gadA

iSSP

Known Region

Unknown Region

Sequence Name
Consensus
4 sequences
Ref.seq
AS1.seq
AS2.seq
AS3.seq

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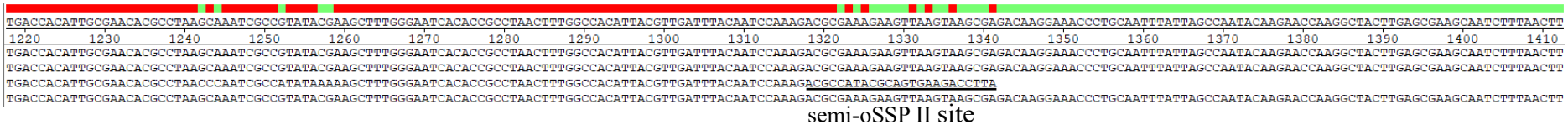
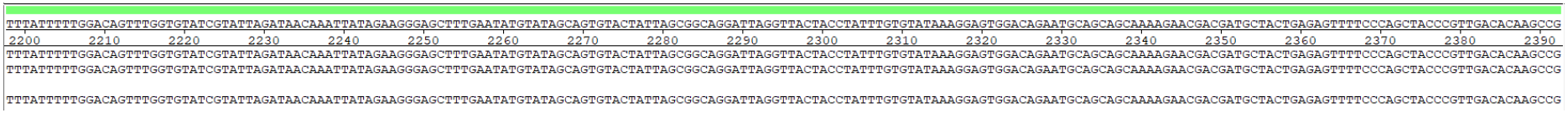
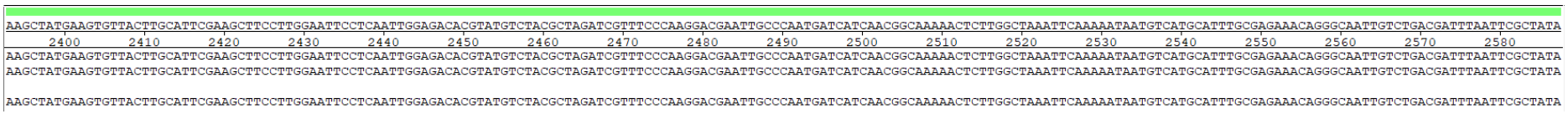
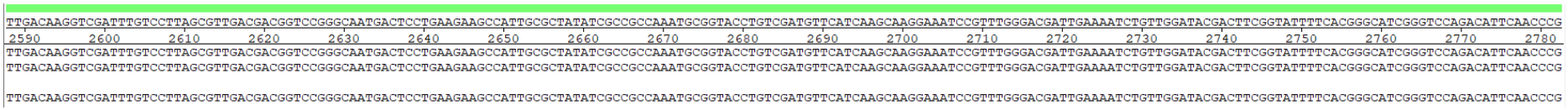
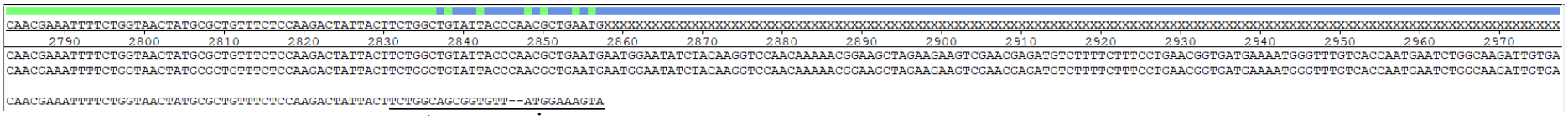


semi-oSSP II site









semi-oSSP III site

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7750      7760      7770      7780      7790      7800      7810      7820      7830      7840      7850      7860      7870      7880      7890      7900      7910      7920      7930      7
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semi-oSSP I site

gadR

iSSP

Known Region

Unknown Region

Sequence Name
Consensus
4 Sequences
Ref.seq
RS1.seq
RS2.seq
RS3.seq

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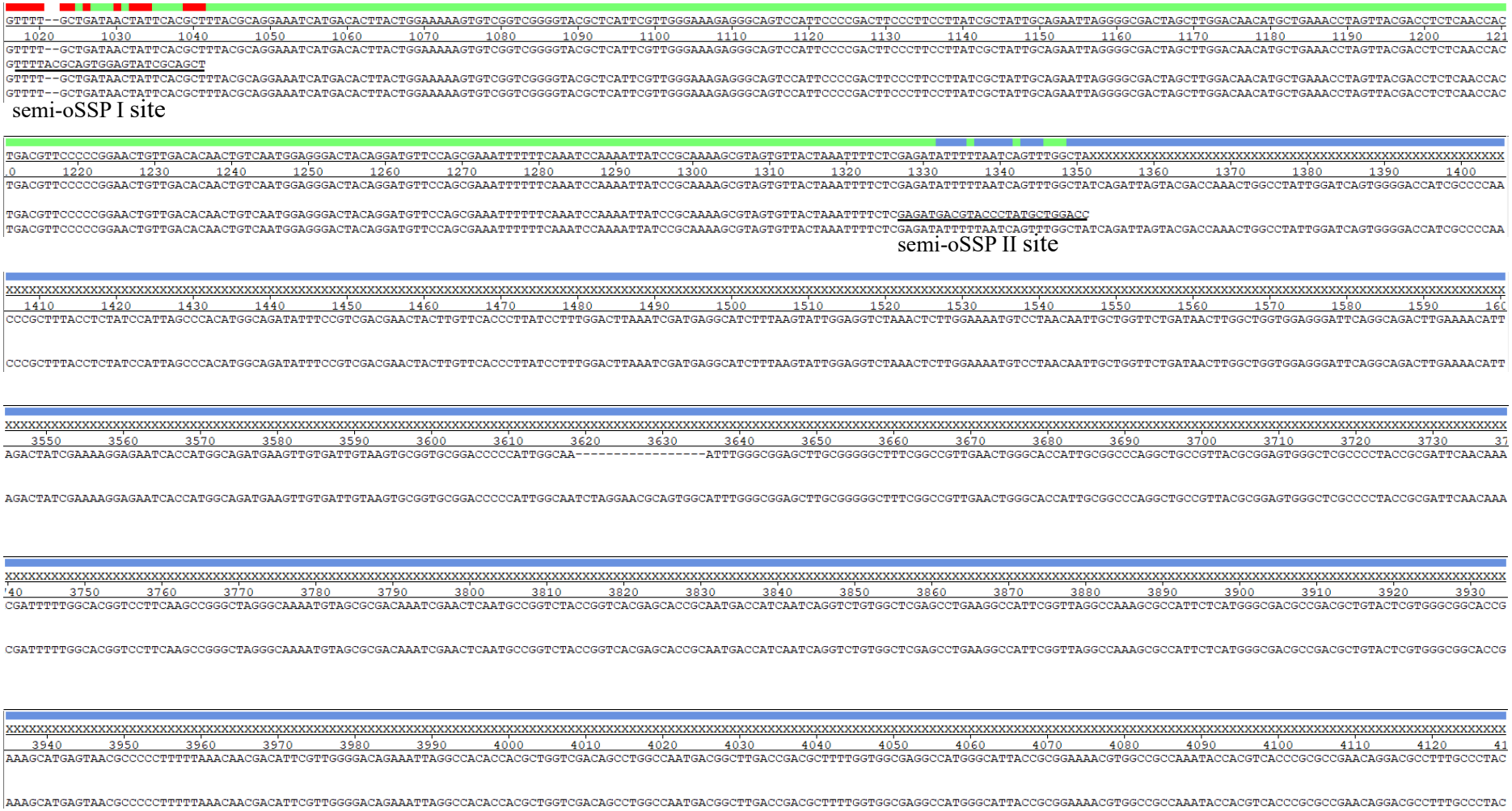
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semi-oSSP II site

semi-oSSP III site

hyg

iSSP

Known Region

Unknown Region

Sequence Name
Consensus
6 Sequences
Ref.seq
H1.seq
H2.seq
H3.seq
H4.seq
H5.seq

semi-oSSP I site

semi-oSSP II site

semi-oSSP II site

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semi-oSSP III site

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```
1600      1610      1620      1630      1640      1650      1660      1670      1680      1690      1700      1710      1720      1730      1740      1750      1760      1770      1780      179
CGGCATGATGGCTGGAGCAATCTGCTCATGAGTGAGGCCGATGGCGTCCTTTGCTCGGAAGAGTATGAAGATGAACAAAGCCCTGAAAAGATTATCGAGCTGTATGCGGAGTGCATCAGGCTCTTTCACTCCATCGACATATCGGATTGTCCCTATACGAATAGCTTAGACAGCCGCTTAGCCGAATTGGATTAC
CGGCATGATGGCTGGAGCAATCTGCTCATGAGTGAGGCCGATGGCGTCCTTTGCTCGGAAGAGTATGAAGATGAACAAAGCCCTGAAAAGATTATCGAGCTGTATGCGGAGTGCATCAGGCTCTTTCACTCCATCGACATATCGGATTGTCCCTATACGAATAGCTTAGACAGCCGCTTAGCCGAATTGGATTAC
```

```
0         1800      1810      1820      1830      1840      1850      1860      1870      1880      1890      1900      1910      1920      1930      1940      1950      1960      1970      1980
TTACTGAATAACGATCTGGCCGATGTGGATTGCGAAAACTGGGAAGAAGACACTCCATTTAAAGATCCGCGCGAGCTGTATGATTTTTTAAAGACGGAAAAGCCCGAAGAGGAACTTGTCTTTTCCCACGGCGACCTGGGAGACAGCAACATCTTTGTGAAAGATGGCAAAGTAAGTGGCTTTATTGATCTTGGC
TTACTGAATAACGATCTGGCCGATGTGGATTGC--AG-TTCGGTCCGGGTCAGTTC---TAA
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