

chrI

N2 chrI:4520897-4569931 vs. vc2010 chrI:4574828-4688011

Zoom: 125 : 1

Word length: 10

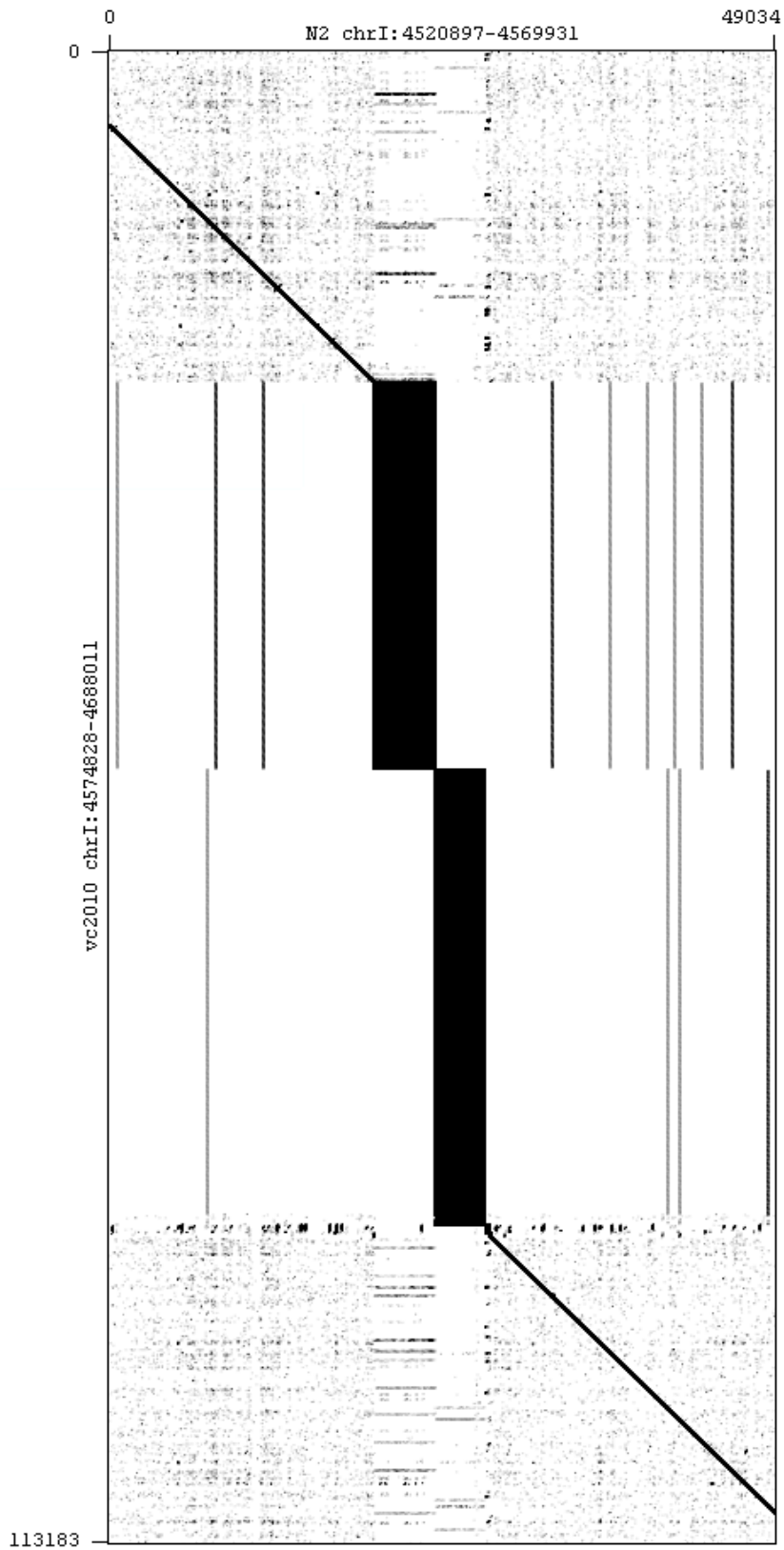
GC ratio seq1: 0.3566

Window size: 0

GC ratio seq2: 0.3829

Matrix: DNA

Program: Gepard (1.40 final)



The horizontal axis shows the N2 genome (ce10) around the gap.

The vertical axis presents the corresponding region in VC2010 genome.

chrII

N2 chrII:14310000-14360000 vs. vc2010 chrII:14495913-14586327

Zoom: 100 : 1

Word length: 10

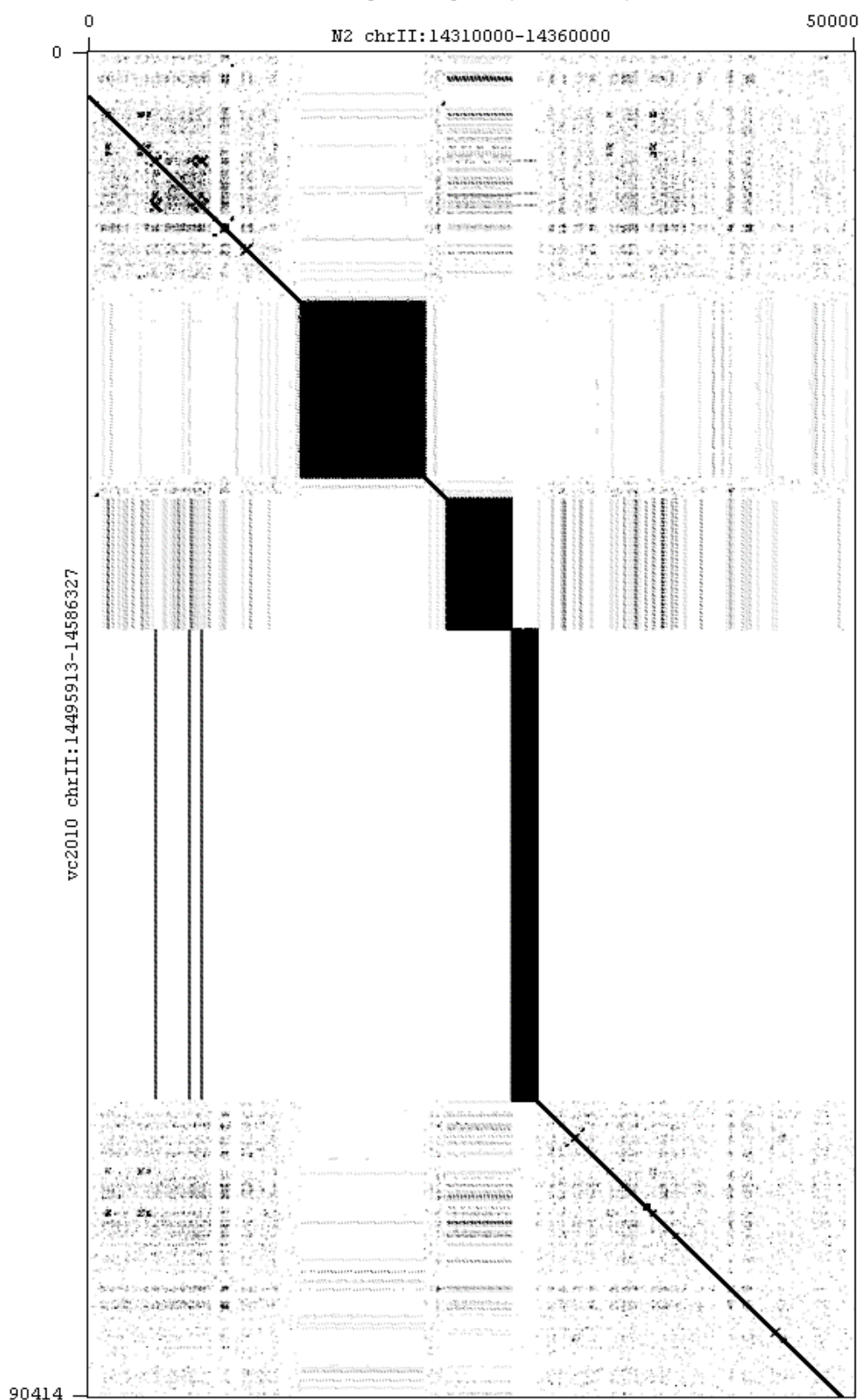
Window size: 0

Matrix: DNA

GC ratio seq1: 0.3528

GC ratio seq2: 0.3662

Program: Gepard (1.40 final)

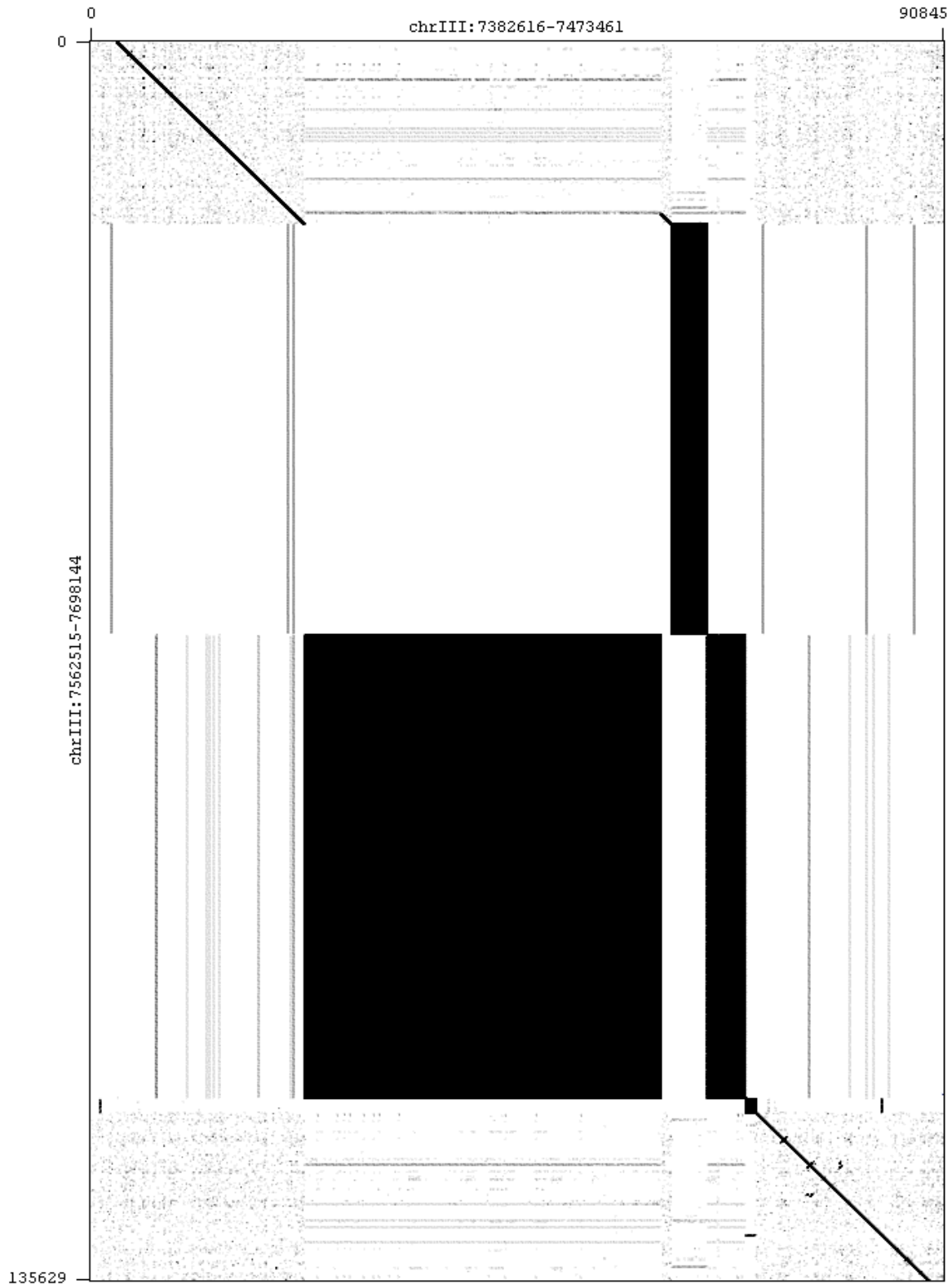


The horizontal axis shows the N2 genome (ce10) around the gap.

The vertical axis presents the corresponding region in VC2010 genome.

chrIII

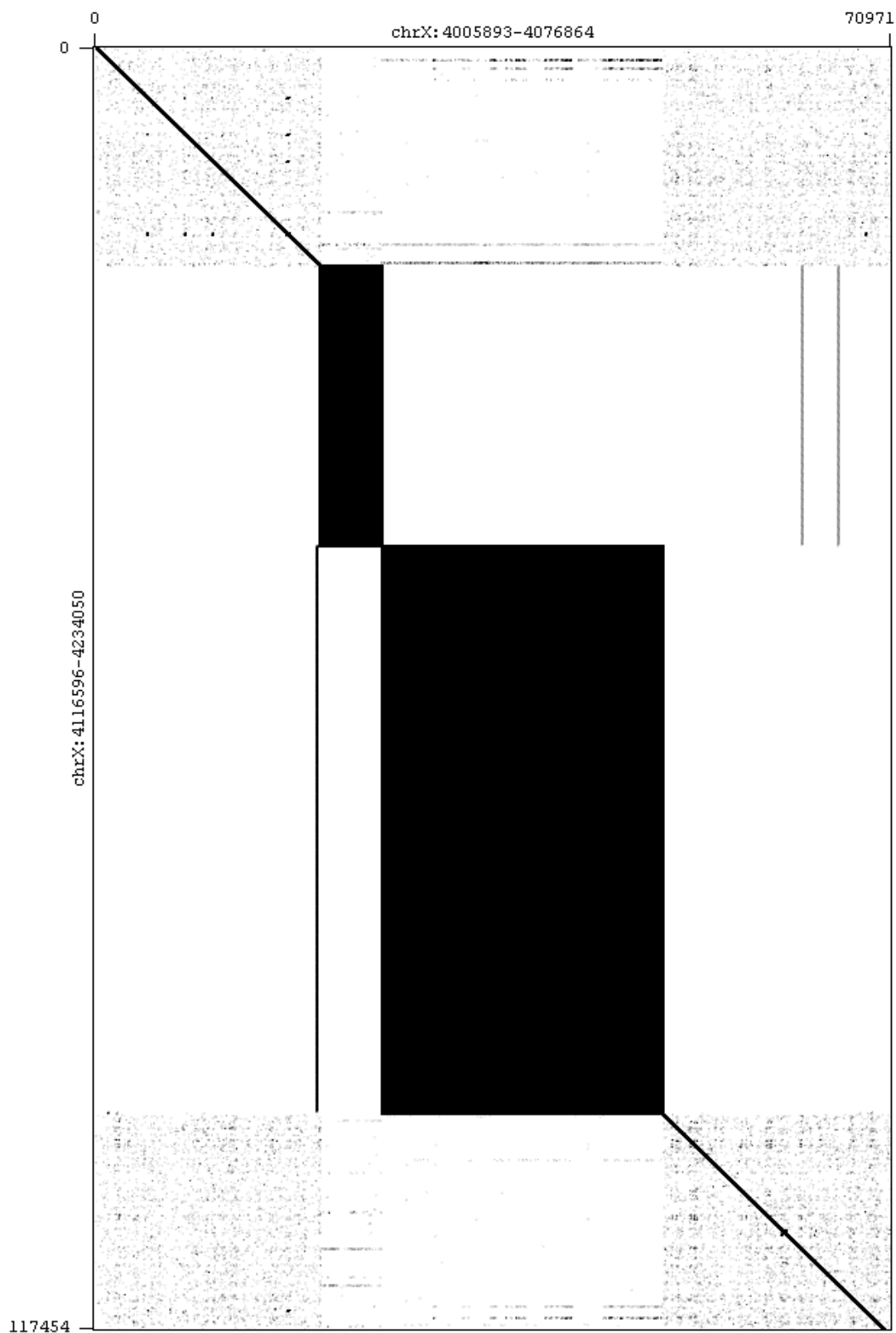
chrIII:7382616-7473461 vs. chrIII:7562515-7698144
Zoom: 149 : 1
Word length: 10 GC ratio seq1: 0.3583
Window size: 0 GC ratio seq2: 0.3762
Matrix: DNA Program: Gepard (1.40 final)



The horizontal axis shows the N2 genome (ce10) around the gap.
The vertical axis presents the corresponding region in VC2010 genome.

chrX 4M

chrX:4005893-4076864 vs. chrX:4116596-4234050
Zoom: 129 : 1
Word length: 10 GC ratio seq1: 0.3343
Window size: 0 GC ratio seq2: 0.3123
Matrix: DNA Program: Gepard (1.40 final)



The horizontal axis shows the N2 genome (ce10) around the gap.

The vertical axis presents the corresponding region in VC2010 genome.

chrX 5M

chrX:5026355-5076355 vs. chrX:5183208-5291274

Zoom: 119 : 1

Word length: 10

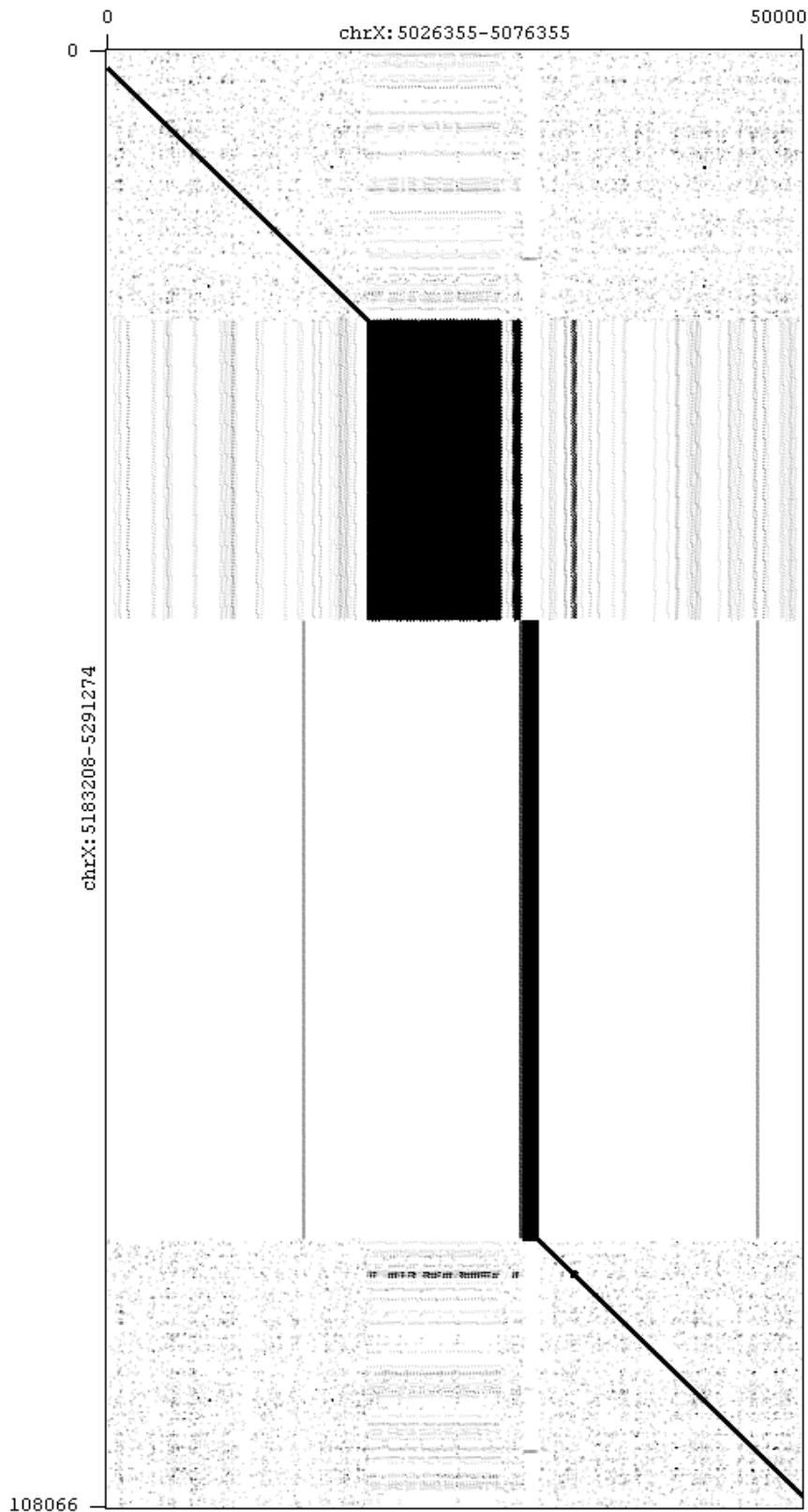
Window size: 0

Matrix: DNA

GC ratio seq1: 0.3318

GC ratio seq2: 0.3551

Program: Gepard (1.40 final)



The horizontal axis shows the N2 genome (ce10) around the gap.

The vertical axis presents the corresponding region in VC2010 genome.