

# Result Annotations

Each annotation has the following qualifiers:

- *area\_average\_threshold* — average window threshold in the area (i.e.  $total\_threshold / windows\_number$ )
- *total\_threshold* — sum of all window thresholds in the area
- *windows\_number* — number of windows in the area

■ dna_flex	144..156
area_average_threshold	14.672
total_threshold	58.689
windows_number	4



Using the [DNA Graphs Package](#) you can see the flexibility graph of a DNA sequence.