| Matrix ID | Name | Score  | Relative Score | Sequence ID | Start | End  | Strand | Predicted          |
|-----------|------|--------|----------------|-------------|-------|------|---------|--------------------|
| MA0050.1  | IRF1 | 12.4031| 0.891287       | SUCNR1      | 1536  | 1547 | -       | aataacgaa          |
| MA0050.1  | IRF1 | 12.132 | 0.885245       | SUCNR1      | 1833  | 1844 | +       | gaaaataaa          |
| MA0050.2  | IRF1 | 15.3143| 0.858548       | SUCNR1      | 1822  | 1842 | -       | ttttatttt          |
| MA0050.1  | IRF1 | 10.6525| 0.852264       | SUCNR1      | 1482  | 1493 | -       | taaagggaa          |
| MA0050.1  | IRF1 | 10.2148| 0.842508       | SUCNR1      | 1495  | 1506 | -       | aaaactgaa          |
| MA0050.2  | IRF1 | 13.7635| 0.840798       | SUCNR1      | 1828  | 1848 | -       | tgaggtttt          |
| MA0050.1  | IRF1 | 10.0175| 0.83811        | SUCNR1      | 36    | 47   | -       | caaaaatgat         |
| MA0050.2  | IRF1 | 13.3312| 0.835849       | SUCNR1      | 1816  | 1836 | -       | tttcacattt         |
| MA0050.1  | IRF1 | 9.09761| 0.817605       | SUCNR1      | 1827  | 1838 | +       | gaatctgaa          |
| MA0050.1  | IRF1 | 9.03716| 0.816258       | SUCNR1      | 1664  | 1675 | +       | taaaatgaa          |
| MA0050.1  | IRF1 | 8.73889| 0.809609       | SUCNR1      | 282   | 293  | +       | gaaagagga          |
| MA0050.2  | IRF1 | 10.9076| 0.808106       | SUCNR1      | 1413  | 1433 | -       | agttatattt         |
| MA0050.2  | IRF1 | 10.2678| 0.800783       | SUCNR1      | 1491  | 1511 | +       | tttaaatttt         |
sequence
uaac
uagc
cacattcacatt
uaac
uaat
tatattcacatt
uagc
cacattatatatatatatataatataatataatttttttttatagt