percentage of bases (500kb windows)

C. elegans chromosome I position (Mb)
Percentage of bases (500kb windows) against C. elegans chromosome IV position (Mb).
percentage of bases (500kb windows)

C. elegans chromosome V position (Mb)
percentage of bases (500kb windows)

C. elegans chromosome X position (Mb)
percentage of bases (500kb windows)

C. elegans chromosome II position (Mb)

"exon.II"
percentage of bases (500kb windows)

C. elegans chromosome III position (Mb)

"exon.III"
percentage of exons in orthologs (500kb windows)

C. elegans chromosome I position (Mb)

"ortholog.I"
percentage of exons in orthologs (500kb windows)

C. elegans chromosome II position (Mb)
percentage of exons in orthologs (500kb windows)

C. elegans chromosome III position (Mb)
percentage of exons in orthologs (500kb windows)

C. elegans chromosome IV position (Mb)
percentage of exons in orthologs (500kb windows) vs C. elegans chromosome X position (Mb)
The graph shows the percentage of bases (500kb windows) in C. elegans chromosome IV position (Mb). The x-axis represents the C. elegans chromosome IV position in Mb, and the y-axis represents the percentage of bases. The line labeled "repeat.IV" indicates the distribution of repeat elements across the chromosome.
percentage of bases (500kb windows)

C. elegans chromosome V position (Mb)

"repeat.V"
percentage of bases (500kb windows)

C. elegans chromosome X position (Mb)

"repeat.X"
number of telomeric repeats (500kb windows)

C. elegans chromosome I position (Mb)
number of telomeric repeats (500kb windows)

C. elegans chromosome III position (Mb)

"telomere.III"
number of telomeric repeats (500kb windows)

C. elegans chromosome IV position (Mb)
number of telomeric repeats (500kb windows)

C. elegans chromosome X position (Mb)