

## Part 1: Pseudo-code of operators

### [Mutation]

**SwappedMutation** (parent, child)

```
{
int subLength; // swapped part length
int pos1; // part 1 start index
int pos2; // part 2 start index

for (int i = 0; i < numberOfGenes; i++)
{
    if (i < pos1) // i before part 1
    {
        // Copy parent to child;
    }else if (i >= pos1 && i < pos1 + subLength) // i is in part 1
    {
        // Copy parent's part2 to child;
    }else if (i >= pos1 + subLength && i < pos2) // i is between part 1 and part 2
    {
        // Copy parent to child;
    }else if (i >= pos2 && i < pos2 + subLength) // i is in part 2
    {
        // Copy parent's part1 to child;
    }
    else
    {
        // Copy parent to child;
    }
}
}
```

**ReciprocalMutation** (parent, child)

```
{
    int pos1; // first element
    int pos2; // second element
    // make sure pos1 is less than pos2
    if (pos1 > pos2)
    {
        // Exchange pos1 & pos2;
    }
    for (int i = 0; i < numberOfGenes; i++)
    {
        if (i < pos1)
        {
            // copy parent to child
        }else if (i == pos1)
        {
            // copy parent's pos2 to child
        }else if (i > pos1 && i < pos2)
        {
            // copy parent to child
        }else if (i == pos2)
        {
            // copy parent's pos1 to child
        }
        else
        {
            // copy parent to child
        }
    }
}
```

### DisplacementMutation ( parent, child)

```
{
    int pos1; // cut-part start position
    int pos2; // cut-part end position
    // make sure pos1 is less then pos2
    if (pos1 > pos2)
    {
        // exchange pos1 & pos2
    }
    int subLength = pos2 - pos1;
    // determine insert position
    int insertPos = randomizer.Next(numberOfGenes - subLength);
    int minPos = Math.Min(pos1, insertPos);
    for (int i = 0; i < numberOfGenes; i++)
    {
        if (minPos == pos1)
        {
            if(i < pos1)
            {
                // copy parent to child
            }else if (i >= pos1 && i < insertPos)
            {
                // copy parent to child (except cut-part)
            }else if (i >= insertPos && i < insertPos + subLength)
            {
                // copy cut-part to child
            }
            else
            {
                // copy parent to child
            }
        }
        else
        {
            if (i < insertPos)
            {
                // copy parent to child
            }else if (i >= insertPos && i < insertPos + subLength)
```

```
{  
    // copy cut-part to child  
}else if (i >= insertPos + subLength && i < pos2)  
{  
    // copy parent to child (except cut-part)  
}  
else  
{  
    // copy parent to child  
}  
}  
}  
}
```

**InsertionMutation** ( parent, child)

```
{
    int pos1; // select a cut-part
    int pos2; // select a pos to insert cut-part
    for (int i = 0; i < numberOfGenes; i++)
    {
        if (i < pos1)
        {
            // copy parent to child
        }else if (i > pos2)
        {
            // copy parent to child
        }else if (i == pos1)
        {
            // copy parent to child's pos2
        }else
        {
            // copy parent to child orderly
        }
    }
}
```

**InversionMutation** ( parent, child)

```
{
    int pos1; // inverse-part start position
    int pos2; // inverse-part end position
    // make sure pos1 is less then pos2
    if (pos1 > pos2)
    {
        // exchange pos1 & pos2
    }
    for (int i = 0; i < numberOfGenes; i++)
    {
        // copy parent to child
    }

    for (int i = pos1; i < pos2; i++)
    {
        // fill in number inversely into child (overwrite original one)
    }
}
```

## [Crossover]

**PartialMappedCrossover** (father, mother, child1, child2)

```
{
    int pos1 = randomizer.Next(numberOfGenes);
    int pos2 = randomizer.Next(numberOfGenes);
    // make sure pos1 is less than pos2
    if (pos1 > pos2)
    {
        // exchange pos1 & pos2
    }
    // prepare proto-child
    for (int i = 0; i < numberOfGenes; i++)
    {
        if (i >= pos1 && i < pos2)
        {
            temp1[i] = chromosomes[motherIdx][i];
            temp2[i] = chromosomes[fatherIdx][i];
        }
        else
        {
            temp1[i] = chromosomes[fatherIdx][i];
            temp2[i] = chromosomes[motherIdx][i];
        }
    }
    // prepare map
    int cutLength = pos2 - pos1;
    int[][] map = new int[cutLength][];
    for (int i = 0; i < cutLength; i++)
    {
        map[i] = new int[2];
        for (int j = 0; j < 2; j++)
        {
            // initialize map member
        }
    }
    for (int i = 0; i < cutLength; i++)
    {
```

```

        // fill in map
    }
    // rearrange map
    for (int i = 0; i < cutLength; i++)
    {
        for (int j = i+1; j < cutLength; j++)
        {
            if (map[i][0] == map[j][1])
            {
                map[j][1] = map[i][1];
                map[i][0] = map[i][1] = -1; // erase useless(repeated) numbers
            }
            if (map[i][1] == map[j][0])
            {
                map[j][0] = map[i][0];
                map[i][0] = map[i][1] = -1; // erase useless(repeated) numbers
            }
        }
    }
    // prepare child
    for (int i = 0; i < numberOfGenes; i++)
    {
        if (i >= pos1 && i < pos2)
        {
            // copy cut-part to childs
        }
        else
        {
            // prepare child1
            if (IsContain(temp1[i], map))
            {
                for (int j = 0; j < cutLength; j++)
                {
                    if (temp1[i] == map[j][0])
                    {
                        chromosomes[child1Idx][i] = map[j][1]; // copy from map
                        break;
                    }
                }
            }
        }
    }

```



```

    }
}
else
{
    // copy from father
}
// prepare child2
if (IsContain(temp2[i], map))
{
    for (int j = 0; j < cutLength; j++)
    {
        if (temp2[i] == map[j][1])
        {
            chromosomes[child2Idx][i] = map[j][0]; // copy from map
            break;
        }
    }
}
else
{
    // copy from mother
}
}
}
}

```

**OrderCrossover** (father, mother, child1, child2)

```
{
    int pos1 = randomizer.Next(numberOfGenes);
    int pos2 = randomizer.Next(numberOfGenes);
    // make sure pos1 is less than pos2
    if (pos1 > pos2)
    {
        // exchange pos1 & pos2
    }
    // copy parent to proto-child
    for (int i = 0; i < numberOfGenes; i++)
    {
        temp1[i] = chromosomes[fatherIdx][i];
        temp2[i] = chromosomes[motherIdx][i];
    }
    // tag father's gene position with respect to mother's gene (by setting num = -1)
    for (int i = 0; i < numberOfGenes; i++)
    {
        if (i >= pos1 && i < pos2)
        {
            for (int j = 0; j < numberOfGenes; j++)
            {
                if (temp1[i] == temp2[j]) temp2[j] = -1;
            }
        }
    }
    // prepare child 1
    for (int i = 0; i < numberOfGenes; i++)
    {
        if (i < pos1)
        {
            for (int j = 0; j < numberOfGenes; j++)
            {
                if (temp2[j] != -1)
                {
                    chromosomes[child1Idx][i] = temp2[j]; // copy to child 1
                    temp2[j] = -1; // turn off this number
                    break;
                }
            }
        }
    }
}
```

```

        }
    }
}
}else if (i >= pos1 && i < pos2)
{
    chromosomes[child1Idx][i] = temp1[i];
}else
{
    for (int j = 0; j < numberOfGenes; j++)
    {
        if (temp2[j] != -1)
        {
            chromosomes[child1Idx][i] = temp2[j];
            temp2[j] = -1;
            break;
        }
    }
}
}
// copy parent to proto-child again
for (int i = 0; i < numberOfGenes; i++)
{
    temp1[i] = chromosomes[fatherIdx][i];
    temp2[i] = chromosomes[motherIdx][i];
}
// tag mother's gene position with respect to father's gene (by setting num
= -1)
for (int i = 0; i < numberOfGenes; i++)
{
    if (i >= pos1 && i < pos2)
    {
        for (int j = 0; j < numberOfGenes; j++)
        {
            if (temp2[i] == temp1[j]) temp1[j] = -1;
        }
    }
}
// prepare child 2
for (int i = 0; i < numberOfGenes; i++)

```

```

{
    if (i < pos1)
    {
        for (int j = 0; j < numberOfGenes; j++)
        {
            if (temp1[j] != -1)
            {
                chromosomes[child2Idx][i] = temp1[j]; // copy to child 2
                temp1[j] = -1; // turn off this number
                break;
            }
        }
    }
    else if (i >= pos1 && i < pos2)
    {
        chromosomes[child2Idx][i] = temp2[i];
    }
    else
    {
        for (int j = 0; j < numberOfGenes; j++)
        {
            if (temp1[j] != -1)
            {
                chromosomes[child2Idx][i] = temp1[j];
                temp1[j] = -1;
                break;
            }
        }
    }
}

```

**PositionBasedCrossover** (father, mother, child1, child2)

```
{
    // copy parent to proto-child
    for (int i = 0; i < numberOfGenes; i++)
    {
        temp1[i] = chromosomes[fatherIdx][i];
        temp2[i] = chromosomes[motherIdx][i];
        temp3[i] = chromosomes[fatherIdx][i];
        temp4[i] = chromosomes[motherIdx][i];
    }
    // determine number of selected genes
    int numOfSelectedGenes = randomizer.Next(numberOfGenes/3);
    int[] arrayPos = new int[numOfSelectedGenes];
    for (int i = 0; i < numOfSelectedGenes; i++)
    {
        arrayPos[i] = -1; // initialize array position index to -1
    }
    for (int i = 0; i < numOfSelectedGenes; i++)
    {
        int pos1 = randomizer.Next(numberOfGenes);
        if (!IsContain2(pos1, arrayPos))
        {
            arrayPos[i] = pos1; // add number to arrayPos
        }
        else
        {
            continue;
        }
        for (int j = 0; j < numberOfGenes; j++)
        {
            if (temp1[pos1] == temp2[j]) temp2[j] = -1; // turn off this number
            if (temp4[pos1] == temp3[j]) temp3[j] = -1; // turn off this number
        }
    }
    // prepare child 1
    for (int i = 0; i < numberOfGenes; i++)
    {
        if (IsContain2(i, arrayPos))
```

```

{
    chromosomes[child1Idx][i] = temp1[i];
}
else
{
    for (int j = 0; j < numberOfGenes; j++)
    {
        if (temp2[j] != -1)
        {
            chromosomes[child1Idx][i] = temp2[j];
            temp2[j] = -1;
            break;
        }
    }
}
// prepare child 2
for (int i = 0; i < numberOfGenes; i++)
{
    if (IsContain2(i, arrayPos))
    {
        chromosomes[child2Idx][i] = temp4[i];
    }
    else
    {
        for (int j = 0; j < numberOfGenes; j++)
        {
            if (temp3[j] != -1)
            {
                chromosomes[child2Idx][i] = temp3[j];
                temp3[j] = -1;
                break;
            }
        }
    }
}
}

```

**OrderBasedCrossover** (father, mother, child1, child2)

```
{
    // copy parent to proto-child
    for (int i = 0; i < numberOfGenes; i++)
    {
        temp1[i] = chromosomes[fatherIdx][i];
        temp2[i] = chromosomes[motherIdx][i];
        temp3[i] = chromosomes[fatherIdx][i];
        temp4[i] = chromosomes[motherIdx][i];
    }
    // determine number of selected genes
    int numOfSelectedGenes = randomizer.Next(numberOfGenes / 3);
    int[] arrayPos = new int[numOfSelectedGenes];
    int[] arrayPos2 = new int[numOfSelectedGenes];
    for (int i = 0; i < numOfSelectedGenes; i++)
    {
        // initialize arrayPos and arrayPos2
        arrayPos[i] = -1;
        arrayPos2[i] = -1;
    }
    for (int i = 0; i < numOfSelectedGenes; i++)
    {
        int pos1 = randomizer.Next(numberOfGenes);
        if (!IsContain2(pos1, arrayPos))
        {
            arrayPos[i] = pos1;
            for (int j = 0; j < numberOfGenes; j++)
            {
                if (temp1[pos1] == temp2[j])
                {
                    temp2[j] = -1; // turn off this number
                    arrayPos2[i] = j;
                }
                if (temp4[j] == temp3[pos1])
                {
                    temp3[pos1] = -1; // turn off this number
                }
            }
        }
    }
}
```

```

    }
    else
    {
        continue;
    }
}
Array.Sort(arrayPos);
Array.Sort(arrayPos2);

int count = 0;
for (int i = 0; i < numOfSelectedGenes; i++)
{
    if (arrayPos[i] != -1)
    {
        count = i;
        break;
    }
}

int count2 = 0;
for (int i = 0; i < numOfSelectedGenes; i++)
{
    if (arrayPos2[i] != -1)
    {
        count2 = i;
        break;
    }
}
// prepare child 1
for (int i = 0; i < numberOfGenes; i++)
{
    if (IsContain2(i, arrayPos2))
    {
        chromosomes[child1Idx][i] = temp1[arrayPos[count]];
        count++;
    }
    else
    {

```



```

for (int j = 0; j < numberOfGenes; j++)
{
    if (temp2[j] != -1)
    {
        chromosomes[child1Idx][i] = temp2[j];
        temp2[j] = -1;
        break;
    }
}
}

// prepare child 2
for (int i = 0; i < numberOfGenes; i++)
{
    if (IsContain2(i, arrayPos))
    {
        chromosomes[child2Idx][i] = temp4[arrayPos2[count2]];
        count2++;
    }
    else
    {
        for (int j = 0; j < numberOfGenes; j++)
        {
            if (temp3[j] != -1)
            {
                chromosomes[child2Idx][i] = temp3[j];
                temp3[j] = -1;
                break;
            }
        }
    }
}
}

```

**CycleCrossover** ( father, mother, child1, child2)

```
{
    // copy parent to proto-child
    for (int i = 0; i < numberOfGenes; i++)
    {
        temp1[i] = chromosomes[fatherIdx][i];
        temp2[i] = chromosomes[motherIdx][i];
        temp5[i] = -1;
        temp6[i] = -1;
    }
    temp5[0] = 0; // cycle1 index container
    temp6[0] = temp1[0]; // cycle1 container
    temp6[1] = temp2[0];

    // find cycle1 in parents
    for (int i = 1; i < numberOfGenes; i++)
    {
        for (int j = 0; j < numberOfGenes; j++)
        {
            if (temp6[i] == temp1[j])
            {
                temp6[i + 1] = temp2[j];
                temp5[i] = j;
                break;
            }
        }
        // check cycle is completed or not
        if (temp6[i + 1] == temp6[0])
        {
            break;
        }
    }
    // prepare child 1
    for (int i = 0; i < numberOfGenes; i++)
    {
        if (IsContain2(i, temp5))
        {
            //Copy father to child1
```

```
    }  
    else  
    {  
        // copy mother to child1  
    }  
}  
// prepare child 2  
for (int i = 0; i < numberOfGenes; i++)  
{  
    if (IsContain2(i, temp5))  
    {  
        // copy mother to child2  
    }  
    else  
    {  
        // copy father to child2  
    }  
}  
}
```

**SubtourCrossover** ( father , mother, child1, child2)

```
{
    // determine subtour position
    int pos1 = randomizer.Next(numberOfGenes);
    int pos2 = randomizer.Next(numberOfGenes);
    // make sure pos1 is less then pos2
    if (pos1 > pos2)
    {
        // exchange pos1 & pos2
    }
    // determine sublength
    int subLength = pos2 - pos1;
    // copy parent to proto-child
    for (int i = 0; i < numberOfGenes; i++)
    {
        temp1[i] = chromosomes[fatherIdx][i];
        temp2[i] = chromosomes[motherIdx][i];
        temp3[i] = -1;
    }
    // find father's subtour corresponding position in mother
    for (int i = 0; i < numberOfGenes; i++)
    {
        if (i >= pos1 && i < pos2)
        {
            for (int j = 0; j < numberOfGenes; j++)
            {
                if (temp1[i] == temp2[j])
                {
                    temp3[i] = j;
                    break;
                }
            }
        }
    }
}

Array.Sort(temp3);

// check mother's subtour is reasonable or not
```

```

bool result = true;
for (int i = 0; i < numberOfGenes-1; i++)
{
    if (temp3[i] != -1)
    {
        if (temp3[i+1] != (temp3[i] + 1))
        {
            result = false;
            break;
        }
    }
}

int startIndex = -1;
if (result == true)
{
    // find startIndex in mother
    for (int i = 0; i < numberOfGenes; i++)
    {
        if (temp3[i] != -1)
        {
            startIndex = temp3[i];
            break;
        }
    }

    // prepare child 1
    for (int i = 0; i < numberOfGenes; i++)
    {
        if (i >= pos1 && i < pos2)
        {
            chromosomes[child1Idx][i] = chromosomes[motherIdx][startIndex];
            startIndex++;
        }
        else
        {
            chromosomes[child1Idx][i] = chromosomes[fatherIdx][i];
        }
    }
}

```

```

    }

    // find startIndex in mother again
    for (int i = 0; i < numberOfGenes; i++)
    {
        if (temp3[i] != -1)
        {
            startIndex = temp3[i];
            break;
        }
    }

    int pos3 = pos1;
    // prepare child 2
    for (int i = 0; i < numberOfGenes; i++)
    {
        if (i >= startIndex && i < startIndex+subLength)
        {
            chromosomes[child2Idx][i] = chromosomes[fatherIdx][pos3];
            pos3++;
        }
        else
        {
            chromosomes[child2Idx][i] = chromosomes[motherIdx][i];
        }
    }
    else
    {
        for (int i = 0; i < numberOfGenes; i++)
        {
            // copy father and mother to child directly
        }
    }
}

```

## Part 2: Comments on Job Assignment Problem Solved by Binary- and Permutation-Encoded GA Solvers

### [Mutation comparison by experiments]

PMX + inversion

26 26 28 28 30

PMX + insertion

26 30 27 27 27

PMX + displacement

26 26 26 28 26

PMX + reciprocal exchange

26 26 26 26 26

PMX + swapped

26 26 29 30 26

Conclusion : “Reciprocal exchange” has average best solution (minimum objective value).

### [Crossover comparison by experiments]

PMX + reciprocal exchange

26 26 26 26 26

OX + reciprocal exchange

26 26 26 26 26

PBX + reciprocal exchange

26 26 26 26 26

OBX + reciprocal exchange

30 26 28 29 26

CX + reciprocal exchange

26 26 26 26 26

STX + reciprocal exchange

26 26 28 26 26

PMX + inversion

26 26 26 30 26

OX + inversion

26 26 26 26 27

PBX + inversion

28 29 31 30 31

OBX + inversion

30 31 31 30 27

CX + inversion

26 28 32 30 30

STX + inversion

26 26 28 31 27

Conclusion: PMX & OX are relatively good at finding best solution, while OBX is relatively bad.