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# Comparative Study of Molecular Orbitals of Ruthenium (II) Chloride and Ruthenium (II) Iodide Based on Molecular Mechanics

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## ABSTRACT

*We have studied the molecular orbitals of ruthenium halides, in order to study the extent of contribution of 4d, 5s and 5p orbitals in the formation of molecular orbitals. The 3D modeling and geometry optimization of the ruthenium halides have been done by CAChe software using molecular mechanics with EHT option. Eigenvector analysis shows that  $4dx^2-y^2$  and  $4dxy$  orbitals of ruthenium play a major role in bonding between ruthenium and halogens, 5s orbital is next and 5p orbitals have a negligible role. There is a difference in energy levels of s and p orbitals of chloride and iodide are 0.1691 and 0.7472 respectively. The overlap population analysis shows that the nonbonding orbitals are present in 6<sup>th</sup> and 7<sup>th</sup> molecular orbitals in both. No molecular orbital is formed by only two atomic orbitals. All molecular orbitals have contribution from many atomic orbitals; the difference is only in extent of involvement.*

**Keywords:** Ruthenium (II) chloride, ruthenium (II) iodide, sd hybridization, population analysis, overlap population analysis, eigenvector, eigenvalues.

## INTRODUCTION

In the recent years Landis presented the result of DFT calculation of transition metal hydride [1-3]. He also gave the results of an NBO analysis of the transition metal-hydrogen bonds, which show dominantly sdn hybridized bond orbitals and negligible np participation [1]. However, there is a serious technical flaw in the analysis. The NBO method requires preselection of those orbitals, which are considered as valence orbitals, and may become occupied in the population analysis. In the last few decades, there has been a phenomenal advancement in theoretical inorganic chemistry. Commercial programs incorporating the latest methods have become widely available, and are capable of providing more information about molecular orbitals with a simple input of chemical formula. The focus of attention has

been on computational transition-metal chemistry [4, 5]. This is largely due to the successful employment of gradient corrected density functional theory in calculating molecule; particularly of the heavier atoms [6-9] and in the use of small-core relativistic effective core potential [10-12] which set the stage for calculation of geometries, bond energies, and chemical reaction and other important properties of transition metal compounds with impressive accuracy [9, 13, 14]. Application of molecular mechanics to organometallic and transition metal compounds is growing [15]. Molecular orbital parameters such as eigenvalues, eigenvectors and overlap matrix are well calculated with this method. In this paper we present the comparative study of ruthenium (II) chloride and ruthenium (II) iodide based on eigenvalues, eigenvector, overlap matrix, population analysis and overlap population analysis, in order to study the extent of contribution of 4d, 5s and 5p orbitals in the formation of molecular orbitals. Such a quantitative study will provide correct information about the involvement of 5p orbital of ruthenium in bonding.

### MATERIALS AND METHODS

The study materials of this paper are ruthenium (II) chloride and ruthenium (II) iodide. The 3D modeling and geometry optimization of the halide have been done by CAChe software using molecular mechanics with EHT option. Eigenvalues, eigenvectors and overlap matrix values have been obtained with the same software, using the same option. With the help of these values, eigenvector analysis, magnitude of contribution of atomic orbital in MO formation and population analysis have been made and discussed. The method adopted for various calculations is based on the following principles.

The molecular orbitals are formed by the linear combination of basis functions. Most molecular quantum-mechanical methods (such as- SCF, CI etc.) begin the calculation with the choice of a basis functions  $\chi_r$ , which are used to express the MOs  $\phi_i$  as  $\phi_i = \sum_r c_{ri} \chi_r$  ( $c$  = coefficient of  $\chi$ ,  $r$  = number of atomic orbital,  $i$  = molecular orbital number). The use of an adequate basis set is an essential requirement for the calculation. The basis functions are usually taken as AOs. Each AO can be represented as a linear combination of one or more Slater-type orbitals (STOs) [15, 25, 26]. Each molecular orbital  $\phi_i$  is expressed as  $\phi_i = \sum_r c_{ri} \chi_r$ , where, the  $\chi_r$ 's are the STO basis functions. Here we use the STO-6G basis set (which is contracted Gaussian) [27-30] for the SCF calculation.

The coefficients in linear combination for each molecular orbital being found by solution of the Roothaan equation [31]. The most efficient way to solve the Roothaan equation is to use matrix-algebra methods. In matrix-algebra methods, the matrix elements are computed [32], and the secular equation is solved to give the set of orbital energies (i.e. eigenvalues). These orbital energies [33-35] are used to solve Roothaan equations for the set of coefficients (i.e. eigenvectors) giving a set of molecular orbitals. The calculations are done using a computer. By the above calculation, the values of orbital energies (eigenvalues) and eigenvectors (coefficients) have been calculated.

A widely used method to analyze SCF wave function is population analysis, introduced by Mulliken [36, 37]. He proposed a method that apportions the electrons of an  $n$ -electron

molecule into net populations  $n_r$  in the basis functions  $\chi_r$  and overlap populations  $n_{r-s}$  for all possible pairs of basis functions.

For the set of basis functions  $\chi_1, \chi_2, \dots, \chi_b$ , each molecular orbital  $\phi_i$  has the form  $\phi_i = \sum_s c_{si} \chi_s = c_{1i} \chi_1 + c_{2i} \chi_2 + \dots + c_{bi} \chi_b$ . For simplicity, we shall assume that the  $c_{si}$ 's and  $\chi_s$ 's are real. The probability density associated with one electron in  $\phi_i$  is (s and b are the number of the atomic orbital other than r)

$$|\phi_i|^2 = c_{1i}^2 \chi_1^2 + c_{2i}^2 \chi_2^2 + \dots + 2c_{1i}c_{2i} \chi_1 \chi_2 + 2c_{1i}c_{3i} \chi_1 \chi_3 + 2c_{2i}c_{3i} \chi_2 \chi_3 + \dots$$

Integrating this equation over three-dimensional space and using the fact that  $\phi_i$  and the  $\chi_s$ 's are normalized, we get

$$1 = c_{1i}^2 + c_{2i}^2 + \dots + 2c_{1i} c_{2i} S_{12} + 2c_{1i} c_{3i} S_{13} + 2c_{2i} c_{3i} S_{23} + \dots \quad (3)$$

Where the S's are overlap integrals:  $S_{12} = \int \chi_1 \chi_2 dv_1 dv_2$ , etc. Mulliken proposed that the terms in (3) be apportioned as follows. One electron in the molecular orbital  $\phi_i$  contributes  $c_{1i}^2$  to the net population in  $\chi_1$ ,  $c_{2i}^2$  to the net population in  $\chi_2$ , etc., and contributes  $2c_{1i}c_{2i}S_{12}$  to the overlap population between  $\chi_1$  and  $\chi_2$ ,  $2c_{1i} c_{3i} S_{13}$  to the overlap population between  $\chi_1$  and  $\chi_3$ , etc.

Let there be  $n_i$  electrons in the molecular orbital  $\phi_i$  ( $n_i = 0, 1, 2$ ) and let  $n_{r,i}$  and  $n_{r-s,i}$  symbolize the contributions of electrons in the molecular orbital  $\phi_i$  to the net population in  $\chi_r$  and to the overlap population between  $\chi_r$  and  $\chi_s$ , respectively. We have

$$n_{r,i} = n_i c_{ri}^2,$$

$$n_{r-s,i} = n_i (2c_{ri} c_{si} S_{rs})$$

Based on the above principle, the contribution of electrons in each occupied molecular orbital has been calculated with the help of eigenvector values and also calculated overlap population analysis for distinguishing the bonding, nonbonding and antibonding nature of molecular orbital.

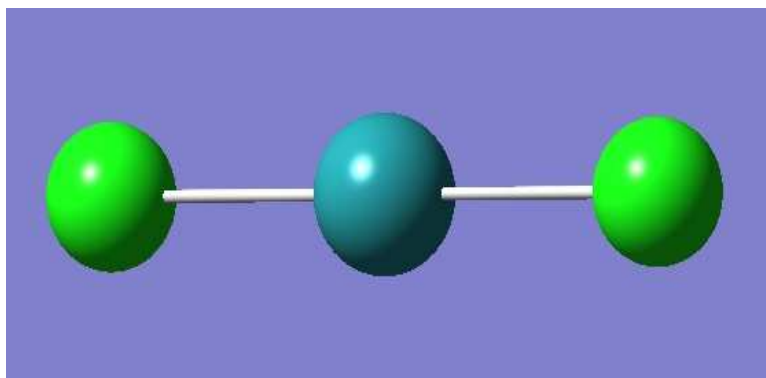
## RESULTS AND DISCUSSION

Ruthenium (II) chloride and ruthenium (II) iodide is triatomic molecule, having the following (Fig. 1.1, 1.2) optimized geometry [16, 17] as obtained from molecular mechanics [18-21] method.

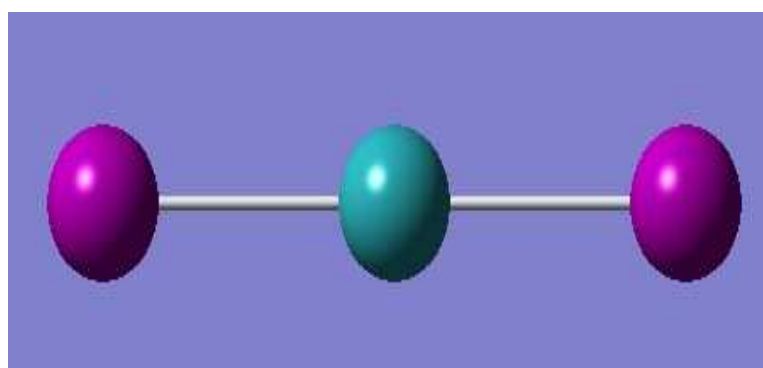
The MOs of these halides (chloride and iodide) are formed by linear combination of 9 ruthenium orbitals and 4 orbitals from each halogen as detailed below-

$$\begin{aligned} \text{Ru-1} &= 5s, 5p_x, 5p_y, 5p_z, 4d_{x^2-y^2}, 4d_z^2, 4d_{xy}, 4d_{xz}, 3d_{yz} &= 9 \\ \text{X-2} &= ns, n p_x, n p_y, n p_z &= 4 \\ \text{X-3} &= ns, n p_x, n p_y, n p_z &= 4 \\ &&\text{Total} = 17 \end{aligned}$$

where X = Cl or I; n = 3 for Cl and n = 5 for I.



**Fig.1.1: Optimized geometry of Ruthenium (II) Chloride.**



**Fig.1.2: Optimized geometry of Ruthenium (II) Iodide.**

In order to examine the contribution of various atomic orbitals in the formation of molecular orbitals the LCAO has been studied. The 17 AOs give LCAO approximations to the 17 MOs of ruthenium (II) halides. The various AOs are represented by  $\chi$  and MOs by  $\phi$ .  $\chi_1$  to  $\chi_9$  are 5s, 5px, 5py, 5pz,  $4dx^2-y^2$ ,  $4dz^2$ , 4dxy, 4dxz, 4dyz, respectively and  $\chi_{10}$  to  $\chi_{13}$  and  $\chi_{14}$  to  $\chi_{17}$  are ns, np<sub>x</sub>, np<sub>y</sub>, np<sub>z</sub> for X-2 and X-3, respectively are atomic orbitals of halides.

The eigenvalues of 17 MOs ( $\phi_1$  to  $\phi_{17}$ ) of ruthenium (II) chloride are -0.9810, -0.9696, -0.5934, -0.5824, -0.5824, -0.5476, -0.5476, -0.5326, -0.5271, -0.5271, -0.4986, -0.4726, -0.4726, -0.2118, -0.2118, 0.1413 and 0.6063, respectively and of ruthenium (II) iodide are -0.6884, -0.6679, -0.5565, -0.5565, -0.5560, -0.5476, -0.5476, -0.4843, -0.4811, -0.4700, -0.4700, -0.4521, -0.4521, -0.2336, -0.2336, -0.0681 and 0.2251, respectively. The coefficients of  $\chi$  are the eigenvector and overlap matrix which has been taken from Table-1.1, Table- 1.2 and Table- 2.1, Table- 2.2 respectively.

**Table 1.1: Eigenvector values of molecular orbitals of Ruthenium (II) chloride.**

| Atom | AOs<br>( $\chi$ )                | Eigenvector values or coefficients of Atomic Orbitals |         |         |         |         |         |         |         |         |         |         |         |         |         |         |         |         |
|------|----------------------------------|---|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|
|      |                                  | MO-1  | MO-2    | MO-3    | MO-4    | MO-5    | MO-6    | MO-7    | MO-8    | MO-9    | MO-10   | MO-11   | MO-12   | MO-13   | MO-14   | MO-15   | MO-16   | MO-17   |
| Ru-1 | 5s                               | -0.1029   | 0.0000  | 0.0966  | -0.0000 | 0.0000  | 0.0000  | 0.0000  | -0.0000 | -0.0000 | 0.0000  | -0.4603 | -0.0000 | -0.0000 | -0.0000 | -0.0000 | -1.1218 | 0.0000  |
|      | 5px                              | -0.0000   | -0.0688 | -0.0000 | 0.0000  | -0.0000 | 0.0000  | 0.0000  | -0.1681 | -0.0014 | 0.0000  | 0.0000  | -0.0000 | -0.0000 | 0.0111  | 0.0000  | 0.0000  | 1.4554  |
|      | 5py                              | 0.0000  | -0.0007 | 0.0000  | 0.0000  | -0.0000 | -0.0000 | -0.0000 | -0.0018 | 0.1271  | -0.0006 | -0.0000 | 0.0000  | 0.0000  | -1.0225 | -0.0003 | 0.0000  | 0.0158  |
|      | 5pz                              | -0.0000   | -0.0000 | -0.0000 | 0.0000  | 0.0000  | 0.0000  | -0.0000 | 0.0000  | 0.0006  | 0.1271  | -0.0000 | -0.0000 | 0.0000  | 0.0003  | -1.0226 | -0.0000 | -0.0000 |
|      | 4dx <sup>2</sup> -y <sup>2</sup> | -0.1049   | -0.0000 | 0.4628  | -0.0027 | 0.0161  | -0.0000 | -0.4999 | 0.0000  | 0.0000  | -0.0000 | 0.6669  | 0.0003  | 0.0146  | -0.0000 | -0.0000 | -0.4071 | 0.0000  |
|      | 4dz <sup>2</sup>                 | 0.0606  | -0.0000 | -0.2673 | 0.0000  | -0.0000 | -0.0000 | -0.8660 | -0.0000 | -0.0000 | 0.0000  | -0.3851 | -0.0000 | -0.0000 | 0.0000  | 0.0000  | 0.2351  | -0.0000 |
|      | 4dxy                             | -0.0023   | 0.0000  | 0.0100  | 0.1234  | -0.7415 | -0.0000 | -0.0108 | -0.0000 | -0.0000 | 0.0000  | 0.0145  | -0.0131 | -0.6721 | -0.0000 | -0.0000 | -0.0088 | -0.0000 |
|      | 4dxz                             | 0.0000  | -0.0000 | -0.0000 | 0.7416  | 0.1234  | 0.0108  | -0.0000 | -0.0000 | -0.0000 | -0.0000 | 0.0000  | 0.6722  | -0.0131 | 0.0000  | -0.0000 | -0.0000 | 0.0000  |
|      | 4dyz                             | 0.0000  | 0.0000  | 0.0000  | 0.0080  | 0.0013  | -0.9999 | 0.0000  | -0.0000 | 0.0000  | 0.0000  | -0.0000 | 0.0073  | -0.0001 | -0.0000 | 0.0000  | -0.0000 | -0.0000 |
| Cl-2 | 3s                               | -0.6626   | -0.6829 | -0.1572 | 0.0000  | -0.0000 | -0.0000 | -0.0000 | 0.0675  | 0.0000  | -0.0000 | -0.0028 | -0.0000 | -0.0000 | 0.0000  | 0.0000  | 0.3728  | -0.5297 |
|      | 3px                              | -0.0142   | -0.0187 | -0.4916 | -0.0007 | 0.0043  | -0.0000 | 0.0000  | 0.6403  | -0.0074 | 0.0000  | 0.2812  | -0.0001 | -0.0064 | -0.0028 | -0.0000 | -0.6084 | 0.6516  |
|      | 3py                              | -0.0002   | -0.0002 | -0.0053 | 0.0660  | -0.3966 | -0.0000 | -0.0000 | 0.0069  | 0.6796  | -0.0032 | 0.0030  | 0.0115  | 0.5894  | 0.2616  | 0.0001  | -0.0066 | 0.0071  |
|      | 3pz                              | 0.0000  | -0.0000 | -0.0000 | 0.3967  | 0.0660  | 0.0000  | -0.0000 | 0.0000  | 0.0032  | 0.6796  | 0.0000  | -0.5894 | 0.0115  | -0.0001 | 0.2616  | 0.0000  | -0.0000 |
| Cl-3 | 3s                               | -0.6626   | 0.6829  | -0.1572 | -0.0000 | 0.0000  | 0.0000  | -0.0000 | -0.0675 | 0.0000  | -0.0000 | -0.0028 | 0.0000  | 0.0000  | 0.0000  | 0.0000  | 0.3728  | 0.5297  |
|      | 3px                              | 0.0142  | -0.0187 | 0.4916  | 0.0007  | -0.0043 | -0.0000 | 0.0000  | 0.6403  | -0.0074 | 0.0000  | -0.2812 | 0.0001  | 0.0064  | -0.0028 | -0.0000 | 0.6084  | 0.6516  |
|      | 3py                              | 0.0002  | -0.0002 | 0.0053  | -0.0660 | 0.3966  | -0.0000 | -0.0000 | 0.0069  | 0.6796  | -0.0032 | -0.0030 | -0.0115 | -0.5894 | 0.2616  | 0.0001  | 0.0066  | 0.0071  |
|      | 3pz                              | 0.0000  | 0.0000  | -0.0000 | -0.3967 | -0.0660 | 0.0000  | -0.0000 | 0.0000  | 0.0032  | 0.6796  | 0.0000  | 0.5894  | -0.0115 | -0.0001 | 0.2616  | 0.0000  | 0.0000  |

**Table 1.2: Eigenvector values of molecular orbitals of Ruthenium (II) iodide.**

| Atom | AOs<br>( $\chi$ )                | Eigenvector values or coefficients of Atomic Orbitals |         |         |         |         |         |         |         |         |         |         |         |         |         |         |         |         |
|------|----------------------------------|---|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|
|      |                                  | MO-1  | MO-2    | MO-3    | MO-4    | MO-5    | MO-6    | MO-7    | MO-8    | MO-9    | MO-10   | MO-11   | MO-12   | MO-13   | MO-14   | MO-15   | MO-16   | MO-17   |
| Ru-1 | 5s                               | 0.1682  | 0.0000  | -0.0000 | -0.0000 | 0.0083  | -0.0000 | -0.0000 | 0.0000  | 0.4973  | -0.0000 | 0.0000  | 0.0000  | -0.0000 | 0.0000  | 0.0000  | -1.0160 | 0.0000  |
|      | 5px                              | 0.0000  | -0.1190 | 0.0000  | 0.0000  | 0.0000  | 0.0000  | -0.0000 | 0.2102  | -0.0000 | 0.0000  | -0.0013 | 0.0000  | -0.0000 | -0.0109 | -0.0000 | 0.0000  | 1.3059  |
|      | 5py                              | -0.0000   | -0.0013 | 0.0000  | 0.0000  | -0.0000 | -0.0000 | 0.0000  | 0.0023  | -0.0000 | -0.0005 | 0.1194  | -0.0000 | 0.0000  | 1.0072  | 0.0023  | 0.0000  | 0.0142  |
|      | 5pz                              | -0.0000   | 0.0000  | -0.0000 | 0.0000  | 0.0000  | 0.0000  | 0.0000  | 0.0000  | 0.0000  | 0.1194  | 0.0005  | -0.0000 | -0.0000 | 0.0023  | -1.0073 | -0.0000 | 0.0000  |
|      | 4dx <sup>2</sup> -y <sup>2</sup> | 0.2522  | -0.0000 | -0.0002 | -0.0202 | -0.6232 | 0.0000  | 0.4999  | -0.0000 | -0.4917 | 0.0000  | -0.0000 | -0.0001 | 0.0081  | 0.0000  | 0.0000  | -0.3287 | 0.0000  |
|      | 4dz <sup>2</sup>                 | -0.1457   | -0.0000 | -0.0000 | -0.0000 | -0.3599 | 0.0000  | 0.8660  | 0.0000  | 0.2839  | -0.0000 | 0.0000  | 0.0000  | -0.0000 | -0.0000 | -0.0000 | 0.1898  | -0.0000 |
|      | 4dxy                             | 0.0055  | 0.0000  | 0.0079  | 0.9309  | -0.0135 | 0.0000  | 0.0108  | 0.0000  | -0.0107 | 0.0000  | -0.0000 | 0.0037  | -0.3745 | 0.0000  | 0.0000  | -0.0071 | -0.0000 |
|      | 4dxz                             | 0.0000  | 0.0000  | -0.9311 | 0.0079  | -0.0000 | 0.0108  | -0.0000 | -0.0000 | 0.0000  | -0.0000 | -0.0000 | 0.3745  | 0.0037  | 0.0000  | -0.0000 | 0.0000  | 0.0000  |
|      | 4dyz                             | 0.0000  | -0.0000 | -0.0101 | 0.0001  | 0.0000  | -0.9999 | 0.0000  | 0.0000  | -0.0000 | 0.0000  | 0.0000  | 0.0041  | 0.0000  | -0.0000 | 0.0000  | 0.0000  | 0.0000  |
| I-2  | 5s                               | 0.6043  | -0.6703 | 0.0000  | 0.0000  | 0.2911  | 0.0000  | 0.0000  | -0.1031 | -0.0636 | 0.0000  | -0.0000 | -0.0000 | 0.0000  | -0.0000 | -0.0000 | 0.3208  | -0.4207 |
|      | 5px                              | -0.0470   | -0.0091 | -0.0000 | -0.0023 | 0.3536  | 0.0000  | 0.0000  | 0.6191  | -0.4060 | 0.0000  | 0.0075  | -0.0001 | -0.0064 | -0.0022 | 0.0000  | -0.5813 | 0.6031  |
|      | 5py                              | -0.0005   | -0.0001 | 0.0018  | 0.2067  | 0.0038  | 0.0000  | 0.0000  | -0.0067 | -0.0040 | -0.0027 | 0.6879  | -0.0068 | 0.6787  | -0.2026 | -0.0005 | -0.0063 | 0.0065  |
|      | 5pz                              | -0.0000   | -0.0000 | -0.2076 | 0.0018  | 0.0000  | 0.0000  | 0.0000  | -0.0000 | 0.0000  | 0.6880  | 0.0027  | -0.6787 | -0.0068 | -0.0005 | 0.2026  | 0.0000  | -0.0000 |
| I-3  | 5s                               | 0.6043  | 0.6703  | -0.0000 | -0.0000 | 0.2911  | 0.0000  | 0.0000  | 0.1031  | -0.0637 | 0.0000  | -0.0000 | 0.0000  | -0.0000 | -0.0000 | -0.0000 | 0.3208  | 0.4207  |
|      | 5px                              | 0.4070  | -0.0091 | 0.0000  | 0.0023  | -0.3536 | -0.0000 | -0.0000 | -0.6191 | 0.4060  | 0.0000  | -0.0075 | -0.0001 | 0.0074  | 0.0022  | 0.0000  | 0.5813  | 0.6031  |
|      | 5py                              | 0.0005  | -0.0001 | -0.0018 | -0.2076 | -0.0038 | 0.0000  | 0.0000  | -0.0067 | 0.0044  | -0.0027 | -0.6879 | 0.0068  | -0.6787 | -0.2026 | -0.0005 | 0.0063  | 0.0065  |
|      | 5pz                              | -0.0000   | 0.0000  | 0.2076  | -0.0018 | 0.0000  | 0.0000  | 0.0000  | 0.0000  | 0.0000  | 0.6880  | 0.0027  | 0.6787  | 0.0068  | -0.0005 | 0.2026  | 0.0000  | 0.0000  |

**Table 2.1: Overlap matrix (Overlap integrals values) of Ruthenium (II) chloride.**

| AOs                                      | 5s<br>(Ru-1) | 5px<br>(Ru-1) | 5py<br>(Ru-1) | 5pz<br>(Ru-1) | 4dx <sup>2</sup> -y <sup>2</sup><br>(Ru-1) | 4dz <sup>2</sup><br>(Ru-1) | 4dxy<br>(Ru-1) | 4dxz<br>(Ru-1) | 4dyz<br>(Ru-1) | 3s<br>(Cl-2) | 3px<br>(Cl-2) | 3py<br>(Cl-2) | 3pz<br>(Cl-2) | 3s<br>(Cl-3) | 3px<br>(Cl-3) | 3py<br>(Cl-3) | 3pz<br>(Cl-3) |  |
|--|--------------|---------------|---------------|---------------|--|----------------------------|----------------|----------------|----------------|--------------|---------------|---------------|---------------|--------------|---------------|---------------|---------------|--|
| 5s (Ru-1)                                | 1.0000       |               |               |               |  |                            |                |                |                |              |               |               |               |              |               |               |               |  |
| 5px (Ru -1)                              | -0.0000      | 1.0000        |               |               |  |                            |                |                |                |              |               |               |               |              |               |               |               |  |
| 5py (Ru -1)                              | -0.0000      | -0.0000       | 1.0000        |               |  |                            |                |                |                |              |               |               |               |              |               |               |               |  |
| 5pz (Ru -1)                              | 0.0000       | 0.0000        | 0.0000        | 1.0000        |  |                            |                |                |                |              |               |               |               |              |               |               |               |  |
| 4dx <sup>2</sup> -y <sup>2</sup> (Ru -1) | 0.0000       | 0.0000        | -0.0000       | 0.0000        | 1.0000                                     |                            |                |                |                |              |               |               |               |              |               |               |               |  |
| 4dz <sup>2</sup> (Ru -1)                 | -0.0000      | -0.0000       | -0.0000       | 0.0000        | 0.0000                                     | 1.0000                     |                |                |                |              |               |               |               |              |               |               |               |  |
| 4dxy (Ru -1)                             | -0.0000      | 0.0000        | 0.0000        | 0.0000        | -0.0000                                    | 0.0000                     | 1.0000         |                |                |              |               |               |               |              |               |               |               |  |
| 4dxz (Ru 1)                              | 0.0000       | 0.0000        | 0.0000        | 0.0000        | 0.0000                                     | 0.0000                     | 0.0000         | 1.0000         |                |              |               |               |               |              |               |               |               |  |
| 4dyz (Ru -1)                             | 0.0000       | 0.0000        | 0.0000        | 0.0000        | 0.0000                                     | -0.0000                    | 0.0000         | -0.0000        | 1.0000         |              |               |               |               |              |               |               |               |  |
| 3s (Cl-2)                                | 0.2270       | 0.3406        | 0.0037        | 0.0000        | 0.0994                                     | -0.0574                    | 0.0022         | 0.0000         | 0.0000         | 1.0000       |               |               |               |              |               |               |               |  |
| 3px (Cl-2)                               | -0.3205      | -0.3862       | -0.0060       | 0.0000        | -0.1310                                    | 0.0757                     | -0.0038        | 0.0000         | 0.0000         | -0.0000      | 1.0000        |               |               |              |               |               |               |  |
| 3py (Cl-2)                               | -0.0035      | -0.0060       | 0.1707        | 0.0000        | -0.0034                                    | 0.0008                     | 0.0923         | 0.0000         | 0.0000         | -0.0000      | 0.0000        | 1.0000        |               |              |               |               |               |  |
| 3pz (Cl-2)                               | 0.0000       | 0.0000        | 0.0000        | 0.1708        | 0.0000                                     | 0.0000                     | 0.0000         | 0.0923         | 0.0010         | 0.0000       | 0.0000        | 0.0000        | 1.0000        |              |               |               |               |  |
| 3s (Cl-3)                                | 0.2270       | -0.3406       | -0.0037       | 0.0000        | 0.0994                                     | -0.0574                    | 0.0022         | 0.0000         | 0.0000         | 0.0004       | -0.0027       | -0.0000       | 0.0000        | 1.0000       |               |               |               |  |
| 3px (Cl-3)                               | 0.3205       | -0.3862       | -0.0060       | 0.0000        | 0.1310                                     | -0.0757                    | 0.0038         | 0.0000         | 0.0000         | 0.0027       | -0.0108       | -0.0001       | 0.0000        | 0.0000       | 1.0000        |               |               |  |
| 3py (Cl-3)                               | 0.0035       | -0.0060       | 0.1707        | 0.0000        | 0.0034                                     | -0.0008                    | -0.0923        | 0.0000         | 0.0000         | 0.0000       | -0.0001       | 0.0011        | 0.0000        | -0.0000      | -0.0000       | 1.0000        |               |  |
| 3pz (Cl-3)                               | 0.0000       | 0.0000        | 0.0000        | 0.1708        | 0.0000                                     | 0.0000                     | 0.0000         | -0.0923        | -0.0010        | 0.0000       | 0.0000        | 0.0000        | 0.0011        | 0.0000       | 0.0000        | 0.0000        | 1.0000        |  |

**Table 2.2: Overlap matrix (Overlap integrals values) of Ruthenium (II) iodide.**

| AOs                                      | 5s      | 5px     | 5py     | 5pz    | 4dx <sup>2</sup> -y <sup>2</sup> | 4dz <sup>2</sup> | 4dxy    | 4dxz    | 4dyz    | 5s      | 5px     | 5py     | 5pz    | 5s      | 5px     | 5py    | 5pz    |  |
|--|---------|---------|---------|--------|----------------------------------|------------------|---------|---------|---------|---------|---------|---------|--------|---------|---------|--------|--------|--|
|  | (Ru-1)  | (Ru-1)  | (Ru-1)  | (Ru-1) | (Ru-1)                           | (Ru-1)           | (Ru-1)  | (Ru-1)  | (Ru-1)  | (I-2)   | (I-2)   | (I-2)   | (I-2)  | (I-3)   | (I-3)   | (I-3)  | (I-3)  |  |
| 5s (Ru-1)                                | 1.0000  |         |         |        |                                  |                  |         |         |         |         |         |         |        |         |         |        |        |  |
| 5px (Ru -1)                              | -0.0000 | 1.0000  |         |        |                                  |                  |         |         |         |         |         |         |        |         |         |        |        |  |
| 5py (Ru -1)                              | -0.0000 | -0.0000 | 1.0000  |        |                                  |                  |         |         |         |         |         |         |        |         |         |        |        |  |
| 5pz (Ru -1)                              | 0.0000  | 0.0000  | 0.0000  | 1.0000 |                                  |                  |         |         |         |         |         |         |        |         |         |        |        |  |
| 4dx <sup>2</sup> -y <sup>2</sup> (Ru -1) | 0.0000  | 0.0000  | -0.0000 | 0.0000 | 1.0000                           |                  |         |         |         |         |         |         |        |         |         |        |        |  |
| 4dz <sup>2</sup> (Ru -1)                 | -0.0000 | -0.0000 | -0.0000 | 0.0000 | 0.0000                           | 1.0000           |         |         |         |         |         |         |        |         |         |        |        |  |
| 4dxy (Ru -1)                             | -0.0000 | 0.0000  | 0.0000  | 0.0000 | -0.0000                          | 0.0000           | 1.0000  |         |         |         |         |         |        |         |         |        |        |  |
| 4dxz (Ru 1)                              | 0.0000  | 0.0000  | 0.0000  | 0.0000 | 0.0000                           | 0.0000           | 0.0000  | 1.0000  |         |         |         |         |        |         |         |        |        |  |
| 4dyz (Ru -1)                             | 0.0000  | 0.0000  | 0.0000  | 0.0000 | 0.0000                           | -0.0000          | 0.0000  | -0.0000 | 1.0000  |         |         |         |        |         |         |        |        |  |
| 5s (I-2)                                 | 0.1844  | 0.2783  | 0.0030  | 0.0000 | 0.0748                           | -0.0432          | 0.0016  | 0.0000  | 0.0000  | 1.0000  |         |         |        |         |         |        |        |  |
| 5px (I-2)                                | -0.2793 | -0.3722 | -0.0053 | 0.0000 | -0.1156                          | 0.0667           | -0.0032 | -0.0000 | -0.0000 | -0.0000 | 1.0000  |         |        |         |         |        |        |  |
| 5py (I-2)                                | -0.0030 | -0.0053 | 0.1185  | 0.0000 | -0.0026                          | 0.0007           | 0.0605  | -0.0000 | 0.0000  | -0.0000 | 0.0000  | 1.0000  |        |         |         |        |        |  |
| 5pz (I-2)                                | 0.0000  | 0.0000  | 0.0000  | 0.1185 | 0.0000                           | 0.0000           | 0.0000  | 0.0605  | 0.0007  | 0.0000  | 0.0000  | 0.0000  | 1.0000 |         |         |        |        |  |
| 5s (I-3)                                 | 0.1844  | -0.2783 | -0.0030 | 0.0000 | 0.0748                           | -0.0432          | 0.0016  | 0.0000  | 0.0000  | 0.0001  | -0.0006 | -0.0000 | 0.0000 | 1.0000  |         |        |        |  |
| 5px (I-3)                                | 0.2793  | -0.3722 | -0.0053 | 0.0000 | 0.1156                           | -0.0667          | 0.0032  | -0.0000 | -0.0000 | 0.0006  | -0.0024 | -0.0000 | 0.0000 | 0.0000  | 1.0000  |        |        |  |
| 5py (I-3)                                | 0.0030  | -0.0053 | 0.1185  | 0.0000 | 0.0026                           | -0.0007          | -0.0605 | -0.0000 | 0.0000  | 0.0000  | -0.0000 | 0.0002  | 0.0000 | -0.0000 | -0.0000 | 1.0000 |        |  |
| 5pz (I-3)                                | 0.0000  | 0.0000  | 0.0000  | 0.1185 | 0.0000                           | 0.0000           | 0.0000  | -0.0605 | -0.0007 | 0.0000  | 0.0000  | 0.0000  | 0.0002 | 0.0000  | 0.0000  | 0.0000 | 1.0000 |  |



In order to examine the extent of involvement of 4d, 5s and 5p orbitals in the formation of molecular orbitals the values of coefficient of these orbitals have been added to see the total involvement in all the eleven MOs ( $\phi_1$ - $\phi_{11}$ ). The summation values of 4dxy, 4dxz, 4dx<sup>2</sup>-y<sup>2</sup>, 5s, 5px, 5py, and 5pz of ruthenium (II) chloride are 0.9025, 0.8758, 1.7533, 0.6598, 0.2383, 0.1302 and 0.1277, respectively and that of ruthenium (II) iodide are 0.9738, 0.9498, 1.8874, 0.6738, 0.3305, 0.1230 and 0.1199, respectively. The nonbonding orbitals 4dz<sup>2</sup> and 4dyz are excluded. It is clearly indicated that the maximum involvement is of 4dx<sup>2</sup>-y<sup>2</sup> orbital and the minimum of 5pz orbital in both halides. In ruthenium (II) chloride the value of 5p orbitals are in the range 0.2383 to 0.1277 which is very low in comparison to d orbitals (dxy, dxz, dx<sup>2</sup>-y<sup>2</sup>) which is in the range 1.7533 to 0.8758 and the value for 5s is 0.6598; whereas in ruthenium (II) iodide the value of 5p orbitals are between 0.3305 to 0.1199 which is very low in comparison to d orbitals (dxy, dxz, dx<sup>2</sup>-y<sup>2</sup>) which is in the range 1.8874 to 0.9498. The value for 5s is 0.6738. So the involvement of 5p orbitals is negligible in both in comparison of 4d orbitals. The extent of involvement of 4d, 5s and 5p orbitals of ruthenium in the formation of MOs in the chloride and iodide is well demonstrated by the graph (Fig-2) drawn between the orbitals and the summation values of their coefficients. The graph showing below clearly shows that the involvement of p orbitals is negligible. The summation values are highest in case of iodide and lowest in chloride. It is perhaps on this account the splitting of d orbitals is maximum in iodide and minimum in chloride.

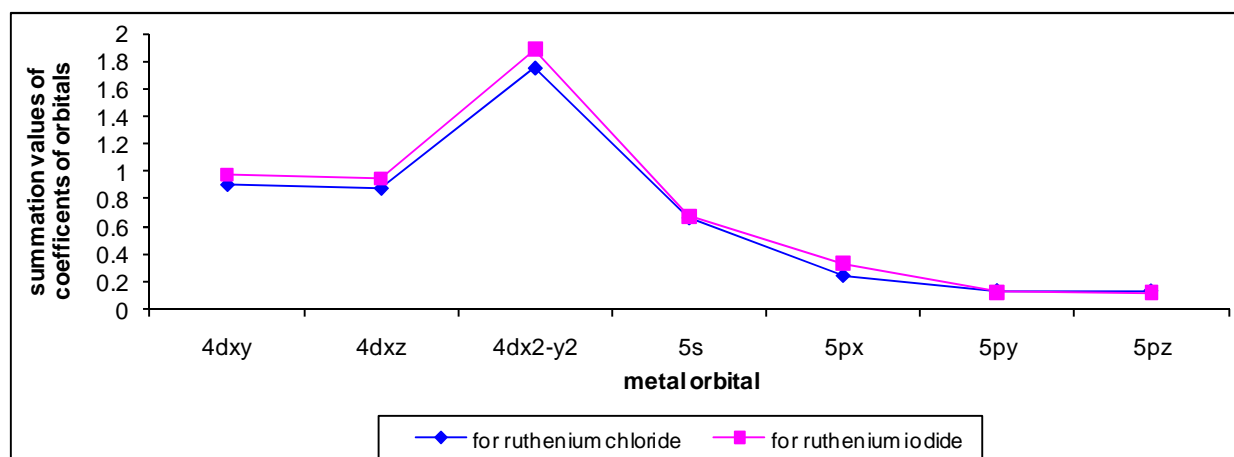


Fig- 2 Trend of extent of involvement of metal orbital in the formation of MOs of RuCl<sub>2</sub> & RuI<sub>2</sub>

### 3.1. Population Analysis:

The contributions of electrons in each occupied MO are calculated by using the population analysis method, introduced by Mulliken. This method apportions the electrons of n-electron molecule into net population  $n_r$  in the basis function  $\chi_r$ .

Let there be  $n_i$  electrons in the MO  $\phi_i$  ( $n_i = 0, 1, 2$ ) and let  $n_{r,i}$  symbolize the contribution of electrons in the MO  $\phi_i$  to the net population in  $\chi_r$ . We have

$$n_{r,i} = n_i c_{ri}^2 \quad (1)$$

where,  $c_{ri}$  is the coefficient of atomic orbitals for the  $i^{\text{th}}$  MO ( $r=1-17$ ).

Equation- 1 has been solved for 22 electrons of 11 molecular orbitals. Two electrons in the 1<sup>st</sup> MO to 11<sup>th</sup> MO have been considered. The six molecular orbitals having no electron are left

over. The data relating to  $c_{ri}$  have been taken from Table 1.1, 1.2. The results of solution of equation-1 are included in Table 3.1, 3.2 which enlists the contribution of electrons in molecular orbitals under two sections- major and minor. It is evident that major contribution is from 4d and 5s orbital. The p orbitals have negligible contribution. The details of contribution are as below:

**Table 3.1: Contribution of electrons in MO of Ruthenium (II) chloride.**

| MO. No | $n_i$ | Major contribution                      |                          | Minor contribution          |                          |
|--------|-------|---|--------------------------|-----------------------------|--------------------------|
|        |       | Basis function ( $\chi_r$ )             | $n_{r,i} = n_i c_{ri}^2$ | Basis function ( $\chi_r$ ) | $n_{r,i} = n_i c_{ri}^2$ |
| 1      | 2     | 5s (Ru 1)                               | 0.0211                   | 4dz <sup>2</sup> (Ru 1)     | 0.0073                   |
|        |       | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | 0.0220                   |                             |                          |
|        |       | 3s (Cl 2)                               | 0.8780                   |                             |                          |
|        |       | 3s (Cl 3)                               | 0.8780                   |                             |                          |
| 2      | 2     | 3s (Cl 2)                               | 0.9327                   | 5px (Ru 1)                  | 0.0094                   |
|        |       | 3s (Cl 3)                               | 0.9327                   |                             |                          |
| 3      | 2     | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | 0.4283                   | 5s (Ru 1)                   | 0.0186                   |
|        |       | 4dz <sup>2</sup> (Ru 1)                 | 0.1428                   | 3s (Cl 2)                   | 0.0494                   |
|        |       | 3px (Cl 2)                              | 0.4833                   | 3s (Cl 3)                   | 0.0494                   |
|        |       | 3px (Cl 3)                              | 0.4833                   |                             |                          |
| 4      | 2     | 4dxz (Ru 1)                             | 1.0999                   | 4dxy (Ru 1)                 | 0.0304                   |
|        |       | 3pz (Cl 2)                              | 0.3147                   | 4py (Cl 2)                  | 0.0087                   |
|        |       | 3pz (Cl 3)                              | 0.3147                   | 4py (Cl 3)                  | 0.0087                   |
| 5      | 2     | 4dxy (Ru 1)                             | 1.0996                   | 4dxz (Ru 1)                 | 0.0304                   |
|        |       | 3py (Cl 2)                              | 0.3145                   | 3pz (Cl 2)                  | 0.0087                   |
|        |       | 3py (Cl 3)                              | 0.3145                   | 3pz (Cl 3)                  | 0.0087                   |
| 6      | 2     | 4dyz (Ru 1)                             | 1.9996                   |                             |                          |
| 7      | 2     | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | 0.4998                   |                             |                          |
|        |       | 4dz <sup>2</sup> (Ru 1)                 | 1.4999                   |                             |                          |
| 8      | 2     | 5px (Ru 1)                              | 0.0565                   | 3s (Cl 2)                   | 0.0091                   |
|        |       | 3px (Cl 2)                              | 0.8199                   | 3s (Cl 3)                   | 0.0091                   |
|        |       | 3px (Cl 3)                              | 0.8199                   |                             |                          |
| 9      | 2     | 5py (Ru 1)                              | 0.0323                   |                             |                          |
|        |       | 3py (Cl 2)                              | 0.9237                   |                             |                          |
|        |       | 3py (Cl 3)                              | 0.9237                   |                             |                          |
| 10     | 2     | 5pz (Ru 1)                              | 0.0323                   |                             |                          |
|        |       | 3pz (Cl 2)                              | 0.9237                   |                             |                          |
|        |       | 3pz (Cl 3)                              | 0.9237                   |                             |                          |
| 11     | 2     | 5s (Ru 1)                               | 0.4237                   |                             |                          |
|        |       | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | 0.8895                   |                             |                          |
|        |       | 4dz <sup>2</sup> (Ru 1)                 | 0.2866                   |                             |                          |
|        |       | 3px (Cl 2)                              | 0.1581                   |                             |                          |
|        |       | 3px (Cl 3)                              | 0.1581                   |                             |                          |

Besides contribution of electrons the Mulliken's method is also used for evaluating overlap population, in order to distinguish bonding, nonbonding and antibonding molecular orbitals. This method allocates proportionally the overlap population  $n_{r-s}$  for all possible pairs of basis functions. Which is shown by the equation-2.

$$n_{r-s,i} = n_i (2c_{ri} c_{si} S_{rs}) \quad (2)$$

Where,  $c_{ri}$  = the coefficient of atomic orbitals for one atom.

$c_{si}$  = the coefficient of atomic orbitals for other atom  
and  $S_{rs}$  = the overlap integral between the two AOs (one of an atom and one of other atom).

**Table 3.2: Overlap populations of 1st MO of Ruthenium (II) iodide.**

| $n_i$                     | AOs                                     | $c_{ri}$ | AOs      | $c_{si}$ | $S_{rs}$ | $n_{r-s,i} = n_i(2c_{ri} \cdot c_{si} \cdot S_{rs})$ |
|---------------------------|---|----------|----------|----------|----------|--|
| 2                         | 5s(Ru 1)                                | 0.1682   | 5s(I 2)  | 0.6043   | 0.1844   | 0.0749   |
| 2                         | 5s(Ru 1)                                | 0.1682   | 5px(I 2) | -0.0470  | -0.2793  | 0.0088   |
| 2                         | 5s(Ru 1)                                | 0.1682   | 5py(I 2) | -0.0005  | -0.0030  | 0.0000   |
| 2                         | 5s(Ru 1)                                | 0.1682   | 5s(I 3)  | 0.6043   | 0.1844   | 0.0749   |
| 2                         | 5s(Ru 1)                                | 0.1682   | 5px(I 3) | 0.0470   | 0.2793   | 0.0088   |
| 2                         | 5s(Ru 1)                                | 0.1682   | 5py(I 3) | 0.0005   | 0.0030   | 0.0000   |
| 2                         | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | 0.2522   | 5s(I 2)  | 0.6043   | 0.0748   | 0.0455   |
| 2                         | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | 0.2522   | 5px(I 2) | -0.0470  | -0.1156  | 0.0054   |
| 2                         | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | 0.2522   | 5py(I 2) | -0.0005  | -0.0026  | 0.0000   |
| 2                         | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | 0.2522   | 5s(I 3)  | 0.6043   | 0.0748   | 0.0455   |
| 2                         | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | 0.2522   | 5px(I 3) | 0.0470   | 0.1156   | 0.0054   |
| 2                         | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | 0.2522   | 5py(I 3) | 0.0005   | 0.0026   | 0.0000   |
| 2                         | 4dz <sup>2</sup> (Ru 1)                 | -0.1457  | 5s(I 2)  | 0.6043   | -0.0432  | 0.0152   |
| 2                         | 4dz <sup>2</sup> (Ru 1)                 | -0.1457  | 5px(I 2) | -0.0470  | 0.0667   | 0.0018   |
| 2                         | 4dz <sup>2</sup> (Ru 1)                 | -0.1457  | 5py(I 2) | -0.0005  | 0.0007   | 0.0000   |
| 2                         | 4dz <sup>2</sup> (Ru 1)                 | -0.1457  | 5s(I 3)  | 0.6043   | -0.0432  | 0.0152   |
| 2                         | 4dz <sup>2</sup> (Ru 1)                 | -0.1457  | 5px(I 3) | 0.0470   | -0.0667  | 0.0018   |
| 2                         | 4dz <sup>2</sup> (Ru 1)                 | -0.1457  | 5py(I 3) | 0.0005   | -0.0007  | 0.0000   |
| 2                         | 4dxy(Ru 1)                              | 0.0055   | 5s(I 2)  | 0.6043   | 0.0016   | 0.0000   |
| 2                         | 4dxy(Ru 1)                              | 0.0055   | 5px(I 2) | -0.0470  | -0.0032  | 0.0000   |
| 2                         | 4dxy(Ru 1)                              | 0.0055   | 5py(I 2) | -0.0005  | 0.0605   | 0.0000   |
| 2                         | 4dxy(Ru 1)                              | 0.0055   | 5s(I 3)  | 0.6043   | 0.0016   | 0.0000   |
| 2                         | 4dxy(Ru 1)                              | 0.0055   | 5px(I 3) | 0.0470   | 0.0032   | 0.0000   |
| 2                         | 4dxy(Ru 1)                              | 0.0055   | 5py(I 3) | 0.0005   | -0.0605  | 0.0000   |
| 2                         | 5s(I 2)                                 | 0.6043   | 5s(I 3)  | 0.6043   | 0.0001   | 0.0001   |
| 2                         | 5s(I 2)                                 | 0.6043   | 5px(I 3) | 0.0470   | 0.0006   | 0.0000   |
| 2                         | 5px(I 2)                                | -0.0470  | 5s(I 3)  | 0.6043   | -0.0006  | 0.0000   |
| 2                         | 5px(I 2)                                | -0.0470  | 5px(I 3) | 0.0470   | -0.0024  | 0.0000   |
| 2                         | 5py(I 2)                                | -0.0005  | 5py(I 3) | 0.0005   | 0.0002   | 0.0000   |
| $\sum n_{r-s,i} = 0.3033$ |   |          |          |          |          |  |

It is evident from equation-2 that for overlap population analysis of MOs of a molecule, we need eigenvector values (coefficients), values of overlap matrix (overlap integrals) and number of electrons in each MO. The eigenvector and overlap integral values for halides of ruthenium have been taken from Table-1.1, 1.2 and Table-2.1, 2.2 respectively and the number of electrons is taken as two for 1<sup>st</sup> to 11<sup>th</sup> MOs and zero for 12<sup>th</sup> to 17<sup>th</sup> MO. With these values Table 4 is constructed for overlap-population contributions  $n_{r-s,i}$  of one molecular orbital. This table has 7 columns, defined as below. There will be 17 such tables for 17 MO but only 11 tables for each halide are constructed, because remaining six which have no electrons are left over. In such a way there will be 22 tables for all the two halides.

Column 1 – number of electron  $n_i$

Column 2, 4 – atomic orbitals of ruthenium and halogen.

Column 3 – coefficients of AOs of one atom ( $c_{ri}$ )

Column 5 – coefficients of AOs of other atom ( $c_{si}$ )

Column 6 – overlap integral between two AOs of different atoms ( $S_{rs}$ )

Column 7 – overlap population contribution  $n_{r-s,i}$ .

The possible overlaps between the various AOs of metal and halogens in each molecular orbital will be 88, as detailed below–

8 overlaps – 5s AO of ruthenium with ns, npx, npy, npz AOs of X-2 and X-3.

8 overlaps – 5px AO of ruthenium with ns, npx, npy, npz AOs of X-2 and X-3.

8 overlaps – 5py AO of ruthenium with ns, npx, npy, npz AOs of X-2 and X-3.

8 overlaps – 5pz AO of ruthenium with ns, npx, npy, npz AOs of X-2 and X-3.

8 overlaps –  $4dx^2-y^2$  AO of ruthenium with ns, npx, npy, npz AOs of X-2 and X-3.

8 overlaps –  $4dz^2$  AO of ruthenium with ns, npx, npy, npz AOs of X-2 and X-3.

8 overlaps – 4dxy AO of ruthenium with ns, npx, npy, npz AOs of X-2 and X-3.

8 overlaps – 4dxz AO of ruthenium with ns, npx, npy, npz AOs of X-2 and X-3.

8 overlaps – 4dyz AO of ruthenium with ns, npx, npy, npz AOs of X-2 and X-3.

4 overlaps – ns AO of X-2 with ns, npx, npy, npz AO of X-3.

4 overlaps – npx AO of X-2 with ns, npx, npy, npz AO of X-3.

4 overlaps – npy AO of X-2 with ns, npx, npy, npz AO of X-3.

4 overlaps – npz AO of X-2 with ns, npx, npy, npz AO of X-3.

Total– 88 overlaps

For the study of overlap population we have to construct eleven tables for each halide, each having 88 possible overlaps but while building up the table we have dropped the values of zero eigenvector value (Table 1.1, 1.2), hence each table of overlap-population contribution differs in its number of orbitals. For obtaining the values of overlap-population contributions ( $n_{r-s,i}$ ) we have to discuss each table separately, but for brevity we here discuss Table 4 for I<sup>st</sup> MO of ruthenium chloride.

#### **Ruthenium chloride:**

This table has 31 possible overlaps; out of which 24 provide coefficient values of ruthenium orbitals and 7 for Cl-2, in column 3 that are  $c_{ri}$ . Column-5 is for coefficient value  $c_{si}$ , for both the chlorines. Up to 24, both the chlorines are involved and for remaining seven only Cl-3. Column-6 is overlapping integral  $S_{rs}$  and exhibits the magnitude of overlap between the AOs represented in column-2 and 4. The values are self explanatory for indicating the magnitude.

#### **Ruthenium iodide**

This table has 29 possible overlaps; out of which 24 provide coefficient values of ruthenium orbitals and 5 for I-2, in column 3 that are  $c_{ri}$ . Column-5 is for coefficient value  $c_{si}$ , for both the chlorines. Up to 24, both the iodides are involved and for remaining five only I-3. Column-6, is overlap integral  $S_{rs}$  and exhibits the magnitude of overlap between the AOs represented in column-2 and 4. The values are self explanatory for indicating the magnitude.

The overlap population analysis also shows negligible involvement of 5p orbitals of ruthenium. It has earlier been suggested that much smaller radius of the 4d orbital than the 5s orbital makes the involvement of 5s orbital dominant contribution in the bonding<sup>[22, 23]</sup>. This hypothesis is the central theme of a recent text book of transition-metal chemistry by Gerloch and Constable<sup>[24]</sup>. While the importance of the valence ns and (n-1) d functions for the

description for transition metal bond is undisputed, the status of the empty np orbital is controversially discussed.

**Table 4: Overlap populations of I<sup>st</sup> MO of Ruthenium (II) chloride.**

| n <sub>i</sub>            | AOs                                     | c <sub>ri</sub> | AOs       | c <sub>si</sub> | S <sub>rs</sub> | n <sub>r-si</sub> = n <sub>i</sub> (2c <sub>ri</sub> .c <sub>si</sub> .S <sub>rs</sub> ) |
|---------------------------|---|-----------------|-----------|-----------------|-----------------|--|
| 2                         | 5s(Ru 1)                                | -0.1029         | 3s(Cl 2)  | -0.6626         | 0.2270          | 0.06190  |
| 2                         | 5s(Ru 1)                                | -0.1029         | 3px(Cl 2) | -0.0142         | -0.3205         | -0.00187   |
| 2                         | 5s(Ru 1)                                | -0.1029         | 3py(Cl 2) | -0.0002         | -0.0035         | 0.00000  |
| 2                         | 5s(Ru 1)                                | -0.1029         | 3s(Cl 3)  | -0.6626         | 0.2270          | 0.06190  |
| 2                         | 5s(Ru 1)                                | -0.1029         | 3px(Cl 3) | 0.0142          | 0.3205          | -0.00187   |
| 2                         | 5s(Ru 1)                                | -0.1029         | 3py(Cl 3) | 0.0002          | 0.0035          | 0.00000  |
| 2                         | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | -0.1049         | 3s(Cl 2)  | -0.6626         | 0.0994          | 0.02763  |
| 2                         | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | -0.1049         | 3px(Cl 2) | -0.0142         | -0.1310         | -0.00078   |
| 2                         | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | -0.1049         | 3py(Cl 2) | -0.0002         | -0.0034         | 0.00000  |
| 2                         | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | -0.1049         | 3s(Cl 3)  | -0.6626         | 0.0994          | 0.02763  |
| 2                         | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | -0.1049         | 3px(Cl 3) | 0.0142          | 0.1310          | -0.00078   |
| 2                         | 4dx <sup>2</sup> -y <sup>2</sup> (Ru 1) | -0.1049         | 3py(Cl 3) | 0.0002          | 0.0034          | 0.00000  |
| 2                         | 4dz <sup>2</sup> (Ru 1)                 | 0.0606          | 3s(Cl 2)  | -0.6626         | -0.0574         | 0.00921  |
| 2                         | 4dz <sup>2</sup> (Ru 1)                 | 0.0606          | 3px(Cl 2) | -0.0142         | 0.0757          | -0.00026   |
| 2                         | 4dz <sup>2</sup> (Ru 1)                 | 0.0606          | 3py(Cl 2) | -0.0002         | 0.0008          | 0.00000  |
| 2                         | 4dz <sup>2</sup> (Ru 1)                 | 0.0606          | 3s(Cl 3)  | -0.6626         | -0.0574         | 0.00921  |
| 2                         | 4dz <sup>2</sup> (Ru 1)                 | 0.0606          | 3px(Cl 3) | 0.0142          | -0.0757         | -0.00026   |
| 2                         | 4dz <sup>2</sup> (Ru 1)                 | 0.0606          | 3py(Cl 3) | 0.0002          | -0.0008         | 0.00000  |
| 2                         | 4dxy(Ru 1)                              | -0.0023         | 3s(Cl 2)  | -0.6626         | 0.0022          | 0.00001  |
| 2                         | 4dxy(Ru 1)                              | -0.0023         | 3px(Cl 2) | -0.0142         | -0.0038         | 0.00000  |
| 2                         | 4dxy(Ru 1)                              | -0.0023         | 3py(Cl 2) | -0.0002         | 0.0923          | 0.00000  |
| 2                         | 4dxy(Ru 1)                              | -0.0023         | 3s(Cl 3)  | -0.6626         | 0.0022          | 0.00001  |
| 2                         | 4dxy(Ru 1)                              | -0.0023         | 3px(Cl 3) | 0.0142          | 0.0038          | 0.00000  |
| 2                         | 4dxy(Ru 1)                              | -0.0023         | 3py(Cl 3) | 0.0002          | -0.0923         | 0.00000  |
| 2                         | 3s(Cl 2)                                | -0.6626         | 3s(Cl 3)  | -0.6626         | 0.0004          | 0.00070  |
| 2                         | 3s(Cl 2)                                | -0.6626         | 3px(Cl 3) | 0.0142          | 0.0027          | -0.00010   |
| 2                         | 3px(Cl 2)                               | -0.0142         | 3s(Cl 3)  | -0.6626         | -0.0027         | -0.00010   |
| 2                         | 3px(Cl 2)                               | -0.0142         | 3px(Cl 3) | 0.0142          | -0.0108         | 0.00000  |
| 2                         | 3px(Cl 2)                               | -0.0142         | 3py(Cl 3) | 0.0002          | -0.0001         | 0.00000  |
| 2                         | 3py(Cl 2)                               | -0.0002         | 3px(Cl 3) | 0.0142          | -0.0001         | 0.00000  |
| 2                         | 3py(Cl 2)                               | -0.0002         | 3py(Cl 3) | 0.0002          | 0.0011          | 0.00000  |
| $\sum n_{r-si} = 0.19218$ |   |                 |           |                 |                 |  |

Our results indicate that involvement of np orbital in transition metal bond is negligible and the main role is played by ns and by (n-1) d orbital. Landis<sup>[1-3]</sup> has also emphatically denied the involvement of np orbital in hybridization. He has supported sd hybridization and has based his observation on the bond angles. The idealized sd hybridization has been shown to have angles of 90<sup>0</sup>. This is because the energy curves are a function of the bond angles and have two minima one below 90<sup>0</sup> and one above 90<sup>0</sup>. The bond angles also support the Landis concept.

The column-7 of Table 4 enlists the values of overlap population, derived from the equation –2. The sum of the values of overlap-populations decides whether the MO in a covalent molecule is bonding, nonbonding or antibonding. If the sum of this inter atomic overlap population contribution is substantially positive, the MO is bonding; if substantially negative, the MO is antibonding and if zero or near zero, the MO is nonbonding. Table 4

indicates that the sum of overlap- population contribution in first MO of RuCl<sub>2</sub> is 0.19218 which is positive indicating or supporting the bonding nature of MO.

Similarly the sum of overlap population for the 11 MO in each halide has been worked out and the results are tabulated in Table 5 below:

| Table 5                                     |   |          |               |
|---|---|----------|---------------|
| Nature of occupied MOs of RuCl <sub>2</sub> |   |          |               |
| MO. No                                      | Sum of overlap population contribution ( $\sum n_{r-s,i}$ ) |          | Nature of MOs |
| 1   | 0.19218   | Positive | Bonding       |
| 2   | 0.12479   | Positive | Bonding       |
| 3   | 0.22345   | Positive | Bonding       |
| 4   | 0.22263   | Positive | Bonding       |
| 5   | -0.21482  | Negative | Antibonding   |
| 6   | 0.00000   | Zero     | Nonbonding    |
| 7   | 0.00000   | Zero     | Nonbonding    |
| 8   | 0.28493   | Positive | Bonding       |
| 9   | 0.06038   | Positive | Bonding       |
| 10  | 0.12005   | Positive | Bonding       |
| 11  | 0.07327   | Positive | Bonding       |
| Nature of occupied MOs of RuI <sub>2</sub>  |   |          |               |
| 1   | 0.3033  | Positive | Bonding       |
| 2   | 0.0888  | Positive | Bonding       |
| 3   | 0.0934  | Positive | Bonding       |
| 4   | 0.0934  | Positive | Bonding       |
| 5   | 0.1329  | Positive | Bonding       |
| 6   | 0.00000   | Zero     | Nonbonding    |
| 7   | 0.00000   | Zero     | Nonbonding    |
| 8   | 0.3358  | Positive | Bonding       |
| 9   | 0.3062  | Positive | Bonding       |
| 10  | 0.0781  | Positive | Bonding       |
| 11  | 0.0781  | Positive | Bonding       |

The overlap population analysis as presented in Table 5 shows that the nonbonding electrons are present in 6<sup>th</sup> and 7<sup>th</sup> molecular orbitals in both RuCl<sub>2</sub> and RuI<sub>2</sub>. The non bonding orbital is degenerate in all the cases. The eigenvector analysis as presented in Table-1.1, 1.2 indicates that these orbitals are 4dyz and 4dz<sup>2</sup>.

From the above discussion it is clear that no molecular orbital is formed by only two atomic orbitals. All molecular orbitals have contribution of many basis functions or atomic orbitals; as a result every molecular orbital has a definite shape having contribution from many basis functions.

## CONCLUSION

- Eigenvector analysis shows that 4dx<sup>2</sup>-y<sup>2</sup> and 4dxy orbitals of ruthenium play a major role in bonding between ruthenium and halides, 5s orbital is next and 4p orbitals have a negligible role. This supports the Landis observation and concept of sd hybridization.
- s and p orbitals of halogen are involved in bonding with ruthenium. There is a difference in energy levels of s and p orbitals are 0.1691 in chloride and 0.7472 in iodide.

- The overlap population analysis shows that the nonbonding orbitals are present in 6<sup>th</sup> and 7<sup>th</sup> molecular orbitals in both.
- No molecular orbital is formed by only two atomic orbitals. All molecular orbitals have contribution from many atomic orbitals; the difference is only in extent of involvement.

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