Ideos

14 September, 2024

Contents

Queries]
nuccomp	1
fast_win	1
Ideo function	

Queries

query	Length
45S	9382
26S	1544
5.8S	157
18S	1552
5S	259
CS-237_satelite	557
$CsatSD_centromere_237bp$	237
${\tt CsatSD_subtelomere_370bp}$	370

nuccomp

 $fast_win$

Ideo function

Time difference of 7.731123 secs

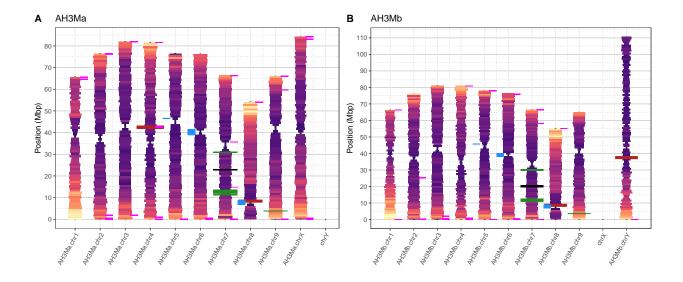


Figure 1: Ideogram of the *Cannabis sativa* genome. Each chromosome is presented as a set of 1 Mbp windows where the width is proportional to the abundance of the 'CpG' motif (wide is low abundance, narrow is high abundance) and colored by gene density (cool or blue colors are low gene density and hot or yellow colors are high gene density). The 45S, 26S, 18S, and 5.8S probes are in firebrick red (#B22222), the 5S probes are in black, Cannabinoids (CBCAS, CBDAS, THCAS, OAC) are in forest green. The probe 'CsatSD_centromere_237bp' is in Dodger blue (#1E90FF), the probe 'CsatSD_centromere_370bp' is in magenta (#FF00FF).