Collation Adjustments for DUCET

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The UTC has just agreed (UTC 131-A27) to add the following contractions to DUCET:

```
OFB2 OF71 ; [.255A.0020.0002.0FB2][.2570.0020.0002.0F71]
OFB3 OF71 ; [.255D.0020.0002.0FB3][.2570.0020.0002.0F71]
```

I believe these may be expressed as a tailoring

```
& \u0fb2\u0f71 = \u0fb2\u0f71
& \u0fb3\u0f71 = \u0fb3\u0f71
```

I have tested the collation of a range of Tibetan vowels and found that these additions changed the collation orders of some perfectly valid Tibetan sequences:

```
0FBC 0F72 and 0FB2 0F71 0F72
0FBC 0F74 and 0FB2 0F71 0F74
0FB3 0F72 and 0FB3 0F71 0F72
0FB3 0F74 and 0FB3 0F71 0F74
```

Disappointingly, neither of the tests defined by CollationTest_NON_IGNORABLE.txt and CollationTest_SHIFTED.txt detected these transpositions.

I therefore propose that another 4 tailorings be added to undo this accidental re-ordering and re-establish their previous ordering:

```
& \u0fb2\u0f71\u0f72 = \u0fb2\u0f71\u0f72
& \u0fb2\u0f71\u0f74 = \u0fb2\u0f71\u0f74
& \u0fb3\u0f71\u0f72 = \u0fb3\u0f71\u0f72
& \u0fb3\u0f71\u0f74 = \u0fb3\u0f71\u0f74
```

These should be applied before the two new tailorings.

The intention is that if no other changes were made to DUCET, these should add the following entries to DUCET:

```
        0FB2
        0F73
        ; [.255A.0020.0002.0FB2][.2572.0020.0002.0F73]

        0FB2
        0F71
        0F72
        ; [.255A.0020.0002.0FB2][.2572.0020.0002.0F73]

        0FB2
        0F75
        ; [.255A.0020.0002.0FB2][.2576.0020.0002.0F75]

        0FB3
        0F71
        0F74
        ; [.255A.0020.0002.0FB2][.2576.0020.0002.0F75]

        0FB3
        0F73
        ; [.255D.0020.0002.0FB3][.2572.0020.0002.0F73]

        0FB3
        0F71
        0F72
        ; [.255D.0020.0002.0FB3][.2572.0020.0002.0F73]

        0FB3
        0F75
        ; [.255D.0020.0002.0FB3][.2576.0020.0002.0F75]

        0FB3
        0F71
        0F74
        ; [.255D.0020.0002.0FB3][.2576.0020.0002.0F75]
```

In particular, any collation-based search for U+0F73 should continue to find it in the sequence <U+0FB2, U+0F73>.