

6 HsMhc1\_fl\_(5) 7 HsMhc1\_fl\_(6) result panel

1 HsMhc1\_fl 2 HsMhc1\_fl\_(1) 3 HsMhc1\_fl\_(2) 4 HsMhc1\_fl\_(3) 5 HsMhc1\_fl\_(4)

HsMhc1\_fl\_(1) incomplete HsMhc1\_fl\_(1) selected

| Name          | Match-ratio | Query length (aa) | Number of Contigs |
|---------------|-------------|-------------------|-------------------|
| HsMhc1_fl_(1) | 98          | 1939              | 1                 |

Search details search details

Target file genomes\_ucsc/Homo\_sapiens\_v18\_chromosome.fasta

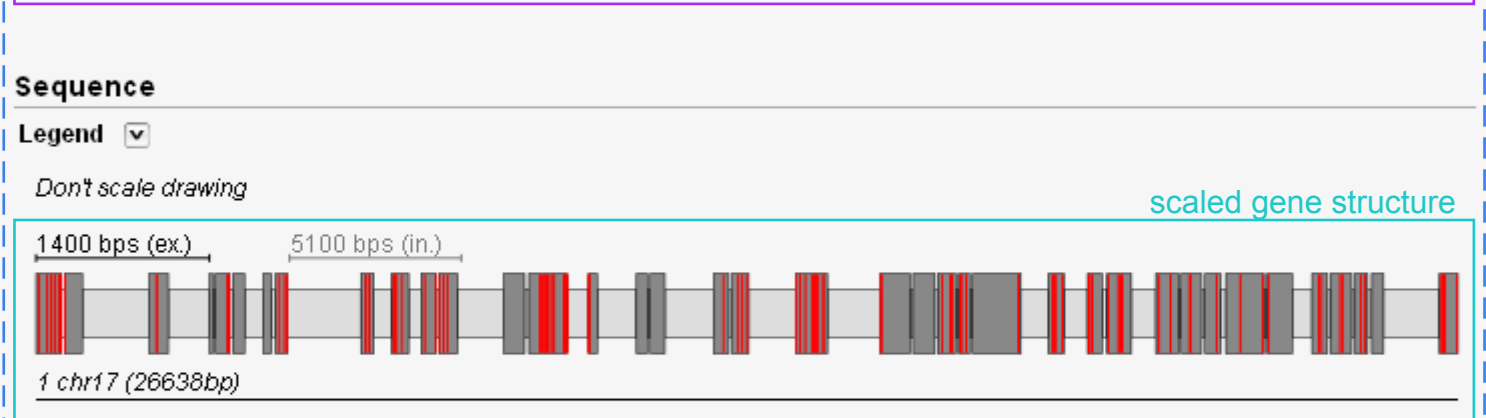
Scpio version Scpio v1.4 (20100627-unreleased)

Search time Thu Jul 1 14:59:21 2010

Targets

Target Name chr17

| Target Number | Status     | Reason     | Target Location            | Pro_len | Matches | Mismatches | Target Strand | Identity | Score |
|---------------|------------|------------|----------------------------|---------|---------|------------|---------------|----------|-------|
| 1             | incomplete | mismatches | -10391962 ...<br>-10365324 | 1939    | 1838    | 101        |               | 94.8%    | 0.896 |



For clarity introns have been scaled down by a factor of 3.7

Statistics

Exons: 38, Introns: 37

Contigs: 1

result views