monte Carlo)

Single monomeric chlorine oxide species were exposed to an Isobaric-Isothermal **GIBBS** ensemble (GEMC) simulation, applying Grand Canonical Monte Carlo (GCMC) methodology. This represents a statistical mechanical periodic ensemble at constant T and P, where the number of molecular entities (introduced as single molecular periodic models) N , is preserved throughout each simulation cycle. It is frequently referred to as an NPT ensemble [7] but in this instance, with a stepwise increment of single species to a maximum pre-defined population. The objective being to reach a sustained but minimum potential energy, which ultimately should present the optimum model size. The initial Grand Scale periodic cell dimensions were permitted to be adjusted as well. NPT ensembles were constructed, permitting between five to one hundred molecules in a single global periodic environment.

The Potential Energy (U_i) for the ensemble model was determined at each optimum step, followed by further additions of species and the trajectory energy graphically displayed to determine the minimum (optimum) internal energy point. This technique was used to mimic chemically realistic inter- and intra-molecular interactions in a confined chemical space (at the molecular dynamics level of simulation), as opposed to merely analysing them as single molecular structures. The optimum

ensemble model step was extracted and subjected further to **VASP-6.2.1** [2] and **MOPAC-2016** [3] refinement and analyses, to obtain thermochemical properties.

Table-3.1: *GIBBS* (GEMC) Conditions Applied for GCMC Simulation

<i>GIBBS</i> Execution Conditions	
Initialize: Non-Bond	Lennard Jones (6-9)
Electrostatic Sums:	Ewald
ensemble: Fixed (\$) T,a P	NPT (298K, 1 Atm.)
Steps (Per species addition):	1000
Periodic initial Cell size:	10x10x10 Å
Internal moves:	0.2 Å
Pivoting around centre:	0.2 Å

The process flow that was adopted to run *GIBBS* Monte Carlo and the parameters activated at each stage, are tabulated below:

Initialise
Non-bond Lennard Jones (6-9)
Electrostatics: Ewald
Random seed: -1

Single Phased properties
ensemble: NVT
temperature: \$T

Run	
N- Steps	1000
Intervals	
Writing the main averages	1000
Writing the configuration	10000
Adjusting displacement amplitude	20000
Writing the restart file	-1
Steps between checking for stopping	-1
Writing intermediate configurations	10
Writing the trajectory file	10

Move Probabilities	
Internal moves	
Internal moves	0.2
internal rotation	0.2
pivoting around the centre	0.2
reptation of linear molecules	0
concerted rotation	0
double bridging	0
internal double rebridging	0
stretching (atomic translation)	0
Insertion test insertions and deletions	
	0
Rigid Moves	
rigid translation:	0.2
rigid translation and rotation	0
rigid rotation	0
Exchange two molecules	0

Figure-3.1: MedeA GIBBS Monte Carlo (GEMC) Process Flow

3.4 Gaussian-16 (Single molecular optimization)

The structures of all the species in this study were first optimised using *Hybrid Density Functional Theory* (DFT) applying a small basis set of: 6-311++G (3d2f,3p2d). The B3LYP method (Becke's three-parameter nonlocal exchange functional with the correlation functional of Lee, Yang, and Parr) using basis set (cc-pv5z) was then applied to determine optimum species energies.

3.4.1 DFT Approach -- Gaussian

Three (3) options are presented to compute DFT frontier molecular orbitals:

- Unrestricted Hartree Fock (UHF) method
- Restricted Hartree Fock (RHF) method
- Restricted Open Shell Hartree Fock (ROHF) approach

UHF uses a separate orbital for each electron, even if they are paired (in this study this was applied to ions, excited states and radicals) [9]. Energies and optimisation of geometries were performed at DFT (Density Functional Theory) levels, using Pople's higher basis set (aug-cc-pv5z) and Dunning's correlation.

3.4.2 DFT and Ab Initio structure refinement

The *Gaussian-16* software supports a non-periodic (Molecular) model environment. All single model species were prepared with the *GaussView-6* software [10], [11]. For *Gaussian-16* calculations, theory level CCSD (T) has shown to provide for the most accurate thermochemical properties [10], but is computationally resource demanding, especially when combined with higher basis sets [12]. Computational conditions applied for *Gaussian-16* analyses, are presented in Table- 3.2

Table-3.2: Executional Conditions Applied for *Gaussian* Analysis

<i>Gaussian</i> Execution Conditions	Selection
Optimization Conditions:	OPT selected plus FREQUENCY
Temperature:	298.15K
Pressure:	1 atm
Spin polarization: Restricted Open Shell HF	ROHF : Charged and Radical Species
Spin polarization: Unrestricted Hartree Fock	UHF : Open shell, Radical, Charged species
Spin polarization: Restricted Hartree Fock	RHF : Neutral Species
Theory:	Hartree Fock
basis set:	cc-pV5Z
Charge:	To specify for charged species
Theory:	DFT
Functional:	B3LYP, CCSD
basis set:	cc-pV5Z
Spin:	Selected

It was furthermore required to apply an additional calculation step (externally applied to *Gaussian* derived electronic energies [13]) to obtain the Heats of Formation (ΔH_f 298) and is briefly outlined below and in the Appendix.

A similar approach was followed to determine Gibbs Free Energy of Formation (ΔG_f 298).

3.4.3 Heats of Formation Calculation (Transformed *Gaussian-16* Computations)

The enthalpies of formation of the species was calculated at **0K** [$\Delta H_f(0K)$] and **298K** [$\Delta H_f(0K)$]

First calculate $\Delta H_f(M, \mathbf{0K})$ for each molecule [14]:

$$\begin{aligned}\Delta H_f(M, \mathbf{0K}) &= \sum_{atoms} x \Delta H_f(X, 0K) - \sum D_0(M) \\ &= \sum_{atoms} x \Delta H_f(X, 0K) - [\sum x \epsilon_0(X) - \epsilon_0(M) - \epsilon_{ZPE}(M)]\end{aligned}$$

Where:

$$M = \text{Molecule}$$

X	=	Element which makes up <i>Molecule</i> (M)
x	=	The number of atoms of X in M
$\sum D_0(M)$	=	Atomization energy of the molecule
(ϵ_0, M)	=	Total energies of the molecule
$\epsilon_{ZPE}(M)$	=	Zero-point energy of the molecule [14]

The $\Delta H_f(M, \mathbf{0K})$ energy obtained was then used to calculate $\Delta H_f(M, \mathbf{298K})$ for each molecule [14]:

$$\Delta H_f(M, \mathbf{298K}) = \Delta H_f(M, 0K) + [\Delta H_{fM}(298 K) - \Delta H_f(M, 0K)] - \sum_{atoms x} [\Delta H_{fx}(298 K) - \Delta H_{fx}(0K)]$$

The complete suite of ΔH_f calculations, using the energies of species obtained at the B3LYP (cc-PV5Z) level of theory, is outlined in the Appendix section (A4) of this document as a working example, to fully perceive the calculations undertaken.

3.5 MOPAC-2016 computations

MOPAC-2016 (Version: 20.302W) was used to compute Heats of Formation of the single species selected for this study. Relevant charges were assigned to ionic species and subjected for refinement as non-periodic structures. *MOPAC* is a semi-empirical quantum chemistry program and is based on neglect of diatomic differential overlap (NDDO) approximation as defined by Dewar and Thiel [15].

Seven (7) different semi-empirical Hamiltonians are in *MOPAC-2016* namely:

Austin Model 1 (AM1)

Recife Model 1 (RM1)

Parametric Method 3 (PM3)

Parametric Method 6 (PM 6)

Parametric Method 7 (PM7)

Modified Neglect of Diatomic Overlap (MNDO)

Modified Neglect of Diatomic Overlap with D Atomic Orbitals (MNDOD)

- MNDO was among the earliest semi-empirical NDDO methods developed for quantum calculations. The key benefit of MNDO over former methods, was that parameter values were enhanced to emulate molecular properties instead of atomic properties [16]
- AM1 was developed as an improvement of MNDO, since it has the ability to replicate hydrogen bonds and estimate accurate activation energies for reactions [17]
- RM1 parameters were developed to reproduce experimental properties: such as ionization potentials, geometry, enthalpies of formation, electronic charges and dipole moments [18]

- PM3 theory is based on the equation and formalism similar to the AM1 method. PM3 functionality methodically defines intermolecular hydrogen bonding in small polar molecules [19]
- MNDOD replaced and improved on the Modified Intermediate Neglect of Diatomic Overlap (MINDO) method.
- PM6 is a re-parameterized version of the NDDO method.
- PM7 is improved form the PM6 method of the neglect of diatomic differential overlap (NDDO) theory, developed for large systems. Atomic parameters were enhanced with a ~10% reduction in average errors in organic compounds [15].

MOPAC-2016 computational steps applied, are presented in Table-3.3

Table-3.3: MOPAC conditions applied for analysis

<i>MOPAC Execution Conditions</i>	
Hamiltonians applied:	AM1/MNDO/MNDOD PM3/PM6/RM1/PM7
SCF convergence:	0.01 kcal/mol
Structure Optimize:	Normal
Wave function:	Automatic
Periodic Cell:	Optimize
Starting Hessian:	Automatic
Convergence:	Normal
Iterations:	10 000
Initial Temp:	50K
Final Temp:	300K

The overall workflow used to execute *MedeA (MOPAC-2016)* is presented in Figure-3.2.

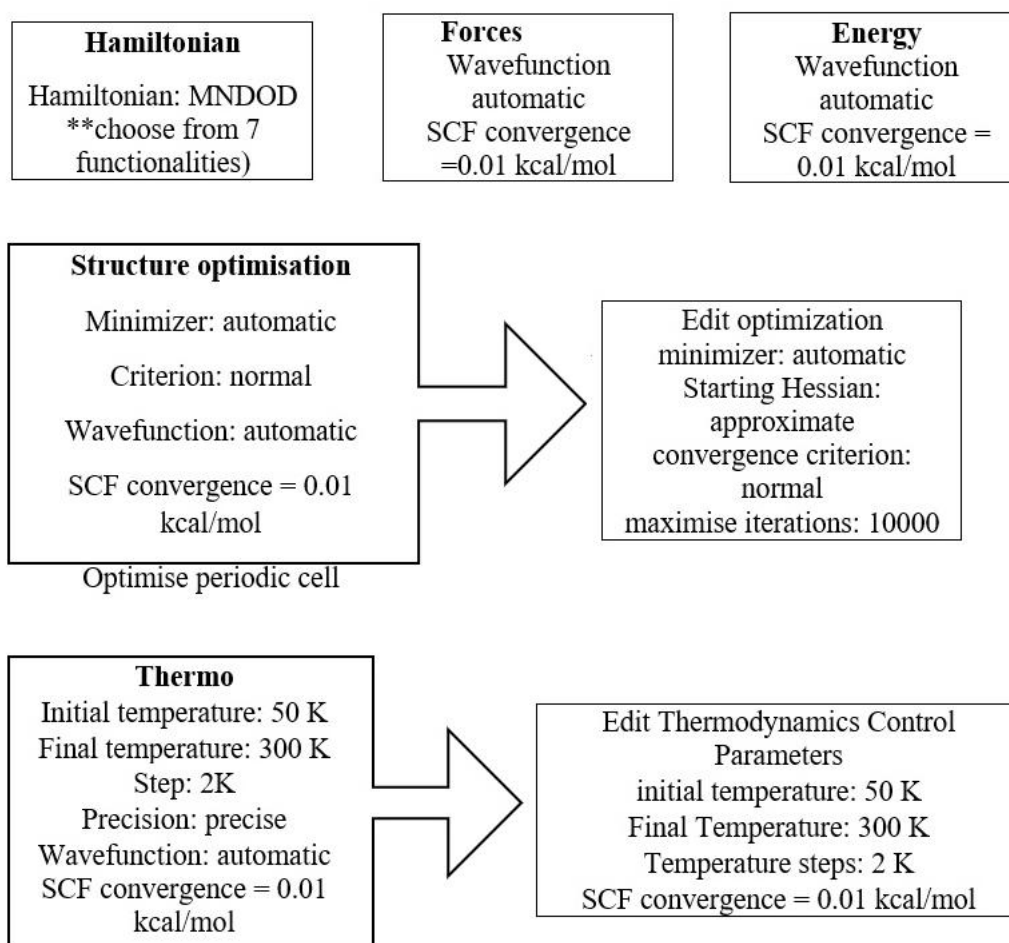


Figure-3.2: Medea/MOPAC-2016

Single neutral periodic species were first optimised with *Medea-3-3.1*/(*VASP-6.2.1*) and the resultant model then exposed to the single point meta-generalized gradient approximation (Meta-GGA) scan method. Both computations used the same parameters with the exception of the calculation module where the DFT Exchange Correlation (GGA-BLYP) was used for optimisation and the GGA-SCAN functional was used for single point analysis.

3.6 *VASP-6.2.1* Computations

VASP-6.2.1 was used to compute the thermochemistry of neutral and radical single molecular species (confined to periodic space) with all atomic positions relaxed and unit-cell dimensions refined, as pre-condition for refinement of the larger ensemble models. No *van der Waals* dispersion corrections were applied. Computations were undertaken at the DFT theory level, applying both LDA (*Linear Density Approximation*) and Exchange Correlations (GGA-BLYP) and (GGA-PBE) for potential approximations, applied for full structural refinement at group symmetry *PI(1)*. Energies of Formation were determined with a Projector Augmented Wave (PAW) plane-wave cut-off of 400 eV, increased to compensate for cell-shape and cell-volume optimization. All *VASP-6.2.1* calculations were employed with the same input parameters, presented in Table-3.4.

Table-3.4: VASP-6.2.1 Conditions Applied for DFT Analysis
(Remaining Conditions taken as default)

VASP Execution Conditions	Single Molecule	Ensemble Models
Theory: Functional	DFT	DFT
Exchange Correlation: <i>Becke-3-LYP</i>	GGA-BLYP	GGA-BLYP
Perdue Burke Ernzerhof	GGA-PBE	GGA-PBE
Cell size:	15x15x15 Å	<i>GIBBS</i> optimized model
Cell Optimize:	No	No
Volume Change:	No	No
Relax Atoms:	Yes	No
Energy of Formation:	Yes	Yes
Augmented PAW plane-wave cut off:	280-400 eV	280-400 eV
Brillouin zone (k-mesh) spacing:	0.5 Å ⁻¹	0.5 Å ⁻¹
Mesh points:	2x2x2 Å ⁻¹	2x2x2 Å ⁻¹
Smearing: Methfessel-Paxton (width)	0.2 eV	0.2 eV
Energy response surface: Optimized	Conjugate Gradient (CG)	Conjugate Gradient (CG)
Vd Waals Dispersion:	No	No
SCF convergence:	1x 10 ⁻⁵ eV	1x 10 ⁻⁵ eV
Max. iterations:	120	120

The parameter flowchart applied for *MedeA/VASP-6.2.1* computations are presented in Figure-3.4 and Figure-3.5.

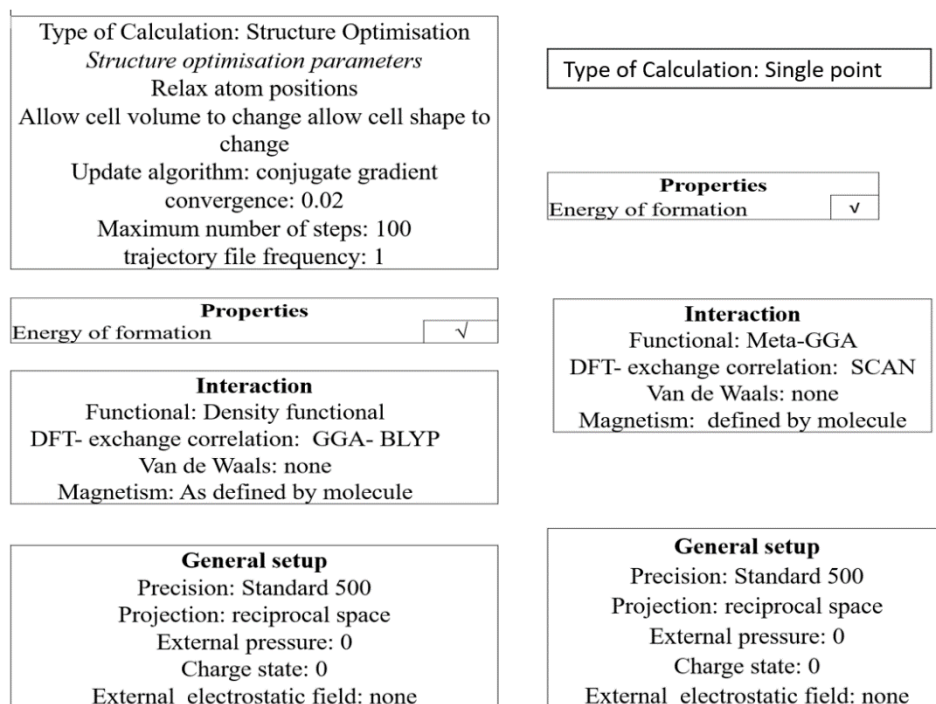


Figure-3.3: Calculation Module of VASP-6

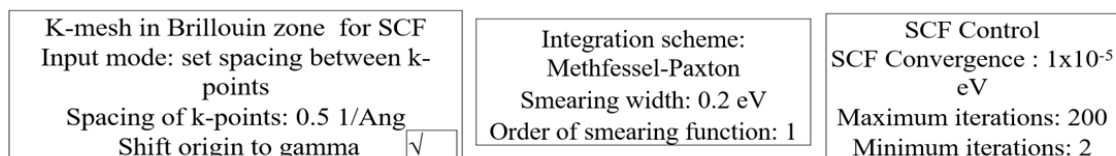


Figure-3.4: SCF Module of VASP-6

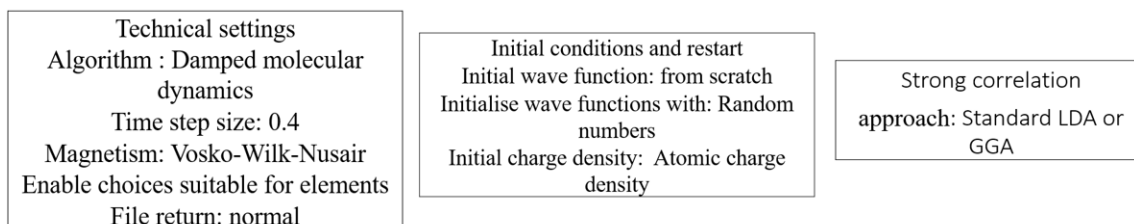


Figure-3.5: Advanced Restart Module of VASP-6

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CHAPTER 4: COMPUTATIONAL RESULTS: SINGLE MOLECULES

4.1 Introduction

The results presented in this Chapter, reflect on the correlation between the different software systems to derive at Heats of Formation, as one measure of conformity to optimum molecular geometries and effective electronic structure. The optimum structural property for each species was important to apply to a Grand Ensemble Canonical Simulation but scaled to atomic structural representations applying empirical force field parameters.

4.2 *Gaussian* Computations

Ab Initio open literature investigations utilised a collection of computational software systems such as: *Gaussian*, *Molpro*, *OpenMP* (parallel version of Mihaly Kallay's general coupled cluster code *MRCC* [1]). Application of these computational software systems are all compute resource demanding and come with innumerable challenges and usually require extended cluster computational hardware. These factors probably justify why only a selected few ClO₂ species have been selected and studied by computational means up to now.

Chlorine oxide species present a significant class of organic molecules, described by complex covalent bonding (and in some instances dative bonding) between highly electronegative atoms. Additionally, some molecules have unpaired electrons (resulting in open shell geometries) which usually result in strong lone-pair repulsions, or in cases of hydrogen derivative species, in strong intermolecular hydrogen bonding. The O-Cl bond creates a challenge for single referenced electron correlation methods, as these species are usually dominated by severe non-dynamical correlation (NDC) effects [1] and spin contamination, with resultant polarization.

Structural properties required to calculate geometries, spectroscopic properties and energy characteristics of molecules, are obtained from theoretical methods utilising one-electron basis set theory.

Many researchers have discussed some of the complexities associated with these studies, especially in relation to the calculation of Enthalpies of Formation and typical atmospheric kinetic models [2], [3], [1].

Heats of formation (ΔH_f) were calculated for single species (identified in chapter 2 from open literature), using DFT/B3LYP unrestricted level of theory, with Prof John Pople's basis set (cc-pV5Z) programmed into the *Gaussian-16* program.

- An external (to *Gaussian*) calculation step was required to derive at Heats of Formation and Free Energies of Formation [4]. Tables-A4.1 and A4.2 in the Appendix of this manuscript outlines the steps to follow in deriving at final Heats of Formation and Free Energies of Formation, from all *Gaussian-16* simulations for chlorine oxide species.

- Tabulated Heats of Formation derived with the *Gaussian-16* software are listed in Table-4.4
- Literature referenced, alternative techniques utilized to derive at Heats of Formation for most of the $C\ell O_2$ species, are listed in Table-4.1 (below)

Table-4.1: Literature referenced alternative computational and experimental methods used to obtain ΔH_f energies. (The structure of $C\ell_2O_3$ and its isomer $C\ell O C\ell O_2$ was referenced to Clark and Francisco [5])

Investigator		
($C\ell O$)	ΔH_f (kcal/mol)	Method
Jr. M. W. Chase	24.19 [6]	Determined experimentally
Abramowitz and Chase	24.27 [7]	Determined experimentally by spectroscopy
Karton et al.	24.28 [1]	W8 theory
Ventura et al.	24.88 [2]	G4 theory
Ruscic et al.	24.31 [8] [9] [10]	ATcT thermochemical network (TN)
($O C\ell O$)	ΔH_f (kcal/mol)	Method
Xu et al.	23.70 [11]	G2M//B3LYP/6-311+G(3df,2p)
Abramowitz and Chase	23.17 [7]	Determined experimentally by spectroscopy
Karton et al.	23.67 [12]]	W8 theory
Ventura et al.	27.2 [2]	Ab initio method: CCSD(T) aug-cc-pV6Z
Ruscic et al.	24.36 [8] [9] [10]	ATcT thermochemical network (TN)
($C\ell O O$)	ΔH_f (kcal/mol)	Method
Zhu and Lin	23.8 [13]	G2M//B3LYP/6-311G(3df, 2p)
Abramowitz and Chase	21.5 [7]	Determined experimentally by spectroscopy
Karton et al.	24.30 [12]	W8 theory
Ventura et al.	24.30 [2]	Ab initio method: CCSD(T) aug-cc-pV6Z
Ruscic et al.	24.56 [8] [9] [10]	ATcT thermochemical network (TN)

Table-4.1 continued

Investigator	ΔH_f (kcal/mol)	Method
(HOCl)		
Zhu and Lin	-17 [13]	G2M//B3LYP/6-311G(3df. 2p)
Ventura et al.	-18 [2]	Ab initio method: CCSD(T) aug-cc-pV6Z
Karton et al.	-18 [12]	W4 theory
Ruscic et al.	-18.36 [8] [9] [10]	ATcT thermochemical network (TN)
(ClO ₃)		
Zhu and Lin	47.2 [13]	G2M//B3LYP/6-311G(3df. 2p)
Sicre and Cobos	49.3 [14]	G3M//B3LYP/6-311G(3df. 2p)
Sicre and Cobos	57.0 [14]	G1 and isodesmic reaction
Karton et al.	50.9 [12]	W4 theory
Ruscic et al.	44.46 [8] [9] [10]	ATcT thermochemical network (TN)
(ClO ₄)		
Sicre and Cobos	65.2 [14]	G2M//B3LYP/6-311G(3df. 2p)
Sicre and Cobos	66.4 [14]	Ab initio: B3LYP/6-311+G(3d2f)
Ruscic at al.	57.6 [8] [9] [10]	ATcT thermochemical network (TN)
(ClOCl)		
Jr. M. W. Chase	22 [13]	Determined experimentally
Karton et al.	18.82 [12]	W4 theory
Abramowitz and Chase.	31.77 [7]	Determined experimentally by spectroscopy
Klobas and Wilmouth.	31.03 [15]	Determined from the slope of the van't Hoff plot in the third law analysis.
(Cl ₂ O ₃)		
Clark and Francisco	46.2 [5]	G1 and G2 theory
	44.9 [5]	
Sicre and Cobos	37.8 [14]	Ab initio techniques: B3LYP 6-311+G(3d2f)
	53.9 [14]	G2 theory
Burkholder et al.	35.8 [16]	Determined with UV spectroscopy
(HOClO)		
Karton et al.	5.0 [12]	W4 theory
Clark and Francisco	11.9 [5]	G2 theory
Ruscic at al.	4.94 [8] [9] [10]	ATcT thermochemical network (TN)
(HOClOO)		
Xu et al.	0.1 [11]	G2M//B3LYP/6-311+G(3df.2p)
Karton et al.	-2.6 [1]	W4 theory
Ruscic at al.	-1.10 [8] [9] [10]	ATcT thermochemical network (TN)
(HOOCLO)		
Xu et al.	20.6 [11]	G2M//B3LYP/6-311+G(3df.2p)
Ruscic at al.	21.54 [8] [9] [10]	ATcT thermochemical network (TN)

Table-4.1 continued

Investigator		
(HOOC ℓ O ₂)	ΔH_f (kcal/mol)	Method
Xu et al.	0.1 [11]	G2M//B3LYP/6-311+G(3df.2p)
Colussi and Grela	-4.2 [17]	Determined with UV spectroscopy
Ruscic et al.	16.55 [8] [9] [10]	ATcT thermochemical network (TN)
(HOClO ₃)	ΔH_f (kcal/mol)	Method
Xu et al.	3.2 [11]	G2M//B3LYP/6-311+G(3df.2p)
Clark and Francisco	10.80 [5]	G2 theory
Colussi and Grela	-1.5 [17]	Determined with UV spectroscopy
Ruscic et al.	0.43 [8] [9] [10]	ATcT thermochemical network (TN)

Unrestricted Hartree Fock (UHF) spin polarization restriction was applied on radical and charged species, with *Restricted Hartree Fock* (RHF) applied to neutral species. *Restricted Open Shell Hartree Fock* (ROHF) ruling was used for some charged species in an attempt to compensate for spin polarization.

Although a significant group of chlorine oxide related species exist, a few have been studied computationally, possibly owing to the complexity of their bonding character. Many researchers have attempted to compute the thermochemical properties of a few chlorine oxide species [1] [2] [5][6] [7] citing discrepancies in their energy values. A list of reported energies and the methods used to derive at them, are presented below in Table-4.2.

- The ΔH_f of twenty (20) SINGLE species were considered unacceptable, since these values show a difference in energy, exceeding 8.5 kcal/mol correlated against Literature Reference ΔH_f energies. This equates to thirty-three (33) percent of the species producing unacceptable results.
- The computed ΔH_f of thirteen (13) species showed differences in energy, between the calculated and the reference energies that lie between 3.5 - 8.5 kcal/mol hence twenty-two (22) percent produced fair results.
- Accurate results were obtained for thirty-five (35) percent of the species
- The *Gaussian* (DFT/B3LYP) theory supported by the (*cc-pV5Z*) basis set, was found by many researchers to be sufficiently accurate, but noted the (*cc-pV6Z*) basis set to be more acceptable for these compounds [2]
- **Gaussian-09** supports computations with the (*cc-pV6Z*) basis set on 3rd row elements [18]. However, this basis set is not supported in the later *Gaussian-16* [18].

- *Gaussian-16* theory level CCSD (T) gave the most accurate thermochemical properties [2], but found to be computationally resource demanding, especially when combined with higher basis sets [19].

4.3 MOPAC Computations

The Heats of Formation (ΔH_f) for the set of sixty (60) species were calculated, applying each of the seven (7) *MOPAC* Hamiltonians. Different ΔH_f energies were obtained for each compound. The author of the *MOPAC* program, Dr James (Jim) Stewart [20] very kindly corrected the number of orbitals assigned, to deal with effective electron population and charge distribution in each ionic species. Although reference ΔH_f of a few species are unknown, the result (applying the different Hamiltonians) corresponding to the most sterically correct geometry were finally accepted.

The ΔH_f results for the studied set of species computed with *MOPAC-2016* are tabulated in Table 4.2 and were used in determining Free Energies of Formation, Entropy and Heat Capacity properties.

Table-4.2: Calculated Heats of Formation (kcal/mol) with MOPAC-2016, applying all the supported Hamiltonians for single species (in ‘vacuum’). Entries in bold were selected as Heats of Formation. Entries closest to reference values were selected as the final Heat of Formation values.

Molecule Formula	Reference	Hamiltonian (ΔH_f kcal/mol)						
		AM1	MNDO	MNDOD	PM3	PM6	PM7	RM1
ClO	24.31 [8]	31.09	34.84	17.73	8.27	25.47	21.57	12.83
[ClO] ⁻	-28.26 [8]	-15.27	-7.78	-35.81	-32.98	-24.00	-31.23	-20.98
[ClO] ⁺	274.84 [8]	321.72	334.48	316.78	238.35	284.77	276.94	279.73
ClOO	24.56 [8]	44.45	22.24	23.28	13.51	19.07	7.35	-1.15
[ClOO] ⁻	-59.28 [8]	-42.04	-44.04	-43.15	-40.56	-32.58	-41.39	-53.36
[ClOO] ⁺	286.71 [8]	277.39	299.61	293.61	264.05	249.24	259.35	272.89
ClOCl	18.62 [8]	19.45	31.26	16.02	-16.25	8.53	4.89	9.1
[ClOCl] ⁻	-35.28 [8]	-39.81	-30.51	-36.72	-53.84	-60.37	-61.23	-57.16
[ClOCl] ⁺	269.14 [8]	284.81	302.43	261.74	213.39	237.23	247.28	244.88
ClClO	31.81 [8]	55.03	73.16	28.63	9.18	37.78	40.13	17.28
[ClClO] ⁻	-44.11 [8]	-32.32	-39.71	-43.69	-62.98	-56.57	-49.02	-75.53
[ClClO] ⁺	270.09 [8]	285.23	307.67	264.74	234.68	256.85	271.85	240.95
ClClO ₂	29.16 [8]	132.97	170.87	38.44	-55.80	28.98	46.02	19.42
ClOOO	53.29 [8]	42.54	45.51	41.66	4.03	9.07	10.83	30.75
[ClOOO] ⁻	-25.00 [8]	-13.25	-6.47	-4.21	-35.61	-9.43	-53.81	-25.06
Cl(O)O ₂	63.37 [8]	63.42	63.96	62.96	74.32	83.95	61.23	63.47
[Cl(O)O ₂] ⁻	-25.22 [8]	-8.12	-12.12	-10.55	-12.99	-35.34	-53.81	-14.77
ClO ₃	44.46 [8]	153.42	171.64	38.79	-21.19	39.31	53.10	158.13
[ClO ₃] ⁻	-50.83 [8]	72.95	130.44	-55.19	-155.81	-65.04	-59.67	-41.97
[ClO ₃] ⁺	297.67 [8]	497.19	530.66	300.82	300.82	324.37	339.03	386.74

Table-4.2 Continued

Molecule Formula	Reference	Hamiltonian (ΔH_f kcal/mol)						
		AM1	MNDO	MNDOD	PM3	PM6	PM7	RM1
ClO ₄	57.60 [8]	219.47	249.03	79.10	46.14	136.89	92.82	163.42
[ClO ₄] ⁻	-64.98 [8]	166.46	238.88	-80.38	-128.15	-72.82	-82.73	2.47
ClOClO	41.80 [8]	82.39	117.00	36.38	-4.37	38.19	33.41	42.73
ClOOCl	31.37 [15]	33.69	41.70	36.71	2.33	16.00	14.64	18.80
ClOClO ₂	37.80 [16]	150.69	212.90	38.45	-83.71	22.17	26.44	48.41
ClOClO ₃	37.40 [14]	267.68	325.67	50.07	-112.63	54.44	49.52	66.97
ClO ₂ ClO ₂	N/R	275.27	350.88	103.16	-112.63	50.72	74.49	68.02
ClO ₂ -O- ClO ₂	N/R	283.17	397.84	41.27	-198.43	18.51	22.32	49.18
ClOOClO ₃	69 [14]	227.59	303.20	73.94	-20.74	78.4	68.99	111.43
ClO ₂ -O- ClO ₃	72.40 [14]	403.53	509.39	47.01	-119.90	44.94	37.45	154.79
ClO ₃ -O- ClO ₃	86.20 [14]	526.16	623.36	57.12	-26.25	76.7	59.59	260.46
ClO ₂ -O-O- ClO ₂ [22]	N/R	214.46	363.24	82.18	-143.74	62.83	79.42	105.24
ClO ₃ -O-O- ClO ₃ [22]	111.40 [22] ^s	128.12	585.97	108.37	44.32	137.16	128.12	182.76
[ClO ₃ ClO ₃] ⁻² [18]	N/R	212.66	578.86		-120.01	44.91	37.45	106.06
OCtO	24.36 [8]	97.57	142.95	25.47	43.1	80.35	41.21	90.43
[OCtO] ⁻	-25.74 [8]	22.37	54.39	-33.48	-36.85	-16.07	-23.14	2.2
[OCtO] ⁺	262.13 [8]	385.36	418.23	270.55	188.61	266.34	279.2	284.22
OCtOO	54.31 [8]	3.33	18.82	1.71	4.02	26.69	44.72	-8.60
[OCtOO] ⁻	-8.19 [8]	-22.07	2.48	-25.07	-23.26	1.52	-12.15	-24.39
OCtClO ₂	46.20 [5]	196.34	243.10	82.73	-83.71	22.18	27.21	50.13
[ClOH ₂] ⁺	194.66 [8]	207.65	201.43	197.10	183.5	201.92	200	207.75
HOCl	-18.36 [8]	-21.78	-15.68	-15.67	-34.30	-17.81	-23.08	-20.93
[HOCl] ⁻	47.38 [8]	-48.80	-61.54	-9.02	-55.81	-65.41	-63.2	-60.86
[HOCl] ⁺	237.86 [8]	244.01	253.76	219.18	201.4	215.87	220.85	224.08
HOClO	4.94 [8]	35.04	69.25	-1.73	-25.15	9.49	6.77	12.76
[HOClO] ⁻	-37.31 [8]	-22.66	-13.62	-34.19	-48.81	-49.98	-47.06	-42.32
[HOClO] ⁺	236.82 [8]	265.87	293.57	221.29	177.25	222.4	223.62	233.62
HOClO ₂	10.90 [23]	106.10	165.03	-3.27	-108.27	-12.98	-10.91	5.86
HOClO ₃	0.43 [8]	223.17	277.8	3.71	-26.48	16.03	7.49	110.80
[HOClO ₃] ⁺	0.43 [8]	401.01	430.41	304.78	273.23	303.74	325.2	344.25

Table-4.2 Continued

Molecule Formula	Reference	Hamiltonian (ΔH_f kcal/mol)						
		AM1	MNDO	MNDOD	PM3	PM6	PM7	RM1
HOOC ℓ	282.36 [8]	9.72	9.23	5.81	-13.83	-1.6	0.47	-0.68
[HOOC ℓ] ⁻	-0.32 [8]	-39.22	-61.72	-56.94	-52.75	-53.14	-52.6	-58.96
[HOOC ℓ] ⁺	-8.12 [8]	251.3	259.09	252.98	227.53	223.78	230.67	244.23
HOOC ℓ O	27.30 [23]	49.08	82.04	17.76	-7.17	30.06	23.08	22.05
HOOC ℓ O ₂	16.55 [8]	114.9	174.77	19.82	-80.24	19.25	21.63	35.03
HOOC ℓ O ℓ	14.00 [23]	13.6	14.86	11.88	-6.34	7.94	3.80	1.65
HOOC ℓ O ℓ O	22.70 [5]	29.39	29.21	27.10	5.81	24.78	17.94	11.62
HCC ℓ O	33.24 [8]	43.63	76.22	21.77	10.50	27.70	30.43	34.08
HCC ℓ O ₂	45.65 [8]	121.11	177.07	31.72	-70.49	37.56	49.06	47.7
HCC ℓ O ₃	57.10[23]	106.1	302.02	45.70	-11.39	73.01	77.41	5.86

*N/R - not reported

[§] Indicates that the $\Delta H_f(298K)$ of C ℓ O₃-O-O-C ℓ O₃ is calculated from Energies by Beltran et al., (1999) [22]

Additional properties i.e., Spin Multiplicity, Point groups, Entropy and Heat Capacities (Cp) at 298K, were calculated with *MOPAC-2016* are listed in Table 4.3. These chemical properties are required as supporting data for the population of the *FactSage* database (chapter 7).

Table-4.3: Entropy, Heat capacities, Point Groups and Multiplicities

Molecule Formula	S (J/mol K)	Cp (J/mol K)	Multiplicity (Spin State)	Point Group
C ℓ O	215.09	30.78	2	C* ν
[C ℓ O] ⁻	215.64	31.01	1	C* ν
[C ℓ O] ⁺	213.25	21.66	1	C* ν
C ℓ OO	256.64	40.54	2	C2v
[C ℓ OO] ⁻	265.18	46.01	1	C2v
[C ℓ OO] ⁺	258.75	35.69	1	C2v
C ℓ OCC ℓ	264.16	43.33	1	C2v
[C ℓ OCC ℓ] ⁻	279.21	48.37	2	C2v
[C ℓ OCC ℓ] ⁺	262.36	42.76	2	C2v
C ℓ CC ℓ O	272.67	45.19	1	Cs
[C ℓ CC ℓ O] ⁻	293.01	49.86	2	Cs
[C ℓ CC ℓ O] ⁺	275.27	47.94	2	Cs
C ℓ CC ℓ O ₂	296.4	61.96	1	Cs

Table-4.3 continued

Molecule Formula	S	Cp	Multiplicity	Point Group
ClOOO	297.01	52.81	2	C1
[ClOOO] ⁻	286.06	56.29	1	C1
Cl(O)O ₂	323.26	63.22	2	C2v
[Cl(O)O ₂] ⁻	300.27	62.35	1	C2v
ClO ₃	280.33	62.65	2	D3h
[ClO ₃] ⁻	264.82	56.18	1	D3h
[ClO ₃] ⁺	273.42	35.56	1	D3h
ClO ₄	308.9	79.68	2	C2v
[ClO ₄] ⁻	265.7	63.11	1	Td
ClOClO	302.40	62.52	1	C1
ClOOCl	289.38	57.09	1	C2
ClOClO ₂	308.11	69.63	1	Cs
ClOClO ₃	335.93	86.60	1	C1
ClO ₂ ClO ₂	340.46	96.51	1	D2h
ClO ₂ -O-ClO ₂	332.84	99.84	1	D2h
ClOOClO ₃	359.09	102.89	1	C1
ClO ₂ -O-ClO ₃	367.43	111.22	1	C1
ClO ₃ -O-ClO ₃	369.09	126.94	1	D3h
[ClO ₃ ClO ₃] ⁻²	403.99	126.56	1	D3d
ClO ₂ -O-O-ClO ₂	403.86	129.85	1	C2h
ClO ₃ -O-O-ClO ₃	471.17	164.93	1	C1
OClO	250.18	38.97	2	C2v
[OClO] ⁻	253.39	42.54	1	C2v
[OClO] ⁺	272.24	32.61	1	C2v
OClOO	368.09	67.23	2	Cs
[OClOO] ⁻	302.31	65.09	1	Cs
OClClO ₂	340.18	77.86	1	Cs
[ClOH ₂] ⁺	246.18	27.56	2	C2v
HOCl	236.16	36.05	1	Cs
[HOCl] ⁺	235.36	35.62	2	Cs
HOClO	263.60	47.48	1	Cs
[HOClO] ⁻	283.68	57.80	2	Cs
[HOClO] ⁺	271.55	52.30	2	Cs
HOClO ₂	288.18	62.57	1	Cs

Table-4.3 continued

Molecule Formula	S (J/mol K)	Cp (J/mol K)	Multiplicity (Spin State)	Point Group
HOClO ₃	307.07	74.37	1	Cs
[HOClO ₃] ⁺	312.33	80.74	2	Cs
HOOCℓ	263.45	49.09	1	Cs
[HOOCℓ] ⁻	270.30	52.15	2	Cs
[HOOCℓ] ⁺	266.73	51.32	2	Cs
HOOCℓO	293.57	65.99	1	C1
HOOCℓO ₂	322.75	82.36	1	C1
HOOCℓ	286.64	58.79	1	C1
HOOCℓO	329.09	84.55	1	C1
HClO	237.13	38.579	1	Cs
HClO ₂	253.22	47.44	1	C2v
HClO ₃	264.06	50.98	1	C3v

MOPAC Hamiltonians: MNDOD, PM6 and PM7, accounted for 35.60 %, 22 % and 25 %, accurate ΔH_f energies respectively. Final (geometry optimized) MOPAC Heats of Formation energies are reported jointly in Table-4.4, with the Gaussian (single) molecule energies.

MOPAC-2016 surprisingly, offered excellent results:

- Approximately 21 % of the calculated ΔH_f obtained were between 6 to 10 kcal/mol higher than the reference values.
- The calculated ΔH_f of five species calculated (highlighted in yellow in Table- 4.5) were beyond the limit of the reference values and classified as unacceptable
- The remainder of the calculated ΔH_f were within 5 kcal/mol of the reference ΔH_f and acceptable.
- Finite ΔH_f energies for $[\text{ClO}_3\text{ClO}_3]^{-2}$, $\text{ClO}_2\text{-O-ClO}_2$, ClO_2ClO_2 and $\text{ClO}_2\text{-O-O-ClO}_2$ could not be confirmed with no reference ΔH_f energies being available.

4.4 VASP Computation (including all consolidated results)

The ΔH_f energies of neutral *single* models, were calculated using VASP-6.2.1 applying two functionals: GGA-BLYP and GGA-PBE. All neutral single species exposed to VASP-6.2.1 calculations, were confined to periodic cell conditions under Space Group symmetry restrictions of P1(1).

Periodic cell parameters assigned were increased to 15x15x15 Å to ensure a true single molecular environment persists. The GGA-BLYP functional produced accurate energies and the calculated

(single molecule) VASP-6.2.1 ΔH_f energies are presented in *Table-4.4* which also summarises **all** results obtained for single molecule species (from the MOPAC Software)

Table-4.4: Heats of Formation (ΔH_f 298K) of single chlorine oxide species calculated with the *Gaussian-16* (DFT B3LYP/cc-pV5Z) and MOPAC-2016 software

Species	Literature Reference ΔH_f (298K)	<i>Gaussian-16</i> (DFT B3LYP, cc-pV5Z) ΔH_f (298K)	MOPAC-2016 (derived) ΔH_f (298K)	VASP-6.2.1 GGA-BLYP ΔH_f (298K)	VASP-6.2.1 GGA-PBE ΔH_f (298K)
ClO [8]	24.31 [8]	29.71	25.47	25.91	26.62
[ClO] ⁻ [8]	-28.26 [8]	-23.46	-24.00		
[ClO] ⁺ [8]	274.84 [8]	278.18	276.94		
ClOO [8]	24.56 [8]	21.50	23.28	23.74	23.06
[ClOO] ⁻ [8]	-59.28 [8]	-54.75 #	-53.36		
[ClOO] ⁺ [8]	286.71 [8]	299.79	293.61		
ClOCl [8]	18.62 [8]	21.03	19.45	16.13	16.27
[ClOCl] ⁻ [8]	-35.28 [8]	-35.26	-36.72		
[ClOCl] ⁺ [8]	269.14 [8]	269.03	261.74		
ClClO [8]	31.81 [8]	34.64	37.80	23.94	21.79
[ClClO] ⁻ [8]	-44.11 [8]	-54.10 #	-43.69		
[ClClO] ⁺ [8]	270.09 [8]	275.83	271.85		
ClClOO [8] *# [ClClO ₂]	29.16 [8]	36.79	28.96	9.75	5.83
ClOOO [8]	53.29 [8]	67.80	45.51	65.8	66.06
[ClOOO] ⁻ [8]	-25.00 [8]	-21.93	-25.06		
ClO(O)O [8]*# [Cl(O)O ₂]	63.37 [8]	69.46	63.47		
[ClO(O)O] ⁻ [8]*# [Cl(O)O ₂] ⁻	-25.22 [8]	-21.95	-35.34		
ClO ₃ [8]	44.46 [8]	47.79	39.31	72.17	61.87
[ClO ₃] ⁻ [8]	-50.83 [8]	-35.31	-55.19		
[ClO ₃] ⁺ [8]	297.67 [8]	329.73	300.82		
[ClO ₄] [8]	57.60 [8]	74.64	46.14	62.99	45.53
[ClO ₄] ⁻ [8]	-64.98 [8]	-42.11	-72.83		
ClOClO [8]	41.80 [8]	54.98	38.19	15.54	13.52
ClOOCl [8]	31.37 [15]	38.67	33.69	31.38	10.86
ClOClO ₂ [8]	37.80 [16]	48.86 **	38.45	40.46	34.81
ClOClO ₃ [8]	37.40 [14]	65.77	49.52		
ClOOClOO [21]* [ClO ₂ ClO ₂]	N/R	69.78	50.72	21.63	16.32
ClO ₂ -O-ClO ₂ [21]	N/R	81.21	41.27	104.5	90.07

Table-4.4 Continued

Species	Literature Reference ΔH_f (298K)	Gaussian-16 (DFT B3LYP, cc-pV5Z) ΔH_f (298K)	MOPAC-2016 (derived) ΔH_f (298K)	VASP-6.2.1 GGA-BLYP		VASP-6.2.1 GGA-PBE	
				ΔH_f (298K)			
Cl ₂ O ₅ [14] ** [ClOOClO ₃]	69 [14]	72.17	73.94	49.87	33.22		
Cl ₂ O ₆ [14] ** [ClO ₂ -O-ClO ₃]	72.4 [14]	85.24 *	47.01	28.78	15.45		
Cl ₂ O ₇ [14]** ClO ₃ -O-ClO ₃	86.2 [14]	89.64 *	76.70	77.53	44.22		
ClO ₂ -O-O-ClO ₂ [21]	N/R	104.17 **	79.42	105.61	25.24		
Cl ₂ O ₈ [22] ** ClO ₃ -O-O-ClO ₃	111.40 [22] [§]	118.45	108.37				
[ClO ₃ -ClO ₃] ⁻² [21]	N/R	105.74	106.06				
OClO [8]	24.36 [8]	23.89	25.47	21.04	66.92		
[OClO] ⁻ [8]	-25.74 [8]	-16.29	-23.14				
[OClO] ⁺ [8]	262.13 [8]	278.24	266.34				
OClOO [8]	54.31 [8]	51.63	44.72	36.12	33.65		
[OClOO] ⁻ [8]	-8.19 [8]	-9.89	-12.15				
OClClO ₂ [8]	46.2 [5]	146.67 #	50.13	43.07	35.69		
[ClOH ₂] ⁺ [8]	194.66 [8]	202.38	197.10				
HOCl [8]	-18.36 [8]	-16.43 *	-17.81	-18.04	-19.28		
[HOCl] ⁻ [8]	47.38 [8]	54.25	-9.02 **				
[HOCl] ⁺ [8]	237.86 [8]	239.06	244.01				
HOClO [8]	4.94 [8]	13.01	6.77	6.15	1.9		
[HOClO] ⁻ [8]	-37.31 [8]	-57.33 #	-34.19				
[HOClO] ⁺ [8]	236.82 [8]	238.93	233.62				
HOCl(O)O [8] ** [HOClO ₂]	10.90 [23]	11.18 **	-3.27	13.7	7.84		
[HOClO ₃] [8]	0.43 [8]	12.13 **	3.71	9.09	-10.44		
[HOClO ₃] ⁺ [8]	0.43 [8]	325.73 #	273.23				
HOOC _l [8]	282.36 [8]	2.60 *	-0.68	2.97	0.97		
[HOOC _l] ⁻ [8]	-0.32 [8]	-10.60	-39.22				
[HOOC _l] ⁺ [8]	-8.12 [8]	253.27 #	244.23				
HOOC _l O [23]	27.30 [23]	27.78 *	23.08	27.49	21.73		
HOOC _l (O)O [8]** [HOOC _l O ₂]	16.55 [8]	37.26 **	19.25	5.08	-37.37		
HOOC _l O _l [8]	14.00 [23]	27.44 *	11.88	3.39	-0.53		
HOOC _l O _l [5]	22.70 [5]	28.18	24.78	12.92	8.31		
HClO [8]	33.24 [8]	33.89 *	30.45	31.05	28.15		
HClO ₂ [8]	45.65 [8]	48.31 *	49.06	88.98	81.31		
HClO ₃ [23]	57.1 [23]	46.07 *	45.70	42.49	25.82		

N/R - not reported

** Highlights the naming convention adopted for this study

* Restricted: DFT/B3LYP/cc-pv5z

**Restricted open shell. DFT/B3LYP/cc-pv5z

Unrestricted: DFT/B3LYP/6-311++G(3d2f,3p2d)

§ Indicates that the ΔH_f (298K) of $\text{ClO}_3\text{-O-O-ClO}_3$ is calculated from Energies by Beltran et al., 1999 [22]

Some discrepancies observed for both *Gaussian* and *MOPAC* single molecule species, are highlighted as bold text. Comparison of ΔH_f obtained with three computational programs i.e., *Gaussian-16*, *MOPAC-2016* and *VASP-6.2.1* resulted in variable results for the species:



Further investigations are required to elucidate these variations, probably related to spin polarization and/or improved atom potentials.

4.5 Conclusion

A noticeable variation in spin-states were observed for higher-order ClO_2 species structures, applying both the *Gaussian* and *MOPAC* software programs. This was specifically prevalent for species hosting a higher oxygen content, which resembles a complex array of internal open shell connectivity and spin polarization. This must not be conceived as flaws in the software algorithms, but rather considered as a result of slight alterations in electronic structural configurations, adopted by the software algorithms. Species of this nature (with intrinsic radical and charge dissipation character) are particularly vulnerable, to produce varying spin conditions. Both *Gaussian-16* and *MOPAC-2016* performed well in terms of the computational determination of the Heat of Formation, which could comfortably be utilised in the subsequent extraction of chemistry reaction schemes.

4.6 References

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CHAPTER 5: ENSEMBLE MODEL COMPUTATIONS

5.1 Introduction

The first critical and significantly important aspect in simulation at the force field level of molecular space definition, is the specification of suitable empirical force field parameters. The *MedeA/GIBBS* software features a built-in algorithm to assign appropriate force field parameters to a predefined model (periodic or molecular) and is based on molecular geometry and α - β bonding from every atom, within a specified cut off distance from each atom. Cut off distances of 3.5Å are typically permitted around each atom. Atom type assignments associated with immediate 3D-neighbours are therefore effectively considered. For organic structures all facets of torsion; bond angles (typical); bond lengths; long-distance interactions; coulomb forces; bond character/type (to restrain bond distance); bond increments; and vd Waals interactions are accounted for and have to be assigned as a completed set, prior to any simulation. This represents a molecular mechanical representation at the atomic-level [1], side stepping the *Ab Initio* details of electron-electron and electron-nucleon interactions.

The force field concept relates to classical mechanics and is a collection of equations and associated parameters, designed to reproduce molecular geometry and selected properties, normally derived from a vast set of known structures [2]. The force field method uses a set of empirical formulae to portray the interatomic interactions in an averaged fashion [3]. This method is used as an excellent averaged molecular structural representation tool, when electron (or charge) distribution is not considered or required and can successfully describe properties of interest and as a result are computationally less demanding.

The technique is sufficiently accurate and powerful, to optimize a molecular (or periodic) structure to within an acceptable energy regime, for quantum chemistry to take over.

5.2 MedeA GIBBS ensemble (NPT)

Panagiotopoulos [4] proposed the concept of Gibbs ensemble Monte Carlo (GEMC) simulations in 1987, which entailed a method of directly computing the phase coexistence properties of either pure fluids or mixtures [4]. The technique is dependent on a confined model space and is undertaken in a periodic (Grand Cell) environment. GEMC has an advantage over other conventional methods since two-phase equilibrium properties can be calculated without requiring a Free Energy value [4]. In this case, each chlorine oxide species was built as a periodic model, force field parameters assigned and executed as a single-phase system in a GEMC ensemble simulation.

5.2.1 Chemical Approach

Single monomeric chlorine oxide species were exposed to an isobaric-isothermal Gibbs ensemble (GEMC) simulation, applying Grand Canonical Monte Carlo (GCMC) methodology at constant temperature (T) & pressure (P). The number of molecular entities N is preserved throughout each

5.2.2.2 Bond Increments parameters

Bond increments describe the charge distribution formed during the bonding of two atoms. The assignment of the atomic charges of a molecule into bond increments, creates a set of general rules, based on the type of atoms in the bond [6].

I and J are the atom types of the bonded atoms [8]. Delta IJ and Delta JI (presented as partial atomic charges measured in elementary charge unit e) refers to the change in atomic charges depending on the direction of the charge i.e., I to J or from J to I [8].

Bond increments parameters are presented in Table 5.2.

Table-5.2: Bond Increment Parameters

I	J	Delta-IJ	Delta-JI	Force field reference
cl4	o_1r	0.3864	-0.6364	pcff file
cl	o	0.3864	-0.6364	This work
h*	o	0.4100	-0.4100	pcff file
cl	o_1r	0.3864	-0.6364	This work

Entries for bond increments: **cl [o]** and **cl [o_1r]** constants have been borrowed from the *cvff* force field set for: **cl4 [o_1r]**. No other constants have been published to support this study. Bond lengths for **cl [o]** = 1.7 Å, **cl4 [o_1r]** = 1.4 Å and **cl [o_1r]** = 1.6 Å [7].

5.2.2.3 Quartic bond parameters

The quartic bond-stretching potential is presented by:

$$E = K_2 * (R - R_0)^2 + K_3 * (R - R_0)^3 + K_4 * (R - R_0)^4$$

Where:

R is the current bond length and *R*₀ the reference bond length in angstroms.

*K*₂, *K*₃ and *K*₄ are the coefficients for the quadratic, cubic and quartic terms in units of kcal/mol Å⁻², kcal mol Å⁻³ and kcal/mol Å⁻⁴ respectively.

I and *J* are the atom types of the bonded atoms [8].

Quartic bond parameters are presented in Table 5.3.

Table-5.3: Quartic Bond Parameters

I	J	R 0	K2	K3	K4	Force field Reference
cl	o	1.6500	307.0632	0.0	0.0	cvff file
h	o	0.9600	493.8480	0.0	0.0	cvff file
cl4	o_1r	1,4523	419.3650	-838.7299	978.5182	cvff file
cl	o_1r	1.679	419.3650	-838.7299	978.5182	This work
cl4	oh	1.641	419.3650	-838.7299	978.5182	This work
o	o	1.2080	833.6868	0,00	0.00	cvff file
o	o_1r	1.3603	833.6868	0,00	0.00	This work
o_1r	o_1r	1.3227	833.6868	0,00	0.00	This work

- Entries for the: **cl [o_1r]** and **cl4 [oh]** constants have been extracted from the **cl4 [o_1r]** entry in *pcff+*. The relevant bond lengths (R0) were obtained from literature [7].
- Entries for the: **o [o_1r]** and **o_1r [o_1r]** constants have been extracted from the o - o entry in *pcff+*. The relevant bond lengths (R0) were obtained from literature [9].

5.2.2.4 Quartic bond angle parameters

The quartic angle is defined as follows in *pcff+*:

$$\Delta = \theta - \theta_0$$

$$E = K2 * \Delta^2 + K3 * \Delta^4$$

$$\Delta^3 + K4 * \Delta^4$$

Where:

Δ is used to represent Delta, the changed angle value

θ_0 is the equilibrium value of the angle.

Theta (θ) is the current bond angle, Theta0 (θ_0) is the reference bond angle in degrees.

The usual half factor is included in K [10].

K_2 , K_3 and K_4 are quadratic force constants in units of: kcal/mol rad⁻².

I, J, K are the atom types involved in the angle.

Quartic angle parameters are presented in Table 5.4.

Table-5.4: Quartic Angle Parameters

I	J	K	Theta0	K2	K3	K4	Force field reference
o1=	s2=	o1=	119.300	115.2627	-35.6278	-26.1261	cvff file
o	cl	o	115.400	115.2627	-35.6278	-26.1261	This work
*	p	*	109.5000	45.0000	0.0	0.0	pcff file
o	cl	oh	109.5000	45.0000	0.0	0.0	This work
*	o	*	109.5	60.000	0.0	0.0	pcff file
cl	o_1r	cl4	111,9	60.000	0.0	0.0	This work
cl4	o_1r	o_1r	110,3	60.000	0.0	0.0	This work
cl4	o_1r	o	110.4700	60.000	0.0	0.0	This work
cl	o	o_1r	111,74	60.000	0.0	0.0	This work
cl4	o_1r	cl4	119,11	60.000	0.0	0.0	This work
cl4	oh	ho	105	60.000	0.0	0.0	This work
o_1r	cl4	o_1r	112.2363	108.1226	-24.1496	-23.3346	pcff file
o_1r	cl4	oh	104,2	108.1226	-24.1496	-23.3346	This work
o	cl	o_1r	104,6	108.1226	-24.1496	-23.3346	This work

- Constants from the SO₂ (cvff file entry) were borrowed for the **o-cl-o** quartic angle and was found to be an appropriate chemical fit for ClO species.
- The actual quartic angle from literature [7] was substituted for Theta0.
Constants and Theta0 from the * [p] * entry were borrowed for the **o – cl - oh** quartic angle. cl was substituted for ‘p’. ‘o’ and ‘oh’ were substituted for *.
Constants from the * [o] * entry were borrowed for the following quartic angle entries:
 - cl [o_1r] cl4 o_1r was substituted for ‘o’. cl and cl4 were substituted for *
 - cl4 [o_1r] cl4 o_1r was substituted for ‘o’. cl4 and cl4 were substituted for *
 - cl4 [o_1r] o_1r o_1r was substituted for ‘o’. cl4 and o_1r were substituted for *
 - cl4 [o_1r] o o_1r was substituted for ‘o’. cl4 and o were substituted for *
 - cl [o] o-1r cl and o_1r were substituted for *
 - cl [oh] ho oh was substituted for ‘o’. cl was substituted for *
 - cl [o_1r] cl4: The quartic angle for cl - o – cl bond was extracted from literature [7] and was substituted for θ_0
 - cl4 [o_1r] o_1r: The quartic angle for cl - o – o bond was extracted from literature [11] and was substituted for θ_0
 - cl4 [o_1r] o: The quartic angle for cl - o – o bond was extracted from literature [9] and was substituted for θ_0

- cl [o] o-1r: The quartic angle for cl - o – o bond was extracted from literature [9] and was substituted for θ_0
- cl [oh] ho: The quartic angle for cl -o – h bond was extracted from literature [7] and was substituted for θ_0

Constants from the o_1r [cl4] o_1r entry were borrowed for the following Quartic angles:

- a) o [1r – cl4] oh ! oh was substituted for ‘o_1r’
- b) o [cl] o_1r ! o was substituted for ‘o_1r’
- c) o_1r [cl4] oh ! Quartic angle, from literature [11] and was substituted for θ_0
- d) o [cl – o] 1r ! Quartic angle was extracted from literature [9] was substituted for θ_0

5.2.2.5 Wilson out of plane parameters

The Wilson out-of-plane potential is defined according to the angle between one bond from the central atom and the plane defined by the other two bonds [8]. The format of the out-of-plane potential (Wilson definition) section is:

$$E = K * (\text{Chi} - \text{Chi}0)^2$$

Where:

I, J, K and L are the atom types of the four atoms involved in the out-of-plane term.

J being the central atom: Chi (χ_0) is the reference angle in degrees [8].

Wilson out of plane parameter are presented in Table 5.5.

Table-5.5: Wilson out of Plane Parameters

I	J	K	L	K-Chi	Chi0	Force field Reference
*	p	*	*	0.000	0.000	cvff file
o	cl	o	o	0.000	0.000	This work
o	cl	o	oh	0.000	0.000	This work
o	cl	o	h	0.000	0.000	This work
o	cl	o	cl	0.000	0.000	This work
cl4	o_1r	o_1r	cl4	0.000	0.000	This work
o_1r	cl4	o_1r	cl4	0.000	0.000	This work
cl	o	o_1r	cl4	0.000	0.000	This work

The constants from entry * -**p**- * - * were used for all Wilson out of plane entries, **cl** was substituted for **p**. The relevant atoms were substituted for *.

5.2.2.6 Torsion parameters

The three-term cosine expansion of the torsion potential is:

$$E = \sum_{(n=1,3)} \{V(n) * [1 - \cos (n*\Phi - \Phi_0(n))]\}$$

Where:

V_1 , V_2 and V_3 are the barrier heights in kcal/mol

Φ (ϕ) is the current torsion angle.

Φ_{01} , Φ_{02} and Φ_{03} are the reference torsion angles in degrees.

(The reference angles are usually 0° or 180°)

(The atoms are bonded to each other in the order *I-J-K-L*) [10].

Torsion parameters are presented in Table 5.6.

Table-5.6: Torsion Parameters

I	J	K	L	V(1)	Phi1(0)	V(2)	Phi2(0)	V(3)	Phi3(0)	Force field Reference
*	c	o	*	0	0	0	0	0.13	0	cvff file
o	cl	o	o	0	0	0	0	0.13	0	This work
o	cl	o	h	0	0	0	0	0.13	0	This work
o	cl	oh	ho	0	0	0	0	0.13	0	This work
o_1r	cl4	o_1r	cl	0	0	0	0	0.13	0	This work
o_1r	cl4	o_1r	cl4	0	0	0	0	0.13	0	This work
o_1r	cl4	o_1r	o	0	0	0	0	0.13	0	This work
o_1r	cl4	oh	ho	0	0	0	0	0.13	0	This work
o	cl	o_1r	cl4	0	0	0	0	0.13	0	This work
o	cl	o	oh	0	0	0	0	0.13	0	This work
o	cl	oh	o	0	0	0	0	0.13	0	This work
o	cl	o	cl	0	0	0	0	0.13	0	This work
o	cl	o	o_1r	0	0	0	0	0.13	0	This work
o_1r	cl4	o_1r	o_1r	0	0	0	0	0.13	0	This work
*	o	o	*	0	0	0	0	1	0	cvff file
cl4	o_1r	o_1r	cl4	0	0	0	0	1	0	This work
cl	o	o_1r	cl4	0	0	0	0	1	0	This work

The constants from * - cl - o - * and * - o - o - * in *pcff+* were used for torsion entries derived for the chlorine species.

5.3 Heats of Formation

5.3.1 Ensemble Models -- Potential Energy of Chlorine Species

GEMC Simulations were undertaken for all neutral and radical species, assuming that a good approximation of molecular geometries has been achieved, irrespective of the uneven electron population for radicals. The potential energy (U_i) for each ensemble model was determined at each step of addition of molecular units, followed by a dynamic step and the minimized energy extracted.

A typical GIBBS (*GEMC*) energy profile for Cl_2O_2 is graphically presented in Figure-5.1 to demonstrate the effective inflection point, when the interaction of a sufficient number of molecules have reached a global equilibrium.

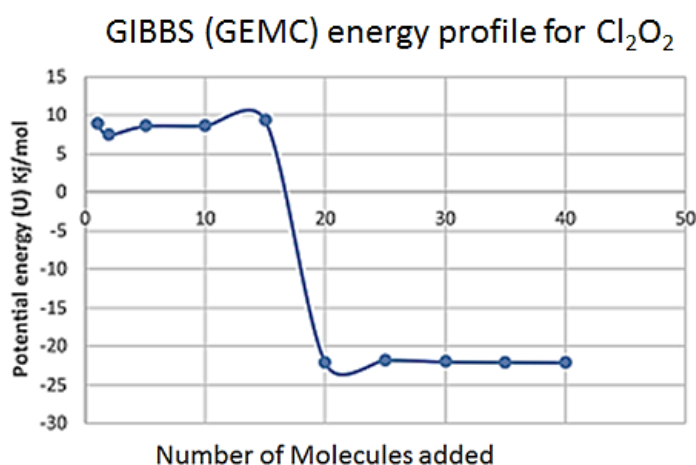


Figure-5.1: *GEMC* Potential Energy profile for Cl_2O_2

The model ensemble energies were recorded at:

- a) The minimum point
- b) Before the minimum point
- c) Beyond the minimum Internal Energy

Neutral species structures were successfully computed under *GEMC* with the exception of HClO , HClO_2 and HClO_3 due to lacking force field parameters.

It must further be noted that applying *MOPAC* Hamiltonians to describe an optimum model status, may differ with **single-molecular** models (in vacuum or periodic) and rely on optimum basis sets to effectively describe an **ensemble model**.

Table-5.7 displays the *MOPAC* calculated ΔH_f energies for the ensemble models. Reference ΔH_f energies are listed as well. In cases where no reference ΔH_f was available, the more consistent energy obtained through different Hamiltonians was reported.

In Table-5.7 the ensemble model energies noted as: “Model before minimum (U)”, “Model at minimum (U)” and “Model after minimum (U)” are shown. These are listed to demonstrate that a global minimum had been reached. Optimum ΔH_f energies from selected *MOPAC* Hamiltonians, are given in the last column.

Table-5.7: Heat of Formation of ensemble neutral models applying the seven Hamiltonians in MOPAC-2016

Ensemble model Energy nodal points	Literature Ref. ΔH_f (kcal/mol)	HEAT OF FORMATION (kcal/mol)							ΔH_f (Calculated)		
		AM1	MNDO	MNDOD	PM3	PM6	PM7	RM1	Model	Functional	ΔH_f (kcal/mol)
OCtO before minimum (U)	24.36	151.31	218.44	72.68	47.97	83.88	83.49	91.26	Single Molecule	MNDOD	25.47
OCtO at minimum (U)		92.06	104.09	20.13	-109.64	4.85	-13.7	-9.66			
OCtO after minimum (U)		41.56	84.09	21.51	-64.49	34.29	-11.09	16.62	ensemble Model		20.13
CtOOCt before minimum (U)	31.38	30.4	87.26	27.62	17.91	23.1	18.85	15.64	Single Molecule	AM1	33.69
CtOOCt at minimum (U)		30.48	33.36	35.36	18.62	21.22	10.08	13.56			
CtOOCt after minimum (U)		29.45	32.27	33.88	17.05	19.91	9.47	12.79	ensemble Model		30.48
CtO before minimum (U)	24.31	31.05	35.85	18.98	5.81	21.61	15.6	7.23	Single Molecule	PM6	25.47
CtO at minimum (U)		41.64	48.48	21.87	-24.47	5.73	8.18	11.66			
CtO after minimum (U)		34.26	50.35	22.81	-14.95	15.34	6.35	9.14	ensemble Model		MNDOD
CtOCtO ₂ before minimum (U)	37.4	154.97	215.95	55.61	-74.72	26.31	32.35	51.75	Single Molecule	MNDOD	37.59
CtOCtO ₂ at minimum (U)		144.69	215.39	39.25	-167.11	-13.23	-23.35	6.34			
CtOCtO ₂ after minimum (U)		143.07	214.92	40.16	-187.73	-17.67	-27.37	9.65	ensemble Model		39.25
CtO ₂ -O-CtO ₂ before minimum (U)	N/R	284.64	398.04	40.83	-198.48	18.06	22.26	50.42	Single Molecule	RM1	50.55
CtO ₂ -O-CtO ₂ at minimum (U)		268.89	400.56	27.67	-381.76	-53.16	-75.27	-2.92			
CtO ₂ -O-CtO ₂ after minimum (U)		268.36	396.55	27.17	-392.01	-64.15	-79.88	-6.2	ensemble Model		50.42

Table-5.7 Continued

Ensemble model. Energy nodal points	Literature Ref. ΔH_f (kcal/mol)	HEAT OF FORMATION (kcal/mol)							ΔH_f (Calculated)		
		AM1	MNDO	MNDOD	PM3	PM6	PM7	RM1	Model	Functional	ΔH_f (kcal/mol)
$\text{ClO}_2\text{-O-O-ClO}_2$ before minimum (U)	N/R	111.44	197.15	40.35	-72.25	31.78	36.10	52.88	single molecule	RM1	79.42
$\text{ClO}_2\text{-O-O-ClO}_2$ at minimum (U)		104.97	150.25	37.26	-163.38	-4.55	-9.21	23.46	ensemble model	AM1	104.97
$\text{ClO}_2\text{-O-O-ClO}_2$ after minimum (U)		102.55	153.47	36.89	-158.55	-4.99	-8.56	25.31			
ClOOCLO_3 before minimum (U)	69.00	193.21	236.12	73.95	-199.46	48.78	65.84	129.36	single molecule	PM7	73.84
ClOOCLO_3 at minimum (U)		171.61	252.84	83.87	-371.14	53.66	10.89	11.96	ensemble model		65.84
ClOOCLO_3 after minimum (U)		158.03	249.98	84.47	-359.16	47.16	55.19	27.74			
$\text{ClO}_2\text{-O-ClO}_3$ before minimum (U)	72.4	276.34	347.19	87.40	13.02	67.15	58.41	156.56	single molecule	MNDOD	47.01
$\text{ClO}_2\text{-O-ClO}_3$ at minimum (U)		284.29	358.71	92.12	12.30	67.28	60.18	157.11	ensemble model	PM6	67.28
$\text{ClO}_2\text{-O-ClO}_3$ after minimum (U)		283.84	357.55	92.37	10.85	67.22	59.65	157.25			
$\text{ClO}_3\text{-O-ClO}_3$ before minimum (U)	82.6	135.89	261.66	56.45	-260.12	78.94	83.52	33.03	single molecule	PM6	77.47
$\text{ClO}_3\text{-O-ClO}_3$ at minimum (U)		150.33	261.66	91.78	-310.25	33.56	-5.52	53.61	ensemble model		78.94
$\text{ClO}_3\text{-O-ClO}_3$ after minimum (U)		160.17	261.66	65.69	-324.62	34.88	-8.48	49.11			
ClClO before minimum (U)	31.84	55.88	78.87	50.01	18.98	44.74	38.43	34.93	single molecule	MNDOD	28.63
ClClO at minimum (U)		55.44	83.91	54.73	18.64	43.58	34.08	34.41	ensemble model	RM1	34.41
ClClO after minimum (U)		56.58	82.00	52.48	19.28	44.85	36.56	35.28			
ClOO before minimum (U)	24.56	5.56	20.78	23.28	12.02	0.74	7.36	0.60	single molecule	MNDOD	23.28
ClOO at minimum (U)		33.71	50.53	28.26	18.95	21.89	15.91	0.40	ensemble model		28.26
ClOO after minimum (U)		5.19	24.18	28.52	13.57	-4.44	-5	-36.79			
HOCl before minimum (U)	-18.36	-13.71	3.25	0.03	-31.1	-27	-28.18	-20.71	single molecule	PM6	-17.81
HOCl at minimum (U)		-16.57	-0.39	-3.40	-32.76	-18.56	-31.99	-22.44	ensemble model		-18.56
HOCl after minimum (U)		-16.19	1.49	-0.44	-32.09	-17.51	-31.19	-21.68			

Table-5.7 Continued		HEATS of FORMATION (kcal/mol)							ΔH_f (calculated)		
Ensemble model Energy nodal points	Literature Ref. ΔH_f (kcal/mol)	AM1	MNDO	MNDOD	PM3	PM6	PM7	RM1	Model	Functional	ΔH_f (kcal/mol)
HOClO before minimum (U)	4.94	50.01	75.94	11.18	-10.42	21.31	14.43	29.56	single molecule	PM7	6.77
HOClO at minimum (U)		52.05	85.54	23.18	-11.19	18.9	6.49	28.78	ensemble model		6.49
HOClO after minimum (U)		52.44	86.12	25.76	-7.95	22.08	8.52	30.63			
HOClO ₃ before minimum (U)	0.43	220.98	281.3	1.17	-126.05	-9.49	-25.44	28.42	single molecule	MNDOD	3.71
HOClO ₃ at minimum (U)		28.42	282.58	3.17	-143.86	-16.06	-30.61	26.02	ensemble model		3.17
HOClO ₃ after minimum (U)		81.54	287.76	3.17	-110.97	-11.02	-36.04	73.5			
HOClO ₂ before minimum (U)	1.1	146.79	193.2	-2.82	-14.77	44.4	-10.18	98.57	single molecule	MNDOD	-3.27
HOClO ₂ at minimum (U)		144.9	198.6	-11.67	-11.25	46.15	-66.3	99.02	ensemble model		-2.82
HOClO ₂ after minimum (U)		141.66	195.54	-11.32	-22.53	38.45	42.84	93.79			
HOOCtO ₂ before minimum (U)	16.56	29.78	174.26	16.04	-163.7	-8.4	100.99	5.64	single molecule	PM6	19.25
HOOCtO ₂ at minimum (U)		63.52	175.04	19.33	-179.74	-18.81	-29.29	1.17	ensemble model	MNDOD	19.33
HOOCtO ₂ after minimum (U)		35.75	174.91	18.63	-183.26	-22.65	-33.78	3.22			
HOOCt before minimum (U)	-0.32	-0.88	0.91	3.94	-18.27	-1.6	-5.07	-9.32	single molecule	RM1	-0.68
HOOCt at minimum (U)		-3.56	3.94	4.65	-19.01	-4.83	-17.56	-23.33	ensemble model	AM1	-3.56
HOOCt after minimum (U)		-4.1	5.19	6.28	-19.04	-5.73	-18.4	-21.61			

Table-5.7 Continued

Ensemble model Energy nodal points	Literature Ref. ΔH_f (kcal/mol)	HEATS of FORMATION (kcal/mol)							ΔH_f (calculated)		
		AM1	MNDO	MNDOD	PM3	PM6	PM7	RM1	Model	Functional	ΔH_f (kcal/mol)
HOOOCl before minimum (U)		14.5	15.02	12.55	-4.73	7.89	3.70	1.12	single molecule	MNDOD	11.88
HOOOCl at minimum (U)	10.11	10.68	19.89	21.80	-4.49	1.75	-9.77	-5.37			
HOOOCl after minimum (U)		10.9	20.15	22.84	-5.34	0.85	-10.46	-9.51	ensemble model	AM1	10.68
ClO ₃ -O-O-ClO ₃ before minimum (U)		428.47	514.59	132.79	75.26	116.31	107.13	225.37	single molecule	MNDOD	108.37
ClO ₃ -O-O-ClO ₃ at minimum (U)	111.40	430.80	519.64	126.35	66.72	107.45	100.03	223.79			
ClO ₃ -O-O-ClO ₃ after minimum (U)		432.53	522.22	130.02	68.53	109.60	102.64	223.43	ensemble model	PM6	107.45
ClOCl before minimum (U)		19.52	31.39	16.29	-16.12	8.63	4.78	9.91	single molecule		19.45
ClOCl at minimum (U)	18.64	19.73	37.35	29.9	-14.13	7.79	-5.91	-8.92		AM1	
ClOCl after minimum (U)		19.48	36.6	29.55	-14.16	6.47	-6.05	-12.53	ensemble model		19.73
ClOClO before minimum (U)		40.67	58.77	18.76	-1.73	19.57	16.76	20.2	single molecule	PM6	38.19
ClOClO at minimum (U)	39.77	38.48	60.19	23.07	-39.03	4.78	4.16	8.12			
ClOClO after minimum (U)		37.77	59.33	21.97	-21.21	6.25	4.03	9.06	ensemble model	AM1	38.48

Table-5.7 Continued

Ensemble model. Energy nodal points	Literature Ref. ΔH_f (kcal/mol)	HEAT OF FORMATION (kcal/mol)							ΔH_f (Calculated)		
		AM1	MNDO	MNDOD	PM3	PM6	PM7	RM1	Model	Functional	ΔH_f (kcal/mol)
CtOOO before minimum (U)	53.32	42.65	46.29	41.71	3.83	8.47	12.92	32.68	single molecule	MNDO	45.51
CtOOO at minimum (U)		41.35	48.32	50	6.86	2.21	1.56	7.88	ensemble model	MNDOD	50
CtOOO after minimum (U)		41.93	71.23	70.42	18.06	40.91	28.27	26.19			
OCtCtO ₂ before minimum (U)	46.2	127.51	243.82	83.73	-83.46	43.96	97.34	49.71	single molecule	RM1	50.13
OCtCtO ₂ at minimum (U)		110.92	245.76	52.86	-	-8.77	-13.24	2.74	ensemble model	MNDOD	52.86
OCtCtO ₂ after minimum (U)		142.52	226.62	52.67	-	-6.96	-24.39	10.67			
CtOCtO ₃ before minimum (U)	37.4	211.01	262.07	58.84	7.15	37.93	29.62	106.24	single molecule	PM7	49.49
CtOCtO ₃ at minimum (U)		211.78	261.91	61.21	8.08	39	30.43	107.32	ensemble model	PM6	39
CtOCtO ₃ after minimum (U)		212.09	263.52	61.86	8.75	39.43	31.15	107.58			
CtO ₂ CtO ₂ before minimum (U)	N/R	132.01	166.67	51.62	-46.3	22.6	21.51	60.74	single molecule	RM1	68.02
CtO ₂ CtO ₂ at minimum (U)		66.55	145.42	104.12	-117.22	-8.09	-11.98	11.84	ensemble model	AM1	66.55
CtO ₂ CtO ₂ after minimum (U)		68.11	154.04	24.57	-113.87	-9.69	-11.87	106.43			
CtO ₄ before minimum (U)	57.6	233.59	237.17	35.74	-87.59	70.83	61.15	147.87	single molecule	PM3	46.14
CtO ₄ at minimum (U)		227.73	243.25	77.65	-99.56	64.97	48.92	125.1	ensemble model	PM7	48.92
CtO ₄ after minimum (U)		228.67	244.04	78.66	-95.81	114.7	48.23	120.66			
CtO ₃ before minimum (U)	44	153.68	154.08	38.84	-23.37	48.62	53.19	81.31	single molecule	PM6	39.31
CtO ₃ at minimum (U)		99.55	216.84	20.17	-166.07	1.04	-16.84	41.52	ensemble model	RM1	41.52
CtO ₃ after minimum (U)		151.65	164.93	27.25	-136.88	-4.2	-16.94	50.98			

Optimum GEMC **ensemble** models extracted at the energy intervals outlined above, were subjected to *MOPAC-2016* computation to determine the ΔH_f Heats of Formation.

ΔH_f energies for **ensemble** models and **single** species using *MOPAC* analysis for ClOCl , ClOClO , ClOClO_2 , HOClO_3 , HOCl , HOClO , HOClO_2 and HOOOCl produced similar results.

- The majority of the ensemble model analyses offered overall Heat of Formation values similar to reference values when compared to single species energies for: ClOO , ClClO , ClOOO , ClO_3 , ClOOCl , ClOClO_3 , $\text{ClO}_2\text{-O-ClO}_3$, OClOO , $\text{ClO}_3\text{-O-ClO}_3$, HOClO , HOCl , HOClO_2 and HOOOCl
- A few single species i.e., ClO , ClClO_2 , Cl(O)O_2 , ClO_4 , ClOOClO_3 and OClO were the exception, producing results that matched reference values more closely than ensemble models.

A summary of the single molecular and ensemble model *MOPAC-2016* Heats of Formation obtained from ensemble model simulations, are presented in Table-5.8 below.

Table-5.8: Heat of Formation Derived from ensemble Model Space (GEMC Derived)
Calculated with MOPAC-2016.

ensemble Models	ΔH_f (calculated)			Literature Ref. ΔH_f (kcal/mol)
	MODEL Space	MOPAC Hamiltonian	ΔH_f (kcal/mol)	
ClO	single	PM6	25.47	24.31 [12]
	ensemble	MNDOD	21.87	
ClOO	single	MNDOD	23.28	24.56 [12]
	ensemble	MNDO	24.18	
ClOCl	single	AM1	19.45	18.62 [12]
	ensemble	AM1	19.73	
ClClO	single	MNDOD	28.64	31.84 [12]
	ensemble	RM1	34.41	
ClClO_2	single	PM6	28.96	29.16 [12]
	ensemble	MNDOD	22.34	
ClOOO	single	MNDO	45.54	53.29 [12]
	ensemble	MNDOD	50.00	

Table-5.8 Continued

Ensemble Models	ΔH_f (calculated)			Literature Ref. ΔH_f (kcal/mol)
	MODEL Space	MOPAC Hamiltonian	ΔH_f (kcal/mol)	
Cl(O)O ₂	single	MNDOD	63.00	63.37 [12]
	ensemble	MNDOD	66.14	
ClO ₃	single	PM6	39.31	44.30 [12]
	ensemble	RM1	41.52	
ClO ₄	single	PM3	50.54	57.60 [12]
	ensemble	PM7	48.92	
ClOClO	single	PM6	38.19	41.80[12]
	ensemble	AM1	38.48	
ClOOCl	single	AM1	33.69	31.37 [13]
	ensemble	AM1	30.48	
ClOClO ₂	single	MNDOD	38.45	37.80 [14]
	ensemble	MNDOD	39.25	
ClOClO ₃	single	PM7	49.52	37.40 [14]
	ensemble	PM6	39.00	
ClO ₂ ClO ₂	single	MNDOD	103.16	Not reported
	ensemble	AM1	66.55	
ClO ₂ -O-ClO ₂	single	RM1	50.55	Not reported
	ensemble	RM1	50.42	
ClOOClO ₃	single	PM7	69.23	69.00 [14]
	ensemble	PM7	65.84	
ClO ₂ -O-ClO ₃	single	MNDOD	46.97	72.40 [15]
	ensemble	PM6	67.28	
ClO ₃ -O-ClO ₃	single	PM6	77.66	82.60 [15]
	ensemble	PM6	78.94	
ClO ₂ -O-O-ClO ₂	single	RM1	79.42	Not reported
	ensemble	AM1	104.97	
ClO ₃ -O-O-ClO ₃	single	PM6	108.37	111.40 [9] ^s
	ensemble	PM6	107.45	
OClO	single	MNDOD	25.47	24.36 [12]
	ensemble	MNDOD	20.13	

Table-5.8 Continued

Ensemble Models	ΔH_f (calculated)			Literature Ref. ΔH_f (kcal/mol)
	MODEL Space	MOPAC Hamiltonian	ΔH_f (kcal/mol)	
OClOO	single	PM6	26.69	54.3 [12]
	ensemble	MNDOD	53.73	
OClClO ₂	single	RM1	50.13	46.2 [14]
	ensemble	RM1	49.71	
HOCl	single	PM6	-17.81	-18.36 [12]
	ensemble	PM6	-18.56	
HOClO	single	PM7	0.22	4.94 [12]
	ensemble	PM7	6.49	
HOClO ₂	single	MNDOD	-3.18	-1.10 [12]
	ensemble	MNDOD	-2.82	
HOClO ₃	single	MNDOD	3.71	0.43 [12]
	ensemble	MNDOD	3.17	
HOOC _l	single	RM1	-0.68	-0.32 [12]
	ensemble	AM1	-0.88	
HOOC _l O	single	RM1	22.05	21.54 [12]
	ensemble	MNDOD	22.7	
OOClO ₂	single	PM6	19.27	16.55 [12]
	ensemble	MNDOD	19.33	
HOOC _l	single	MNDOD	11.88	10.11 [12]
	ensemble	AM1	10.68	
HOOC _l O _l	single	PM6	24.78	22.7 [14]
	ensemble	PM6	21.58	

[§] Indicates that the ΔH_f (298K) of ClO₃-O-O-ClO₃ is calculated from energies by Beltran et al. (1999) [9].

Ensemble models subjected to *MOPAC-2016* analyses consistently, produced acceptable results. Over 60 % producing acceptable results compared to energies obtained for their single molecule counterparts.

No reference ΔH_f energies for ClO₂-O-ClO₂ and ClO₂-O-O-ClO₂ were available from the Open Literature. Calculated values of 50.42 kcal/mol and 104.97 kcal/mol were assigned respectively for these ensemble analyses. These energies were consistently obtained, by applying a number of Hamiltonians (*MOPAC*) which produced similar results or a result closest in value to their single

molecules counterparts. The energy of the single molecule analysis was selected based on which Hamiltonian produced the most sterically correct structure.

5.3.2 *GEMC Results*

ΔH_f energies obtained applying the GEMC ensemble model approach, consistently produced results that matched reference values and should in general be applied to derive at effective chemical properties.

5.4 Conclusion

Application of the GEMC (GCMC) approach in calculation the Heats of Formation for this group of C{O2 species has proven to be successful. But the technique relies on an appropriate set of force field parameters. Extending the molecular/compound description (even in vacuum space) validates the importance of a multiple molecular ensemble. In this study, the extraction of confirmed Heats of Formation proved invaluable. Extending the technique to other compounds and substances, cannot be claimed to offer the same result. It will however be beneficial in extending the force field tables to include parameters for ions and radicals, specifically for a selection of species supporting open shell electron configurations.

A Species-based level of thermodynamics can now be undertaken with confidence.

5.5 References

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CHAPTER 6: BONDING CHARACTER OF CHLORINE OXIDE

6.1 Introduction

The ClO group of species, exhibits a unique bonding combination of chlorine (a strong electronegative atom) with oxygen (a paramagnetic element). The atomic stoichiometry of having to consider these two elements in the construction of ClO₂ and its subspecies, including the isometric extension into larger subspecies with higher oxygen atom content, poses a complex bonding character to decipher. Effective atom electron potentials pose a further complication for quantum chemical analysis, with basis sets approximating (assumed) near acceptable structural geometries. The quantum chemical analyses undertaken, demonstrated a clear ability of the software systems applied (*Gaussian*, *MOPAC* and *VASP*), to produce promising results in structural geometries, and thermochemical properties which are agreement with available referenced values. This verification step helps one to carry out the species level water interactions outline in chapter 7 with confidence.

6.2 Variation in Heats of Formation

A selected few species are displayed in Table-6.1 depicting their bond distances and atomic partial charges, demonstrating the intrinsic charge distributions in these complex compounds.

Table-6.1: Selected few species extracted, to demonstrate their varying partial charge distributions (Mulliken) and bonding character in relation to their molecular spin states. Atom connectivity is presented in parenthesis. (A species naming convention has been adopted to portray bond sequencing)

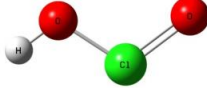

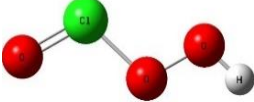
Species	Molecular Structure	Atomic Partial Charges	Spin State	Bond Distances (Å) (Atom Connectivity)
HOClO		O1: -0.26 Cl: 0.34 O2: -0.33 H: 0.24	singlet	Cl-O1 = 1.51 Cl-O2 = 1.77 O2-H = 0.98 (O1-Cl-O2-H)
ClOOCl		Cl1: 0.05 O1: -0.03 O2: -0.08 Cl2: 0.06	singlet	Cl1-O1 = 1.82 O1-O2 = 1.32 (Bridge) Cl2-O2 = 1.82 (Cl-O1-O2-Cl)
HOOClo		O3: -0.33 Cl: 0.53 O2: -0.22 O1: -0.32 H: 0.34	singlet	Cl-O3 = 1.51 Cl-O2 = 1.84 O2-O1 = 1.46 O1-H = 0.98 (O3-Cl-O2-O1-H)

Table-6.1 continued

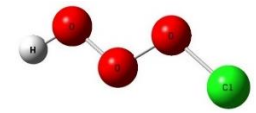
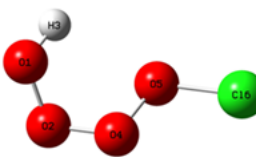
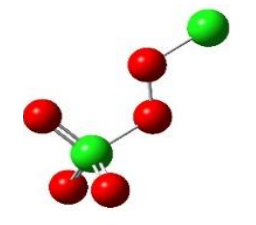
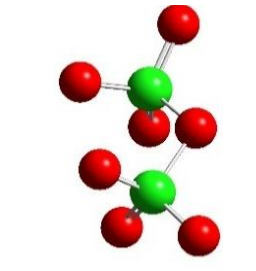
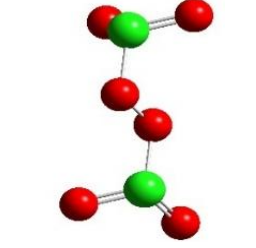
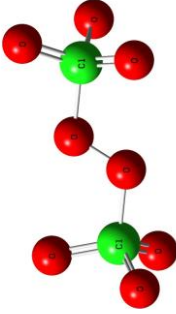
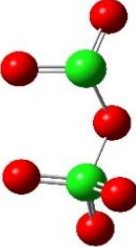

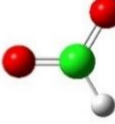
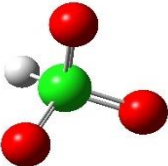
Species	Molecular Structure	Atomic Partial Charges	Spin State	Bond Distances (Å) (Atom Connectivity)
HOOC ℓ		Cl: 0.1 O3: -0.09 O2: -0.07 O1: -0.27 H: 0.34	singlet	Cl-O3 = 1.81 O3-O2 = 1.35 (Bridge) O2-O1 = 1.48 O1-H = 0.98 (O3-Cl-O2-O1-H)
HOOOOCl		Cl: 0.24 O4: -0.25 O3: -0.021 O2: 0.022 O1: -0.54 H: 0.59	singlet (triplet)	Cl-O4 = 1.63 O4-O3 = 1.75 O3-O2 = 1.26 (Bridge) O2-O1 = 1.57 O1-H = 0.98 (Cl-O4-O3-O2-O1-H)
ClOOClO ₃		Cl1: 0.93 O1: -0.11 O2: -0.30 O3: -0.32 O4: -0.25 O5: 0.061 Cl2: -0.0056	singlet	Cl1-O1 = 1.65 O1-O2 = 1.44 (Bridge) Cl2-O2 = 1.95 Cl2-O3 = 1.44 Cl2-O4 = 1.47 Cl2-O5 = 1.44 (Cl1-O1-O2-Cl2-O3{O4}{O5})
ClO ₃ -O-ClO ₃		Cl1: 0.99 O1: -0.28 O2: -0.28 O3: -0.28 O4: -0.26 Cl2: -0.99 O5: -0.30 O6: -0.28 O7: -0.28	singlet	Cl1-O1 = 1.44 Cl1-O2 = 1.44 Cl1-O3 = 1.48 Cl1-O4 = 1.90 Cl2-O4 = 1.94 Cl2-O5 = 1.42 Cl2-O6 = 1.42 Cl2-O7 = 1.46 (Cl1-O1{O2}{O3}-O4-Cl2-O5{O6}{O7})
ClO ₂ -O-O-ClO ₂		Cl1: 0.28 O1: -0.20 O2: -0.21 O3: 0.13 Cl2: 0.28 O4: 0.13 O5: -0.21 O6: -0.20	singlet	Cl1-O1 = 1.46 Cl1-O2 = 1.54 Cl1-O3 = 1.97 O3-O4 = 1.32 (Bridge) Cl2-O4 = 1.97 Cl2-O5 = 1.46 Cl2-O6 = 1.54 (O1{O2}-Cl1-O3-O4-Cl2{O5}{O6})

Table-6.1 continued

Species	Molecular Structure	Atomic Partial Charges	Spin State	Bond Distances (Å) (Atom Connectivity)
$\text{ClO}_3\text{-O-O-ClO}_3$		Cl1: 1.02 O1: -0.14 O5: -0.30 O6: -0.29 O7: -0.30 Cl2: 1.02 O2: -0.14 O8: -0.29 O9: -0.30 O10: -0.30	singlet	Cl1-O5 = 1.37 Cl1-O6 = 1.37 Cl1-O7 = 1.374 Cl1-O1 = 1.64 O1-O2 = 1.36 (Bridge) Cl2-O2 = 1.64 Cl2-O8 = 1.37 Cl3-O9 = 1.37 Cl4-O10 = 1.37 (Cl1-O5{O6}{O7}-O1-O2-Cl2-O8{O9}{O10})
$\text{ClO}_2\text{-O-ClO}_3$		Cl1: 0.97 O1: -0.28 O2: -0.25 Cl2: 0.78 O3: -0.26 O4: -0.35 O5: -0.29 O6: -0.31	singlet	Cl1-O1 = 1.41 Cl1-O2 = 1.43 Cl1-O3 = 1.82 Cl2-O3 = 1.93 Cl2-O4 = 1.90 Cl2-O5 = 1.43 Cl2-O6 = 1.77 (Cl1-O1{O2}-O3-Cl2-O4{O5}{O6})
HClO		Cl: 0.41 O: -0.47 H: 0.064	singlet	Cl-O = 1.56 Cl-H = 1.32 (H-Cl-O)
HClO_2		Cl: 0.32 O1: -0.46 O2: -0.46 H: 0.14	singlet	Cl-O1 = 1.47 Cl-O2 = 1.47 Cl-H = 1.36 (H-Cl-O1{O2})
HClO_3		Cl1: -0.94 O1: -0.33 O2: -0.33 O3: -0.33 H: 0.058	singlet (doublet)	Cl-O1 = 1.42 Cl-O2 = 1.42 Cl-O3 = 1.42 Cl-H = 1.33 (H-Cl-O1{O2}{O3})

A comprehensive collection of partial charges derived for all chlorine oxide species in this study, can be found in Table-A6 (APPENDIX).

The hydrogen terminated species, HOOOCl , HOOCl , HOClO and HOClO all display varying atomic partial charges at the chlorine positions, ranging from 0.96ϵ to 0.07ϵ --- the larger partial charges observed for chlorine bounded by two oxygen atoms. This large variation can be ascribed to varying bonding character, adopted with the central and neighbouring oxygen atoms. The assignment of

two possible spin-states (singlet and triplet) for HOOOOCl , can be tied to this vast variation in partial charge distribution. For HOOOOCl , HOOCOCl and ClOOCl a bridging double bond is observed (1.26 Å, 1.35 Å and 1.32 Å respectively) between the central two oxygen atoms, clearing the one major paramagnetic contribution, leaving chlorine and/or oxygen atoms on either end, to share a bond in a closed shell condition (as a dative bond). This can result in either a triplet for HOOOOCl or a singlet state, in which case a partial double-bond charge distribution has to be at play with the two atoms neighbouring the central double bond.

It is furthermore significant to observe the large partial charges (**0.97** and **0.86**) for the two chlorine atoms in the two structures (HOClO and $\text{HOOC}l\text{O}$ respectively) flanked on either side by oxygen atoms. The unique assignment of a singlet spin state for both these species, displaying a reduced bond distance of 1.51 Å (a typical Cl-O bond distance) between their terminal oxygen atoms and the chlorine atoms (considered as double bonds) may dictate the residual charge distribution on the inner oxygen atoms.

The complex construction of electron sharing and resulting bond character, is demonstrated in Table-6.1 for these species and may be the reason for their vulnerability to dissociate in media such as water, imposing a dipole moment of 1.85 Debye, through dipole-dipole interaction and charge transfer. This large bond distance observed between oxygen, bonded to two chlorine atoms, noted as 'bridge' bonding is noticeable in all structures, clearly assigned to single bond character.

Figure-6.1 presents a composite graph of all Heats of Formation values derived. empirical construction of the ionic species, supported by force field structure conditioning, were not possible and these were calculated as single species entities. Varying spin conditions were applied in all cases, also considering different basis sets. Table-4.3 (chapter 4) lists the observed spin-states.

Singlet and doublet spin conditions were considered during the calculations.

(ClOOClO₃) as Cl1-O1-O2-Cl2-O3{O4}{O5} which has two types of O-Cl bonds, i.e., a **bridged bond** between Cl2-O2 atoms at 1.95 Å, a bond distance of ±1.4 Å for the **terminal Cl2-O3{O4}{O5}** bonds and 1.65 Å between the single-valent bonded terminal Cl1-O1.

HClO, HClO₂ and HClO₃ are hypervalent configurations [1] (solely due to their terminal oxygen atoms) whose instability is attributed to an electropositive charged hypervalent halogen (Table-6.1), being associated to the electropositive H-atom, which appears to affect the relative stability order and is attributed to the degree of valency on the halogen [1].

Lee et al., [2] and suggested that the multi-valency of a halogen is due to one, two, or three lone-pair valence electrons achieving significant degrees of *d*-character, resulting in multiple *pd*-hybrid halogen bonds. He examined ClXO₂, and HXO type molecules (X = Cl) and found that these molecules undergo *pd* hybridization. This hybridization process entails *p* - *d* orbital promotion of lone-pair electrons. This imparts the hypervalent character to the chlorine molecule thereby allowing the formation of more than one bond thus increasing the electron count. This type of hypervalent bonding for a halogen is characterized by a large ionic component [2]. This large ionic component imparts high thermodynamic stability to -XO_z, and HXO type molecules.

Lee et al., proved that “-XO_z are very stable while the -XO species usually are less stable than the normal valent isomer -OX” (z = number of oxygen molecules and X = Cl) [2]. This change in stability in the *sp* hybridisation process occurs since the *d* shell becomes partially occupied which lowers its energy. This lower energy equates to more stability. This stability trend is clearly seen in the mid-section of the Figure-6.1 which represents -XO₂ type molecules such as ClOOCl, ClO₂-O-O-ClO₂, ClO₂-O-ClO₃ and the HXO type molecules such as HOOClo, HOOCl, HOOOCl, HClO, HClO₂ and HClO₃. The higher stability of the mentioned chlorine oxides is represented by the low Heats of Formation values.

Single Cl-O bonds are ionic (dative) in nature and as stated earlier, their thermodynamic stability is affected by various factors, one being electronegativity. Examination of the atomic charges of Cl in Cl=O bonds (Table-6.1) reveals that the poly-oxide species possess highly electropositive charged chlorine atoms. This electro-positivity indicates that these species are prone to chemical attack from radicals and most likely explains why the Cl=O bonds are easily broken and break up into small components becoming available for use in other chemical processes.

6.3 Conclusion

ClO₂ compounds and its associated sub-species display a peculiar bonding character. The combination of strong electronegativity (chlorine) in the presence (directly bonded) to elements with an unpaired electron configuration (oxygen) and in this case, enclosed into relatively small molecular structures, results in a complicated bonding arrangement. This is further demonstrated by the thermochemical instability of some of the species (indicated by their high Heat of Formation Figure-6.1) in forming excess amounts of O₂, Cl⁻, Cl₂, [ClO₂]⁻ and [ClO₃]⁻ which ultimately, could play a major role in pathogen control.

6.4 References

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CHAPTER 7: SPECIES BASED THERMODYNAMICS IN AQUEOUS MEDIUM

7.1 Objective

The ultimate objective was to demonstrate the optimum conditions for specific species identified, to persist in an aqueous medium, to play a role in pathogen control in water treatment.

- Extract the main chemical reaction schemes
- Identify the ClO_2 species behavior in an aqueous medium

7.2 Species-based Thermodynamics

FactSage is an amalgamation of two popular thermochemistry computational software packages namely: FACT-Win (formerly *F*A*C*T*) and *CHEMSAGE* (formerly *SOLGASMIX*) [1]. The initial programs were used to calculate thermochemical properties for pure substances and ideal gases. *FactSage* contains two types of thermochemical databases i.e.:

- A pure substances database
- A solution database

The “*Facility for the Analysis of Chemical Thermodynamics*” (FACT) solution databases is updated frequently via globally funded research programs [1].

The Gibbs energy minimisation module of *FactSage*, ‘EQUILIB’ which determines the concentration of compounds in an equilibrium state, was used to predict the sustainability of the studied chlorine species in aqueous medium at standard temperature and pressure. All chlorine species (with confirmed and/or attainable Heats of Formation for ‘unknowns’) were added to the *FactSage* compound database. Selected thermochemical properties C_p , ΔH_f and S derived in this study by semi-empirical and *Ab Initio* supported techniques for single molecules (ensuring correlation with ensemble derived energies) were populated into the compound database. Each species was reacted with H_2O in a mole ratio of water: species of 1:1 and the resultant product species recorded. Resultant mole fractions for products, lower than 10^{-5} were discarded. The product species that resulted from reaction with H_2O is tabulated in the Appendix, Table-A7.

The following product species were found to be prevalent in most aqueous reactions: O_2 , $[\text{ClOH}_2]^+$, Cl^- , and $[\text{ClO}_4]^-$. These species persist in the aqueous environment and act as precursors for further chlorine oxide interactions.

7.3 Populating the *FactSage* Software Database

The *FactSage* database was populated with: *GIBBS* Free Energies, Heats of Formation and Heat Capacities, of all chlorine oxide species studied – listed in Table-7.1

NOTE:

The *FactSage* software (in its current status -- Sept. 2022) applying the *EQUILIB* module, does not support the option to invoke (select) ionic species, but will report on their contribution (prior populating the database via the *COMPOUND* Module).

This imposed a restriction on the analysis of comprehensive chlorine oxide and sub-species analyses, which prompted the alternative approach of exposing each individual (neutral) species to water, in a 1:1 mol ratio. A detailed list of a predominant species, resulting from interaction of single species with water analyses are listed in Table-A7 (Appendix). This was used to derive at a comprehensive set of reaction schemes, in demonstrating the significant formation of specifically O_2 , Cl^\cdot , $[\text{ClO}_2]^\cdot$, $[\text{ClO}_3]^\cdot$ as species known, to be involved in pathogen control.

The individual Free Energy of Formation ($\Delta\mathbf{G}_f$) of each species in aqueous medium could not be obtained with *FactSage*. *FactSage* provides the overall Free Energy of Formation of each species with water. Hence, the Standard Free Energy of Formation ($\Delta\mathbf{G}_f$) of each species was either:

- Obtained from literature.
- Calculated using *Ab Initio* electronic structure results (*Gaussian-16*) as outlined by Ochterski, J W [2].

The *Gaussian-16* software does not offer $\Delta\mathbf{H}_f$ and $\Delta\mathbf{G}_f$ energies as a standard output property. A separate external $\Delta\mathbf{H}_f$ and $\Delta\mathbf{G}_f$ energies calculation was required. The calculation step was extracted from a publication by J. W. Ochterski [2] and the results presented in Table-S7.

Table-7.1: Calculated Gibbs Free Energies of Formation and Free Heat of Formation using values obtained with *Gaussian-16*

Species	ΔH_f (OK) (kcal/mol)	ΔH_f (298K) (kcal/mol)	ΔG_f (298K) (kcal/mol)	Species	ΔH_f (OK) (kcal/mol)	ΔH_f (298K) (kcal/mol)	ΔG_f (298K) (kcal/mol)
ClO	29.77	29.715	22.07	OCtO	24.50	23.89	7.45
[ClO] ⁻	-23.43	-23.46	-31.37	[OCtO] ⁻	-15.96	-16.30	-32.48
[ClO] ⁺	278.19	278.18	270.94	[OCtO] ⁺	278.90	278.24	261.31
ClOO	21.62	21.50	6.87	OCtOO	51.67	51.63	30.17
[ClOO] ⁻	-54.57	-54.75	-69.30	[OCtOO] ⁻	-8.70	-9.87	-33.85
[ClOO] ⁺	300.13	299.79	284.57	OCtClO ₂	147.45	146.67	113.47
ClOCl	21.46	21.03	5.11	[ClOH ₂] ⁺	204.09	202.38	179.42
[ClOCl] ⁻	-34.99	-35.26	-50.19	HOCl	-15.73	-16.44	-31.02
[ClOCl] ⁺	269.53	269.03	253.32	[HOCl] ⁻	54.74	54.25	38.96
ClClO	35.05	34.64	19.18	[HOCl] ⁺	239.76	239.06	224.84
[ClClO] ⁻	-53.42	-54.10	-69.56	HOCtO	14.44	13.01	-11.04
[ClClO] ⁺	276.12	275.83	260.96	[HOCtO] ⁻	-55.76	-57.33	-80.78
ClClO ₂	37.86	36.79	11.04	[HOCtO] ⁺	240.42	238.93	215.1
ClOOO	68.41	67.8	43.48	HOCtO ₂	13.26	11.17	-23.15
[ClOOO] ⁻	-20.62	-21.94	-47.99	HOCtO ₃	14.84	12.13	-32.56
Cl(O)O ₂	69.75	69.46	46.56	[HOCtO ₃] ⁺	327.65	325.73	283.97
[Cl(O)O ₂] ⁻	-20.62	-21.94	-47.99	HOOCt	4.06	2.60	-21.57
ClO ₃	49.16	47.79	21.47	[HOOCt] ⁻	-9.90	-10.60	-32.12
[ClO ₃] ⁻	-34.08	-35.31	-61.76	[HOOCt] ⁺	254.85	253.29	229.36
[ClO ₃] ⁺	330.74	329.73	303.30	HOOCtO	29.56	27.78	-5.72
ClO ₄	76.78	74.64	37.12	HOOCtO ₂	40.10	37.26	-7.35
[ClO ₄] ⁻	-40.15	-42.11	-79.79	HOOOCt	29.47	27.44	-6.50
ClOClO	56.06	54.98	29.40	HOOOCt	30.21	28.18	-14.83
ClOOCl	39.29	38.67	13.96	HClO	34.60	33.89	19.39
ClOClO ₂	50.56	48.83	12.93	HClO ₂	49.88	48.30	23.82
ClOClO ₃	67.92	65.77	19.98	HClO ₃	48.51	46.07	11.05
ClO ₂ ClO ₂	73.08	69.78	21.80	H ₂ O	-49.76	-50.44	-64.83
ClO ₂ -O-ClO ₂	84.34	81.21	24.39	CO ₂	-410.17	-410.02	-426.77
ClOOClO ₃	75.18	72.17	15.45	C ₂ H ₆	144.51	140.65	85.19
ClO ₂ -O-ClO ₃	89.75	85.25	16.03	OH(g)			-33.15 [3]
ClO ₃ -O-ClO ₃	93.30	89.64	13.61	H ⁺			-6.28 [4]
ClO ₂ -O-O-ClO ₂	107.20	104.17	37.96	HCl			69.90 [5]
ClO ₃ -O-O-ClO ₃	124.17	118.45	28.50	[HOO] ⁻			369.50 [6]
[ClO ₃ ClO ₃] ⁻²	110.20	105.74	36.14	Cl ⁻			-131.20 [3]

7.4 GIBBS Free Energy of Reactions

In an attempt to verify the major species in an aqueous medium with chlorine oxides achieved with *FactSage*, a possible reaction scheme is presented in Figure-7.1. The evaluation of ClO_2 behaviour in an aqueous medium was considered. The Standard Free Energy of Formation of the individual species (Table-7.1) was used to determine the overall reaction *GIBBS* Free Energy of Formation of the participating reactions.

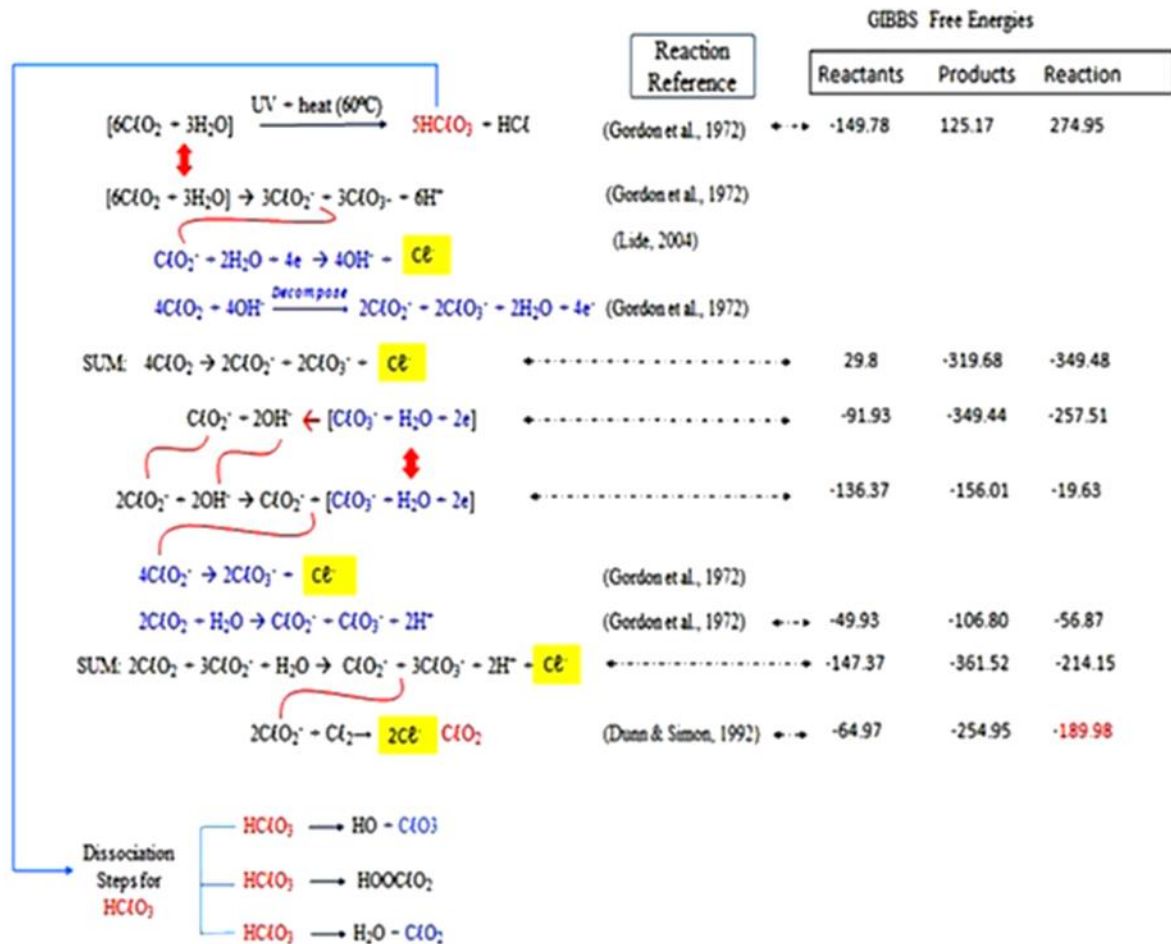


Figure-7.1: Sequence of the main reactions, some of their inter-dependencies and the distinct formation of Cl^- , $[\text{ClO}_2]^-$ and $[\text{ClO}_3]^-$ in an aqueous medium
References [8,9,10] refer to the reaction sequences.

The reaction scheme considers the overall ClO_2 breakdown products released in an aqueous medium taking into consideration decomposition and dissociation reactions. Neutral and alkaline aqueous medium reaction were considered. The final reaction products of Cl^- , $[\text{ClO}_2]^-$, $[\text{ClO}_3]^-$ verifies the major species identified by *FactSage* with the exception of O_2 . The decomposition products of ClO_2 in sunlight is O_2 and Cl_2 [7], however all major species identified are verified in the reaction scheme. The majority of reaction *GIBBS* Free Energies (ΔG_f) reveal negative values, indicating that reactions are occurring spontaneously.

7.5 Conclusion

- Progressive partial regeneration of ClO_2 following consecutive reactions, with gradual depletion into other sub-species.
- The results obtained, identifies the following species: O_2 , Cl^- , $[\text{ClO}_2]^-$, $[\text{ClO}_3]^-$ to be the main products in a large proportion of the reactions.
- This is further demonstrated by the thermochemical instability of some of the species (indicated by their high Heat of Formation values) in Chapter 6, Figure-6.1 in forming excess amounts of O_2 , Cl^- , Cl_2 , $[\text{ClO}_2]^-$ and $[\text{ClO}_3]^-$ which in turn, could play a major role in pathogen control.
- It was observed (Appendix Table-A7) that all chlorine oxide species containing more than one chlorine atom (displaying lower Heats of Formation) resulted in the highest product components of: O_2 and Cl^- as basic precursors for further interactions.

7.6 References

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CHAPTER 8: CONCLUSION

8.1 Conclusion

This study was undertaken both as a theoretical and investigational computational approach. It exemplifies the core of this study in support of water purification treatment, applying ClO_2 as an oxidative agent.

The endeavour to create a *GIBBS* (GEMC) ensemble simulation environment (Chapter 5), has confirmed that acceptable thermochemical properties can be derived. This also offered optimum molecular geometries, in conjunction with subsequent minimization steps, applied during *Ab Initio* analysis – a joint approach. Not only can electronic properties be derived but Heat Capacity, Entropy and Free Energies as well. However, it is imperative for the *GIBBS* (GEMC) Canonical simulations to be supported by a proper set of force field parameters.

The ultimate objectives to identify the major species derived from ClO_2 through a computational approach have achieved all facets of:

- a) optimum molecular geometries
- b) deriving thermochemical properties
- c) the ability to generate chemical reaction schemes.

The outcome of this work should lead to further studies in scaling these thermochemical observations into a finite element approach, to determine micro-scale physical interaction in quantifying macro-scale coagulation, in particular the interactions with fibrous material (NOM).

Considering the global water scarcity, this work will be useful, to understand the full benefits (health and economic) of ClO_2 in water purification processes, justifying its use in the water treatment industry.

8.2 Future work

A meso-scale investigation could utilize the results from this study, applied to various engineering initiatives to quantify the impact on inline treatment facilities.

- a) Possible improvement in NOM removal upon the addition of ClO_2 as a pre-treatment agent was observed, which primed the investigation into the effects of ClO_2 species (as oxidant) have on aqueous NOM and coagulation [1], [2].
- b) ClO_2 also appears to alter the chemical/physical state of dissolved and suspended solids, allowing for removal in a clarifier [1] [2]. These observations substantiate a probe into zeta-potential investigations of NOM particles once exposed to ClO_2 .
- c) A further study to identify the chemical species with the most appropriate ability to kill pathogens, is important and should be included in further studies. This implies an extensive quantum chemical resolution of the highest rate of charge transfer on an extreme model size,

between species and the environment. This can be accomplished through finite element analysis (an engineering feat), having the correct (meso-scale) charge dissipation of the large (molecular) model.

8.3 References

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APPENDIX

A4 - SUPPLEMENTARY DATA FOR CHAPTER 4

Tables-A4-1 and A4-2: Calculation steps [7] to extract Heat of Formation, derived from *Gaussian-16* applying **B3LYP** theory.

Table A4-1: Example of Heat of Formation calculation derived from molecular electronic energies, calculated with output from *Gaussian-16*

	OCℓO	O	Cl	H	Units
ϵ_0 (B3LYP)	-610.582863	-75.100483	-460.181573	-0.502428	Hartree
ϵ_{ZPE}	0.005783	0.000000	0.000000	0.000000	Hartree
E_{tot}	0.00894	0.001416	0.001416	0.001416	Hartree
H_{corr}	0.009884	0.002360	0.002360	0.002360	Hartree
G_{corr}	-0.019255	-0.013508	-0.013508	-0.010654	Hartree

Table-A4-2: Experimental enthalpies of formation of elements (kcal/mol) from JANAF tables [1]

Elements	$\Delta H_f(0K)$	$\Delta H_f(298K)$	$\Delta H_f(298K) - \Delta H_f(0K)$
H	51.63	52.64	1.01
O	58.99	60.03	1.04
Cl	28.59	29.69	1.10

a) First calculate $\Delta H_f(M, 0K)$ for each molecule: [2]

$$\Delta H_f(M, 0K) = \sum_{atoms} x \Delta H_f(X, 0K) - \sum D_0(M)$$

$$\begin{aligned} \Delta H_f(M, 0K) &= \sum_{atoms} x \Delta H_f(X, 0K) - [\sum x \epsilon_0(X) - \epsilon_0(M) - \epsilon_{ZPE}(M)] \\ &= \sum_{atoms} x \Delta H_f(X, 0K) - [627.51 \sum x \epsilon_0(X) - \epsilon_0(M) - \epsilon_{ZPE}(M)] \end{aligned}$$

Where: M = Molecule

X = Elements which make up M

x = Number of atoms of X in M

$\sum D_0(M)$ = Atomization energy of the molecule

$\epsilon_{ZPE}(M)$ = Zero Point energy of the species

$\epsilon_0(M)$ = Total energy of the species

H_{corr} = Thermal correction to Enthalpy

E_{tot} = Total electronic energy

G_{corr} = Correction to the Gibbs Free Energy

Multiply by 627.51 = Conversion (Hartree to kcal)

b) $\Delta H_f(M, 298K)$ for each molecule can be obtained as:

$$\Delta H_f(M, 298K) = \Delta H_f(M, 0K) + [H^\circ_M(298K) - H^\circ_M(0K)] \\ - \sum_{atoms} x [H^\circ_x(298K) - H^\circ_x(0K)]$$

c) $\Delta_f G^\circ(M, 298K)$ is determined as:

$$\Delta_f G^\circ(298K) = \Delta H_f(298K) - 298.15[\sum S_{(X, 298K)} - S_{(M, 298K)}] \quad [2] \\ = \Delta H_f(298K) - 298.15[\sum S_{(X, 298K)} \\ - 627.51(H_{corrM} - G_{corrM})/298.15] \quad (\text{Table-A4.1})$$

Elemental Entropies (cal/mol K):

$$S_{(298K)} Cl = 39.481$$

$$S_{(298K)} H = 27.418$$

$$S_{(298K)} O = 38.494$$

$$S_{(298K)} C = 37.787$$

d) EXAMPLE (OCℓO):

$$H_f(\text{OC}\ell\text{O}, 0K) = (2*58.99 + 28.59) - \quad ! \text{Heats of Formation at } 0K \\ \mathbf{627.51} * \quad ! \text{Conversion: Hartree to kcal/mol} \\ [2(-75.1005) + (-460.1816) \quad ! \sum x \epsilon_0(X) \\ -(-610.583 + 0.005783)] \quad ! \epsilon_0(M) - \epsilon_{ZPE}(M) (\text{Table-A4.2}) \\ = 146.57 - \mathbf{122.124} \\ = \mathbf{24.45} \text{ kcal/mol}$$

e) To calculate the $\Delta H_f(\text{OC}\ell\text{O}, 298K)$:

$$\Delta H_f(\text{OC}\ell\text{O}, 298K) = \mathbf{24.45} + 627.51 * [0.009884 - 0.005783] - (2*1.04 + 1.10) \\ = 24.45 + 2.57342 - 3.18 \\ = \mathbf{23.84} \text{ kcal/mol}$$

f) $\Delta G_f(\text{OC}\ell\text{O}, 298K)$ is determined as:

$$\Delta G_f(\text{OC}\ell\text{O}, 298K) = \mathbf{23.84} - 298.15 [(2*38.494 + 39.481)/1000 \quad ! \text{Convert to kcal} \\ - \mathbf{627.51}(0.009884 - (-0.01926))/298.15] \quad (\text{Table-A4.1}) \\ = \mathbf{23.84} - 298.15 (0.116469 - 0.061338) \\ = \mathbf{7.41} \text{ kcal/mol}$$

A4.3 - Gaussian-16 Thermochemistry outputs

Electronic energies and derived molecular Heats of Formation of the sixty (60) chlorine oxide species.

Table A-4.3: Gaussian DFT/B3LYP/cc-PV5Z output energies (Hartree) and associated Heats of Formation (kcal/mol)

Species	ϵ_0	ϵ_{ZPE}	E_{tot}	H_{corr}	G_{corr}	ΔH (OK)	ΔH (298K)	ΔG (298K)
O (cc-pv5z)	-75.045501	0	-0.00142	-0.00236	0.014952			
Cl (cc-pv5z)	-460.015052	0	-0.00142	-0.00236	0.015677			
H (cc-pv5z)	-0.50142	0	-0.00142	-0.00236	0.010654			
O (6-311++G(3df,2p))	-74.989847	0	0.001416	0.00236	-0.01351			
Cl (6-311++G(3df,2p))	-459.489755	0	0.001416	0.00236	-0.01392			
H (6-311++G(3df,2p))	-0.499995	0	0.001416	0.00236	-0.01568			
ClO	-535.377	0.002787	0.005163	0.006107	-0.01876	29.77	29.715	22.07
[ClO] ⁻	-535.461	0.001987	0.004407	0.005351	-0.01909	-23.43	-23.46	-31.37
[ClO] ⁺	-534.98	0.001753	0.004201	0.005145	-0.02037	278.19	278.18	270.94
ClOO	-610.586	0.004618	0.008553	0.009497	-0.02252	21.62	21.50	6.87
[ClOO] ⁻	-610.707	0.004176	0.007997	0.008941	-0.02321	-54.57	-54.75	-69.30
[ClOO] ⁺	-610.142	0.003875	0.00745	0.008394	-0.0227	300.13	299.79	284.57
ClOCl	-995.618	0.003612	0.007145	0.00809	-0.02235	21.46	21.03	5.11
[ClOCl] ⁻	-995.706	0.001233	0.005028	0.005973	-0.02604	-34.99	-35.26	-50.19
[ClOCl] ⁺	-995.223	0.003982	0.007396	0.00834	-0.02244	269.53	269.03	253.32
Cl ₂ O	-995.597	0.00385	0.007412	0.008357	-0.02282	35.05	34.64	19.18

Table A-4.3 continued

Species	ϵ_0	ϵ_{ZPE}	E_{tot}	H_{corr}	G_{corr}	ΔH (OK)	ΔH (298K)	ΔG (298K)
$[\text{Cl}_2\text{O}]^*$	-995.645	0.002096	0.005615	0.006559	-995.645	-53.42	-54.10	-69.56
$[\text{Cl}_2\text{O}]^+$	-995.212	0.002828	0.006574	0.007518	-995.212	276.12	275.83	260.96
ClClO_2	-1070.79	0.009595	0.013771	0.014715	-1070.79	37.86	36.79	11.04
ClOOO	-685.709	0.007877	0.012692	0.013636	-685.709	68.41	67.8	43.48
$[\text{ClOOO}]^-$	-685.851	0.007116	0.0108	0.011744	-685.851	-20.62	-21.94	-47.99
Cl(O)O_2	-685.707	0.00776	0.01308	0.014024	-685.707	69.75	69.46	46.56
$[\text{Cl(O)O}_2]^-$	-685.851	0.007469	0.011147	0.012091	-685.851	-20.62	-21.94	-47.99
ClO_3	-685.743	0.010472	0.014074	0.015018	-685.743	49.16	47.79	21.47
$[\text{ClO}_3]^-$	-685.875	0.009585	0.013418	0.014362	-685.875	-34.08	-35.31	-61.76
$[\text{ClO}_3]^+$	-685.292	0.00839	0.012558	0.013502	-685.292	330.74	329.73	303.30
ClO_4^*	-760.893	0.01041	0.014433	0.015377	-760.893	76.78	74.64	37.12
$[\text{ClO}_4]^-$	-761.083	0.014186	0.018493	0.019438	-761.083	-40.15	-42.11	-79.79
ClOClO	-1070.76	0.00723	0.01139	0.012334	-1070.76	56.06	54.98	29.40
ClOOCl	-1070.79	0.006504	0.011388	0.012332	-1070.79	39.29	38.67	13.96
ClOClO_2^*	-1145.97	0.010652	0.015435	0.01638	-1145.97	50.56	48.83	12.93
ClOClO_3	-1221.14	0.016573	0.022351	0.023295	-1221.14	67.92	65.77	19.98
ClO_2ClO_2	-1221.13	0.0143	0.018233	0.019177	-0.01505	73.08	69.78	21.80
$\text{ClO}_2\text{O-ClO}_2$	-1296.31	0.018927	0.024792	0.025736	-0.01268	84.34	81.21	24.39
$\text{ClO}_2\text{O-ClO}_3$	-1371.51	0.027295	0.032621	0.033565	-0.00339	89.75	85.25	16.03
ClOOClO_3	-1296.33	0.01996	0.026008	0.026952	-0.01163	75.18	72.17	15.45

Table A-4.3 continued

Species	ϵ_0	ϵ_{ZPE}	E_{tot}	H_{corr}	G_{corr}	ΔH (OK)	ΔH (298K)	ΔG (298K)
ClO_3O-ClO_3	-1446.7	0.030374	0.038703	0.039647	-0.00473	93.30	89.64	13.61
$ClO_2O-O-ClO_2$	-1371.47	0.023365	0.031038	0.031982	-0.00977	107.20	104.17	37.96
$ClO_3O-O-ClO_3$	-1521.85	0.037317	0.044028	0.044972	0.004479	124.17	118.45	28.50
$[ClO_3-ClO_3]^{-2}$	-1371.47	0.028141	0.033538	0.034483	-0.00186	110.20	105.74	36.14
$OClO$	-610.583	0.005783	0.00894	0.009884	-0.01926	24.50	23.89	7.45
$[OClO]^-$	-610.645	0.003267	0.006849	0.007793	-0.02175	-15.96	-16.30	-32.48
$[OClO]^+$	-610.178	0.006108	0.009185	0.010129	-0.01822	278.90	278.24	261.31
$OClOO$	-685.733	0.004705	0.010421	0.011365	-0.02806	51.67	51.63	30.17
$[OClOO]^-$	-685.829	0.00475	0.008655	0.009599	-0.02582	-8.70	-9.87	-33.85
$OClClO_2^*$	-1145.81	0.005881	0.012201	0.013145	-0.02631	147.45	146.67	113.47
$[ClOH_2]^+$	-536.291	0.025091	0.028046	0.028991	0.002482	204.09	202.38	179.42
$HOCl$	-536.045	0.013162	0.016105	0.017049	-0.00978	-15.73	-16.44	-31.02
$[HOCl]^-$	-535.926	0.006934	0.010224	0.011168	-0.01455	54.74	54.25	38.96
$[HOCl]^+$	-535.637	0.012962	0.01592	0.016864	-0.01055	239.76	239.06	224.84
$HOClO$	-611.193	0.015261	0.018722	0.019666	-0.01037	14.44	13.01	-11.04
$[HOClO]^{-*}$	-611.066	0.011127	0.014348	0.015292	-0.01570	-55.76	-57.33	-80.78
$[HOClO]^+$	-610.834	0.015946	0.019318	0.020262	-0.01012	240.42	238.93	215.1
$HOClO_2$	-686.393	0.019286	0.023362	0.024307	-0.00765	13.26	11.17	-23.15
$HOClO_3$	-761.593	0.027229	0.031957	0.032902	-0.00083	14.84	12.13	-32.56

Table A-4.3 continued

Species	ϵ_0	ϵ_{ZPE}	E_{tot}	H_{corr}	G_{corr}	ΔH (OK)	ΔH (298K)	ΔG (298K)
[HOClO ₃] ⁺	-761.084	0.016512	0.022501	0.023446	-0.01494	327.65	325.73	283.97
HOCl* [*]	-611.212	0.017467	0.020866	0.02181	-0.00805	4.06	2.60	-21.57
[HOCl] ⁻	-611.228	0.011698	0.016318	0.017262	-0.0168	-9.90	-10.60	-32.12
[HOCl] ⁺	-610.812	0.016787	0.020038	0.020982	-0.00924	254.85	253.29	229.36
HOClO* [*]	-686.369	0.020306	0.024864	0.025809	-0.00746	29.56	27.78	-5.72
HOClO ₂ * [*]	-761.551	0.024822	0.029341	0.030285	-0.00357	40.10	37.26	-7.35
HOOC ₂ Cl	-686.37	0.021388	0.025538	0.026482	-0.00608	29.47	27.44	-6.50
HOOCOC ₂ Cl	-761.564	0.022424	0.028236	0.02918	-0.00723	30.21	28.18	-14.83
HClO	-535.961	0.009811	0.012762	0.013706	-0.01325	34.60	33.89	19.39
HClO ₂ * [*]	-611.137	0.015657	0.018874	0.019818	-0.00952	49.88	48.30	23.82
HClO ₃	-686.341	0.022859	0.026355	0.027299	-0.00356	48.51	46.07	11.05

Basis sets applied:

DFT B3LYP/(cc-PV5Z)

DFT/B3LYP 6-311++G/(3df,2p)

DFT/B3LYP/(aug-cc-pv5z)

The predominant basis set used was DFT/B3LYP (cc-PV5Z)

DFT/B3LYP 6-311G computations highlighted with *

A6 -SUPPLEMENTARY INFORMATION FOR CHAPTER 6

A6.1 ATOMIC PARTIAL CHARGES

Mulliken partial charges were obtained using *Gaussian-16* (DFT, B3LYP-cc-pV5Z), *MedeA-3-31* and *MOPAC-2016*.

Bader charges were obtained using *MedeA-3-31/VASP-5.4* software.

Table-A6: Net Atomic, Mulliken and Baden Partial Charges (ND Signifies: Not Determined)

Species	Atomic Partial Charges					
	Atom Label	Mulliken Charge <i>MOPAC-2016</i>	Mulliken Charge <i>Gaussian-16:</i> B3LYP cc-PV5Z	Net atomic Charge <i>MOPAC-2016</i>	Net Atomic Charge <i>Gaussian-16</i> B3LYP cc-PV5Z	Bader Charge Transfer <i>VASP-5.4</i>
ClO	Cl	-0.39	0.08	-0.16	0.02	0.61
	O	-0.38	-0.08	-0.16	0.02	-0.6
[ClO] ⁻	Cl	-0.06	-0.83	-0.21	-0.53	0.02
	O	-0.06	-0.17	-0.79	-0.47	-1.02
[ClO] ⁺	Cl	-0.18	0.79	0.87	0.72	1.38
	O	-0.18	0.21	0.13	0.28	-0.38
OClO	Cl	-0.46	1.14	0.83	0.87	1.66
	O1	-0.42	-0.57	-0.41	-0.43	-0.81
	O2	-0.43	-0.57	-0.41	-0.43	-0.84
[OClO] ⁺	Cl	-0.22	0.89	1.5	1.21	2.28
	O1	-0.21	0.05	-0.25	-0.1	-0.64
	O2	-0.22	0.05	-0.25	-0.1	-0.62
[OClO] ⁻	Cl	-0.14	0.18	0.7	0.4	0.72
	O1	-0.09	-0.59	-0.85	-0.7	-0.86
	O2	-0.09	-0.59	-0.85	-0.7	-0.84
ClOO	Cl	-1.19	0.27	0.02	-0.07	1.66
	O1	-0.22	-0.3	-0.06	0.02	0.81
	O2	-0.1	0.03	0.04	0.05	-0.84
[ClOO] ⁻	Cl	-0.08	-0.26	-0.3	-0.86	-0.62
	O1	-0.13	-0.13	-0.09	0.41	-0.21
	O2	-0.13	-0.62	-0.62	-0.55	-0.17
[ClOO] ⁺	Cl	-0.23	0.49	0.6	0.23	0.58
	O1	-0.17	0.2	0.13	0.78	0.09
	O2	-0.24	0.31	0.27	-0.01	0.34
ClOCl	Cl1	-0.08	0.11	0.06	-0.04	0.19
	Cl2	-0.13	0.11	0.06	-0.04	0.28
	O	-0.16	-0.23	-0.12	0.07	-0.48
[ClOCl] ⁻	Cl1	-0.18	-0.47	-0.39	-0.57	-0.6
	Cl2	-0.39	-0.47	-0.39	-0.57	0.34
	O	-0.25	-0.06	-0.23	0.14	-0.74

Table-A6 continued

Species	Atomic Partial Charges					
	Atom label	Mulliken charge <i>MOPAC-2016</i>	Mulliken charge <i>Gaussian-16:</i> B3LYP cc-PV5Z	Net atomic charge <i>MOPAC-2016</i>	Net atomic charge <i>Gaussian-16</i> B3LYP cc-PV5Z	Bader charge transfer <i>VASP-5.4</i>
[ClOCl] ⁺	Cl1	-3.31	0.51	0.54	0.24	0.64
	Cl2	-2.00	0.51	0.54	0.24	0.71
	O	-0.14	-0.02	-0.09	0.52	-0.34
ClClO	Cl1	-0.13	-0.21	-0.09	-0.24	-0.09
	Cl2	-0.18	0.42	0.57	0.50	0.83
	O	-0.12	-0.22	-0.48	-0.25	-0.74
[ClClO] ⁻	Cl1	-0.33	-0.51	-0.53	-0.92	-0.6
	Cl2	-0.37	0.10	0.15	0.31	0.40
	O	-0.18	-0.59	-0.62	-0.40	-0.79
[ClClO] ⁺	Cl1	-2.35	0.28	0.18	0.09	0.33
	Cl2	-0.43	0.72	0.75	0.86	1.30
	O	-0.15	-0.01	0.06	0.06	-0.62
ClCl(O)O	Cl1	-0.09	-0.06	-0.42	-0.34	-0.20
	Cl2	-0.20	0.66	1.69	1.33	2.03
	O1	-0.19	-0.30	-0.63	-0.50	-0.90
	O2	-0.21	-0.30	-0.63	-0.50	-0.92
ClOOCl	Cl1	-0.12	0.05	0.08	-0.15	0.25
	Cl2	-0.09	0.06	0.08	-0.15	0.26
	O1	-0.14	-0.03	-0.08	0.15	-0.26
	O2	-0.13	-0.08	-0.08	0.14	-0.26
ClOClO	Cl1	-0.15	0.09	0.13	-0.02	0.17
	Cl2	-0.14	0.51	0.86	0.63	1.06
	O1	-0.17	-0.3	-0.39	-0.20	-0.62
	O2	-0.15	-0.3	-0.60	-0.41	-0.61
ClO ₃	Cl	-0.17	0.52	2.77	1.88	1.87
	O1	-0.60	-0.26	-0.92	-0.63	-0.55
	O2	-0.22	0.00	-0.92	-0.63	-0.81
	O3	-0.60	-0.26	-0.92	-0.63	-0.51
[ClO ₃] ⁻	Cl	-0.22	0.13	1.57	0.44	1.15
	O1	-0.15	-0.37	-0.86	-0.48	-0.66
	O2	-0.14	-0.38	-0.86	-0.48	-0.9
	O3	-0.15	-0.38	-0.86	0.48	-0.59
[ClO ₃] ⁺	Cl	-0.25	0.78	1.68	0.15	3.01
	O1	-0.18	0.07	-0.23	0.21	-0.66
	O2	-0.13	0.08	-0.23	0.44	-0.68
	O3	-0.18	0.08	-0.23	0.21	-0.65

Table-A6 continued

Species	Atomic Partial Charges					
	Atom Label	Mulliken Charge <i>MOPAC-2016</i>	Mulliken Charge <i>Gaussian-16:</i> B3LYP cc-PV5Z	Net Atomic Charge <i>MOPAC-2016</i>	Net Atomic Charge <i>Gaussian-16</i> B3LYP cc-PV5Z	Bader Charge Transfer <i>VASP-5.4</i>
ClOOO	Cl	-0.51	0.11	0.01	0.09	0.54
	O1	-0.45	-0.09	-0.02	-0.43	-0.05
	O2	-0.16	0.07	0.00	0.33	0.00
	O3	-0.11	-0.10	0.02	0.01	-0.5
[ClOOO] ⁻	Cl	0.06	-0.17	-0.20	-0.37	-0.53
	O1	-0.12	-0.41	-0.09	-0.39	-0.21
	O2	-0.16	-0.12	-0.10	0.44	0.09
	O3	-0.13	-0.31	-0.61	-0.67	-0.34
ClO(O)O	Cl	-0.62	0.20	-0.07	0.34	0.12
	O1	-0.23	0.21	0.03	-0.17	0.09
	O2	-0.12	-0.20	0.03	-0.01	-0.10
	O3	-0.09	-0.20	0.02	-0.17	-0.11
[ClO(O)O] ⁻	Cl	-0.07	-0.83	-0.45	0.16	-0.35
	O1	-0.17	0.33	0.64	-0.55	0.00
	O2	-0.18	-0.25	-0.59	-0.06	-0.31
	O3	-0.20	-0.25	-0.6	0.55	-0.33
OClOO	Cl	-0.17	0.08	0.33	-0.03	0.84
	O1	-0.10	-0.08	-0.15	0.02	-0.65
	O2	-0.19	0.00	-0.16	0.02	-0.13
	O3	-0.10	0.00	-0.02	-0.02	-0.06
[OClOO] ⁻	Cl	-0.14	0.33	0.51	-0.73	0.29
	O1	-0.08	-0.56	-0.79	0.05	-0.78
	O2	-0.13	-0.13	-0.11	0.19	-0.23
	O3	-0.10	-0.64	-0.61	-0.51	-0.25
ClO ₄	Cl	-0.16	1.08	2.98	1.18	3.45
	O1	-4.03	-0.27	-1.01	-0.14	-0.84
	O2	-0.27	-0.27	-0.66	-0.46	-0.84
	O3	-0.29	-0.27	-0.68	-0.14	-0.92
	O4	-0.28	-0.27	-0.65	-0.46	-0.84

Table-A6 continued

Species	Atomic Partial Charges					
	Atom Label	Mulliken Charge <i>MOPAC-2016</i>	Mulliken Charge <i>Gaussian-16:</i> B3LYP cc-PV5Z	Net Atomic Charge <i>MOPAC-2016</i>	Net Atomic Charge <i>Gaussian-16</i> B3LYP cc-PV5Z	Bader Charge Transfer <i>VASP-5.4</i>
[ClO ₄] ⁻	Cl	-0.19	0.96	2.77	1.63	3.29
	O1	-0.16	-0.46	-0.94	-0.8	-1.08
	O2	-0.11	-0.46	-0.94	-0.81	-1.07
	O3	-0.09	-0.46	-0.94	-0.21	-1.08
	O4	-0.12	-0.59	-0.94	-0.81	-1.06
ClOClO ₂	Cl1	-0.17	0.68	0.95	1.11	1.68
	Cl2	-0.13	0.21	0.17	-0.22	0.66
	O1	-0.15	-0.3	-0.45	-0.64	-0.85
	O2	-0.15	-0.29	-0.42	-0.34	-0.8
	O3	-0.12	-0.3	-0.25	0.09	-0.69
OClClO ₂	Cl1	-0.17	0.53	0.88	1.17	1.59
	Cl2	-0.14	0.07	0.09	0.02	0.63
	O1	-0.21	-0.27	-0.3	-0.52	-0.78
	O2	-0.22	-0.28	-0.38	-0.54	-0.78
	O3	-0.2	0.05	-0.29	-0.12	-0.67
ClO ₂ ClO ₂	Cl1	-0.21	0.33	1.01	1.05	1.68
	Cl2	-0.21	0.33	1.01	-0.53	1.62
	O1	-0.19	-0.16	-0.51	-0.53	-0.84
	O2	-0.21	-0.16	-0.51	-0.53	-0.82
	O3	-0.19	-0.16	-0.51	1.05	-0.8
	O4	-0.21	-0.16	-0.51	-0.53	-0.83
ClOClO ₃	Cl1	-0.2	0.52	2.33	2.09	3.19
	Cl2	-0.13	-0.26	0.31	0.05	0.33
	O1	-0.17	-0.26	-0.7	-0.57	-0.98
	O2	-0.1	-0.26	-0.69	-0.57	-1.01
	O3	-0.17	-0.26	-0.65	-0.61	-0.98
	O4	-0.15	0.52	-0.6	-0.39	-0.55

Table-A6 continued

Species	Atomic Partial Charges					
	Atom Label	Mulliken Charge <i>MOPAC-2016</i>	Mulliken Charge <i>Gaussian-16:</i> B3LYP cc-PV5Z	Net Atomic Charge <i>MOPAC-2016</i>	Net Atomic Charge <i>Gaussian-16</i> B3LYP cc-PV5Z	Bader Charge Transfer <i>VASP-5.4</i>
ClOOClO ₃	Cl1	0.2	0.93	2.35	-0.52	2.89
	Cl2	-0.14	-0.0056	0.17	0.1	0.19
	O1	-0.15	-0.11	-0.51	2.29	-0.16
	O2	-0.13	-0.3	-0.68	0.01	-0.92
	O3	-0.13	-0.32	-0.66	-0.62	-0.91
	O4	-0.1	-0.25	-0.71	-0.64	-0.95
	O5	-0.16	0.06	0.04	-0.61	-0.13
ClO ₂ -O-ClO ₂	Cl1	0.2	0.93	2.35	-0.52	2.89
	Cl2	-0.14	-0.01	0.17	0.1	0.19
	O1	-0.15	-0.11	-0.51	2.29	-0.16
	O2	-0.13	-0.3	-0.68	0.01	-0.92
	O3	-0.13	-0.32	-0.66	-0.62	-0.91
	O4	-0.1	-0.25	-0.71	-0.64	-0.95
	O5	-0.16	0.06	0.04	-0.61	-0.13
ClO ₂ -O-ClO ₃	Cl1	-0.14	0.97	1.04	2.24	3.2
	Cl2	-0.12	0.78	0.72	1.71	2.21
	O1	-0.18	-0.28	-0.41	-1.15	-0.65
	O2	-0.12	-0.25	-0.21	-0.55	-0.88
	O3	-0.14	-0.26	-0.23	-0.49	-0.86
	O4	-0.13	-0.35	-0.29	-0.6	-1.02
	O5	-0.11	-0.29	-0.3	-0.6	-1
O6	-0.09	-0.31	-0.33	0.55	-1	
ClO ₂ -O-O-ClO ₂	Cl1	-0.22	0.278	1.89	1.02	0.8
	Cl2	-0.22	0.278	1.88	1.02	0.8
	O1	-0.15	-0.195	-0.51	-0.42	-0.05
	O2	-0.15	-0.212	-0.51	-0.44	-0.06
	O3	-0.2	0.13	-0.7	-0.16	-0.07
	O4	-0.18	0.13	-0.67	-0.16	-0.68
	O5	-0.18	-0.212	-0.67	-0.42	-0.68
O6	-0.2	-0.195	-0.71	-0.44	-0.05	

Table-A6 continued

Species	Atomic Partial Charges					
	Atom Label	Mulliken Charge <i>MOPAC-2016</i>	Mulliken Charge <i>Gaussian-16:</i> B3LYP cc-PV5Z	Net Atomic Charge <i>MOPAC-2016</i>	Net Atomic Charge <i>Gaussian-16</i> B3LYP cc-PV5Z	Bader Charge Transfer <i>VASP-5.4</i>
ClO ₃ -O-ClO ₃	Cl1	-0.2	0.99	2.32	2.19	3.21
	Cl2	-0.2	0.99	2.32	2.19	3.22
	O1	-0.19	-0.28	-0.61	-0.56	-0.96
	O2	-0.17	-0.28	-0.58	-0.61	-0.97
	O3	-0.19	-0.3	-0.62	-0.56	-0.96
	O4	-0.11	-0.26	-1.03	-0.93	-0.58
	O5	-0.18	-0.3	-0.61	-0.56	-0.98
	O6	-0.12	-0.28	-0.58	-0.61	-1
	O7	-0.17	-0.28	-0.61	-0.56	-0.97
ClO ₃ -O-O-ClO ₃	Cl1	-0.20	1.022	2.23	2.47	1.87
	Cl2	-0.20	1.022	2.22	2.47	1.85
	O1	-0.18	-0.139	-0.34	-0.47	-0.09
	O2	-0.17	-0.139	-0.32	-0.47	-0.04
	O5	-0.15	-0.297	-0.65	-0.64	-0.89
	O6	-0.17	-0.289	-0.57	-0.69	-0.84
	O7	-0.19	-0.296	-0.65	-0.67	-0.11
	O8	-0.18	-0.289	-0.60	-0.69	-0.85
	O9	-0.09	-0.296	-0.67	-0.67	-0.03
	O10	-0.18	-0.297	-0.65	-0.64	-0.86
[ClOH ₂] ⁺	Cl	-0.15	0.3	0.41	0.66	0.52
	O	-0.09	-0.16	-0.22	-0.5	-1.1
	H1	0	0.43	0.41	0.42	0.81
	H2	0	0.43	0.41	0.42	0.79
HOCl	Cl	-0.12	0.04	0.1	0.03	0.24
	O	-0.14	-0.37	-0.41	-0.28	-0.9
	H	ND	0.33	0.31	0.29	0.66
[HOCl] ⁻	Cl	-0.35	-0.27	-0.64	-0.95	-0.53
	O	-0.22	0.211	-0.42	-0.5	-1.06
	H	ND	-0.941	0.056	0.444	0.59

Table-A6 continued

Species	Atomic Partial Charges					
	Atom Label	Mulliken Charge <i>MOPAC-2016</i>	Mulliken Charge <i>Gaussian-16:</i> B3LYP cc-PV5Z	Net Atomic Charge <i>MOPAC-2016</i>	Net Atomic Charge <i>Gaussian-16</i> B3LYP cc-PV5Z	Bader Charge Transfer <i>VASP-5.4</i>
[HOCl] ⁺	Cl	-0.62	0.63	0.62	0.49	0.93
	O	-0.17	-0.02	0.02	0.09	-0.68
	H	ND	0.4	0.35	0.42	0.76
HOClO	Cl	-0.14	0.34	0.83	0.64	1.1
	O1	-0.12	-0.26	-0.6	-0.38	-0.89
	O2	-0.12	-0.33	-0.56	-0.52	-1
	H	ND	0.24	0.33	0.25	0.69
[HOClO] ⁻	Cl	-0.31	-0.34	-0.47	-0.18	-1.11
	O1	-0.17	-0.25	-0.36	-0.11	-0.86
	O2	-0.2	-0.38	-0.3	-0.68	0.41
	H	ND	-0.03	0.13	0.03	0.56
[HOClO] ⁺	Cl	-0.39	0.84	0.65	0.24	1.77
	O1	-0.17	-0.02	0.17	0.34	-0.75
	O2	-0.48	-0.17	-0.14	0.01	-0.75
	H	ND	0.36	0.32	0.41	0.75
HOOC _l	Cl	-0.11	0.04	-0.06	-0.18	0.3
	O1	-0.14	-0.24	-0.18	-0.29	-0.62
	O2	-0.13	-0.1	0.02	0.23	-0.37
	H	ND	0.31	0.22	0.24	0.68
[HOOC _l] ⁺	Cl	-0.27	0.31	0.59	0.31	0.73
	O1	-0.72	0.11	-0.04	-0.33	-0.38
	O2	-0.25	0.24	0.17	0.48	-0.09
	H	ND	0.34	0.28	0.55	0.75
HOOC _l	Cl	-0.13	0.1	0.01	0.38	0.12
	O1	-0.16	-0.27	-0.04	-0.07	-0.16
	O2	-0.15	-0.07	-0.01	0.27	0
	O3	-0.15	-0.09	-0.17	-0.01	-0.61
	H	ND	0.34	0.21	0.43	0.65

Table-A6 continued

Species	Atomic Partial Charges					
	Atom Label	Mulliken Charge <i>MOPAC-2016</i>	Mulliken Charge <i>Gaussian-16:</i> B3LYP cc-PV5Z	Net Atomic Charge <i>MOPAC-2016</i>	Net Atomic Charge <i>Gaussian-16</i> B3LYP cc-PV5Z	Bader Charge Transfer <i>VASP-5.4</i>
HOOC ℓ O	C ℓ	-0.11	0.52	0.48	0.68	0.82
	O1	-0.11	-0.32	-0.20	-0.32	-0.48
	O2	-0.13	-0.22	-0.13	-0.28	-0.22
	O3	-0.14	-0.33	-0.39	-0.39	-0.78
	H	ND	0.34	0.25	0.31	0.66
HO ℓ (O)O	C ℓ	-0.17	0.67	0.94	1.17	1.39
	O1	-0.13	-0.37	-0.3	0.24	-0.71
	O2	-0.13	-0.28	-0.44	-0.49	-0.86
	O3	-0.14	-0.33	-0.46	-0.40	-0.50
	H	ND	0.31	0.26	-0.53	0.68
HO ℓ O ₃	C ℓ	-0.22	0.5	1.88	2.27	3.6
	O1	-0.12	-0.26	-0.45	-0.71	-1.12
	O2	-0.14	-0.17	-0.58	-0.63	-1.06
	O3	-0.12	-0.19	-0.59	-0.66	-1.14
	O4	-0.15	-0.17	0.55	-0.6	-0.96
	H	ND	0.3	0.29	0.32	0.69
[HO ℓ O ₃] ⁺	C ℓ	-0.14	1.01	3.11	1.88	3.62
	O1	-0.32	-0.18	-0.86	-0.30	-0.92
	O2	-0.28	-0.08	-0.54	-0.31	-0.84
	O3	-0.17	-0.09	-0.54	-0.34	-0.91
	O4	-0.32	-0.08	-0.54	-0.31	-0.68
	H	ND	0.41	0.35	0.37	0.74
HOOC ℓ (O)O	C ℓ	-0.2	0.72	1.83	1.18	1.96
	O1	-0.13	-0.28	-0.26	-0.32	-0.53
	O2	-0.14	-0.11	-0.51	-0.10	-0.29
	O3	-0.2	-0.34	-0.68	-0.46	-0.91
	O4	-0.18	-0.32	-0.71	-0.54	-0.91
	H	ND	0.32	0.33	0.24	0.67

Table-A6 continued

Species	Atomic Partial Charges					
	Atom Label	Mulliken Charge <i>MOPAC-2016</i>	Mulliken Charge <i>Gaussian-16:</i> B3LYP cc-PV5Z	Net Atomic Charge <i>MOPAC-2016</i>	Net Atomic Charge <i>Gaussian-16</i> B3LYP cc-PV5Z	Bader Charge Transfer <i>VASP-5.4</i>
HOOCOCℓ	Cℓ	-0.12	0.24	0.02	0.3	0.33
	O1	-0.14	-0.54	-0.18	-0.06	-0.63
	O2	-0.14	-0.022	0.02	0.01	0.05
	O3	-0.16	-0.021	-0.03	-0.24	0
	O4	-0.17	-0.25	-0.04	-0.42	-0.38
	H	ND	0.59	0.22	0.42	0.64
HCℓO	Cℓ	0.455	0.41	0.32	0.24	0.23
	O	-0.5	-0.47	-0.46	-0.35	-0.89
	H	0.045	0.06	0.14	0.11	0.67
HCℓO ₂	Cℓ	-0.23	0.32	1.1	0.617	-0.77
	O1	-0.16	-0.46	-0.43	-0.331	0.98
	O2	-0.23	-0.46	-0.43	-0.331	-0.88
	H	ND	0.14	-0.24	0.045	0.67
HCℓO ₃	Cℓ	0.948	0.94	1.87	1.93	2.18
	O1	-0.11	-0.33	-0.58	-0.62	-0.89
	O2	-0.1	-0.33	-0.58	-0.62	-1.01
	O3	-0.09	-0.33	-0.58	-0.62	-0.96
	H	ND	0.06	-0.14	-0.06	0.68

A7 - SUPPLEMENTARY INFORMATION FOR CHAPTER 7

Major product species extracted from the reaction with H₂O

Table-A7: Major species products -- Exposed to Water

H ₂ O + Reactant (1:1)	Water	Reactant	O ₂	Cl[-]	ClOH ₂ [+]	H ₂ O
	ClO	1	1	0.32	0.32	0.32
ClOO	1	1	0.48	0.24	0.24	0.031222
ClOCl	1	1	0.20	0.40	0.40	0.000013
ClClO	1	1	0.20	0.40	0.40	0.000013
ClClO ₂	1	1	0.33	0.33	0.33	0.000012
ClOOO	1	1	0.58	0.19	0.19	0.031225
Cl(O)O ₂	1	1	0.58	0.19	0.19	0.030783
ClO ₃	1	1	0.58	0.19	0.19	0.031225
ClO ₄	1	1	0.65	0.16	0.16	0.031226
ClOClO	1	1	0.33	0.33	0.33	0.000012
ClOOCl	1	1	0.33	0.33	0.33	0.000012
ClOClO ₂	1	1	0.43	0.29	0.29	0.000012
ClOClO ₃	1	1	0.50	0.25	0.25	0.000012
ClO ₂ ClO ₂	1	1	0.50	0.25	0.25	0.000012
ClO ₂ -O-ClO ₂	1	1	0.56	0.22	0.22	0.000012
ClOOClO ₃	1	1	0.56	0.22	0.22	0.000012
ClO ₂ -O-ClO ₃	1	1	0.60	0.20	0.20	0.000012
ClO ₃ -O-ClO ₃	1	1	0.64	0.18	0.18	0.000012
ClO ₂ -O-O-ClO ₂	1	1	0.60	0.20	0.20	0.000012
ClO ₃ -O-O-ClO ₃	1	1	0.65	0.16	0.16	0.030785
OClO	1	1	0.48	0.24	0.24	0.030781
OClOO	1	1	0.58	0.19	0.19	0.030783
OClClO ₂	1	1	0.43	0.29	0.29	0.000012
HOCl	1	1	0.39	0.39	0.19	0.030761
HOClO	1	1	0.42	0.28	0.28	0.030777
HOClO ₂	1	1	0.54	0.22	0.22	0.030782
HOClO ₃	1	1	0.62	0.18	0.18	0.030784
HOOCl	1	1	0.42	0.28	0.28	0.030777
HOOClO	1	1	0.54	0.22	0.22	0.030782
HOOClO ₂	1	1	0.62	0.18	0.18	0.030784
HOOClO ₃	1	1	0.62	0.18	0.18	0.030784
HOOCOOCl	1	1	0.54	0.22	0.22	0.030782
HOOCOOClO	1	1	0.62	0.18	0.18	0.030784
HOOCOOClO ₂	1	1	0.62	0.18	0.18	0.030784
HOOCOOClO ₃	1	1	0.62	0.18	0.18	0.030784
HClO	1	1	0.19	0.39	0.39	0.030761
HClO ₂	1	1	0.24	0.28	0.28	0.030777
HClO ₃	1	1	0.54	0.22	0.22	0.030782

Appendix References

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- [2] J. W. Ochterski, "Thermochemistry in Gaussian," Gaussian Inc, 2 June 2000. [Online]. Available: www.gaussian.com. [Accessed 8 June 2019].