Dotplots

Anders Krogh

Simple dotplot

```
CIAOl HUMAN Probabl... vs. CIAOl HUMAN Probabl...
Zoom: 1 : 1
Word length: 0
Window size: 1
Matrix: Identity
                       Program: Gepard (1.30)
                                               338
   CIAOl HUMAN Probable cytosolic iron-sul...
```

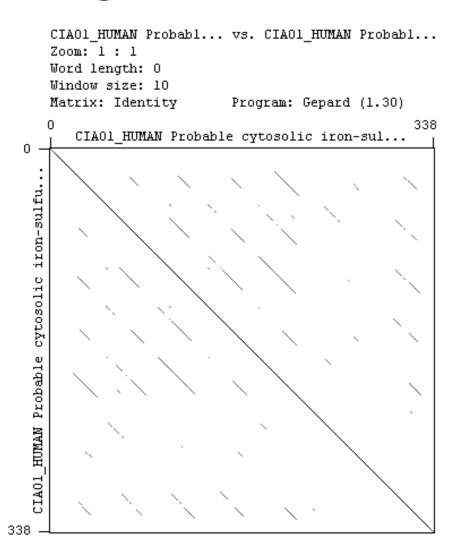
 Make a dot every time the two letters are identical

 This plot is made with gepard (like all the rest)

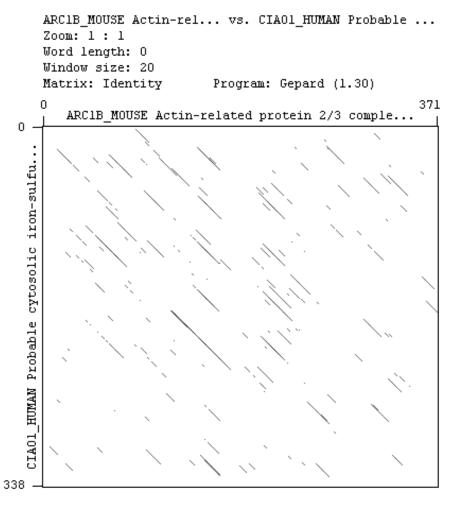
Windowing

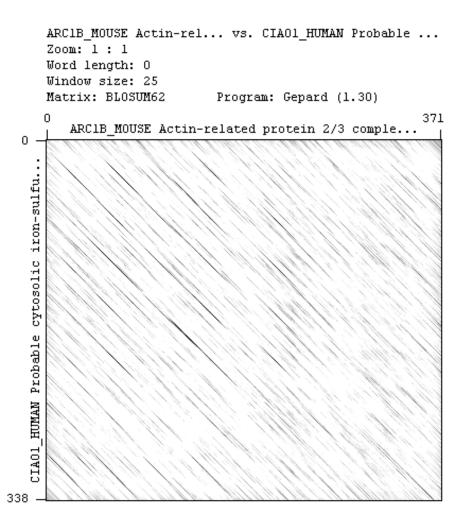
```
...WSPCGNYLASASFDATTCI...
...WSPCGNYDLGSADFDDTTIWK..
```

 Add identical letters in window



Substitution matrix and gray scale





Mouse protein vs human protein, win. 20

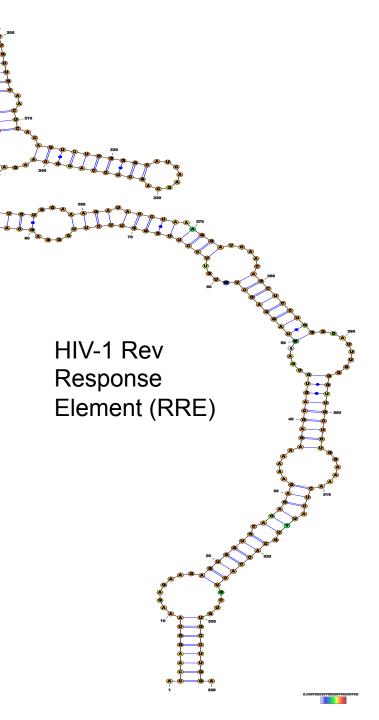
BLOSUM62 and win. size 25

BLOSUM62 matrix

0 - 3 - 30 -2 8 -3 -3 -1 0 -3 -2 -1 -3 -2 5 -1 -3 -1 2 -1 5 0 -2

Inverted repeats etc.

- Genomes have inversions
- Repeats can occur on both strands of DNA
- RNA base-paring looks like inverted repeats

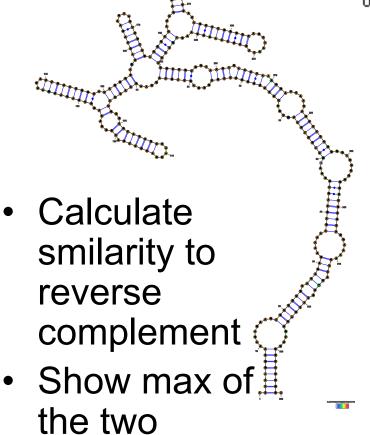


RNA dotplot

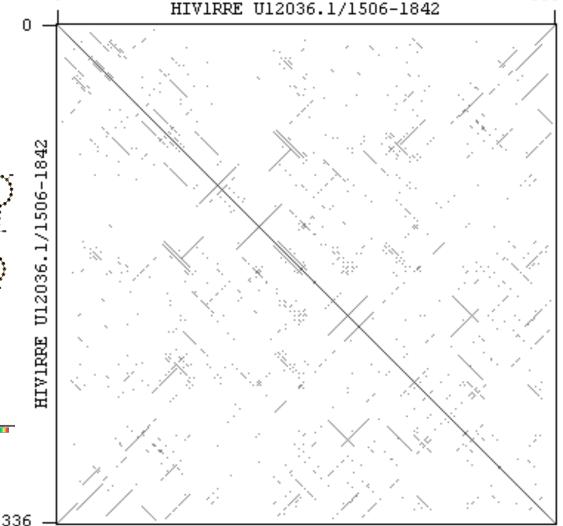
Zoom: 1 : 1
Word length: 0 GC ratio seql: 0.5074
Window size: 20 GC ratio seq2: 0.5074
Matrix: DNA Program: Gepard (1.30)

HIV1RRE U12036.1/1... vs. HIV1RRE U12036.1/1...

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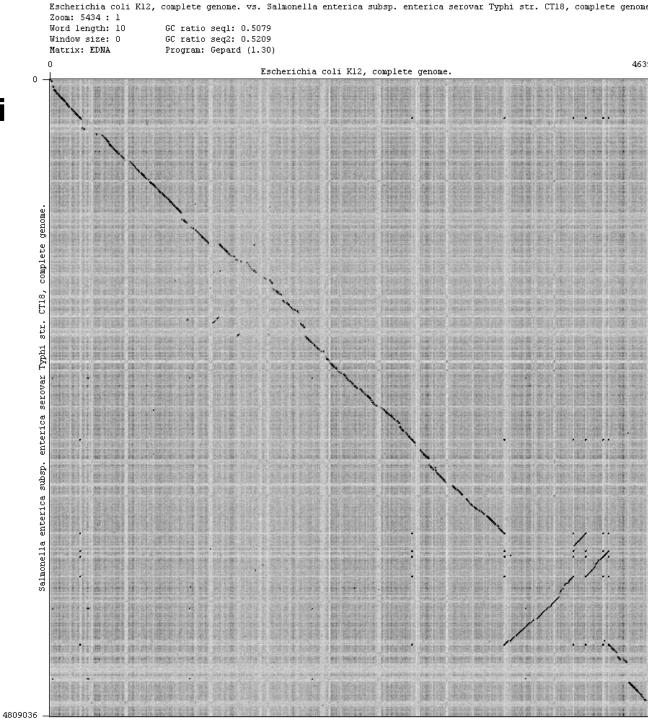
scores



Coli vs Salmonella Typhi

The two are almost colinear

One major region is inverted



Computing dotplots

- 2x10¹³ points in the genome comparison!
- Smart ways to make fast computations
- Dotter
 http://sonnhammer.sbc.su.se/Dotter.html
 is fast
- gepard http://cube.univie.ac.at/gepard is faster

Pairwise alignment

Examples

```
(a)
HBA HUMAN GSAQVKGHGKKVADALTNAVAHVDDMPNALSALSDLHAHKL
          G+ +VK+HGKKV A+++++AH+D++ +++++LS+LH
HBB HUMAN GNPKVKAHGKKVLGAFSDGLAHLDNLKGTFATLSELHCDKL
(b)
HBA HUMAN GSAQVKGHGKKVADALTNAVAHV---D--DMPNALSALSDLHAHKL
          ++ ++++H+ KV + +A ++
                                         +L+ L+++H+ K
LGB2 LUPLU NNPELQAHAGKVFKLVYEAAIQLQVTGVVVTDATLKNLGSVHVSKG
(c)
         GSAQVKGHGKKVADALTNAVAHVDDMPNALSALSD----LHAHKL
HBA HUMAN
          GS++G+DL++H+D+A+ALD++AH+
F11G11.2 GSGYLVGDSLTFVDLL--VAQHTADLLAANAALLDEFPQFKAHQE
```

BLOSUM62 – again

0 -3 -3 0 -2 8 -3 -3 -1 0 -3 -2 -1 -2 -1 1 -2 -1 -3 -2 5 -1 -3 -1 0 -1 -3 -2 2 -1 5 0 -2