

# 第十五章 DNA 数据库

## 第一节 DNA 数据库的动机

最早我设计了数据库一直不想集成在养疗经[17]中进行发布，因为是明文存储，避免文字处理的结果被人肆意使用，因为文字结果没有技术性，也不是算法，花大量时间做这个还不能保密，令我头疼。于是有个最早的动机，用元基来加密。

现在我想把这个数据库的增删改查驱动都用肽展公式来计算实现，那就要完整的语义库，于是我有个强烈的动机就是完善它的基础。满足生化计算的逻辑底层知识来源保障。

## 第二节 DNA 数据库的应用需求

原文: 控制吸收

肽语: PEEOUPPOAVCUPOECAUPEDOAU

肽锁: MMAA

散 列 肽 语 : MMAAPMMAEMMAEMMAOMMAOMMAAUMMAAPMMAAOMMAAAMMAAVMMAAC

MMAAUMMAAPMMAAOMMAAEMMAACMMAAAMMAAUMMAAPMMAAEMMAADMMAAO MMAAAMMAAU

静态密钥: 0. 6/0. 3/0. 5/0. 632

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静态肽展增元概率钥匙 E:

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静态肽展增元概率钥匙 S:

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静态肽展增元：

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时间: 1613805008008

账号随机缓存字符串: ID0. 058573949085271804: 0. 11193692344279549

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开始前序验证:

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开始概率钥匙解析：

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M->SOSOAAOOOSOAAOOSOAAOOSOAAOSOSOAAAOSOAAOOSOAAOOOSOAAOSOSOAAAAS  
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得到原降元元基DNA序列:

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得到新降元元基DNA序列:

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得到原元基DNA序列:

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得到新元基DNA序列:

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SOAAOOSOAAOOSOAAOSOSOAAAASOSOAAE

验证正确?

正确

开始后序验证:

准 备 计 算 元 基 DNA 序 列 :

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P->SOSOAAOOOSOAAOOSOAAOOSOAAOSOSOAAAOSOAAOOSOAAOOOSOAAOSOSOAAAASO  
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 UQIDIIDQUQQUQIUIDIIDIUQQUQQDIDIDIIDIUQQUQQUIDIIDIUQQUQQUQQIDQIDQUQQUQI  
 UQIDQIDQUQQUQQIDIDQIDQUQQUQQUIDQIDQUQIUQQUIDIIDIIDQUQQUQQUQIDIIDIUQQUQ  
 IUIDIIDIUQQUQQIDIDQIDQUQIUQQUQQIDQIDQUQIUQQUIDIIDIUQQUQQUIDIDQIDQUQQUQ  
 IUIDQIDIUQQUQQDIDQIDQUQQUQQDQIDQIDIUQQUQQUQQIDIIDIIDQUQQUQQU

得到原续降元元基DNA序列:

IDQIDIUQIUQIUIDIIDQIDQUQQUQQDIDQIDIUQQUQQUIDQIDQUQQUQQDQIDIIDIUQQUQIUQI  
 DIIDQUQQUQIUIDIIDIUQQUQQDIDIDIIDIUQQUQQUIDIIDIUQQUQQUQQIDQIDQUQQUQIUQI  
 DQIDQUQQUQQIDIDQIDQUQQUQQUIDQIDQUQIUQQUIDIIDIIDQUQQUQQUQIDIIDIUQQUQIUI  
 DIIDQUQQUQQIDIDQIDQUQIUQQUQQIDQIDQUQIUQQUIDIIDIUQQUQQUIDIDQIDQUQQUQIUI  
 DQIDIUQQUQQDIDQIDQUQQUQQDQIDQIDIUQQUQQUQQIDIIDIIDQUQQUQQU

得到后续降元元基DNA序列:

IDQIDIUQIUQIUIDIIDQIDQUQQUQQDIDQIDIUQQUQQUIDQIDQUQQUQQDQIDIIDIUQQUQIUQI  
 DIIDQUQQUQIUIDIIDIUQQUQQDIDIDIIDIUQQUQQUIDIIDIUQQUQQUQQIDQIDQUQQUQIUQI  
 DQIDQUQQUQQIDIDQIDQUQQUQQUIDQIDQUQIUQQUIDIIDIIDQUQQUQQUQIDIIDIUQQUQIUI  
 DIIDQUQQUQQIDIDQIDQUQIUQQUQQIDQIDQUQIUQQUIDIIDIUQQUQQUIDIDQIDQUQQUQIUI  
 DQIDIUQQUQQDIDQIDQUQQUQQDQIDQIDIUQQUQQUQQIDIIDIIDQUQQUQQU

验证正确?

正确

开始整序验证:

准备计算元基DNA序列:

SOSOAAOOOSOAAOOSOAAOOSOAAOSOSOAAAOSOAAOOSOAAOOOSOAAOSOSOAAAASOSO

M->SOSOAAOOOSOAAOOSOAAOOSOAAOOSOAAAOSOAAOOSOAAOOOSOAAOSOSOAAAAS  
OSOAAAOSOAAASOOSOAAOOSOAAOOOSOAAAOSOAAOOSOAAASOOSOAAAASOSOAAOOSOAA  
OOOSOAAOOSOAAOOSOAAOSOSOAAAASOSOAA

P->SOSOAAOOOSOAAOOSOAAOOSOAAOSOSOAAAOSOAAOOSOAAOOOSOAAOSOSOAAAASO  
SOAAAOSOAAASOOSOAAOOSOAAOOSOAAAOSOAAOOSOAAASOOSOAAASOSOAAOOSOAAO  
OOSOAAOOSOAAOOSOAAOSOSOAAAASOSOAA

O->SESSESAAESESSESSESAAESESSESAAESESSESAAESESSESAAAESSESAAESESSESAAESES  
ESSESAAESESSESAAAESSESSESAAAESSESAAESESSESAAESESSESAAESESSESAAAESSES  
AAESESSESAAESESSESAAAESSESSESAAESESSESAAESESSESAAESESSESAAESESSESAAESS  
ESSESAAESESSESAA

A->SESSESVSSESSESSESSESVSSESSESSESVSSESSESSESVSSESSESSESVSSESSESSESVSSES  
SESSESVSSESSESSESSESVSSESSESSESVSSESSESSESVSSESSESSESVSSESSESSESVSSES  
ESSESVSSESSESSESSESVSSESSESSESVSSESSESSESVSSESSESSESVSSESSESSESVSSESSES  
SESVSSESSESSESSESVSSESSESSESVSSESSESSESVSSESSESSESVSSESSESSESVSSESSES  
SESVSSESSESSESSESVSSESSESSESVSSESSESSESVSSESSESSESVSSESSESSESVSSES010000  
010111010011010111100001000011000001100000110000011101011001011100101110101010000111000  
01000011100101011010101000011001011001001101011110100111000111

S->IEQIEIVIVIEIEIEQIEQVQVQIEQIEIVQVQIEQIEQVQVQIEQIEIIEIVQVIVIEIIEQVQVIEIEIIEI  
VQVQIEIEIEIIEIVQVQIEIIEIIEIVQVQVQIEQIEQVQVIVIEQIEQVQVQIEIEQIEQVQVQIEIEQIEQVIV  
QIEIEIIEQVQVQVIEIIEIVQVIEIEIIEQVQVQIEIEQIEQVIVQVQIEQIEQVIVQIEIEIIEIVQVQIEIE  
QIEQVQVIEIEQIEIVQVQIEQIEQVQVQIEQIEQIEIVQVQVQIEIIEQVQVQ

C->IEQIEIVIVIEIEIEQIEQVQVQIEQIEIVQVQIEQIEQVQVQIEQIEIIEIVQVIVIEIIEQVQVIEIEIIEI  
VQVQIEIEIEIIEIVQVQIEIIEIIEIVQVQVQIEQIEQVQVIVIEQIEQVQVQIEIEQIEQVQVQIEIEQIEQVIV  
QIEIEIIEQVQVQVIEIIEIVQVIEIEIIEQVQVQIEIEQIEQVIVQVQIEQIEQVIVQIEIEIIEIVQVQIEIE  
QIEQVQVIEIEQIEIVQVQIEQIEQVQVQIEQIEQIEIVQVQVQIEIIEQVQVQ

E->IDQIDIVIVIUIDIDQIDQVQVQDIDQIDIVQVQUIDQIDQVQVQDQIDIIDIVQVIVIDIIDQVQVUII  
DIIDIVQVQDIDIDIIDIVQVQUIIDIIDIVQVQVQIDQIDQVQVIVIDQIDQVQVQIDIDQIDQVQVQUID  
QIDQVIVQUIDIDIIDQVQVQVIDIIDIVQVUIIDIIDQVQVQIDIDQIDQVIVQVQIDQIDQVIVQUIDIID  
IVQVQUIDIDQIDQVQVUIIDQIDIVQVQDIDQIDQVQVQDQIDQIDIVQVQVQIDIIDQVQVQ

V->IDQIDIUQIUQIUIDIDQIDQUQQUQQDIDQIDIUQQUQQUIDQIDQUQQUQQDQIDIIDIUQQUQI  
UQIDIIDQUQQUQIUIDIIDIUQQUQQDIDIDIIDIUQQUQQUIIDIIDIUQQUQQUQQIDQIDQUQQUQI  
UQIDQIDQUQQUQQIDIDQIDQUQQUQQUIDQIDQUQIUQQUIDIDIIDQUQQUQQUQIDIIDIUQQUQ

IUIDIHDQUQQUQQIDIDQIDQUQIUQQUQQIDQIDQUQIUQQUIDIIDIUQQUQQUIDIDQIDQUQQUQ  
IUIDQIDIUQQUQQDIDQIDQUQQUQQDQIDQIDIUQQUQQUQQIDIHDQUQQUQQ

得到原续降元元基DNA序列:

IDQIDIUQIUQIUIDIDQIDQUQQUQQDIDQIDIUQQUQQUIDQIDQUQQUQQDQIDIIDIUQQUQIUQI  
DIHDQUQQUQIUIDIIDIUQQUQQDIDIDIIDIUQQUQQUIIDIIDIUQQUQQUQQIDQIDQUQQUQIUQI  
DQIDQUQQUQQIDIDQIDQUQQUQQUIDQIDQUQIUQQUIDIDIHDQUQQUQQUQIDIIDIUQQUQIUI  
DIHDQUQQUQQIDIDQIDQUQIUQQUQQIDQIDQUQIUQQUIDIIDIUQQUQQUIDIDQIDQUQQUQIUI  
DQIDIUQQUQQDIDQIDQUQQUQQDQIDQIDIUQQUQQUQQIDIHDQUQQUQQU

得到后续降元元基DNA序列:

IDQIDIUQIUQIUIDIDQIDQUQQUQQDIDQIDIUQQUQQUIDQIDQUQQUQQDQIDIIDIUQQUQIUQI  
DIHDQUQQUQIUIDIIDIUQQUQQDIDIDIIDIUQQUQQUIIDIIDIUQQUQQUQQIDQIDQUQQUQIUQI  
DQIDQUQQUQQIDIDQIDQUQQUQQUIDQIDQUQIUQQUIDIDIHDQUQQUQQUQIDIIDIUQQUQIUI  
DIHDQUQQUQQIDIDQIDQUQIUQQUQQIDQIDQUQIUQQUIDIIDIUQQUQQUIDIDQIDQUQQUQIUI  
DQIDIUQQUQQDIDQIDQUQQUQQDQIDQIDIUQQUQQUQQIDIHDQUQQUQQU

验证正确?

正确

准备整序计算元基DNA序列:

SOSOAAOOOSOA AOOSOA AOOSOA AOSOSOAAAOSOA AOOSOA AOOSOA AOSOSOAAAASOSO  
AAAOSOA ASOOSOA AOOSOA AOOSOA AAAOSOA AOOSOA ASOOSOA AASOSOAAOOSOA AOOO  
SOAAOOSOA AOOSOA AOSOSOAAAASOSOAAE

准备整序计算元基DNA序列:

SOSOAAOOOSOA AOOSOA AOOSOA AOSOSOAAAOSOA AOOSOA AOOSOA AOSOSOAAAASOSO  
AAAOSOA ASOOSOA AOOSOA AOOSOA AAAOSOA AOOSOA ASOOSOA AASOSOAAOOSOA AOOO  
SOAAOOSOA AOOSOA AOSOSOAAAASOSOAAE

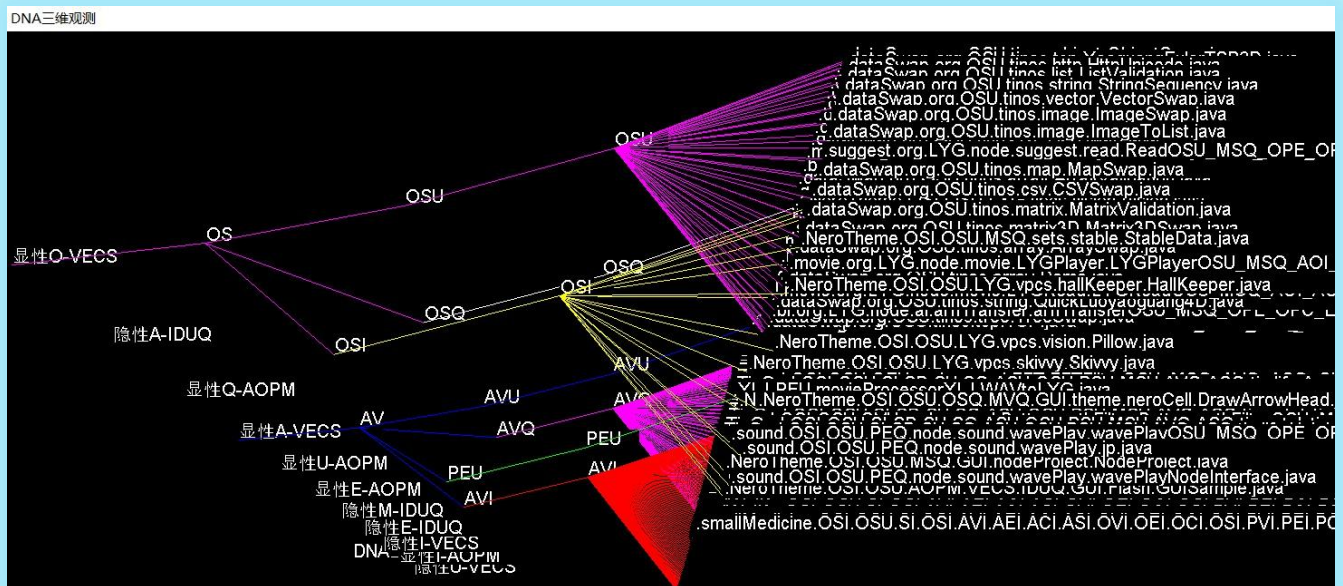
正确

### 第三节 DNA 数据库的具体描述

当数据原子能够完整的 无错的加密与解密,同时,密钥能完整固定不变,密码在解密的过程中能完整还原则为实用.于是我将概率钥匙和密钥对称性进行了离散推导如 **DNA 数据库数据加密**所示,非常的庆幸我有自己的 Socket 流 PLSQL 数据库[4]可以用来做实验,之后我的养疗经[17]将全面采用这种加密方式.

### 第四节 DNA 数据库的应用实现

#### DNA 数据库函数分类



#### DNA 数据库特征隐写

我在这里组织点文字来描述下什么是特征隐写,这里可不是情报学的散列数据加密隐写在大文件里呀,数据库特征隐写,是将每一大段的文字进行元基翻译,然后将元基按三元拆开来统计词汇,这些词汇之后一来用与 DNN 分析,二来做特征索引.

DNA 数据库文件安全 物理加密



之后养疗经[17]的所有文字文件都会变成物理元基加密文件. 用于本地存储, 提高一个安全级别, 因为, 明文文件存储信息非常不安全. 上面这图我做了个对比, 我把笔记原文和用法没有进行解密就显示出来, 大家可以看到明显的区别.

//元基编码加密:

```
String plsql= "setRoot: C: /DetaDB1; "; plsql+=
    "baseName: ZYY; ";
    plsql+= "tableName: zybc: insert; " +
        "columnValue: ID: "+ table. getValueAt(i, 0). toString(). replace(":", "@biky@")+ "; " + "columnValue: 打 分 : "+ table.
        getValueAt(i, 1). toString(). replace(":", "@biky@")+ "; " + "columnValue: 中药名称: "+ table. getValueAt(i, 2). toString().
        replace(":", "@biky@")+ "; " + "columnValue: 笔记原文: "+ new FullDNATokenPDI(). initonSect(table. getValueAt(i, 3).
toString(). replace(":", "@biky@"))+ "; " +
        "columnValue: 功效: "+ new FullDNATokenPDI(). initonSect(table. getValueAt(i, 4). toString(). replace(":", "@biky@"))+ "; "
+
        "columnValue: 风险规避: "+ new FullDNATokenPDI(). initonSect(table. getValueAt(i, 5). toString(). replace(":",
"@biky@"))+ "; " +
        "columnValue: 用量: "+ new FullDNATokenPDI(). initonSect(table. getValueAt(i, 6). toString(). replace(":", "@biky@"))+ "; "
+
        "columnValue: 性味: "+ new FullDNATokenPDI(). initonSect(table. getValueAt(i, 7). toString(). replace(":", "@biky@"))+ "; "
+
        "columnValue: 经脉: "+ new FullDNATokenPDI(). initonSect(table. getValueAt(i, 8). toString(). replace(":", "@biky@"))+ "; "
+
        "columnValue: 中医馆药理: "+ new FullDNATokenPDI(). initonSect(table. getValueAt(i, 9). toString(). replace(":",
"@biky@"))+ "; " +
        "columnValue: 经解: "+ new FullDNATokenPDI(). initonSect(table. getValueAt(i, 10). toString(). replace(":", "@biky@"))+ ";
" +
        "columnValue: 崇源: "+ new FullDNATokenPDI(). initonSect(table. getValueAt(i, 11). toString(). replace(":", "@biky@"))+ ";
" +
        "columnValue: 愚按: "+ new FullDNATokenPDI(). initonSect(table. getValueAt(i, 12). toString(). replace(":", "@biky@"))+ ";
" +
        "columnValue: 搭配: "+ new FullDNATokenPDI(). initonSect(table. getValueAt(i, 13). toString(). replace(":", "@biky@"))+ ";
" +
        "columnValue: 常见药: "+ new FullDNATokenPDI(). initonSect(table. getValueAt(i, 14).
```

```
toString().replace(":", "@biky@"))+ ";
```

```
try {
```

```
    org. ME. plsql. db. plsql. imp. ExecPLSQLImp. ExecPLSQL(plsql, mod);
```

上面是一段我的数据库 SQL, 我目前没有将元基加密集成在数据库工程里, 现在这样调试已经成功. 下面是元基解密函数实例.

```
@SuppressWarnings({ "unchecked", "rawtypes" })
```

```
public Map<String, Object> listToMap(Map<String, Object> dic_yw, Map<String, Object> dic_li,
```

```
    Map<String, Object> dic_hai, Map<String, Object> dic_xz, Map<String, Object> dic_ya,
```

```
    Map<String, Object> dic_jm, Map<String, Object> dic_xw, Map<String, Object> dic_cy,
```

```
    Map<String, Object> dic_jj, Map<String, Object> dic_zf, Map<String, Object> dic_cj,
```

```
    Map<String, Object> dic_yl) {
```

```
    Map<String, Object> dic_map= new ConcurrentHashMap<String, Object>();
```

```
    Map<String, Object> map = null;
```

```
    //for(int i=0; i<)
```

```
    String plsql= "setRoot: C: /DetaDB1; " +
```

```
        "baseName: ZYY; " +
```

```
        "tableName: zybc: select; " +
```

```
        "condition: or: ID|<=|3000; ";
```

```
    //"condition: or: ID|==|2; ";
```

```
try {
```

```
    map= org. ME. plsql. db. plsql. imp. ExecPLSQLImp. ExecPLSQL(plsql, true);
```

```
}catch(Exception e1) { e1.
```

```
    printStackTrace();
```

```
}
```

```
ArrayList list= (ArrayList) map. get("obj");
```

```
Iterator<HashMap<String, Object>> iterator= list. iterator();
```

```
while(iterator. hasNext()) {
```

```
    HashMap<String, Object> hashmap= iterator. next();
```

```
    StringBuilder stringBuilder= new StringBuilder();
```



```

if(hashmap. containsKey("rowValue")) {

    HashMap<String, Object> rowValue= (HashMap<String, Object>) hashmap. get("rowValue");

    String keyName= null;

    if(rowValue. containsKey("中药名称")) {

        HashMap<String, Object> temp= (HashMap<String, Object>) rowValue. get("中药名称");

        keyName= temp. get("columnValue"). toString();

        stringBuilder. append(temp. get("columnValue"). toString());

    }

    if(rowValue. containsKey("常见药")) {

        HashMap<String, Object> temp= (HashMap<String, Object>) rowValue. get("常见药");

        String cj= null== temp. get("columnValue")?"": temp. get("columnValue"). toString();

        cj= new FullDNATokenPDI(). initonDeSect(cj);

        dic_cj. put(keyName, cj. replace("@biky@", ": "));

        stringBuilder. append(cj);

    }

    if(rowValue. containsKey("搭配")) {

        HashMap<String, Object> temp= (HashMap<String, Object>) rowValue. get("搭配");

        String zf= null== temp. get("columnValue")?"": temp. get("columnValue"). toString();

        zf= new FullDNATokenPDI(). initonDeSect(zf);

        dic_zf. put(keyName, zf. replace("@biky@", ": "));

        stringBuilder. append(zf);

    }

    if(rowValue. containsKey("愚按")) {

        HashMap<String, Object> temp= (HashMap<String, Object>) rowValue. get("愚按");

        String ya= null== temp. get("columnValue")?"": temp. get("columnValue"). toString();

        ya= new FullDNATokenPDI(). initonDeSect(ya);

        dic_ya. put(keyName, ya. replace("@biky@", ": "));

        stringBuilder. append(ya);

    }

    if(rowValue. containsKey("崇源")) {

```

```

HashMap<String, Object> temp= (HashMap<String, Object>) rowValue. get("崇源");
String cy= null== temp. get("columnValue")?"" : temp. get("columnValue"). toString();
cy= new FullDNATokenPDI(). initonDeSect(cy);
dic_cy. put(keyName, cy. replace("@biky@", ": "));
stringBuilder. append(cy);
}

if(rowValue. containsKey("经解")) {
    HashMap<String, Object> temp= (HashMap<String, Object>) rowValue. get("经解");
    String jj= null== temp. get("columnValue")?"" : temp. get("columnValue"). toString();
    jj= new FullDNATokenPDI(). initonDeSect(jj);
    dic_jj. put(keyName, jj. replace("@biky@", ": "));
    stringBuilder. append(jj);
}

if(rowValue. containsKey("经脉")) {
    HashMap<String, Object> temp= (HashMap<String, Object>) rowValue. get("经脉");
    String jm= null== temp. get("columnValue")?"" : temp. get("columnValue"). toString();
    jm= new FullDNATokenPDI(). initonDeSect(jm);
    dic_jm. put(keyName, jm. replace("@biky@", ": "));
    stringBuilder. append(jm);
}

if(rowValue. containsKey("性味")) {
    HashMap<String, Object> temp= (HashMap<String, Object>) rowValue. get("性味");
    String xw= null== temp. get("columnValue")?"" : temp. get("columnValue"). toString();
    xw= new FullDNATokenPDI(). initonDeSect(xw);
    dic_xw. put(keyName, xw. replace("@biky@", ": "));
    stringBuilder. append(xw);
}

if(rowValue. containsKey("用量")) {
    HashMap<String, Object> temp= (HashMap<String, Object>) rowValue. get("用量");
    String yl= null== temp. get("columnValue")?"" : temp. get("columnValue"). toString();

```

```

//yl= new FullDNATokenPDI(). initonDeSect(yl);

dic_yl. put(keyName, yl. replace("@biky@", ": "));

stringBuilder. append(yl);

}

if(rowValue. containsKey("风险规避")) {

    HashMap<String, Object> temp= (HashMap<String, Object>) rowValue. get("风险规避");

    String hai= null== temp. get("culumnValue")?"" : temp. get("culumnValue"). toString();

    hai= new FullDNATokenPDI(). initonDeSect(hai);

    dic_hai. put(keyName, hai. replace("@biky@", ": "));

    stringBuilder. append(hai);

}

if(rowValue. containsKey("功效")) {

    HashMap<String, Object> temp= (HashMap<String, Object>) rowValue. get("功效");

    String li= null== temp. get("culumnValue")?"" : temp. get("culumnValue"). toString();

    li= new FullDNATokenPDI(). initonDeSect(li); dic_li.

    put(keyName, li. replace("@biky@", ": "));

    stringBuilder. append(li);

}

if(rowValue. containsKey("笔记原文")) {

    HashMap<String, Object> temp= (HashMap<String, Object>) rowValue. get("笔记原文");

    String yw= null== temp. get("culumnValue")?"" : temp. get("culumnValue"). toString();

    //yw= new FullDNATokenPDI(). initonDeSect(yw);

    dic_yw. put(keyName, yw. replace("@biky@", ": "));

    stringBuilder. append(yw);

}

if(rowValue. containsKey("中医馆药理")) {

    HashMap<String, Object> temp= (HashMap<String, Object>) rowValue. get("中医馆药理");

    String xz= null== temp. get("culumnValue")?"" : temp. get("culumnValue"). toString();

    xz= new FullDNATokenPDI(). initonDeSect(xz);

    dic_xz. put(keyName, xz. replace("@biky@", ": "));

```

```

        stringBuilder.append(xz);
    }
    dic_map.put(keyName, stringBuilder.toString().replace("@biky@", ": "));
}
}
return dic_map;
}

```

上面源码的作用是将文字进行元基化编辑,编辑的过程比较机械,仅仅通过 ASCII 编码进行 char 数字化然后元基编码的逻辑,之后随着语料库的不断完善,我会逐渐将其描述成语义元基编码方式,方便肽展计算.

#### DNA 数据库数据加密

	AOPM	VECS	IDUQ
A -> B	TRUE	TRUE	TRUE
B -> C	TRUE	TRUE	TRUE
A != C	FALSE	FALSE	FASLE
C -> B	TRUE	TRUE	TRUE

今天来组织点文字描述下这个逻辑: A 为待加密文件, B 为肽展公式计算的降元级别, C 为肽展公式计算的增元级别. 通过计算会发现一次降元和一次增元计算后, 文件与结果已经不对应.

#### XOR

	A	B	C
A -> B	AOPM VECS		
B -> C	VECSAOPM		
A != C			
C -> B	MPOA SCEV		

于是我开始进行离散逻辑推导归纳,发现MPOASCEV 的肽展计算能让C 与 B 计算结果吻合. 但 A 还是不等于 C

XNOR

	A	B	C
A -> B	MPOA SCEV		
B -> C	VECSAOPM		
A != C			
C -> B	MPOA SCEV		

而反向的 A 进行 MPOASCEV 肽展计算,A 也是不等于 C, 于是我开始思考是函数的问题还是概率的问题. 于是进行概率钥匙推导计算 Token 元基概率钥匙推导如下:

	A	B	C
A -> B	AOPMVECS	FD	
B -> C		VECSAOPM	FI
A -> B	AOPMVECS	TD	
B -> C		VECSAOPM	TI
C -> B			MPOASCEV TI
B->C		VECSAOPM	TI
C=A -> B	MPOASCEV	TI	

通过测试研究发现, 我得到一个结论,A 到 B 的时候, 已经将元基中的组合拓扑不稳定的元基过滤了, 于是B 到 C 和 C 到 B 的过程是一个循环可破解的过程, 于是我继续跟进测试, 得到健全的列表归纳如下:

Token 元基概率钥匙归纳如下:

DNA		PDE B		PDE C	
Dna	AOPMVECS	TDD	VECSAOPM	TII	
Full dna				MPOASCEV	TDI
Full back			MPOASCEV	TDI	VECSAOPM
dna					TII
Full up			VECSAOPM	TII	MPOASCEV
dna					TDI

在上面表中,2 组概率钥匙 D,I,D 用于服务器端计算,I 用于客户端通讯.  
目前已经在养疗经[17]中实用,效果不错. **T 为带钥匙计算** , **中间的的 DI 为增元降元计算识别** , **右边 DI 为钥匙种类** .

早期的肽展公式我没有单独量化,可以见 git 的 catalytic 最早肽展计算,24 个分类能依次展示,现在我把其概率串量化的版本这里也展示出来,保证演化的完整性.

```
package OSI. OSU. SI. SD. SU. SQ. ASU. OSU. PSU. MSU. AVQ. ASQ. tin. catalytic. procedure. pde;

//注意: 该 文件对应的是罗瑶光先生 DNA 编码 与 计算的两本 国家软著作 思想的编码 实现.

//公安部 与 知识产权委员会 已经备案, 可阅读 相关 著作权 原文 进行逻辑辨别.
```

```
public class PDE_Formular {

    public static TokenPDI tokenPDI;

    public static void main(String[] argv) { Initon
        initonA= new Initon(); initonA. setM(); // 改
        成 O 测试下Initon initonV= new Initon();
        initonV. setS();
        Initon initonS= new Initon(); initonS.
        setI();
```

---

```
Initon initonS1= new Initon(); initonS1. setO();
Initon initonS2= new Initon(); initonS2. setC();
Initon initonS3= new Initon(); initonS3. setU();
Initon initonS4= new Initon(); initonS4. setO();
Initon initonS5= new Initon(); initonS5. setC();
Initon initonS6= new Initon();
initonS6. setI();
Initon initonS7= new Initon(); initonS7. setP();
Initon initonS8= new Initon(); initonS8. setC();
Initon initonS9= new Initon(); initonS9. setU();
Initon initonS10= new Initon(); initonS10.
setP();
Initon initonS11= new Initon(); initonS11.
setC();
Initon initonS12= new Initon();
initonS12. setI();
initonA.  next=   initonV;
initonV.  prev=   initonA;
initonV.  next=   initonS;
initonS.  prev=   initonV;
initonS.  next=   initonS1;
initonS1. prev= initonS;
```

---

```

initonS. next= initonS1; initonS1.
prev= initonS; initonS1. next=
initonS2; initonS2. prev= initonS1;
initonS2. next= initonS3; initonS3.
prev= initonS2; initonS3. next=
initonS4; initonS4. prev= initonS3;
initonS4. next= initonS5; initonS5.
prev= initonS4; initonS5. next=
initonS6; initonS6. prev= initonS5;
initonS6. next= initonS7; initonS7.
prev= initonS6; initonS7. next=
initonS8; initonS8. prev= initonS7;
initonS8. next= initonS9; initonS9.
prev= initonS8; initonS9. next=
initonS10; initonS10. prev=
initonS9; initonS10. next=
initonS11; initonS11. prev=
initonS10; initonS11. next=
initonS12; initonS12. prev=
initonS11;

PDE_Formular pDE_RNA_Formular = new PDE_Formular();
pDE_RNA_Formular.do_PDE_RNA_Formular(initonA);
}

public void do_PDE_RNA_Formular(Initon initon) {
    tokenPDI= new TokenPDI();
    tokenPDI.key= new double[4];

```



---

```

tokenPDI.key[0]= 0. 6;

tokenPDI.key[1]= 0. 3;

tokenPDI.key[2]= 0. 5;

tokenPDI.key[3]= 0. 632;

//初始
Initon InitonPDE= initon;

System. out. print("input: " + InitonPDE. getStore());

while(InitonPDE. hasNext()) {

    InitonPDE= InitonPDE. forwardNext(); System. out.
    print(InitonPDE. getStore());

}

System. out. println();

System. out. println("肽展 降元");

while(InitonPDE. hasPrev()) {

    InitonPDE= InitonPDE. forwardPrev();

}

InitonLinkDNA initonLinkDNA= new InitonLinkDNA();
initonLinkDNA. setInitonLink(InitonPDE);
InitonPDE= new PDE_Decrement_Formular(). PDE_DecrementA(initonLinkDNA);

while(InitonPDE. hasPrev()) {

    InitonPDE= InitonPDE. forwardPrev();

}

System. out. println();

System. out. println("降元A = V + S");

while(InitonPDE. hasNext()) {

    System. out. print(InitonPDE. getStore());

    InitonPDE= InitonPDE. forwardNext();

}

System. out. print(InitonPDE. getStore());

while(InitonPDE. hasPrev()) {

```

---

```

        InitonPDE= InitonPDE. forwardPrev();
    }

    initonLinkDNA. setInitonLink(InitonPDE);

    InitonPDE= new PDE_Decrement_Formular(). PDE_DecrementO(initonLinkDNA);

    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }

    System. out. println();

    System. out. println("降元O = E + S");

    while(InitonPDE. hasNext()) {
        System. out. print(InitonPDE. getStore());
        InitonPDE= InitonPDE. forwardNext();
    }

    System. out. print(InitonPDE. getStore());

    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }

    initonLinkDNA. setInitonLink(InitonPDE);

    InitonPDE= new PDE_Decrement_Formular(). PDE_DecrementP(initonLinkDNA);

    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }

    System. out. println();

    System. out. println("降元P = E + C");

    while(InitonPDE. hasNext()) {
        System. out. print(InitonPDE. getStore());
        InitonPDE= InitonPDE. forwardNext();
    }

    System. out. print(InitonPDE. getStore());

    while(InitonPDE. hasPrev()) {

```

---

```

        InitonPDE= InitonPDE. forwardPrev();
    }

    initonLinkDNA. setInitonLink(InitonPDE);

    InitonPDE= new PDE_Decrement_Formular(). PDE_DecrementM(intonLinkDNA);

    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }

    System. out. println();

    System. out. println("降元M = C + S");

    while(InitonPDE. hasNext()) {
        System. out. print(InitonPDE. getStore());
        InitonPDE= InitonPDE. forwardNext();
    }

    System. out. print(InitonPDE. getStore());

    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }

    initonLinkDNA. setInitonLink(InitonPDE);

    InitonPDE= new PDE_Decrement_Formular(). PDE_DecrementV(intonLinkDNA);

    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }

    System. out. println();

    System. out. println("降元V = U + Q");

    while(InitonPDE. hasNext()) {
        System. out. print(InitonPDE. getStore());
        InitonPDE= InitonPDE. forwardNext();
    }

    System. out. print(InitonPDE. getStore());

    while(InitonPDE. hasPrev()) {

```

---

```

        InitonPDE= InitonPDE. forwardPrev();
    }

    initonLinkDNA. setInitonLink(InitonPDE);

    Initon InitonPDE_COPY= InitonPDE. copyRNA(InitonPDE); System. out.
println();

    System. out. println("1降元概率IU");
    doE_IU(InitonPDE, initonLinkDNA);


    initonLinkDNA. setInitonLink(InitonPDE_COPY); System. out.
println();

    System. out. println("2降元概率DU");
    doE_DU(InitonPDE_COPY, initonLinkDNA);
}

private static void doE_IU(Initon InitonPDE, InitonLinkDNA initonLinkDNA) {
    InitonPDE= new PDE_Decrement_Formular(). PDE_DecrementE_IU(intonLinkDNA, tokenPDI,
false);

    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }

    System. out. println();
    System. out. println("11降元E = I + U");

    while(InitonPDE. hasNext()) {
        System. out. print(InitonPDE. getStore());
        InitonPDE= InitonPDE. forwardNext();
    }

    System. out. print(InitonPDE. getStore());

    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }

    initonLinkDNA. setInitonLink(InitonPDE);

```

---

```

InitonPDE= new PDE_Decrement_Formular(). PDE_DecrementC(intonLinkDNA);

while(InitonPDE. hasPrev()) {
    InitonPDE= InitonPDE. forwardPrev();
}

System. out. println();

System. out. println("11降元C = I + D");

while(InitonPDE. hasNext()) {
    System. out. print(InitonPDE. getStore());
    InitonPDE= InitonPDE. forwardNext();
}

System. out. print(InitonPDE. getStore());

while(InitonPDE. hasPrev()) {
    InitonPDE= InitonPDE. forwardPrev();
}

intonLinkDNA. setInitonLink(InitonPDE);
intonLinkDNA. setInitonLink(InitonPDE);

Initon InitonPDE_COPYSI= InitonPDE. copyRNA(InitonPDE); Initon
InitonPDE_COPYSQ= InitonPDE. copyRNA(InitonPDE); System. out. println();

String s= "1111概率S_";

String si= "1111概率S_I_"; String
sq= "1111概率S_Q_";

System. out. println("1111概率S");

doS(s, InitonPDE, intonLinkDNA);

intonLinkDNA. setInitonLink(InitonPDE_COPYSI); System. out.
println();

System. out. println("1112概率S_I");

doS_I(si, InitonPDE_COPYSI, intonLinkDNA); intonLinkDNA.
setInitonLink(InitonPDE_COPYSQ); System. out. println();

```

---

```

System.out.println("1113概率S_Q");
doS_Q(sq, InitonPDE_COPYSQ, initonLinkDNA);
}

private static void doS_Q(String sq, Initon InitonPDE, InitonLinkDNA initonLinkDNA) { InitonPDE= new
PDE_Decrement_Formular(). PDE_DecrementS_Q(intonLinkDNA, tokenPDI); while(InitonPDE. hasPrev()) {
    InitonPDE= InitonPDE. forwardPrev();
}
System.out.println();

System.out.println(sq+ "降元S = Q");
while(InitonPDE. hasNext()) {
    System.out.print(InitonPDE. getStore());
    InitonPDE= InitonPDE. forwardNext();
}
System.out.print(InitonPDE. getStore());
while(InitonPDE. hasPrev()) {
    InitonPDE= InitonPDE. forwardPrev();
}
intonLinkDNA. setInitonLink(InitonPDE);

//全部 收
System.out.println();
System.out.println(sq+ "肽展 增元");
while(InitonPDE. hasPrev()) {
    InitonPDE= InitonPDE. forwardPrev();
}
intonLinkDNA= new InitonLinkDNA();
intonLinkDNA. setInitonLink(InitonPDE);

```

```

InitonPDE= new PDE_Increment_Formular(). PDE_IncrementV(intonLinkDNA);
while(InitonPDE. hasPrev()) {
    InitonPDE= InitonPDE. forwardPrev();
}
System. out. println();
System. out. println(sq+ "V = U + Q");
while(InitonPDE. hasNext()) {
    System. out. print(InitonPDE. getStore());
    InitonPDE= InitonPDE. forwardNext();
}
System. out. print(InitonPDE. getStore());
while(InitonPDE. hasPrev()) {
    InitonPDE= InitonPDE. forwardPrev();
}
intonLinkDNA. setInitonLink(InitonPDE);
Initon InitonPDE_COPY= InitonPDE. copyRNA(InitonPDE); System. out.
println();
System. out. println(sq+ "概率Increment IU");
doIncrementE_IU(sq, InitonPDE, intonLinkDNA);

intonLinkDNA. setInitonLink(InitonPDE_COPY); System. out.
println();
System. out. println(sq+ "概率Increment DU");
doIncrementE_DU(sq, InitonPDE_COPY, intonLinkDNA);
}

```

```

private static void doS_I(String si, Initon InitonPDE, InitonLinkDNA intonLinkDNA) { InitonPDE= new
PDE_Decrement_Formular(). PDE_DecrementS_I(intonLinkDNA, tokenPDI,
false);
while(InitonPDE. hasPrev()) {

```

---

```

        InitonPDE= InitonPDE. forwardPrev();
    }
    System. out. println();
    System. out. println(si+ "11降元S = I");
    while(InitonPDE. hasNext()) {
        System. out. print(InitonPDE. getStore());
        InitonPDE= InitonPDE. forwardNext();
    }
    System. out. print(InitonPDE. getStore());
    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }
    initonLinkDNA. setInitonLink(InitonPDE);

    //全部 收
    System. out. println();
    System. out. println(si+ "11肽展 增元");
    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }
    initonLinkDNA= new InitonLinkDNA();
    initonLinkDNA. setInitonLink(InitonPDE);
    InitonPDE= new PDE_Increment_Formular(). PDE_IncrementV(initionLinkDNA);
    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }
    System. out. println();
    System. out. println(si+ "11V = U + Q");
    while(InitonPDE. hasNext()) {
        System. out. print(InitonPDE. getStore());

```



---

```

        InitonPDE= InitonPDE. forwardNext();
    }
    System.out.print(InitonPDE. getStore());
    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }
    initonLinkDNA. setInitonLink(InitonPDE);
    Initon InitonPDE_COPY= InitonPDE. copyRNA(InitonPDE); System.out.
println();
    System.out.println(si+ "111概率Increment IU");
    doIncrementE_IU(si, InitonPDE, initonLinkDNA);

    initonLinkDNA. setInitonLink(InitonPDE_COPY); System.out.
println();
    System.out.println(si+ "112概率Increment DU");
    doIncrementE_DU(si, InitonPDE_COPY, initonLinkDNA);
}

private static void doS(String s, Initon InitonPDE, InitonLinkDNA initonLinkDNA) { InitonPDE= new
PDE_Decrement_Formular(). PDE_DecrementS(intonLinkDNA); while(InitonPDE. hasPrev()) {
    InitonPDE= InitonPDE. forwardPrev();
}
    System.out.println();

    System.out.println(s+ "11降元S = I + Q");
    while(InitonPDE. hasNext()) {
        System.out.print(InitonPDE. getStore());
        InitonPDE= InitonPDE. forwardNext();
    }
}

```

---

```

System.out.print(InitonPDE.getStore());

while(InitonPDE.hasPrev()) {
    InitonPDE= InitonPDE.forwardPrev();
}

initonLinkDNA.setInitonLink(InitonPDE);

//全部 收
System.out.println();

System.out.println(s+ "11肽展 增元");

while(InitonPDE.hasPrev()) {
    InitonPDE= InitonPDE.forwardPrev();
}

initonLinkDNA= new InitonLinkDNA();
initonLinkDNA.setInitonLink(InitonPDE);
InitonPDE= new PDE_Increment_Formular().PDE_IncrementV(initonLinkDNA);

while(InitonPDE.hasPrev()) {
    InitonPDE= InitonPDE.forwardPrev();
}

System.out.println();

System.out.println(s+ "11V = U + Q");

while(InitonPDE.hasNext()) {
    System.out.print(InitonPDE.getStore());
    InitonPDE= InitonPDE.forwardNext();
}

System.out.print(InitonPDE.getStore());

while(InitonPDE.hasPrev()) {
    InitonPDE= InitonPDE.forwardPrev();
}

initonLinkDNA.setInitonLink(InitonPDE);

Initon InitonPDE_COPY= InitonPDE.copyRNA(InitonPDE);

```

---

```
System.out.println();
```

```
System.out.println(s+ "111概率Increment IU");
```

```
doIncrementE_IU(s, InitonPDE, initonLinkDNA);
```

```
initonLinkDNA.setInitonLink(InitonPDE_COPY); System.out.
```

```
println();
```

```
System.out.println(s+ "112概率Increment DU");
```

```
doIncrementE_DU(s, InitonPDE_COPY, initonLinkDNA);
```

```
}
```

```
private static void doE_DU(Initon InitonPDE, InitonLinkDNA initonLinkDNA) { InitonPDE= new
```

```
PDE_Decrement_Formular().PDE_DecrementE_DU(intonLinkDNA,
```

```
tokenPDI);
```

```
while(InitonPDE.hasPrev()) {
```

```
    InitonPDE= InitonPDE.forwardPrev();
```

```
}
```

```
System.out.println();
```

```
System.out.println("21降元E = D + U");
```

```
while(InitonPDE.hasNext()) {
```

```
    System.out.print(InitonPDE.getStore());
```

```
    InitonPDE= InitonPDE.forwardNext();
```

```
}
```

```
System.out.print(InitonPDE.getStore());
```

```
while(InitonPDE.hasPrev()) {
```

```
    InitonPDE= InitonPDE.forwardPrev();
```

```
}
```

```
initonLinkDNA.setInitonLink(InitonPDE);
```

```
InitonPDE= new PDE_Decrement_Formular().PDE_DecrementC(intonLinkDNA);
```

```
while(InitonPDE.hasPrev()) {
```

```
    InitonPDE= InitonPDE.forwardPrev();
```

---

```

    }

    System.out.println();

    System.out.println("21降元C = I + D");

    while(InitonPDE.hasNext()) {

        System.out.print(InitonPDE.getStore());

        InitonPDE= InitonPDE.forwardNext();

    }

    System.out.print(InitonPDE.getStore());

    while(InitonPDE.hasPrev()) {

        InitonPDE= InitonPDE.forwardPrev();

    }


    initonLinkDNA.setInitonLink(InitonPDE);

    Initon InitonPDE_COPYSI= InitonPDE.copyRNA(InitonPDE); Initon
    InitonPDE_COPYSQ= InitonPDE.copyRNA(InitonPDE); String s= "2222概率
    S_";

    String si= "2222概率S_I_";

    String sq= "2222概率S_Q_"; System.
    out.println();

    System.out.println("2111概率S");

    doS(s, InitonPDE, initonLinkDNA);


    initonLinkDNA.setInitonLink(InitonPDE_COPYSI); System.out.
    println();

    System.out.println("2112概率S_I");

    doS_I(si, InitonPDE_COPYSI, initonLinkDNA);


    initonLinkDNA.setInitonLink(InitonPDE_COPYSQ); System.out.
    println();

    System.out.println("2113概率S_Q");

```

---

```

doS_Q(sq, InitonPDE_COPYSQ, initonLinkDNA);
}

private static void doIncrementE_DU(String sq, Initon InitonPDE, InitonLinkDNA initonLinkDNA) { initonLinkDNA.
    setInitonLink(InitonPDE);
    InitonPDE= new PDE_Increment_Formular(). PDE_IncrementE_DU(intonLinkDNA);
    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }
    System.out.println();
    System.out.println(sq+ "E = D + U");
    while(InitonPDE. hasNext()) {
        System.out.print(InitonPDE. getStore());
        InitonPDE= InitonPDE. forwardNext();
    }
    System.out.print(InitonPDE. getStore());
    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }
    initonLinkDNA. setInitonLink(InitonPDE);
    InitonPDE= new PDE_Increment_Formular(). PDE_IncrementC(intonLinkDNA);
    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }
    System.out.println();
    System.out.println(sq+ "C = I + D");
    while(InitonPDE. hasNext()) {
        System.out.print(InitonPDE. getStore());
        InitonPDE= InitonPDE. forwardNext();
    }
}

```

---

```

System.out.print(InitonPDE.getStore());

while(InitonPDE.hasPrev()) {
    InitonPDE= InitonPDE.forwardPrev();
}

initonLinkDNA.setInitonLink(InitonPDE);

```

```

initonLinkDNA.setInitonLink(InitonPDE);

Initon InitonPDE_COPYSI= InitonPDE.copyRNA(InitonPDE); Initon
InitonPDE_COPYSQ= InitonPDE.copyRNA(InitonPDE);

```

```

doIncrementS(sq + "EDU_IQ_", InitonPDE, initonLinkDNA);
doIncrementS_I(sq + "EDU_I_", InitonPDE_COPYSI, initonLinkDNA);
doIncrementS_Q(sq + "EDU_Q_", InitonPDE_COPYSQ, initonLinkDNA);
}

```

```

private static void doIncrementE_IU(String s, Initon InitonPDE, InitonLinkDNA initonLinkDNA) { initonLinkDNA.
setInitonLink(InitonPDE);

InitonPDE= new PDE_Increment_Formular().PDE_IncrementE_IU(initonLinkDNA, tokenPDI,
false);

while(InitonPDE.hasPrev()) {
    InitonPDE= InitonPDE.forwardPrev();
}

System.out.println();

System.out.println(s+ "E = I + U");

while(InitonPDE.hasNext()) {
    System.out.print(InitonPDE.getStore());
    InitonPDE= InitonPDE.forwardNext();
}

System.out.print(InitonPDE.getStore());

while(InitonPDE.hasPrev()) {

```

---

```

        InitonPDE= InitonPDE. forwardPrev();
    }

    initonLinkDNA. setInitonLink(InitonPDE);

    InitonPDE= new PDE_Increment_Formular(). PDE_IncrementC(initonLinkDNA);

    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }

    System. out. println();

    System. out. println(s+ "C = I + D");

    while(InitonPDE. hasNext()) {
        System. out. print(InitonPDE. getStore());
        InitonPDE= InitonPDE. forwardNext();
    }

    System. out. print(InitonPDE. getStore());

    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }

    initonLinkDNA. setInitonLink(InitonPDE);

    InitonPDE= new PDE_Increment_Formular(). PDE_IncrementS(initonLinkDNA);

    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }

    initonLinkDNA. setInitonLink(InitonPDE);

    Initon InitonPDE_COPYSI= InitonPDE. copyRNA(InitonPDE); Initon
    InitonPDE_COPYSQ= InitonPDE. copyRNA(InitonPDE);

    doIncrementS(s + "EIU_IQ_", InitonPDE, initonLinkDNA);
    doIncrementS_I(s + "EIU_I_", InitonPDE_COPYSI, initonLinkDNA);
    doIncrementS_Q(s + "EIU_Q_", InitonPDE_COPYSQ, initonLinkDNA);
}

```

---

```

private static void doIncrementS_Q(String s, Initon InitonPDE, InitonLinkDNA initonLinkDNA) { initonLinkDNA.
    setInitonLink(InitonPDE);
    InitonPDE= new PDE_Increment_Formular(). PDE_IncrementS_Q(initonLinkDNA);
    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }
    System. out. println();
    System. out. println(s+ "S = Q");
    while(InitonPDE. hasNext()) {
        System. out. print(InitonPDE. getStore());
        InitonPDE= InitonPDE. forwardNext();
    }
    System. out. print(InitonPDE. getStore());
    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }
    initonLinkDNA. setInitonLink(InitonPDE);
    doIncrementAOPM(s, InitonPDE, initonLinkDNA);
}

```

```

private static void doIncrementS_I(String s, Initon InitonPDE, InitonLinkDNA initonLinkDNA) { initonLinkDNA.
    setInitonLink(InitonPDE);
    InitonPDE= new PDE_Increment_Formular(). PDE_IncrementS_I(initonLinkDNA, tokenPDI);
    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }
    System. out. println();
    System. out. println(s+ "S = I");
    while(InitonPDE. hasNext()) {

```



---

```

        System.out.print(InitonPDE.getStore());
        InitonPDE= InitonPDE.forwardNext();
    }

    System.out.print(InitonPDE.getStore());
    while(InitonPDE.hasPrev()) {
        InitonPDE= InitonPDE.forwardPrev();
    }
    initonLinkDNA.setInitonLink(InitonPDE);
    doIncrementAOPM(s, InitonPDE, initonLinkDNA);
}

private static void doIncrementS(String s, Initon InitonPDE, InitonLinkDNA initonLinkDNA) {
    initonLinkDNA.setInitonLink(InitonPDE);
    InitonPDE= new PDE_Increment_Formular().PDE_IncrementS(initonLinkDNA);
    while(InitonPDE.hasPrev()) {
        InitonPDE= InitonPDE.forwardPrev();
    }
    System.out.println();
    System.out.println(s+ "S = I + Q");
    while(InitonPDE.hasNext()) {
        System.out.print(InitonPDE.getStore());
        InitonPDE= InitonPDE.forwardNext();
    }
    System.out.print(InitonPDE.getStore());
    while(InitonPDE.hasPrev()) {
        InitonPDE= InitonPDE.forwardPrev();
    }
    initonLinkDNA.setInitonLink(InitonPDE);
    doIncrementAOPM(s, InitonPDE, initonLinkDNA);
}

```

---

```

private static void doIncrementAOPM(String s, Initon InitonPDE, InitonLinkDNA initonLinkDNA) { InitonPDE= new
    PDE_Increment_Formular(). PDE_IncrementA(intonLinkDNA); while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }
    System. out. println();
    System. out. println(s+ "A = V + S");
    while(InitonPDE. hasNext()) {
        System. out. print(InitonPDE. getStore());
        InitonPDE= InitonPDE. forwardNext();
    }
    System. out. print(InitonPDE. getStore());
    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }
    initonLinkDNA. setInitonLink(InitonPDE);
    InitonPDE= new PDE_Increment_Formular(). PDE_IncrementO(intonLinkDNA);
    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }
    System. out. println();
    System. out. println(s+ "O = E + S");
    while(InitonPDE. hasNext()) {
        System. out. print(InitonPDE. getStore());
        InitonPDE= InitonPDE. forwardNext();
    }
    System. out. print(InitonPDE. getStore());
    while(InitonPDE. hasPrev()) {
        InitonPDE= InitonPDE. forwardPrev();
    }

```

---

```

    }

    initonLinkDNA.setInitonLink(InitonPDE);

    InitonPDE= new PDE_Increment_Formular(). PDE_IncrementP(intonLinkDNA);

    while(InitonPDE.hasPrev()) {

        InitonPDE= InitonPDE.forwardPrev();

    }

    System.out.println();

    System.out.println(s+ "P = E + C");

    while(InitonPDE.hasNext()) {

        System.out.print(InitonPDE.getStore());

        InitonPDE= InitonPDE.forwardNext();

    }

    System.out.print(InitonPDE.getStore());

    while(InitonPDE.hasPrev()) {

        InitonPDE= InitonPDE.forwardPrev();

    }

    initonLinkDNA.setInitonLink(InitonPDE);

    InitonPDE= new PDE_Increment_Formular(). PDE_IncrementM(intonLinkDNA);

    while(InitonPDE.hasPrev()) {

        InitonPDE= InitonPDE.forwardPrev();

    }

    System.out.println();

    System.out.println(s+ "M = C + S");

    while(InitonPDE.hasNext()) {

        System.out.print(InitonPDE.getStore());

        InitonPDE= InitonPDE.forwardNext();

    }

    System.out.print(InitonPDE.getStore());

    while(InitonPDE.hasPrev()) {

        InitonPDE= InitonPDE.forwardPrev();

```

```
    }  
    initonLinkDNA.setInitonLink(InitonPDE);  
}  
}
```

//上面文件测试结果如下:

input: MSIOCUOCIPCUPCI  
肽展 降元  
降 元  $A = V + S$   
MSIOCUOCIPCUPCI 降 元  
 $O = E + S$   
MSIESCUESCIPCUPCI 降元  
 $P = E + C$   
MSIESCUESCIECCUECCI  
降 元  $M = C + S$   
CSSIESCUESCIECCUECCI 降 元  
 $V = U + Q$   
CSSIESCUESCIECCUECCI  
1降元概率IU  
  
11 降 元  $E = I + U$   
CSSIIUSCUDUSCIDUCCUIUCCI 11降元  
 $C = I + D$   
IDSSIIUSIDUDUSIDIDUIDIDUIUIDIDI  
1111概率S  
  
1111 概 率  $S_{11}$  降 元  $S = I + Q$   
IDIQIQIUIQIDUDUIQIDIDUIDIDUIUIDIDI  
1111概率 $S_{11}$ 肽展 增元

1111 概 率  $S_{11V} = U + Q$   
IDIQIQIUIQIDUDUIQIDIDUIDIDUIUIDIDI  
1111概率 $S_{111}$ 概率Increment IU

1111 概 率  $S_E = I + U$   
IDIQIQIEIQIEEIQIDIEIDIEEIDIDI 1111 概  
率  $S_C = I + D$   
CIQIQIEIQIEEIQCIECIEECI 1111 概 率  
 $S_{EIU\_IQ\_S} = I + Q$   
CSSIESIEESCIECIEECI  
1111概率 $S_{EIU\_IQ\_A} = V + S$   
CSSIESIEESCIECIEECI 1111概率  
 $S_{EIU\_IQ\_O} = E + S$   
CSSIOIEOCIECIEECI  
1111概率 $S_{EIU\_IQ\_P} = E + C$   
CSSIOIEOCIECIEECI  
1111概率 $S_{EIU\_IQ\_M} = C + S$   
MSIOIEOCIECIEECI  
1111 概 率  $S_{EIU\_I\_S} = I$   
CSSSESSEESCSECSEECSS 1111概  
率 $S_{EIU\_I\_A} = V + S$   
CSSSESSEESCSECSEECSS 1111概  
率 $S_{EIU\_I\_O} = E + S$   
CSSSOSEOCSECSEECSS 1111概率  
 $S_{EIU\_I\_P} = E + C$   
CSSSOSEOCSPSEPCS  
1111概率 $S_{EIU\_I\_M} = C + S$   
MSSOSEOMPSEPM  
1111概率 $S_{EIU\_Q\_S} = Q$

CSSIESIEESCIECIEECCI 1111 概率

$$S\_EIU\_Q\_A = V + S$$

CSSIESIEESCIECIEECCI 1111 概率

$$S\_EIU\_Q\_O = E + S$$

CSSIOIEOCIECIEECCI

$$1111 \text{ 概率 } S\_EIU\_Q\_P = E + C$$

CSSIOIEOCIECIEECCI

$$1111 \text{ 概率 } S\_EIU\_Q\_M = C + S$$

MSIOIEOCIECIEECCI

$$1111 \text{ 概率 } S\_112 \text{ 概率 } \text{Increment DU}$$

$$1111 \text{ 概 率 } S\_E = D + U$$

IDIQIUIQIEEIQDIEIDIEIUIDIDI 1111 概

$$\text{率 } S\_C = I + D$$

CIQIUIQIEEIQDIECIEIUCCI 1111 概 率

$$S\_EDU\_IQ\_S = I + Q$$

CSSIIUSIEESCIECIEIUCCI

$$1111 \text{ 概率 } S\_EDU\_IQ\_A = V + S$$

CSSIIUSIEESCIECIEIUCCI 1111 概率

$$S\_EDU\_IQ\_O = E + S$$

CSSIIUSIEOCIECIEIUCCI 1111 概率

$$S\_EDU\_IQ\_P = E + C$$

CSSIIUSIEOCIECIEIUCCI

$$1111 \text{ 概率 } S\_EDU\_IQ\_M = C + S$$

MSIIUSIEOCIECIEIUCCI

$$1111 \text{ 概 率 } S\_EDU\_I\_S = I$$

CSSSSSSUSSEESSCSECSESUCCS 1111 概

$$\text{率 } S\_EDU\_I\_A = V + S$$

CSSSSSSUSSEESSCSECSESUCCS 1111 概

$$\text{率 } S\_EDU\_I\_O = E + S$$

CSSSSSSUSSEOSCSECSOUCCS 1111 概  
率  $S\_EDU\_I\_P = E + C$   
CSSSSSSUSSEOSCSPSOUCCS 1111 概  
率  $S\_EDU\_I\_M = C + S$   
MSSSSSUSSEOSMPSOUCM 1111 概 率  
 $S\_EDU\_Q\_S = Q$   
CISISIIUISIEEISCIECIEIUCCI 1111 概 率  
 $S\_EDU\_Q\_A = V + S$   
CISISIIUISIEEISCIECIEIUCCI 1111 概 率  
 $S\_EDU\_Q\_O = E + S$   
CISISIIUISIEEISCIECIEIUCCI 1111 概 率  
 $S\_EDU\_Q\_P = E + C$   
CISISIIUISIEEISCIPIE IUCCI  
1111概率 $S\_EDU\_Q\_M = C + S$ CISISIIUISIEEISCIPIE IUCCI  
1112概率 $S\_I$   
1001

1111 概 率  $S\_I\_11$  降 元  $S = I$   
IDQIIUIIDUDUQIDIDUIDIDUIUIDIDI  
1111概率 $S\_I\_11$ 肽展 增元

1111 概 率  $S\_I\_11V = U + Q$   
IDQIIUIIDUDVIDIDUIDIDUIUIDIDI  
1111概率 $S\_I\_11$ 概率Increment IU

1111 概 率  $S\_I\_E = I + U$   
IDQIEIIEDVIDIEIDIEEIDIDI 1111  
概 率  $S\_I\_C = I + D$   
CQIEIIEDVCIECIECCI

---

1111 概率  $S\_I\_EIU\_IQ\_S = I + Q$  CQIIEIIEDVCIECIEECI

1111 概率  $S\_I\_EIU\_IQ\_A = V + S$  CQIIEIIEDVCIECIEECI

1111 概率  $S\_I\_EIU\_IQ\_O = E + S$  CQIIEIIEDVCIECIEECI

1111 概率  $S\_I\_EIU\_IQ\_P = E + C$  CQIIEIIEDVCIPIEPCI

1111 概率  $S\_I\_EIU\_IQ\_M = C + S$  CQIIEIIEDVCIPIEPCI

1111 概 率  $S\_I\_EIU\_I\_S = I$

CSSSESEDVCSECSEECCS 1111 概率

$S\_I\_EIU\_I\_A = V + S$

CSSSESEDVCSECSEECCS 1111 概率

$S\_I\_EIU\_I\_O = E + S$

CSSSOSEDVCSECSEECCS 1111 概率

$S\_I\_EIU\_I\_P = E + C$

CSSSOSEDVCPSEPCS

1111 概率  $S\_I\_EIU\_I\_M = C + S$

MSSOSEDVMPSEPM

1111 概率  $S\_I\_EIU\_Q\_S = Q$  CSIIEIIEDVCIECIEECI

1111 概率  $S\_I\_EIU\_Q\_A = V + S$  CSIIEIIEDVCIECIEECI

1111 概率  $S\_I\_EIU\_Q\_O = E + S$  CSIIEIIEDVCIECIEECI

1111 概率  $S\_I\_EIU\_Q\_P = E + C$

CSIIEIIEDVCIPIEPCI

1111 概率  $S\_I\_EIU\_Q\_M = C + S$

MIIEIIEDVCIPIEPCI



1111 概率  $S_I$  112 概率 Increment DU

1111 概 率  $S_I E = D + U$

IDQIIIUIIEDVIDIEIDIEIUIDIDI 1111 概

率  $S_I C = I + D$

CQIIIUIIEDVCIECIEIUCCI 1111 概率

$S_I EDU_IQ_S = I + Q$

CQIIIUIIEDVCIECIEIUCCI

1111 概率  $S_I EDU_IQ_A = V + S$  CQIIIUIIEDVCIECIEIUCCI

1111 概率  $S_I EDU_IQ_O = E + S$

CQIIIUIIEDVCIECIEIUCCI 1111 概率

$S_I EDU_IQ_P = E + C$

CQIIIUIIEDVCIPIEIUCCI

1111 概率  $S_I EDU_IQ_M = C + S$  CQIIIUIIEDVCIPIEIUCCI

1111 概 率  $S_I EDU_I_S = I$

CSSSSUSSEDVCSECSUCCS 1111 概

率  $S_I EDU_I_A = V + S$

CSSSSUSSEDVCSECSUCCS 1111 概

率  $S_I EDU_I_O = E + S$

CSSSSUSSEDVCSECSOUCCS 1111 概

率  $S_I EDU_I_P = E + C$

CSSSSUSSEDVCSPSUCCS 1111 概率

$S_I EDU_I_M = C + S$

MSSSSUSSEDMPSOUCM

1111 概 率  $S_I EDU_Q_S = Q$

CSIIIUIIEDVCIECIEIUCCI 1111 概率

$S_I EDU_Q_A = V + S$

CSIIIUIIEDVCIECIEIUCCI

1111 概率  $S\_I\_EDU\_Q\_O = E + S$

CSIIIUIIEDVVCIECIEIUCCI 1111 概率

$S\_I\_EDU\_Q\_P = E + C$

CSIIIUIIEDVVCIECIEIUCCI

1111 概率  $S\_I\_EDU\_Q\_M = C + S$

MIIIUIIEDVVCIECIEIUCCI

1113 概率  $S\_Q$

1111 概 率  $S\_Q$  降 元  $S = Q$

IDQQI IUQIDUDUIIDIDUIDIDUIUIDIDI

1111 概率  $S\_Q$  肽展 增元

1111 概 率  $S\_Q\_V = U + Q$

IDQQI IVIDUDUIIDIDUIDIDUIUIDIDI

1111 概率  $S\_Q$  概率 Increment IU

1111 概 率  $S\_Q\_E = I + U$

IDQQI IVIEEIIDIEIDIEEIDIDI 1111 概 率

$S\_Q\_C = I + D$  CQQIIVIEEICIECIEECI

1111 概率  $S\_Q\_EIU\_IQ\_S = I + Q$  CQQIIVIEEICIECIEECI

1111 概率  $S\_Q\_EIU\_IQ\_A = V + S$  CQQIIVIEEICIECIEECI

1111 概率  $S\_Q\_EIU\_IQ\_O = E + S$  CQQIIVIEEICIECIEECI

1111 概率  $S\_Q\_EIU\_IQ\_P = E + C$  CQQIIVIEEICIECIEECI

1111 概率  $S\_Q\_EIU\_IQ\_M = C + S$  CQQIIVIEEICIECIEECI

---

1111 概 率  $S\_Q\_EIU\_I\_S = I$

CSSSSVSEESCSECSEEECCS 1111概率

$S\_Q\_EIU\_I\_A = V + S$

CSSSSAEESCSECSEEECCS 1111概率

$S\_Q\_EIU\_I\_O = E + S$

CSSSSAEOCSECSEEECCS

1111概率 $S\_Q\_EIU\_I\_P = E + C$

CSSSSAEOCSPSEPCS

1111概率 $S\_Q\_EIU\_I\_M = C + S$

MSSSAEOMPSEPM

1111概率 $S\_Q\_EIU\_Q\_S = QCSSIIVIEEICIECIEECI$

1111概率 $S\_Q\_EIU\_Q\_A = V + SCSSIIVIEEICIECIEECI$

1111概率 $S\_Q\_EIU\_Q\_O = E + SCSSIIVIEEICIECIEECI$

1111概率 $S\_Q\_EIU\_Q\_P = E + C$

CSSIIVIEEICIEPIEPCI

1111概率 $S\_Q\_EIU\_Q\_M = C + S$

MSIIVIEEICIEPIEPCI

1111概率 $S\_Q$  概率Increment DU

1111 概 率  $S\_Q\_E = D + U$

IDQQIIVIEEIIDIEIDIEIUIDIDI 1111 概

率  $S\_Q\_C = I + D$

CQQIIVIEEICIECIEIUCCI

1111概率 $S\_Q\_EDU\_IQ\_S = I + QCQQIIVIEEICIECIEIUCCI$

1111概率 $S\_Q\_EDU\_IQ\_A = V + SCQQIIVIEEICIECIEIUCCI$

---

1111 概率  $S\_Q\_EDU\_IQ\_O = E + S$

CQQIIVIEEICIECIEIUCCI

1111 概率  $S\_Q\_EDU\_IQ\_P = E + C$  CQQIIVIEEICIECIEIUCCI

1111 概率  $S\_Q\_EDU\_IQ\_M = C + S$  CQQIIVIEEICIECIEIUCCI

1111 概 率  $S\_Q\_EDU\_I\_S = I$

CSSSSVSEESCSECSSESUCCS 1111 概率

$S\_Q\_EDU\_I\_A = V + S$

CSSSSAEESCSECSSESUCCS 1111 概率

$S\_Q\_EDU\_I\_O = E + S$

CSSSSAEOSCSECSOUCCS

1111 概率  $S\_Q\_EDU\_I\_P = E + C$

CSSSSAEOSCPSOUCCS

1111 概率  $S\_Q\_EDU\_I\_M = C + S$

MSSSAEOMPSOUCM

1111 概率  $S\_Q\_EDU\_Q\_S = Q$  CSSIIVIEEICIECIEIUCCI

1111 概率  $S\_Q\_EDU\_Q\_A = V + S$  CSSIIVIEEICIECIEIUCCI

1111 概率  $S\_Q\_EDU\_Q\_O = E + S$  CSSIIVIEEICIECIEIUCCI

1111 概率  $S\_Q\_EDU\_Q\_P = E + C$

CSSIIVIEEICIECIEIUCCI

1111 概率  $S\_Q\_EDU\_Q\_M = C + S$

MSIIVIEEICIECIEIUCCI

2 降元 概率 DU

21 降 元  $E = D + U$

CSSIDUSCUDUSCHIUCCUDUCCI 21 降元 C

$= I + D$

---

IDSSIDUSIDUDUSIDIIUIDIDUDUIDIDI 2111 概率

S

2222 概 率  $S_{11}$  降 元  $S = I + Q$

IDIQIQIDUIQIDUDUIQIDIIUIDIDUDUIDIDI

2222 概率  $S_{11}$  肽展 增元

2222 概 率  $S_{11}V = U + Q$

IDIQIQIDUIQIDUDUIQIDIIUIDIDUDUIDIDI

2222 概率  $S_{111}$  概率 Increment IU 2222 概

率  $S_E = I + U$

IDIQIQIEIQIEEQIDIEIDIEEIDIDI 2222 概

率  $S_C = I + D$

CIQIQIEIQIEEQCIECIEECI 2222 概 率

$S_{EIU\_IQ\_S} = I + Q$

CSSIESIEESCIECIEECI

2222 概率  $S_{EIU\_IQ\_A} = V + S$

CSSIESIEESCIECIEECI 2222 概率

$S_{EIU\_IQ\_O} = E + S$

CSSIOIEOCIECIEECI

2222 概率  $S_{EIU\_IQ\_P} = E + C$

CSSIOIEOCIECIEECI

2222 概率  $S_{EIU\_IQ\_M} = C + S$

MSIOIEOCIECIEECI

2222 概 率  $S_{EIU\_I\_S} = I$

CSSSESSEESCSECSEECSS 2222 概

率  $S_{EIU\_I\_A} = V + S$

CSSSESSEESCSECSEECSS 2222 概

率  $S_{EIU\_I\_O} = E + S$

CSSSOSEOCSECSEECSS 2222 概率

$S_{EIU\_I\_P} = E + C$

CSSSOSEOCSPSEPCS

2222 概 率  $S_{\_EIU\_I\_M} = C + S$

MSSOSEOMPSEPM

2222 概 率  $S_{\_EIU\_Q\_S} = Q$

CSSIESIEESCIECIEECCI 2222 概 率

$S_{\_EIU\_Q\_A} = V + S$

CSSIESIEESCIECIEECCI 2222 概 率

$S_{\_EIU\_Q\_O} = E + S$

CSSIOIEOCIECIEECCI

2222 概 率  $S_{\_EIU\_Q\_P} = E + C$

CSSIOIEOCIECIEECCI

2222 概 率  $S_{\_EIU\_Q\_M} = C + S$

MSIOIEOCIECIEECCI

2222 概 率  $S_{\_112}$  概 率 Increment DU

2222 概 率  $S_{\_E} = D + U$

IDIQIQIEIQIEEQIDHUIDIEEIDIDI 2222 概

率  $S_{\_C} = I + D$

CIQIQIEIQIEEQIUCIEECCI 2222 概 率

$S_{\_EDU\_IQ\_S} = I + Q$

CSSIESIEESCIIUCIEECCI

2222 概 率  $S_{\_EDU\_IQ\_A} = V + S$

CSSIESIEESCIIUCIEECCI 2222 概 率

$S_{\_EDU\_IQ\_O} = E + S$

CSSIOIEOCIIUCIEECCI

2222 概 率  $S_{\_EDU\_IQ\_P} = E + C$

CSSIOIEOCIIUCIEECCI

2222 概 率  $S_{\_EDU\_IQ\_M} = C + S$

MSIOIEOCIIUCIEECCI

2222 概 率  $S_{\_EDU\_I\_S} = I$

---

CSSSSSESSEESSCSSUCSEECCS 2222 概

率  $S_{EDU\_I\_A} = V + S$

CSSSSSESSEESSCSSUCSEECCS 2222 概

率  $S_{EDU\_I\_O} = E + S$

CSSSSSOSSEOSCSSLUCSEECCS 2222 概

率  $S_{EDU\_I\_P} = E + C$

CSSSSSOSSEOSCSSLUCSEPCS 2222 概 率

$S_{EDU\_I\_M} = C + S$

MSSSSOSSEOSMSUMPEM

2222 概 率  $S_{EDU\_Q\_S} = Q$

CISISIEISIEEISCHUCIEECCI 2222概

率 $S_{EDU\_Q\_A} = V + S$

CISISIEISIEEISCHUCIEECCI 2222概

率 $S_{EDU\_Q\_O} = E + S$

CISISIEISIEEISCHUCIEECCI 2222概

率 $S_{EDU\_Q\_P} = E + C$

CISISIEISIEEISCHUCIEPCI 2222概率

$S_{EDU\_Q\_M} = C + S$

CISISIEISIEEISCHUCIEPCI

2112概率 $S_I$

10011111

2222 概 率  $S_{I\_11}$  降 元  $S = I$

IDQQIDUQIDUDUQIDIHUIDIDUDUIDIDI

2222概率 $S_{I\_11}$ 肽展 增元

2222 概 率  $S_{I\_11V} = U + Q$

IDQQIDVIDUDVIDIHUIDIDUDUIDIDI

2222概率 $S_{I\_11}$ 概率Increment IU

2222 概 率  $S_{I\_E} = I + U$

IDQQIDVIDVIDIEIDIEIDIDI

---

2222 概 率  $S\_I\_C = I + D$

CQQCVIEDVCIECIEECI 2222 概率

$S\_I\_EIU\_IQ\_S = I + Q$

CQQCVIEDVCIECIEECI

2222 概率  $S\_I\_EIU\_IQ\_A = V + S$  CQQCVIEDVCIECIEECI

2222 概率  $S\_I\_EIU\_IQ\_O = E + S$

CQQCVIEDVCIECIEECI 2222 概率

$S\_I\_EIU\_IQ\_P = E + C$

CQQCVIEDVCIPIEPCI

2222 概率  $S\_I\_EIU\_IQ\_M = C + S$  CQQCVIEDVCIPIEPCI

2222 概 率  $S\_I\_EIU\_I\_S = I$

CSSCVSEDVCSECSEECSS 2222 概率

$S\_I\_EIU\_I\_A = V + S$

CSSCAEDVCSECSEECSS 2222 概率

$S\_I\_EIU\_I\_O = E + S$

CSSCAEDVCSECSEECSS 2222 概率

$S\_I\_EIU\_I\_P = E + C$

CSSCAEDVCSPSEPCS

2222 概率  $S\_I\_EIU\_I\_M = C + S$

MSCAEDVMPSEPM

2222 概率  $S\_I\_EIU\_Q\_S = Q$  CSSCVIEDVCIECIEECI

2222 概率  $S\_I\_EIU\_Q\_A = V + S$  CSSCVIEDVCIECIEECI

2222 概率  $S\_I\_EIU\_Q\_O = E + S$

CSSCVIEDVCIECIEECI 2222 概率

$S\_I\_EIU\_Q\_P = E + C$

CSSCVIEDVCIPIEPCI



2222 概率  $S\_I\_EIU\_Q\_M = C + S$

MSCVIEDVCIPIEPCI

2222 概率  $S\_I\_112$  概率 Increment DU 2222 概

率  $S\_I\_E = D + U$

IDQQIDVIEDVIDIUIIDIEEIDIDI 2222 概 率

$S\_I\_C = I + D$  CQQCVIEDVCIUCIEECI

2222 概率  $S\_I\_EDU\_IQ\_S = I + Q$  CQQCVIEDVCIUCIEECI

2222 概率  $S\_I\_EDU\_IQ\_A = V + S$  CQQCVIEDVCIUCIEECI

2222 概率  $S\_I\_EDU\_IQ\_O = E + S$

CQQCVIEDVCIUCIEECI 2222 概率

$S\_I\_EDU\_IQ\_P = E + C$

CQQCVIEDVCIUCIEPCI

2222 概率  $S\_I\_EDU\_IQ\_M = C + S$  CQQCVIEDVCIUCIEPCI

2222 概 率  $S\_I\_EDU\_I\_S = I$

CSSCVSEDVCSSUCSEECCS 2222 概率

$S\_I\_EDU\_I\_A = V + S$

CSSCAEDVCSSUCSEECCS 2222 概率

$S\_I\_EDU\_I\_O = E + S$

CSSCAEDVCSSUCSEECCS 2222 概率

$S\_I\_EDU\_I\_P = E + C$

CSSCAEDVCSSUCSEPCS 2222 概率

$S\_I\_EDU\_I\_M = C + S$

MSCAEDVMSUMPEM

2222 概率  $S\_I\_EDU\_Q\_S = Q$

CSSCVIEDVCIUCIEECI

2222 概率  $S\_I\_EDU\_Q\_A = V + S$

---

CSSCVIEDVCIHUCIEECI 2222概率

$S\_I\_EDU\_Q\_O = E + S$

CSSCVIEDVCIHUCIEECI 2222概率

$S\_I\_EDU\_Q\_P = E + C$

CSSCVIEDVCIHUCIEPCI

2222概率 $S\_I\_EDU\_Q\_M = C + S$

MSCVIEDVCIHUCIEPCI

2113概率 $S\_Q$

2222 概 率  $S\_Q$  降 元  $S = Q$

IDQQIDUQIDUDUIIDIIUIDIDUDUIDIDI

2222概率 $S\_Q$ 肽展 增元

2222 概 率  $S\_Q\_V = U + Q$

IDQQIDVIDUDUIIDIIUIDIDUDUIDIDI

2222概率 $S\_Q$ 概率Increment IU 2222 概

率  $S\_Q\_E = I + U$

IDQQIDVIEEIIDIEIDIEIDIDI 2222 概

率  $S\_Q\_C = I + D$

CQQCVIEEICIECIEECI

2222概率 $S\_Q\_EIU\_IQ\_S = I + Q$ CQQCVIEEICIECIEECI

2222概率 $S\_Q\_EIU\_IQ\_A = V + S$ CQQCVIEEICIECIEECI

2222概率 $S\_Q\_EIU\_IQ\_O = E + S$ CQQCVIEEICIECIEECI

2222概率 $S\_Q\_EIU\_IQ\_P = E + C$ CQQCVIEEICIEPCI

2222概率 $S\_Q\_EIU\_IQ\_M = C + S$ CQQCVIEEICIEPCI

2222概率 $S\_Q\_EIU\_I\_S = I$

CSSCVSEESCSECSEECSS

---

2222 概率  $S\_Q\_EIU\_I\_A = V + S$

CSSCAEESCSECSEEECCS

2222 概率  $S\_Q\_EIU\_I\_O = E + S$

CSSCAEOCSECSEEECCS

2222 概率  $S\_Q\_EIU\_I\_P = E + C$

CSSCAEOCSPSEPCS

2222 概率  $S\_Q\_EIU\_I\_M = C + S$

MSCAEOMPSEPM

2222 概率  $S\_Q\_EIU\_Q\_S = QCSSCVIEEICIECIEECI$

2222 概率  $S\_Q\_EIU\_Q\_A = V + SCSSCVIEEICIECIEECI$

2222 概率  $S\_Q\_EIU\_Q\_O = E + SCSSCVIEEICIECIEECI$

2222 概率  $S\_Q\_EIU\_Q\_P = E + C$

CSSCVIEEICIPIEPCI

2222 概率  $S\_Q\_EIU\_Q\_M = C + S$

MSCVIEEICIPIEPCI

2222 概率  $S\_Q$  概率 Increment DU 2222 概

率  $S\_Q\_E = D + U$

IDQQIDVIEEIIDIIUIDIEEIDIDI 2222 概

率  $S\_Q\_C = I + D$

CQQCVIEEICIIUCIEECI

2222 概率  $S\_Q\_EDU\_IQ\_S = I + QCQQCVIEEICIIUCIEECI$

2222 概率  $S\_Q\_EDU\_IQ\_A = V + SCQQCVIEEICIIUCIEECI$

2222 概率  $S\_Q\_EDU\_IQ\_O = E + SCQQCVIEEICIIUCIEECI$

2222 概率  $S\_Q\_EDU\_IQ\_P = E + C$

---

CQQCVIEEICHUCIEPCI

2222 概率  $S\_Q\_EDU\_IQ\_M = C + S$  CQQCVIEEICHUCIEPCI

2222 概 率  $S\_Q\_EDU\_I\_S = I$

CSSCVSEESCSSUCSEECCS 2222 概率

$S\_Q\_EDU\_I\_A = V + S$

CSSCAEESCSSUCSEECCS 2222 概率

$S\_Q\_EDU\_I\_O = E + S$

CSSCAEOCSSUCSEECCS

2222 概率  $S\_Q\_EDU\_I\_P = E + C$

CSSCAEOCSSUCSEPCS

2222 概率  $S\_Q\_EDU\_I\_M = C + S$

MSCAEOMSUMEPM

2222 概率  $S\_Q\_EDU\_Q\_S = Q$  CSSCVIEEICHUCIEECI

2222 概率  $S\_Q\_EDU\_Q\_A = V + S$  CSSCVIEEICHUCIEECI

2222 概率  $S\_Q\_EDU\_Q\_O = E + S$  CSSCVIEEICHUCIEECI

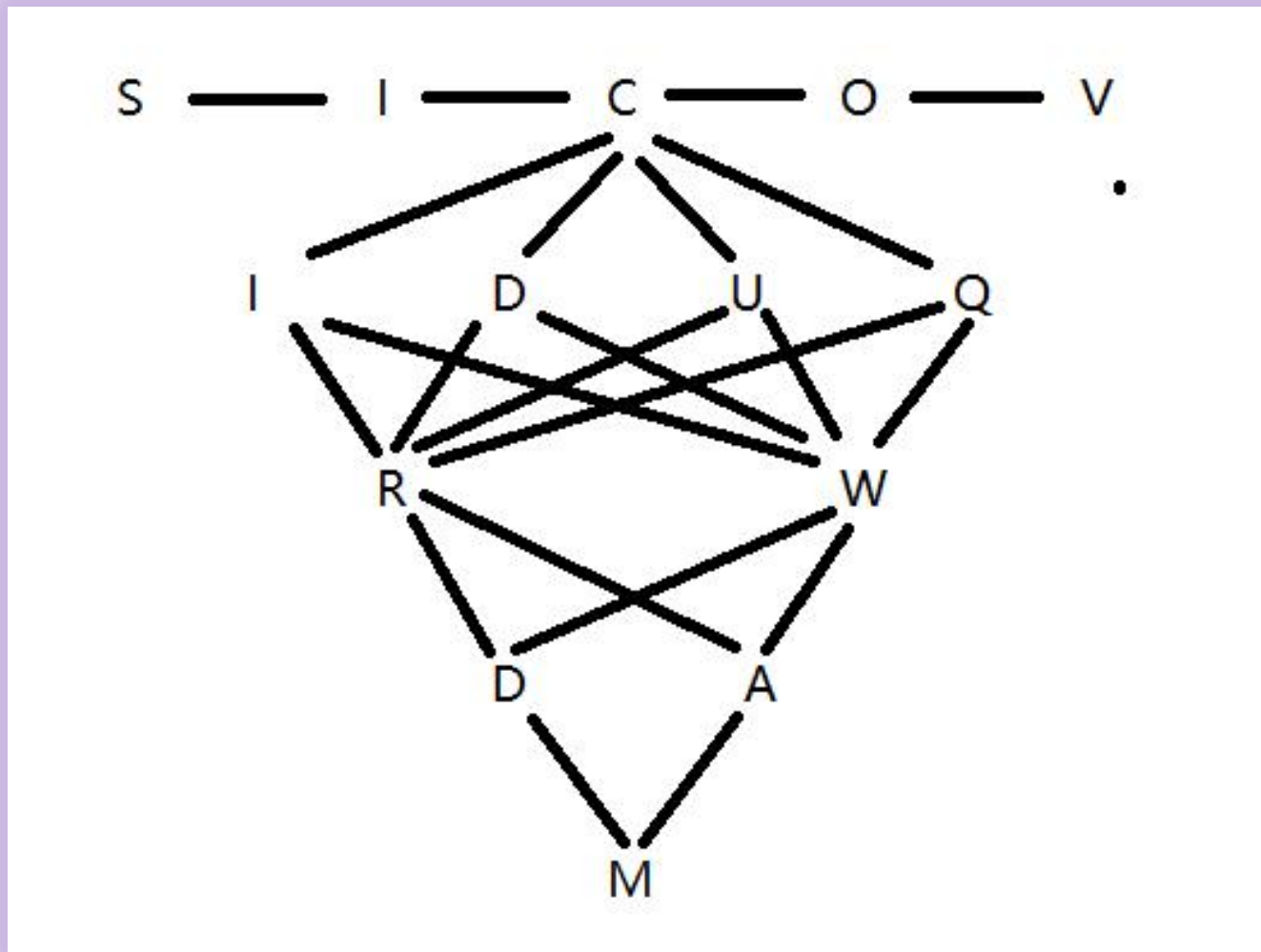
2222 概率  $S\_Q\_EDU\_Q\_P = E + C$

CSSCVIEEICHUCIEPCI

2222 概率  $S\_Q\_EDU\_Q\_M = C + S$

MSCVIEEICHUCIEPCI

通过元基的肽展公式,我设计了两种模式,一种的是增元模式,一种是降元模式,将肽化好的元基表进行比值变换,这里有很多变换算法,变换好后,我开始用肽展降元和增元公式函数进行变换,生成的 TI 概率钥匙用于数据识别. 这个识别的元基串用于服务器端计算. TD 钥匙则用于数据验证. 用于复检.



在数据的SICOV 模式流动下,将控制层进行元基编码和解码进行拆分,然后读写分离,分离后进行函数化在映射和直接寻址间,然后集成在管理系统中.我命名为元基加密组件.我准备分离出来写一个加密解密算法 API,用于各种不同的加密场景.我解释下这些字母:

**S 数据 I 输入 C 控制 O 输出 V 观测 I 增加 D 减少 U 改变 Q 查看 R 读 W 写 D 直接寻址 A 计算内存**

**M 管理器**

//函数实现

```
public static void main(String[] argv) {

    FullDNATokenPDI pDE_RNA_FullFormular= new FullDNATokenPDI();

    @SuppressWarnings("unused")

    String initonKeys= "EIU/0. 6/EDU/0. 4/si/0. 3/sq/0. 7/EIU/0. 5/EDU/0. 5/si/0. 632/sq/0. 368";

    pDE_RNA_FullFormular. key[0]= 0. 6;

    pDE_RNA_FullFormular. key[1]= 0. 3;

    pDE_RNA_FullFormular. key[2]= 0. 5;

    pDE_RNA_FullFormular. key[3]= 0. 632;

    pDE_RNA_FullFormular. text= "控制吸收";

    pDE_RNA_FullFormular. pdw= pDE_RNA_FullFormular. initonSect(pDE_RNA_FullFormular.

text);

    System. out. println("原文: " + pDE_RNA_FullFormular. text);

    //pDE_RNA_FullFormular. pdw= "字典保密: MSIOCUOCIPCUPCI"; String[] lock=

new String[12];

    lock[0] = "A"; lock[3] = "O"; lock[6] = "P"; lock[9] = "M";

    lock[1] = "V"; lock[4] = "E"; lock[7] = "C"; lock[10] = "S";

    lock[2] = "I"; lock[5] = "D"; lock[8] = "U"; lock[11] = "Q"; int i=

(int)(Math. random()* 12)% 12; pDE_RNA_FullFormular. lock+= lock[i];

    i= (int)(Math. random()* 12)% 12;

    pDE_RNA_FullFormular. lock+= lock[i]; i=

(int)(Math. random()* 12)% 12;

    pDE_RNA_FullFormular. lock+= lock[i]; i=

(int)(Math. random()* 12)% 12;

    pDE_RNA_FullFormular. lock+= lock[i];

    for(i= 0; i< pDE_RNA_FullFormular. pdw. length(); i++) {

        pDE_RNA_FullFormular. code+= pDE_RNA_FullFormular. lock + pDE_RNA_FullFormular.
```

```

pdw. charAt(i);

    }

    System.out.println("肽语 : "+ pDE_RNA_FullFormular.pdw); System.out.
println("肽锁 : "+ pDE_RNA_FullFormular.lock); System.out.println("散列肽语:
"+ pDE_RNA_FullFormular.code); pDE_RNA_FullFormular.bys= "0. 6/0. 3/0. 5/0.
632";

    System.out.println("静态密钥: "+ pDE_RNA_FullFormular.bys);

    pDE_RNA_FullFormular.doKeyPress(pDE_RNA_FullFormular.code, pDE_RNA_FullFormular,
false);

    System.out.println("静态肽展降元概率钥匙E: "+ pDE_RNA_FullFormular.pdedeKey); System.out.println("
静态肽展降元概率钥匙S: "+ pDE_RNA_FullFormular.pdedsKey); System.out.println("静态肽展降元: "+
pDE_RNA_FullFormular.pds);

    System.out.println("静态肽展增元概率钥匙E: "+ pDE_RNA_FullFormular.pdeieKey); System.out.println("
静态肽展增元概率钥匙S: "+ pDE_RNA_FullFormular.pdeisKey); System.out.println("静态肽展增元: "+
pDE_RNA_FullFormular.pde);

    pDE_RNA_FullFormular.time= "" + System.currentTimeMillis(); pDE_RNA_FullFormular.cacheId=
"ID" + Math.random() + ": " + Math.random(); System.out.println("时间: "+ pDE_RNA_FullFormular.
time);

    System.out.println("账号随机缓存字符串: "+ pDE_RNA_FullFormular.cacheId);

    pDE_RNA_FullFormular.session_key= pDE_RNA_FullFormular.pde; System.out.
println("Session: "+ pDE_RNA_FullFormular.session_key); System.out.

println("=====
=====");

    System.out.println("开始前序验证: ");

    System.out.println("开始Session解析: "+ pDE_RNA_FullFormular.session_key); System.out.println("
开始概率钥匙解析: "+ pDE_RNA_FullFormular.pdedeKey+
pDE_RNA_FullFormular.pdedsKey

```

---

```
+ pDE_RNA_FullFormular. pdeieKey+ pDE_RNA_FullFormular. pdeisKey);
```

```
FullDNATokenPDI pDE_RNA_FullFormular1= new FullDNATokenPDI(); pDE_RNA_FullFormular1.
pdedeKey= pDE_RNA_FullFormular. pdedeKey. toString(); pDE_RNA_FullFormular1. pdedsKey=
pDE_RNA_FullFormular. pdedsKey. toString(); pDE_RNA_FullFormular1. pdeieKey=
pDE_RNA_FullFormular. pdeieKey. toString(); pDE_RNA_FullFormular1. pdeisKey=
pDE_RNA_FullFormular. pdeisKey. toString();
```

```
pDE_RNA_FullFormular. doKeyUnPress(pDE_RNA_FullFormular. code,
pDE_RNA_FullFormular1, true);

System. out. println();

System. out. println("得到原降元元基DNA序列: "+ pDE_RNA_FullFormular. pds); System. out. println("
得到新降元元基DNA序列: "+ pDE_RNA_FullFormular1. pds); System. out. println("得到原元基DNA序
列: "+ pDE_RNA_FullFormular. pde); System. out. println("得到新元基DNA序列: "+
pDE_RNA_FullFormular1. pde); System. out. println("验证正确? ");
System. out. println(pDE_RNA_FullFormular. pde. equals(pDE_RNA_FullFormular1. pde)? "正确": "失败");
```

```
System. out. println("===== ");

System. out. println("开始后序验证: ");

FullDNATokenPDI pDE_RNA_FullFormular2= new FullDNATokenPDI(); pDE_RNA_FullFormular2.
pdeieKey= pDE_RNA_FullFormular. pdedeKey. toString(); pDE_RNA_FullFormular2. pdeisKey=
pDE_RNA_FullFormular. pdedsKey. toString(); pDE_RNA_FullFormular2. pdedeKey=
pDE_RNA_FullFormular. pdeieKey. toString(); pDE_RNA_FullFormular2. pdedsKey=
pDE_RNA_FullFormular. pdeisKey. toString(); System. out. println("准备计算元基DNA序列: "+
pDE_RNA_FullFormular1. pde); pDE_RNA_FullFormular2.
doSessionKeyUnPress(pDE_RNA_FullFormular1. pde,
```



pDE\_RNA\_FullFormular2, true); System.

out.println();

System.out.println("得到原续降元元基DNA序列: "+ pDE\_RNA\_FullFormular1.pds); System.out.println("得到后续降元元基DNA序列: "+ pDE\_RNA\_FullFormular2.pds); System.out.println("验证正确? ");

System.out.println(pDE\_RNA\_FullFormular1.pds.equals(pDE\_RNA\_FullFormular2.pds)? "正确": "失败");

System.out.println(pDE\_RNA\_FullFormular1.pds.equals(pDE\_RNA\_FullFormular2.pds)? "正确": "失败");

System.out.println("=====

==");

System.out.println("开始整序验证: ");

FullDNATokenPDI pDE\_RNA\_FullFormular3= new FullDNATokenPDI();

pDE\_RNA\_FullFormular3.pdeieKey= pDE\_RNA\_FullFormular.pdeieKey.toString();

pDE\_RNA\_FullFormular3.pdeisKey= pDE\_RNA\_FullFormular.pdeisKey.toString();

pDE\_RNA\_FullFormular3.pdedeKey= pDE\_RNA\_FullFormular.pdeieKey.toString();

pDE\_RNA\_FullFormular3.pdedeKey= pDE\_RNA\_FullFormular.pdeisKey.toString();

System.out.println("准备计算元基DNA序列: "+ pDE\_RNA\_FullFormular1.pde);

pDE\_RNA\_FullFormular3.doFullSessionKeyUnPress(pDE\_RNA\_FullFormular1.pde,

pDE\_RNA\_FullFormular3, true); System.

out.println();

System.out.println("得到原续降元元基DNA序列: "+ pDE\_RNA\_FullFormular1.pds); System.out.println("得到后续降元元基DNA序列: "+ pDE\_RNA\_FullFormular3.pds); System.out.println("验证正确? ");

System.out.println(pDE\_RNA\_FullFormular1.pds.equals(pDE\_RNA\_FullFormular3.pds)? "正确": "失败");

System.out.println(pDE\_RNA\_FullFormular1.pds.equals(pDE\_RNA\_FullFormular3.pds)? "正确": "失败");

---

```

System.out.println("准备整序计算元基DNA序列: "+ pDE_RNA_FullFormular1.pde); System.out.println("
准备整序计算元基DNA序列: "+ pDE_RNA_FullFormular3.pde);
System.out.println(pDE_RNA_FullFormular1.pde.equals(pDE_RNA_FullFormular3.pde)? "正确": "失败");

```

```

}

```

```

private void do_PDE_RNA_FullFormular_FullBack(Initon initon, FullDNATokenPDI
pDE_RNA_FullFormular3
    , boolean bYS)
{ Initon InitonPDE=
initon;

InitonLinkDNA initonLinkDNA= new InitonLinkDNA();

Initon InitonPDE1V= doIncrementV(InitonPDE, initonLinkDNA, pDE_RNA_FullFormular3);
Initon InitonPDE1E= doIncrementE(InitonPDE1V, initonLinkDNA, pDE_RNA_FullFormular3, bYS);
Initon InitonPDE1C= doIncrementC(InitonPDE1E, initonLinkDNA, pDE_RNA_FullFormular3);
Initon InitonPDE1S= doIncrementS(InitonPDE1C, initonLinkDNA, pDE_RNA_FullFormular3, bYS);

Initon InitonPDE1A= doIncrementA(InitonPDE1S, initonLinkDNA, pDE_RNA_FullFormular3);
Initon InitonPDE1O= doIncrementO(InitonPDE1A, initonLinkDNA, pDE_RNA_FullFormular3);
Initon InitonPDE1P= doIncrementP(InitonPDE1O, initonLinkDNA, pDE_RNA_FullFormular3);
Initon InitonPDE2= doIncrementM(InitonPDE1P, initonLinkDNA, pDE_RNA_FullFormular3);
while(InitonPDE2.hasNext()) {
    pDE_RNA_FullFormular3.pde+= InitonPDE2.getStore();
    InitonPDE2= InitonPDE2.next;
}
pDE_RNA_FullFormular3.pde+= InitonPDE2.getStore();
while(InitonPDE2.hasPrev())
{ InitonPDE2= InitonPDE2.
prev;
}

Initon InitonPDEM= doDecrementM(InitonPDE2, initonLinkDNA, pDE_RNA_FullFormular3);

```

```

Initon InitonPDEP= doDecrementP(InitonPDEM, initonLinkDNA, pDE_RNA_FullFormular3);
Initon InitonPDEO= doDecrementO(InitonPDEP, initonLinkDNA, pDE_RNA_FullFormular3);
Initon InitonPDEA= doDecrementA(InitonPDEO, initonLinkDNA, pDE_RNA_FullFormular3);

Initon InitonPDES= doDecrementS(InitonPDEA, initonLinkDNA, pDE_RNA_FullFormular3, bYS);
Initon InitonPDEC= doDecrementC(InitonPDES, initonLinkDNA, pDE_RNA_FullFormular3);
Initon InitonPDEE= doDecrementE(InitonPDEC, initonLinkDNA, pDE_RNA_FullFormular3, bYS);
Initon InitonPDE1= doDecrementV(InitonPDEE, initonLinkDNA, pDE_RNA_FullFormular3);
while(InitonPDE1. hasNext()) {
    pDE_RNA_FullFormular3. pds+= InitonPDE1. getStore();
    InitonPDE1= InitonPDE1. next;
}
pDE_RNA_FullFormular3. pds+= InitonPDE1. getStore();
while(InitonPDE1. hasPrev())
    { InitonPDE1= InitonPDE1.
        prev;
    }
}

```

//下面 9 页 plsql 的 orm 语法我也并入第二卷一开始是为了追梁壁荧写的, 追不到就算了.

```

package org. tinos. language. pletl; import
java. io. File;
import java. util. HashMap; import javax.
swing. JTextPane;
import OSI. OSU. MSQ. sets. stable. StableData; import OSI. OSU.
OEI. PVI. document. load. LoadFile;
import OSI. OSU. OVU. MVQ. GUI. nodeView. NodeShow; import OSI.
OSU. OVU. MVU. GUI. nodeEdit. LinkList; import OSI. OSU. OVU. MVU.
GUI. nodeEdit. LinkNode;
import OSI. OSU. PSQ. OEU. document. neroCell. BootNeroCell; public class
PLETLImpl implements PLETLIntef{

```

---

@Override

```
public boolean doNeroFlow(JTextPane rightBotJTextPane, NodeShow nodeView, LinkList first
    , String documentFlowAddress, HashMap<String, Object> inputMap) {
    //很好的将 《德塔 socket plsql 数据库》 和 《德塔 ETL》变成脑的记忆和计算中枢配合.
    //别急, 这个组合虽然没有自主意识, 但是已经形成了 VPCS 计算神经元的单株 锥形.
```

20200322 罗瑶光

```
try {
    String fileCurrentpath= documentFlowAddress; File file=
    new File(fileCurrentpath);
    if(!file. isFile()) {
        System. out. println(StableData. ATTENSION_RECHOICE); return false;
    }
    LinkNode needDeleteNode= first. first;
    while(needDeleteNode!= null) {
        first. first= first. deletNode(first. first, needDeleteNode. name
            , needDeleteNode. ID, needDeleteNode. primaryKey); if(null==
            needDeleteNode. next) {
            break;
        }
        needDeleteNode= needDeleteNode. next;
    }
    first. first= LoadFile. Load(first. first, nodeView, file, first); BootNeroCell.
    bootCell(first. first, rightBotJTextPane, null);
} catch(Exception loadE) { loadE.
    printStackTrace();
}
return true;
}
```

//大爱如此

---

```
}
```

```
//接口
```

```
package org. tinos. language. plorm;
```

```
import java. util. Map;
```

```
public interface PLORMInterf{
```

```
    public String getPLSQL();
```

```
    public void setPLSQL(String pLSQL);
```

```
    public PLORMInterf withTableCreate(String tableName); public
```

```
    PLORMInterf withTableDelete(String tableName); public PLORMInterf
```

```
    withTableInsert(String tableName); public PLORMInterf
```

```
    withTableUpdate(String tableName); public PLORMInterf
```

```
    withTableSelect(String tableName); public PLORMInterf getCulums();
```

```
    public PLORMInterf startAtRootDir(String rootAddress); public
```

```
    PLORMInterf withBaseName(String baseName); public PLORMInterf
```

```
    withCondition(String conditionType); public PLORMInterf let(String
```

```
    leftSet);
```

```
    public PLORMInterf lessThanAndEqualTo(String compareSet);
```

```
    public PLORMInterf equalTo(String compareSet); public
```

```
    PLORMInterf lessThan(String compareSet); public PLORMInterf
```

```
    greatThan(String compareSet);
```

```
    public PLORMInterf greatThanAndEqualTo(String compareSet);
```

```
    public PLORMInterf notEqualTo(String compareSet) ;
```

```
    public PLORMInterf in(String compareSet) ; public
```

```
    PLORMInterf notIn(String compareSet) ; public
```

```
    PLORMInterf equals(String compareSet);
```

```
    public PLORMInterf notEquals(String compareSet);
```

```
    public PLORMInterf innerJoinWithTable(String baseName, String tableName);
```

```
    public PLORMInterf withRelation(String relationType) ;
```

---

```

    public PLORMInterf as(String compareSet) ;
    public PLORMInterf upTo(String compareSet);
    public PLORMInterf withAggregation(String aggregationType);
    public PLORMInterf changeCulumnName(String newCulumnName, String oldCulumnName);
    public PLORMInterf withCulumnName(String culumnName, String dataType); public PLORMInterf
    withCulumnValue(String culumnName, String culumnValue); public PLORMInterf checkErrors(String
    string);
    public PLORMInterf fixErrors(String string);
    public PLORMInterf finalExec(boolean b) throws Exception;
    public Map<String, Object> returnAsMap();
    public PLORMInterf checkAndFixPlsqlGrammarErrors();
    public PLORMInterf checkAndFixSystemEnvironmentErrors();
}

```

//描述

```

package org. tinos. language. plorm; import
java. util. Map;

import org. ME. plsql. db. plsql. imp. ExecPLSQLImp; public class
PLORMImpl implements PLORMInterf{
    private String PLSQL= ""; private
    String[] PLSQLArray;
    private Map<String, Object> map; public
    String getPLSQL() {
        return PLSQL;
    }

    public void setPLSQL(String pLSQL) { PLSQL=
        pLSQL;
    }
}

```

---

```
public PLORMImpl startAtRootDir(String rootAddress) { PLSQL= Const.
    SET_ROOT+ Const. COLON+ rootAddress
        + Const. SEMICOLON;
    return this;
}

public PLORMImpl withBaseName(String baseName) {
    PLSQL+= Const. SEMICOLON+ Const. BASE_NAME+ Const. COLON
        + baseName;
    return this;
}

public PLORMImpl withTableSelect (String tableName) {
    PLSQL+= Const. SEMICOLON+ Const. TABLE_NAME+ Const. COLON
        + tableName
        + Const. COLON+ Const. SELECT; return
    this;
}

public PLORMImpl withTableCreate(String tableName) {
    PLSQL+= Const. SEMICOLON+ Const. TABLE_NAME+ Const. COLON
        + tableName
        + Const. COLON+ Const. CREATE; return this;
}

public PLORMImpl withTableDelete(String tableName) {
    PLSQL+= Const. SEMICOLON+ Const. TABLE_NAME+ Const. COLON
        + tableName
        + Const. COLON+ Const. DELETE; return
    this;
}

public PLORMImpl withTableInsert(String tableName) {
```

---

```

        PLSQL+= Const. SEMICOLON+ Const. TABLE_NAME+ Const. COLON
            + tableName
            + Const. COLON+ Const. INSERT; return this;
    }

    public PLORMImpl withTableUpdate(String tableName) {
        PLSQL+= Const. SEMICOLON+ Const. TABLE_NAME+ Const. COLON
            + tableName
            + Const. COLON+ Const. UPDATE; return this;
    }

    public PLORMImpl withCondition(String conditionType) {
        PLSQL+= Const. SEMICOLON+ Const. CONDITION+ Const. COLON
            + conditionType;
        return this;
    }

    public PLORMImpl let(String leftSet)
    { PLSQL+= Const. COLON+ leftSet; return
        this;
    }

    public PLORMImpl lessThanAndEqualTo(String compareSet) { PLSQL+= Const.
        LESS_THAN_AND_EQUAL_TO+ compareSet; return this;
    }

    public PLORMImpl equalTo(String compareSet) { PLSQL+=
        Const. EQUAL_TO+ compareSet; return this;
    }

    public PLORMImpl lessThan(String compareSet) { PLSQL+=
        Const. LESS_THAN+ compareSet;

```



---

```
        return this;
    }

    public PLORMImpl greatThan(String compareSet) { PLSQL+=
        Const. GREAT_THAN+ compareSet; return this;
    }

    public PLORMImpl greatThanAndEqualTo(String compareSet) { PLSQL+= Const.
        GREAT_THAN_AND_EQUAL_TO+compareSet; return this;
    }

    public PLORMImpl notEqualTo(String compareSet) { PLSQL+=
        Const. NOT_EQUAL_TO+ compareSet; return this;
    }

    public PLORMImpl in(String compareSet) { PLSQL+=
        Const. IN+ compareSet; return this;
    }

    public PLORMImpl notIn(String compareSet) { PLSQL+=
        Const. NOT_IN+ compareSet; return this;
    }

    public PLORMImpl equals(String compareSet) { PLSQL+=
        Const. EQUALS+ compareSet; return this;
    }

    public PLORMImpl notEquals(String compareSet) { PLSQL+=
        Const. NOT_EQUALS+ compareSet; return this;
    }
}
```

---

```

public PLORMImpl innerJoinWithTable(String baseName, String tableName) { PLSQL+= Const.
    SEMICOLON+ Const. JOIN+ Const. COLON+ baseName
        + Const. COLON+ tableName; return
    this;
}

public PLORMImpl withRelation(String relationType) {
    PLSQL+= Const. SEMICOLON+ Const. RELATION+ Const. COLON
        + relationType; return
    this;
}

public PLORMImpl as(String compareSet) { PLSQL+=
    Const. AS+ compareSet; return this;
}

public PLORMImpl upTo(String compareSet) { PLSQL+=
    Const. UP_TO+ compareSet; return this;
}

public PLORMImpl withAggregation(String aggregationType) { PLSQL+= Const.
    SEMICOLON+ Const. WITH_AGGREGATION
        + Const. COLON+ aggregationType; return this;
}

public PLORMImpl getCulums() {
    PLSQL+= Const. SEMICOLON+ Const. GET_CULUMNS;
    return this;
}

public PLORMImpl changeCulumnName(String newCulumnName, String oldCulumnName) { PLSQL+= Const.
    SEMICOLON+ Const. CHANGES_CULUMN_NAME+ Const. COLON
        + newCulumnName+ Const. COLON+ oldCulumnName;

```

---

```

        return this;
    }

    public PLORMImpl withColumnName(String columnName, String dataType) {
        PLSQL+= Const. SEMICOLON+ Const. CULUMN_NAME+ Const. COLON+ columnName
            + Const. COLON+ dataType; return this;
    }

    public PLORMImpl withColumnValue(String columnName, String columnValue) {
        PLSQL+= Const. SEMICOLON+ Const. CULUMN_VALUE+ Const. COLON+ columnName
            + Const. COLON+ columnValue; return
        this;
    }

    public PLORMInterf exec(boolean b) throws Exception { map=
        ExecPLSQLImp. ExecPLORM(this, true); return this;
    }

    @Override
    public PLORMInterf checkErrors(String string) { return this;
    }

    @Override
    public PLORMInterf fixErrors(String string) { return this;
    }

    @Override
    public PLORMInterf finalExec(boolean b) throws Exception { map=
        ExecPLSQLImp. ExecPLORM(this, true);
        return this;
    }

```

---

```
}
```

```
@Override
```

```
public Map<String, Object> returnAsMap() { return this.
```

```
    map;
```

```
}
```

```
@Override
```

```
public PLORMInterf checkAndFixPlsqlGrammarErrors() {
```

```
    //string to array
```

```
    this.PLSQLArray= PLSQL. split(Const. SEMICOLON);
```

```
    //条件检查 1 过滤 2 修改 3 语义检测
```

```
    //1
```

```
    for(int i= 1; i< PLSQLArray. length; i++) {
```

```
        //1. 1 过滤相同句型
```

```
        //1. 2 过滤无效字符
```

```
        //1. 3 过滤攻击代码
```

```
        if(PLSQLArray[i]. equalsIgnoreCase(PLSQLArray[i- 1]))
```

```
            { PLSQLArray[i]= "";
```

```
        }
```

```
        PLSQLArray[i]= PLSQLArray[i]. replaceAll(">+", ">"); PLSQLArray[i]=
```

```
        PLSQLArray[i]. replaceAll("<+", "<"); PLSQLArray[i]= PLSQLArray[i].
```

```
        replaceAll("\\!+", "!"); PLSQLArray[i]= PLSQLArray[i]. replaceAll("\\~+",
```

```
        "~"); PLSQLArray[i]= PLSQLArray[i]. replaceAll("\\@+", "@");
```

```
        PLSQLArray[i]= PLSQLArray[i]. replaceAll("\\&&+", "&&");
```

```
        PLSQLArray[i]= PLSQLArray[i]. replaceAll("\\\\+", "\\"); PLSQLArray[i]=
```

```
        PLSQLArray[i]. replaceAll("\\[+", "["); PLSQLArray[i]= PLSQLArray[i].
```

```
        replaceAll("\\]+", "]); PLSQLArray[i]= PLSQLArray[i]. replaceAll("\\: +", ":
```

```
        ");
```

---

```

        PLSQLArray[i]= PLSQLArray[i].replaceAll("\\s+", "");
    }

    //2
    //2.1 修改错误比较符号
    //2.2 修改错误语法关键字
    //2.3 修改错误标注符号
    //3
    //3.1 检测是否有关键字前后句段混乱
    //3.2 检测是否有关键字 格式 倒置
    //3.3 检测是否有关键字 句型 倒置
    //return

    String string= "";
    for(int i= 0; i< PLSQLArray. length; i++) {
        string+= PLSQLArray[i]+ Const. SEMICOLON;
    }

    PLSQL= string;
    return this;
}

@Override
public PLORMInterf checkAndFixSystemEnvironmentErrors() { return this;
}
}

```

---

//常量

**package** org. tinos. language. plorm;

**public class** Const{

**public final static** String *SET\_ROOT*= "setRoot";

**public final static** String *BASE\_NAME*= "baseName";

**public final static** String *TABLE\_NAME*= "tableName";

**public final static** String *SELECT*= "select";

**public final static** String *CREATE*= "create";

**public final static** String *DELETE*= "delete";

**public final static** String *INSERT*= "insert";

**public final static** String *UPDATE*= "update";

**public final static** String *CONDITION*= "condition";

**public final static** String *LESS\_THAN\_AND\_EQUAL\_TO*= "<=";

**public final static** String *EQUAL\_TO*= "==";

**public final static** String *LESS\_THAN*= "<";

**public final static** String *GREAT\_THAN*= ">";

**public final static** String *GREAT\_THAN\_AND\_EQUAL\_TO*= ">=";

**public final static** String *NOT\_EQUAL\_TO*= "<>";

**public final static** String *IN*= "|in|";

**public final static** String *NOT\_IN*= "|!in|";

**public final static** String *EQUALS*= "|equal|";

**public final static** String *NOT\_EQUALS*= "|!equal|";

**public final static** String *JOIN*= "join";

**public final static** String *RELATION*= "relation";

**public final static** String *AS*= "|as|";

**public final static** String *UP\_TO*= "|~|";

**public final static** String *WITH\_AGGREGATION*= "aggregation";

**public final static** String *GET\_COLUMNS*= "getCulums";

**public final static** String *CHANGES\_COLUMN\_NAME*= "changeCulumnName";

**public final static** String *COLUMN\_NAME*= "culumnName";

```

    public final static String COLUMN_VALUE= "columnValue";

    public final static String SEMICOLON= "; ";

    public final static String COLON= ": ";

}

```

//ORM 示例 create

```
package org. tinos. language. plorm;
```

```
public class Create{
```

//动机: 准备将下面的`plsql`翻译成`orm`, 省的养疗经[17]的query模式太固定, 上手修改麻烦.

```

//    String plsql= "setRoot: C: /DetaDB; ";

//    plsql+= "baseName: ZYY; ";

//    plsql+= "tableName: zybc: create; " +

//        "columnName: pk: ID: string; " +

//        "columnName: uk: 打分: string; " +

//        "columnName: uk: 中药名称: string; " +

//        "columnName: uk: 笔记原文: string; " +

//        "columnName: uk: 功效: string; " +

//        "columnName: uk: 风险规避: string; " +

//        "columnName: uk: 用量: string; " +

//        "columnName: uk: 性味: string; " +

//        "columnName: uk: 经脉: string; " +

//        "columnName: uk: 中医馆药理: string; " +

//        "columnName: uk: 经解: string; " +

//        "columnName: uk: 崇源: string; " +

//        "columnName: uk: 愚按: string; " +

//        "columnName: uk: 搭配: string; " +

//        "columnName: uk: 常见药: string; ";

```

//这个函数用于确定Deta PLSQL的Root like: 'c: /dsdsd' etc String

```
    PLSQL= "";
```

```
    public String startAtRootDir(String rootAddress) {
```

---

```

        PLSQL="setRoot: "+ rootAddress+ "; ";

        return PLSQL;
    }

    public String withBaseName(String baseName) { PLSQL+=
        "baseName: "+ baseName+ "; "; return PLSQL;
    }

    public String withTableCreate(String tableName) { PLSQL+=
        "tableName: "+ tableName+ ": create; "; return PLSQL;
    }

    public String withCulumnName(String columnName, String keyType, String dataFormat) { PLSQL+=
        "columnName: "+ keyType+ ": "+ columnName+ ": "+ dataFormat+ "; "; return PLSQL;
    }
}

```

---

## Dictionary PLSQL Standard

```

package OSV.VCQ.standard;

import java.io.IOException;
import java.util.ArrayList;
import java.util.List;
import java.util.Map;

import OSI.AOP.MEC.SIQ.plorm.PLORM_E;
import OSI.AOP.MEC.SIQ.plorm.PLORM_C;
//OSV PLSQL 研发, 思想, 设计: 罗瑶光 20210506
public class DictionaryPLSQLStandardDB{
    public List<String> txtToList() throws IOException{
        List<String> dic_list= new ArrayList<>();
        return dic_list;
    }

    public static Map<String, Object> bootORMReadDBInCommonWay(String tabKey) throws IOException{
        Map<String, Object> map= null;
        try {
            PLORM_C orm= new PLORM_E();
            map= orm.startAtRootDir("C:/DetaDB1").withBaseName("ZYY")
                .withTableSelect(tabKey).withCondition("or")
                .let("ID").lessThanAndEqualTo("3000")
                .checkAndFixPlsqlGrammarErrors();//准备完善plsql orm语言 的语法检查函数 和修复函数。
                .checkAndFixSystemEnvironmentErrors();//准备完善plsql orm语言 的系统环境检查函数和修复函
数。

                .finalE(true).returnAsMap();
            //map= org.plsql.db.plsql.imp.E_PLSQLImp.E_PLORM(orm, true);
        }catch(Exception e1) {
            //准备写回滚
            e1.printStackTrace();
        }
        return map;
    }
}

```



```

public static Map<String, Object> bootORMReadDBByRangeRowID(String rootPath, String baseName
    , boolean unTest, String tabKey, String RangeRowCount) throws IOException{
    Map<String, Object> map= null;
    try {
        PLORM_C orm= new PLORM_E();
        map= orm.startAtRootDir(rootPath).withBaseName(baseName)
            .withTableSelect(tabKey).withCondition("or")
            .let("ID").lessThanAndEqualTo(RangeRowCount)
            .checkAndFixPlsqlGrammarErrors()//准备完善plsql orm语言 的语法检查函数 和修复函数。
            .checkAndFixSystemEnvironmentErrors()//准备完善plsql orm语言 的系统环境检查函数和修复函
            .finalE(unTest).returnAsMap();
        //map= org.plsql.db.plsql.imp.E_PLSQLImp.E_PLORM(orm, true);
    }catch(Exception e1) {
        //准备写回滚
        e1.printStackTrace();
    }
    return map;
}

```

数。

//下面这些例子, 本人只是给大家一些更多的参考而已.

```

public static Map<String, Object> bootORMReadDBByLessThanAndEqualTo(String rootPath, String baseName
    , String conditionSubject, String conditionObject, boolean unTest, String tabKey) throws IOException{
    Map<String, Object> map= null;
    try {
        PLORM_C orm= new PLORM_E();
        map= orm.startAtRootDir(rootPath).withBaseName(baseName)
            .withTableSelect(tabKey).withCondition("or")
            .let(conditionSubject).lessThanAndEqualTo(conditionObject)
            .checkAndFixPlsqlGrammarErrors()//准备完善plsql orm语言 的语法检查函数 和修复函数。
            .checkAndFixSystemEnvironmentErrors()//准备完善plsql orm语言 的系统环境检查函数和修复函
            .finalE(unTest).returnAsMap();
        //map= org.plsql.db.plsql.imp.E_PLSQLImp.E_PLORM(orm, true);
    }catch(Exception e1) {
        //准备写回滚
        e1.printStackTrace();
    }
    return map;
}

```

数。

```

public static Map<String, Object> bootPLSQLReadDBInCommonWay(String tabKey) throws IOException{
    Map<String, Object> map= null;
    try {
        String plsql= "setRoot:C:/DetaDB1;" +
            "baseName:ZYY;" +
            "tableName:"+ tabKey +":select;" +
            "condition:or:ID|<=|3000;";
        map= ME.SM.OP.SM.AOP.MEC.SIQ.E.E_PLSQL_E.E_PLSQL(plsql, true);
    }catch(Exception e1) {
        //准备写回滚
        e1.printStackTrace();
    }
    return map;
}
}

```

```

package ME.SM.OP.SM.AOP.MEC.SIQ.E;
import java.util.Map;

import java.util.concurrent.ConcurrentHashMap;

```

---

```

import OSI.AOP.MEC.SIQ.plorm.PLORM_C;
public class E_PLSQL_E {
    public static Map<String, Object> E_PLSQL(String plsql, boolean mod) throws Exception {
        //working for here
        Map<String, Object> output = new ConcurrentHashMap<>();
        //1make container
        output.put("start", "0");
        output.put("countJoins", "0");
        //2make line
        String[] commands = plsql.replace(" ", "").replace("\n", "").split(";");
        String[] acknowledge = null;
        for(String command:commands) {
            acknowledge = command.split(":");
            if(acknowledge[0].equals("setRoot")) {
                PLSQLCommand_E.P_SetRoot(acknowledge, output);
            }
            if(acknowledge[0].equals("baseName")) {
                PLSQLCommand_E.P_BaseName(acknowledge, output);
            }
            if(acknowledge[0].equals("tableName")) {
                PLSQLCommand_E.P_TableName(acknowledge, output);
            }
            if(acknowledge[0].equals("culumnName")) {
                PLSQLCommand_E.P_ListNeedStart(acknowledge, output);
            }
            if(acknowledge[0].equals("changeCulumnName")) {
                PLSQLCommand_E.P_ListNeedStart(acknowledge, output);
            }
            if(acknowledge[0].equals("culumnValue")) {
                PLSQLCommand_E.P_ListNeedStart(acknowledge, output);
            }
            if(acknowledge[0].equals("join")) {
                PLSQLCommand_E.P_Join(acknowledge, output);
            }
            if(acknowledge[0].equals("condition")) {
                PLSQLCommand_E.P_ListNeedStart(acknowledge, output);
            }
            if(acknowledge[0].equals("relation")) {
                PLSQLCommand_E.P_ListNeedStart(acknowledge, output);
            }
            if(acknowledge[0].equals("aggregation")) {
                PLSQLCommand_E.P_ListNeedStart(acknowledge, output);
            }
            if(acknowledge[0].equals("getCulumns")) {
                PLSQLCommand_E.P_ListNeedStart(acknowledge, output);
            }
            output.put("newCommand", acknowledge[0]);
            PLSQLCommand_E.P_E(acknowledge, output, mod);
            output.put("lastCommand", output.get("newCommand"));
        }
        if(null!= acknowledge) {
            if(output.get("start").toString().equals("1")) {
                PLSQLCommand_E.P_E(acknowledge, output, mod);
            }
        }
        System.out.println("1");
        PLSQLCommand_E.P_Check(output.get("newCommand").toString(), output, mod);
        return output;
    }
}

public static Map<String, Object> E_PLORM(PLORM_C orm, boolean b) throws Exception {
    return E_PLSQL(orm.getPLSQL(), true);
}

```

---

```
}
}
```

```
package ME.SM.OP.SM.AOP.MEC.SIQ.E;
import java.io.BufferedReader;
import java.io.File;
import java.io.FileReader;
import java.io.IOException;
import java.math.BigDecimal;
import java.util.ArrayList;
import java.util.HashMap;
import java.util.Iterator;
import java.util.List;
import java.util.Map;
import java.util.concurrent.ConcurrentHashMap;

import MS.OP.SM.AOP.MEC.SIQ.cache.DetaDBBufferCache_M;
import OP.SM.AOP.MEC.SIQ.SM.reflection.Cell;
import OP.SM.AOP.MEC.SIQ.SM.reflection.Row;
import OP.SM.AOP.MEC.SIQ.SM.reflection.Table;
@SuppressWarnings({ "unused", "unchecked"})
public class P_ConditionPLSQL_XCDX_Cache extends P_ConditionPLSQL_XCDX {
    public static void P_Cache(String[] sets, List<Map<String, Object>> output
        , String tableName, String baseName, Map<String, Object> object) {
        Table table= DetaDBBufferCache_M.db.getBase(baseName).getTable(tableName);
        Iterator<String> iterator= table.getRows().keySet().iterator();
        int rowindex=0;
        while(iterator.hasNext()) {
            int count= rowindex++;
            String rowIndex= iterator.next();
            Row row= table.getRow(rowIndex);
            Cell cell=new Cell();
            cell.I_CellValue(rowIndex.replace("row", ""));
            row.putCell("Index", cell);
            if(sets[1].equalsIgnoreCase("<")||sets[1].equalsIgnoreCase("-lt")) {
                double rowCellFromBigDecimal= new BigDecimal(row.getCell(sets[0])
                    .getCellValue().toString()).doubleValue();
                if(rowCellFromBigDecimal< new BigDecimal(sets[2]).doubleValue()
                    && row.containsCell("is_delete_0")) {
                    if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {
                        output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(row));
                        Map<Integer, Boolean> recordRows= (Map<Integer, Boolean>)
object.get("recordRows");
                        recordRows.put(count, true);
                    }
                }
            }
            if(sets[1].equalsIgnoreCase("<=")||sets[1].equalsIgnoreCase("<=")
                ||sets[1].equalsIgnoreCase("-lte")) {
                String set= sets[0];
                Cell setCell= row.getCell(set);
                String cellString= setCell.getCellValue().toString();
                cellString=cellString.isEmpty()? "0": cellString;
                double rowCellFromBigDecimal = new BigDecimal(cellString).doubleValue();
                if(rowCellFromBigDecimal<= new BigDecimal(sets[2]).doubleValue()
                    && row.containsCell("is_delete_0")) {
                    if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {
                        output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(row));
                        Map<Integer, Boolean> recordRows= (Map<Integer, Boolean>)
object.get("recordRows");
```

---

```

        recordRows.put(count, true);
    }
}
}
if(sets[1].equalsIgnoreCase("==")||sets[1].equalsIgnoreCase("=")
    ||sets[1].equalsIgnoreCase("===")) {
    double rowCellFromBigDecimal = new BigDecimal(row.getCell(sets[0])
        .getCellValue().toString()).doubleValue();
    if(rowCellFromBigDecimal == new BigDecimal(sets[2]).doubleValue()
        && row.containsCell("is_delete_0")) {
        if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {
            output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(row));
            Map<Integer, Boolean> recordRows= (Map<Integer, Boolean>)
object.get("recordRows");

            recordRows.put(count, true);
        }
    }
}
if(sets[1].equalsIgnoreCase(">=")||sets[1].equalsIgnoreCase("=>")
    ||sets[1].equalsIgnoreCase("-gte")) {
    double rowCellFromBigDecimal = new BigDecimal(row.getCell(sets[0])
        .getCellValue().toString()).doubleValue();
    if(rowCellFromBigDecimal >= new BigDecimal(sets[2]).doubleValue()
        && row.containsCell("is_delete_0")) {
        if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {
            output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(row));
            Map<Integer, Boolean> recordRows= (Map<Integer, Boolean>)
object.get("recordRows");

            recordRows.put(count, true);
        }
    }
}
if(sets[1].equalsIgnoreCase(">")||sets[1].equalsIgnoreCase("-gt")) {
    double rowCellFromBigDecimal = new BigDecimal(row.getCell(sets[0])
        .getCellValue().toString()).doubleValue();
    if(rowCellFromBigDecimal > new BigDecimal(sets[2]).doubleValue()
        && row.containsCell("is_delete_0")) {
        if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {
            output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(row));
            Map<Integer, Boolean> recordRows= (Map<Integer, Boolean>)
object.get("recordRows");

            recordRows.put(count, true);
        }
    }
}
if(sets[1].equalsIgnoreCase("!=")||sets[1].equalsIgnoreCase("=!")) {
    double rowCellFromBigDecimal = new BigDecimal(row.getCell(sets[0])
        .getCellValue().toString()).doubleValue();
    if(rowCellFromBigDecimal != new BigDecimal(sets[2]).doubleValue()
        && row.containsCell("is_delete_0")) {
        if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {
            output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(row));
            Map<Integer, Boolean> recordRows= (Map<Integer, Boolean>)
object.get("recordRows");

            recordRows.put(count, true);
        }
    }
}
if(sets[1].equalsIgnoreCase("equal") && row.containsCell("is_delete_0")) {
    String rowCellFromString = row.getCell(sets[0]).getCellValue().toString();
    if(rowCellFromString.equalsIgnoreCase(sets[2])) {
        if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {

```



---

```

        public static void P_kernel(String temp, File readDBTableRowIndexColumnFile, File
readDBTableRowIndexFile
        , BufferedReader reader, String DBTableRowIndexPath, List<Map<String, Object>> output,
Row bufferRow
        , Map<String, Object> rowMap) throws IOException {
    String[] columnList = readDBTableRowIndexFile.list();
    NextFile:
        for(String column: columnList) {
            if(column.contains("is_delete")) {
                continue NextFile;
            }
            String DBTableCulumnIndexPath = readDBTableRowIndexFile + "/" + column;
            File readDBTableCulumnIndexPathFile = new File(DBTableCulumnIndexPath);

            Cell cell= new Cell();
            if (readDBTableCulumnIndexPathFile.isDirectory()) {
                //似乎被动了手脚, 20210405 罗瑶光重新检查
                reader = new BufferedReader(new
FileReader(readDBTableCulumnIndexPathFile + "/" + "value.lyg"));
                temp = "";
                String tempString;
                while ((tempString = reader.readLine()) != null) {
                    temp += tempString;
                }
                reader.close();
                rowMap.put(column, temp);
                cell.I_CellValue(temp);
                bufferRow.putCell(column, cell);
            } else {
                rowMap.put(column, null);
                cell.I_CellValue(null);
                bufferRow.putCell(column, cell);
            }
        }
        output.add(rowMap);
    }
}

```

---

```

package ME.SM.OP.SM.AOP.MEC.SIQ.E;
import java.io.BufferedReader;
import java.io.File;
import java.io.FileReader;
import java.io.IOException;
import java.math.BigDecimal;
import java.util.ArrayList;
import java.util.HashMap;
import java.util.Iterator;
import java.util.List;
import java.util.Map;
import java.util.concurrent.ConcurrentHashMap;

import MS.OP.SM.AOP.MEC.SIQ.cache.DetaDBBufferCache_M;
import OP.SM.AOP.MEC.SIQ.SM.reflection.Cell;
import OP.SM.AOP.MEC.SIQ.SM.reflection.Row;
import OP.SM.AOP.MEC.SIQ.SM.reflection.Table;
@SuppressWarnings({ "unused", "unchecked"})
public class P_ConditionPLSQL_XCDX_Map extends P_ConditionPLSQL_XCDX {
    //以后优化成统一对象输出, 不需要再转换。2019-1-15 tin
    public static Map<String, Object> rowToRowMap(Row row) {
        Map<String, Object> culumnMaps= new HashMap<>();
        Map<String, Object> rowMap= new HashMap<>();
    }
}

```

---

```

        Iterator<String> iterator= row.getCells().keySet().iterator();
        while(iterator.hasNext()) {
            String cellName = iterator.next();
            if(!cellName.contains("is_delete")) {
                Cell cell = row.getCell(cellName);
                Map<String, Object> culumnMap = new HashMap<>();
                culumnMap.put("culumnName", cellName);
                culumnMap.put("culumnValue", cell.getCellValue().toString());
                culumnMaps.put(cellName, culumnMap);
            }
        }
        rowMap.put("rowValue", culumnMaps);
        return rowMap;
    }

    public static void P_Map(String[] sets, List<Map<String, Object>> output, String dBTablePath) {
        List<Map<String, Object>> outputTemp = new ArrayList<>();
        Iterator<Map<String, Object>> iterator = output.iterator();
        int rowid = 0;
        while(iterator.hasNext()) {
            Map<String, Object> row = iterator.next();
            Map<String, Object> rowMap = new HashMap<>();
            if(sets[1].equalsIgnoreCase("<")||sets[1].equalsIgnoreCase("-lt")) {
                String rowCellFromString = ((Map<String, Object>)((Map<String,
Object>)(row.get("rowValue"))
                                .get(sets[0]))).get("culumnValue").toString();
                if(new BigDecimal(rowCellFromString).doubleValue() < new
BigDecimal(sets[2]).doubleValue()) {
                    outputTemp.add(row);
                }
            }
            if(sets[1].equalsIgnoreCase("<=")||sets[1].equalsIgnoreCase("<=")
||sets[1].equalsIgnoreCase("-lte")) {
                String rowCellFromString = ((Map<String, Object>)((Map<String,
Object>)(row.get("rowValue"))
                                .get(sets[0]))).get("culumnValue").toString();
                if(new BigDecimal(rowCellFromString).doubleValue() <= new
BigDecimal(sets[2]).doubleValue()) {
                    outputTemp.add(row);
                }
            }
            if(sets[1].equalsIgnoreCase("==")||sets[1].equalsIgnoreCase("=")||sets[1].equalsIgnoreCase("===")) {
                String rowCellFromString = ((Map<String, Object>)((Map<String,
Object>)(row.get("rowValue"))
                                .get(sets[0]))).get("culumnValue").toString();
                if(new BigDecimal(rowCellFromString).doubleValue() == new
BigDecimal(sets[2]).doubleValue()) {
                    outputTemp.add(row);
                }
            }
            if(sets[1].equalsIgnoreCase(">=")||sets[1].equalsIgnoreCase(">=")
||sets[1].equalsIgnoreCase("-gte")) {
                String rowCellFromString = ((Map<String, Object>)((Map<String,
Object>)(row.get("rowValue"))
                                .get(sets[0]))).get("culumnValue").toString();
                if(new BigDecimal(rowCellFromString).doubleValue() >= new
BigDecimal(sets[2]).doubleValue()) {
                    outputTemp.add(row);
                }
            }
            if(sets[1].equalsIgnoreCase(">")||sets[1].equalsIgnoreCase("-gt")) {
                String rowCellFromString = ((Map<String, Object>)((Map<String,

```

---

```

Object>)(row.get("rowValue")))
                                .get(sets[0])).get("columnValue").toString();
                                if(new BigDecimal(rowCellFromString).doubleValue() > new
BigDecimal(sets[2]).doubleValue()) {
                                    outputTemp.add(row);
                                }
                            }
                            if(sets[1].equalsIgnoreCase("!=")||sets[1].equalsIgnoreCase("!=")) {
                                String rowCellFromString = ((Map<String, Object>)((Map<String,
Object>)(row.get("rowValue"))))
                                    .get(sets[0])).get("columnValue").toString();
                                if(new BigDecimal(rowCellFromString).doubleValue() != new
BigDecimal(sets[2]).doubleValue()) {
                                    outputTemp.add(row);
                                }
                            }

                            if(sets[1].equalsIgnoreCase("equal")) {
                                String rowCellFromString = ((Map<String, Object>)((Map<String,
Object>)(row.get("rowValue"))))
                                    .get(sets[0])).get("columnValue").toString();
                                if(rowCellFromString.equalsIgnoreCase(sets[2])) {
                                    outputTemp.add(row);
                                }
                            }

                            if(sets[1].equalsIgnoreCase("!equal")) {
                                String rowCellFromString = ((Map<String, Object>)((Map<String,
Object>)(row.get("rowValue"))))
                                    .get(sets[0])).get("columnValue").toString();
                                if(!rowCellFromString.equalsIgnoreCase(sets[2])) {
                                    outputTemp.add(row);
                                }
                            }

                            if(sets[1].equalsIgnoreCase("in")) {
                                String rowCellFromString = ((Map<String, Object>)((Map<String,
Object>)(row.get("rowValue"))))
                                    .get(sets[0])).get("columnValue").toString();
                                String set = "," + sets[2] + ",";
                                if(set.contains("," + rowCellFromString + ",")){
                                    outputTemp.add(row);
                                }
                            }

                            if(sets[1].equalsIgnoreCase("!in")) {
                                String rowCellFromString = ((Map<String, Object>)((Map<String,
Object>)(row.get("rowValue"))))
                                    .get(sets[0])).get("columnValue").toString();
                                String set = "," + sets[2] + ",";
                                if(!set.contains("," + rowCellFromString + ",")){
                                    outputTemp.add(row);
                                }
                            }
                        }
                    }
                }
            }
        }
    }
}

```



---

```

package ME.SM.OP.SM.AOP.MEC.SIQ.E;
import java.io.BufferedReader;
import java.io.File;
import java.io.FileReader;
import java.io.IOException;
import java.math.BigDecimal;
import java.util.ArrayList;
import java.util.HashMap;
import java.util.Iterator;
import java.util.List;
import java.util.Map;
import java.util.concurrent.ConcurrentHashMap;

import MS.OP.SM.AOP.MEC.SIQ.cache.DetaDBBufferCache_M;
import OP.SM.AOP.MEC.SIQ.SM.reflection.Cell;
import OP.SM.AOP.MEC.SIQ.SM.reflection.Row;
import OP.SM.AOP.MEC.SIQ.SM.reflection.Table;
@SuppressWarnings({ "unused", "unchecked"})
public class P_ConditionPLSQL_XCDX_Table extends P_ConditionPLSQL_XCDX {
//plsql引擎函数获取表开始检查 罗瑶光 20210405 //奇怪了 这是一个没有读 缓存的plsql引擎,我准备对比下history
//object 指令堆栈
//output 数据行
public static void P_Table(String[] sets, List<Map<String, Object>> output
, String DBTablePath, Map<String, Object> object) throws IOException {
    String DBTableRowsPath= DBTablePath + "/rows";
    File fileDBTableRowsPath= new File(DBTableRowsPath);
    if (fileDBTableRowsPath.isDirectory()) {
        String[] rowList= fileDBTableRowsPath.list();
        int count= 0;
        NextRow:
        for(String row: rowList) {
            count++;
            Map<String, Object> rowMap= new HashMap<>();
            String DBTableRowIndexPath= DBTablePath + "/rows/" + row;
            File readDBTableRowIndexFile= new File(DBTableRowIndexPath);
            if (readDBTableRowIndexFile.isDirectory()) {
                String isDelete= DBTableRowIndexPath + "/is_delete_1";
                File isDeleteFile= new File(isDelete);
                if(isDeleteFile.exists()) {
                    continue NextRow;
                }
                String DBTableRowIndexCulumnPath= DBTableRowIndexPath + "/" + sets[0];
                File readDBTableRowIndexCulumnFile= new File(DBTableRowIndexCulumnPath);
                if(readDBTableRowIndexCulumnFile.isDirectory()) {
                    BufferedReader reader= new BufferedReader
                        (new FileReader(readDBTableRowIndexCulumnFile +
"/" + "value.lyg"));

                    String temp= "";
                    String tempString= "";
                    while ((tempString= reader.readLine())!= null) {
                        temp+= tempString;
                    }
                    reader.close();
                    if(temp.isEmpty()) {//增加一行id为空检查, 大家记得给 数据库的id加点值,我lyg的都是
空文件.

                                continue NextRow;
                            }
                    if(sets[1].equalsIgnoreCase("<")|| sets[1].equalsIgnoreCase("-lt")) {
                        if(new BigDecimal(temp.toString()).doubleValue() < new BigDecimal(sets[2].toString()).doubleValue()) {
                            if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {

```

---

```

        Row bufferRow= new Row();
        bufferRow.I_Cells(new ConcurrentHashMap<String, Cell>());
        P_ConditionPLSQL_XCDX_Kernel.P_kernel(row,
        readDBTableRowIndexColumnFile, readDBTableRowIndexFile, reader// 似乎被猫腻哥动了手脚, 我会将手里硬盘数据2年的数据等
        会全部验算
        , row, output, bufferRow, rowMap);output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(bufferRow));
        Map<Integer, Boolean> recordRows = (Map<Integer, Boolean>) object.get("recordRows"); recordRows.put(count, true);

    }

}

if(sets[1].equalsIgnoreCase("<=")||sets[1].equalsIgnoreCase("<=")||sets[1].equalsIgnoreCase("-lte")) {
    if(new BigDecimal(temp.toString()).doubleValue() <= new BigDecimal(sets[2].toString()).doubleValue()) {
        if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {
            Row bufferRow= new Row();
            bufferRow.I_Cells(new ConcurrentHashMap<String, Cell>());
            P_ConditionPLSQL_XCDX_Kernel.P_kernel(row, readDBTableRowIndexColumnFile
            , readDBTableRowIndexFile, reader// 似乎被猫腻哥动了手脚, 我会将手里硬盘数据2年的数据等会全部验算
            , row, output, bufferRow, rowMap);
            output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(bufferRow));
            Map<Integer, Boolean> recordRows = (Map<Integer, Boolean>) object.get("recordRows");
            recordRows.put(count, true);
        }
    }
}

if(sets[1].equalsIgnoreCase("=")||sets[1].equalsIgnoreCase("=")||sets[1].equalsIgnoreCase("===")) {
    if(new BigDecimal(temp.toString()).doubleValue() == new BigDecimal(sets[2].toString()).doubleValue()) {
        if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {
            Row bufferRow= new Row();
            bufferRow.I_Cells(new ConcurrentHashMap<String, Cell>());
            P_ConditionPLSQL_XCDX_Kernel.P_kernel(row, readDBTableRowIndexColumnFile
            , readDBTableRowIndexFile, reader// 似乎被猫腻哥动了手脚, 我会将手里硬盘数据2年的数据等会全部验算
            , row, output, bufferRow, rowMap);
            output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(bufferRow));
            Map<Integer, Boolean> recordRows = (Map<Integer, Boolean>) object.get("recordRows");
            recordRows.put(count, true);
        }
    }
}

if(sets[1].equalsIgnoreCase(">=")||sets[1].equalsIgnoreCase(">=") ||sets[1].equalsIgnoreCase("-gte")) {
    if(new BigDecimal(temp.toString()).doubleValue() >= new BigDecimal(sets[2].toString()).doubleValue()) {
        if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {
            Row bufferRow= new Row();
            bufferRow.I_Cells(new ConcurrentHashMap<String, Cell>());
            P_ConditionPLSQL_XCDX_Kernel.P_kernel(row, readDBTableRowIndexColumnFile
            , readDBTableRowIndexFile, reader// 似乎被猫腻哥动了手脚, 我会将手里硬盘数据2年的数据等会全部验算
            , row, output, bufferRow, rowMap);
            output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(bufferRow));
            Map<Integer, Boolean> recordRows = (Map<Integer, Boolean>) object.get("recordRows");
            recordRows.put(count, true);
        }
    }
}

if(sets[1].equalsIgnoreCase(">")||sets[1].equalsIgnoreCase("-gt")) {
    if(new BigDecimal(temp.toString()).doubleValue() > new BigDecimal(sets[2].toString()).doubleValue()) {
        if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {
            Row bufferRow= new Row();
            bufferRow.I_Cells(new ConcurrentHashMap<String, Cell>());
            P_ConditionPLSQL_XCDX_Kernel.P_kernel(row, readDBTableRowIndexColumnFile
            , readDBTableRowIndexFile, reader// 似乎被猫腻哥动了手脚, 我会将手里硬盘数据2年的数据等会全部验算

```

```

        , row, output, bufferRow, rowMap);
    output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(bufferRow));
    Map<Integer, Boolean> recordRows = (Map<Integer, Boolean>) object.get("recordRows");
    recordRows.put(count, true);
}
}
}
if(sets[1].equalsIgnoreCase("!=")||sets[1].equalsIgnoreCase("!=")) {
    if(new BigDecimal(temp.toString()).doubleValue()!= new BigDecimal(sets[2].toString()).doubleValue()) {
        if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {
            Row bufferRow= new Row();
            bufferRow.I_Cells(new ConcurrentHashMap<String, Cell>());
            P_ConditionPLSQL_XCDX_Kernel.P_kernel(row, readDBTableRowIndexCulumnFile
, readDBTableRowIndexFile, reader// 似乎被猫腻哥动了手脚, 我会将手里硬盘数据2年的数据等会全部验算
, row, output, bufferRow, rowMap);
            output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(bufferRow));
            Map<Integer, Boolean> recordRows = (Map<Integer, Boolean>) object.get("recordRows");
            recordRows.put(count, true);
        }
    }
}
}
if(sets[1].equalsIgnoreCase("equal")) {
    String rowCellFromString = temp.toString();
    if(rowCellFromString.equalsIgnoreCase(sets[2].toString())) {
        if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {
            Row bufferRow= new Row();
            bufferRow.I_Cells(new ConcurrentHashMap<String, Cell>());
            P_ConditionPLSQL_XCDX_Kernel.P_kernel(row, readDBTableRowIndexCulumnFile
, readDBTableRowIndexFile, reader// 似乎被猫腻哥动了手脚, 我会将手里硬盘数据2年的数据等会全部验算
, row, output, bufferRow, rowMap);
            output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(bufferRow));
            Map<Integer, Boolean> recordRows = (Map<Integer, Boolean>) object.get("recordRows");
            recordRows.put(count, true);
        }
    }
}
}
if(sets[1].equalsIgnoreCase("!equal")) {
    String rowCellFromString = temp.toString();
    if(!rowCellFromString.equalsIgnoreCase(sets[2].toString())) {
        if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {
            Row bufferRow= new Row();
            bufferRow.I_Cells(new ConcurrentHashMap<String, Cell>());
            P_ConditionPLSQL_XCDX_Kernel.P_kernel(row, readDBTableRowIndexCulumnFile
, readDBTableRowIndexFile, reader// 似乎被猫腻哥动了手脚, 我会将手里硬盘数据2年的数据等会全部验算
, row, output, bufferRow, rowMap);
            output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(bufferRow));
            Map<Integer, Boolean> recordRows = (Map<Integer, Boolean>) object.get("recordRows");
            recordRows.put(count, true);
        }
    }
}
}
if(sets[1].equalsIgnoreCase("in")) {
    String rowCellFromString = temp.toString();
    String set = "," + sets[2] + ",";
    if(set.contains("," + rowCellFromString + ",")) {
        if(!((Map<Integer, Boolean>)(object.get("recordRows"))).containsKey(count)) {
            Row bufferRow= new Row();
            bufferRow.I_Cells(new ConcurrentHashMap<String, Cell>());
            P_ConditionPLSQL_XCDX_Kernel.P_kernel(row, readDBTableRowIndexCulumnFile
, readDBTableRowIndexFile, reader// 似乎被猫腻哥动了手脚, 我会将手里硬盘数据2年的数据等会全部验算
, row, output, bufferRow, rowMap);
            output.add(P_ConditionPLSQL_XCDX_Map.rowToRowMap(bufferRow));

```



## Deta DNA Index & PLSQL ORM 增删改查 Demo

```
package OSV.ESD.standard;
import java.io.IOException;
import java.util.ArrayList;
import java.util.List;
import java.util.Map;
//import OSI.AOP.MEC.SIQ.plorm.PLORMImpl;
//import OSI.AOP.MEC.SIQ.plorm.PLORMInterf;
//import OSI.OPE.SI.SD.SU.SQ.ASU.OSU.PSU.MSU.AVQ.ASQ.ASU.MPE.procedure.pde.FullDNATokenPDI;
//OSV PLSQL 研发, 思想, 设计: 罗瑶光 20210511
//表删除操作的例子.
public class D_CellStandard{
    public List<String> txtToList() throws IOException{
        List<String> dic_list= new ArrayList<>();
        return dic_list;
    }
    // delete samples
    // tableName:test:delete;
    // condition:or:testCulumn1|<|20:testCulumn2|==|fire;
    // condition:and:testCulumn1|>|100:testCulumn2|==|fire;
    public static Map<String, Object> DeleteCellORM(String rootPath, String baseName
        , String tabKey
        , Map<String, String> rowCells, Boolean initonEncrypt) throws IOException{
        // PLORMInterf orm= new PLORMImpl();
        // orm= orm.startAtRootDir(rootPath)
        //     .withBaseName(baseName)
        //     .withTableInsert(tabKey);
        // condition 不好规范, 稍后. 建议写针对性 delete 语句
        // condition:or:testCulumn1|<|20:testCulumn2|==|fire;
        // condition:and:testCulumn1|>|100:testCulumn2|==|fire;
        return null;
    }
    public static Map<String, Object> DeleteCellPLSQL(String rootPath, String baseName
        , String tabKey
        , Map<String, String> rowCells, Boolean initonEncrypt) throws IOException{
        Map<String, Object> map= null;
        try {
            // String plsqli= "setRoot:"+ rootPath+ ";";
            // plsqli+= "baseName:"+ baseName+ ";";
            // plsqli+= "tableName:"+ tabKey+ ":delete;";
            // condition 不好规范, 稍后. 建议写针对性 delete 语句
            // condition:or:testCulumn1|<|20:testCulumn2|==|fire;
            // condition:and:testCulumn1|>|100:testCulumn2|==|fire;
            // map= OSI.OPE.ME.SM.OP.SM.AOP.MEC.SIQ.imp.E_PLSQLImp.E_PLSQL(plsqli, true);
        }catch(Exception e1) {
            //准备写回滚
        }
    }
}
```

```

        e1.printStackTrace();
    }
    return map;
}
}

```

---

```

package OSV.ESD.standard;
import java.io.IOException;
import java.util.ArrayList;
import java.util.List;
import java.util.Map;
import OSI.AOP.MEC.SIQ.plorm.PLORM_E;
import OSI.AOP.MEC.SIQ.plorm.PLORM_C;
//import OSI.OPE.SI.SD.SU.SQ.ASU.OSU.PSU.MSU.AVQ.ASQ.ASU.MPE.procedure.pde.FullIDNATokenPDI;
//OSV PLSQL 研发, 思想, 设计: 罗瑶光 20210511
//表剔除操作的例子.
public class D_Standard{
    public List<String> txtToList() throws IOException{
        List<String> dic_list= new ArrayList<>();
        return dic_list;
    }
    // drop samples
    // tableName:test:drop;
    public static Map<String, Object> DropCellORM(String rootPath, String baseName
        , String tabKey) throws Exception{
        PLORM_C orm= new PLORM_E();
        orm= orm.startAtRootDir(rootPath)
            .withBaseName(baseName)
            .withTableDrop(tabKey)
            .checkAndFixPlsqlGrammarErrors()
            //准备完善 plsql orm 语言 的语法检查函数 和修复函数。
            .checkAndFixSystemEnvironmentErrors()
            //准备完善 plsql orm 语言 的系统环境检查函数和修复函数。
            .finalE(true);
        return null;
    }
    public static Map<String, Object> DropCellPLSQL(String rootPath, String baseName
        , String tabKey) throws IOException{
        Map<String, Object> map= null;
        try {
            String plsql= "setRoot:"+ rootPath+ ";";
            plsql+= "baseName:"+ baseName+ ";";
            plsql+= "tableName:"+ tabKey+ ":drop;";
            map= ME.SM.OP.SM.AOP.MEC.SIQ.E.E_PLSQL_E.E_PLSQL(plsql, true);
        }catch(Exception e1) {
            //准备写回滚
            e1.printStackTrace();
        }
    }
}

```

```

    }
    return map;
}
}

```

---

```

package OSV.ESI.standard;
import java.io.IOException;
import java.util.ArrayList;
import java.util.Iterator;
import java.util.List;
import java.util.Map;
import OSI.AOP.MEC.SIQ.plorm.PLORM_E;
import OSI.AOP.MEC.SIQ.plorm.PLORM_C;
import OSI.OPE.SI.SD.SU.SQ.ASU.OSU.PSU.MSU.AVQ.ASQ.ASU.MPE.procedure.pde.FullDNATokenPDI;
//OSV PLSQL 研发, 思想, 设计: 罗瑶光 20210510
//插入一行完整的数据.
//我没有测试, 现在养疗经还用不到, 优先级稍后.
//稍后会考虑不完整的插入情况 赋值 empty 或 null.
public class IU_CellStandard{
    public List<String> txtToList() throws IOException{
        List<String> dic_list= new ArrayList<>();
        return dic_list;
    }
    public static Map<String, Object> InsertCellORM(String rootPath, String baseName
        , String tabKey
        , Map<String, String> rowCells, Boolean initonEncrypt) throws IOException{
        Iterator<String> iterators= rowCells.keySet().iterator();
        Map<String, Object> map= null;
        try {
            PLORM_C orm= new PLORM_E();
            orm= orm.startAtRootDir(rootPath)
                .withBaseName(baseName)
                .withTableInsert(tabKey);
            while(iterators.hasNext()) {
                String string= iterators.next();
                if(initonEncrypt) {
                    orm= orm.withCulumnValue(string
                        , new FullDNATokenPDI().initonSect
                        (rowCells.get(string).replace(":", "@Tin@")));
                }else {
                    orm= orm.withCulumnValue(string, rowCells.get(string).replace(":"
                        , "@Tin@"));
                    //我稍后会思考: 符号怎么进行实体化,省的要变通配符代替.
                }
            }
            orm.checkAndFixPlsqlGrammarErrors()
            //准备完善 plsql orm 语言 的语法检查函数 和修复函数。

```

```

        .checkAndFixSystemEnvironmentErrors()
        //准备完善 plsql orm 语言 的系统环境检查函数和修复函数。
        .finalE(true);
        //map= org.plsql.db.plsql.imp.E_PLSQLImp.E_PLORM(orm, true);
    }catch(Exception e1) {
        //准备写回滚
        e1.printStackTrace();
    }
    return map;
}

public static Map<String, Object> InsertCellPLSQL(String rootPath, String baseName
    , String tabKey
    , Map<String, String> rowCells, Boolean initonEncrypt) throws IOException{
    Iterator<String> iterators= rowCells.keySet().iterator();
    Map<String, Object> map= null;
    try {
        String plsql= "setRoot:"+ rootPath+ ";";
        plsql+= "baseName:"+ baseName+ ";";
        plsql+= "tableName:"+ tabKey+ ":insert;";
        while(iterators.hasNext()) {
            String string= iterators.next();
            if(initonEncrypt) {
                plsql+= "columnValue:"+ string+ ":"
                    + new FullDNATokenPDI().initonSect
                    (rowCells.get(string).replace(":", "@Tin@"))+ ";";
            }else {
                plsql+= "columnValue:"+ string+ ":"
                    + rowCells.get(string).replace(":", "@Tin@")+ ";";
            }
        }
        map= ME.SM.OP.SM.AOP.MEC.SIQ.E.E_PLSQL_E.E_PLSQL(plsql, true);
    }catch(Exception e1) {
        //准备写回滚
        e1.printStackTrace();
    }
    return map;
}
}

```

---

```

package OSV.ESU.standard;
import java.io.IOException;
import java.util.ArrayList;
import java.util.List;
import java.util.Map;
import OSI.AOP.MEC.SIQ.plorm.PLORM_E;
import OSI.AOP.MEC.SIQ.plorm.PLORM_C;
//OSV PLSQL 研发, 思想, 设计: 罗瑶光 20210507

```



//选中一个 cell 进行改变

```
public class U_CellStandard{
```

```
    public List<String> txtToList() throws IOException{
```

```
        List<String> dic_list= new ArrayList<>();
```

```
        return dic_list;
```

```
    }
```

// 稍后我会把下面这个 2 个模式变成 ORM UNSQL, 函数是很好写, 但是我在思考

//or and 太多 怎么进行规范传参..

// 我的动机很简单, 就是养疗经数据表可以直接数据库操作修改.

// 见 DNA 编码与计算第一卷 308 页

// update samples

// tableName:test:update;

// condition:or:testCulumn1|<|20:testCulumn2|==|fire;

// condition:and:testCulumn1|>|100:testCulumn2|==|fire;

// culumnValue:date0:19850525;

// culumnValue:date1:19850526;

//

// update samples tableName:test:update;

// condition:or:testCulumn1|<|20:testCulumn2|==|fire;

// condition:and:testCulumn1|>|100:testCulumn2|==|fire;

// join:backend:utest;

// condition:and:uCulumn3|<|20;

// relation:and:testCulumn1|==|uCulumn1:testCulumn2|!=|uCulumn2;

// culumnValue:date0:19850525;

// culumnValue:date1:19850526;

```
public static Map<String, Object> UpdateCellORM(String tabKey, String rowId
```

```
    , String cellName
```

```
    , String cellValue) throws IOException{
```

```
    Map<String, Object> map= null;
```

```
    try {
```

```
        PLORM_C orm= new PLORM_E();
```

```
        orm.startAtRootDir("C:/DetaDB1").withBaseName("ZYY")
```

```
        .withTableUpdate(tabKey).withCondition("or")
```

```
        .let("ID").equalTo(rowId)
```

```
        .checkAndFixPlsqlGrammarErrors()//准备完善 plsql orm 语言 的语法检查函数 和修复函数。
```

```
        .checkAndFixSystemEnvironmentErrors()//准备完善 plsql orm 语言 的系统环境检查函数和修复函数。
```

```
        .withCulumnValue(cellName, cellValue)
```

```
        .finalE(true);
```

```
        //map= org.plsql.db.plsql.imp.E_PLSQLImp.E_PLORM(orm, true);
```

```
    }catch(Exception e1) {
```

```
        //准备写回滚
```

```
        e1.printStackTrace();
```

```
    }
```

```
    return map;
```

```
}
```

```
public static Map<String, Object> UpdateCellORMByRowId(String rootPath
```

```
    , String baseName, boolean unTest
```

```
    , String tabKey, String rowId, String cellName, String cellValue)
```

```

        throws IOException{
Map<String, Object> map= null;
try {
    PLORM_C orm= new PLORM_E();
    orm.startAtRootDir(rootPath).withBaseName(baseName)
    .withTableUpdate(tabKey).withCondition("or")
    .let("ID").equalTo(rowId)
    .checkAndFixPlsqlGrammarErrors();//准备完善 plsql orm 语言 的语法检查函数 和修复函数。
    .checkAndFixSystemEnvironmentErrors();//准备完善 plsql orm 语言 的系统环境检查函数和修复函数。
    .withCulumnValue(cellName, cellValue)
    .finalE(unTest);
    //map= org.plsql.db.plsql.imp.E_PLSQLImp.E_PLORM(orm, true);
} catch (Exception e1) {
    //准备写回滚
    e1.printStackTrace();
}
return map;
}

public static Map<String, Object> UpdateCellORMByEquals(String rootPath
    , String baseName
    , boolean unTest, String tabKey, String conditionSubject
    , String conditionObject
    , String cellName, String cellValue) throws IOException{
Map<String, Object> map= null;
try {
    PLORM_C orm= new PLORM_E();
    orm.startAtRootDir(rootPath).withBaseName(baseName)
    .withTableUpdate(tabKey).withCondition("or")
    .let(conditionSubject).equalTo(conditionObject)
    .checkAndFixPlsqlGrammarErrors();//准备完善 plsql orm 语言 的语法检查函数 和修复函数。
    .checkAndFixSystemEnvironmentErrors();//准备完善 plsql orm 语言 的系统环境检查函数和修复函数。
    .withCulumnValue(cellName, cellValue)
    .finalE(unTest);
    //map= org.plsql.db.plsql.imp.E_PLSQLImp.E_PLORM(orm, true);
} catch (Exception e1) {
    //准备写回滚
    e1.printStackTrace();
}
return map;
}

// update samples tableName:test:update;
// condition:or:testCulumn1|<|20:testCulumn2|==|fire;
// condition:and:testCulumn1|>|100:testCulumn2|==|fire;
// join:backend:utest;
// condition:and:uCulumn3|<|20;
// relation:and:testCulumn1|==|uCulumn1:testCulumn2|!=|uCulumn2;
// culumnValue:date0:19850525;
// culumnValue:date1:19850526;
// String plsql= "setRoot:C:/DetaDB1;" +

```

```
// "baseName:ZYY;" +
// "tableName:"+ tabKey+ ":update;" +
// "condition:or:ID|==|rowId;" +
// "columnValue:columnName:cellValue;";
// condition:"我似乎没有设计 rowid columnid 的数字选项 函数,稍后补充下":19850526;
// condition 可以用 uid in 或者 uid== 来实现.这样会导致计算变慢,所以 设计 rowid columnid 的数字比较选项
//是有必要的.
```

```
public static Map<String, Object> UpdateCellPLSQL(String tabKey, String rowId
    , String cellName
    , String cellValue) throws IOException{
    Map<String, Object> map= null;
    try {
        String plsqli= "setRoot:C:/DetaDB1;" +
            "baseName:ZYY;" +
            "tableName:"+ tabKey+ ":update;" +
            "condition:or:ID|==|" + rowId+ ";" +
            "columnValue:"+cellName+ ":" + cellValue+ ";";
        map= ME.SM.OP.SM.AOP.MEC.SIQ.E.E_PLSQL_E.E_PLSQL(plsqli, true);
    }catch(Exception e1) {
        //准备写回滚
        e1.printStackTrace();
    }
    return map;
}
}
```