

deepBlockAlign v1.0

Alignment parameters:

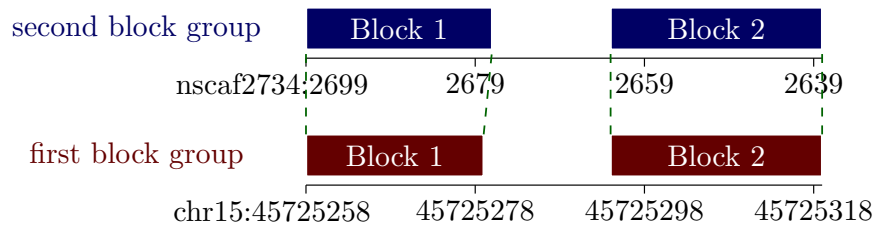
```
[Input]
date and time          Sun, 03 Feb 2013 01:08:34 +0100
first block group      BrainSknshraRep2_cluster_3891
first file             /home/projects/users/sachin/frontier/putative_mirna//tmp1500query.tmp
second block group     bg_67
second file            /home/projects/users/sachin/frontier/bgs/mirbase.all.map.bgs.flagged.sig

[Block group alignment]
distance weight        1
block weight           1
gap penalty            -1

[Block alignment]
gap initialization penalty -2
gap extension penalty    -1
threshold for expression difference 1
match score             1
mismatch score          -1
profile shape difference penalty 1

[Block group information]
Entropy (first block group) 1.53
Reads (first block group) 19.00
% Unique reads (first block group) 100
Entropy (second block group) 1.23
Reads (second block group) 161.00
% Unique reads (second block group) 100
```

Block group alignment: BrainSknshraRep2-cluster-3891-hsa-mir-147b(+) with bg-67-bmo-mir-993b-MI0010625(-) [Score: 0.81]



Block alignment:

