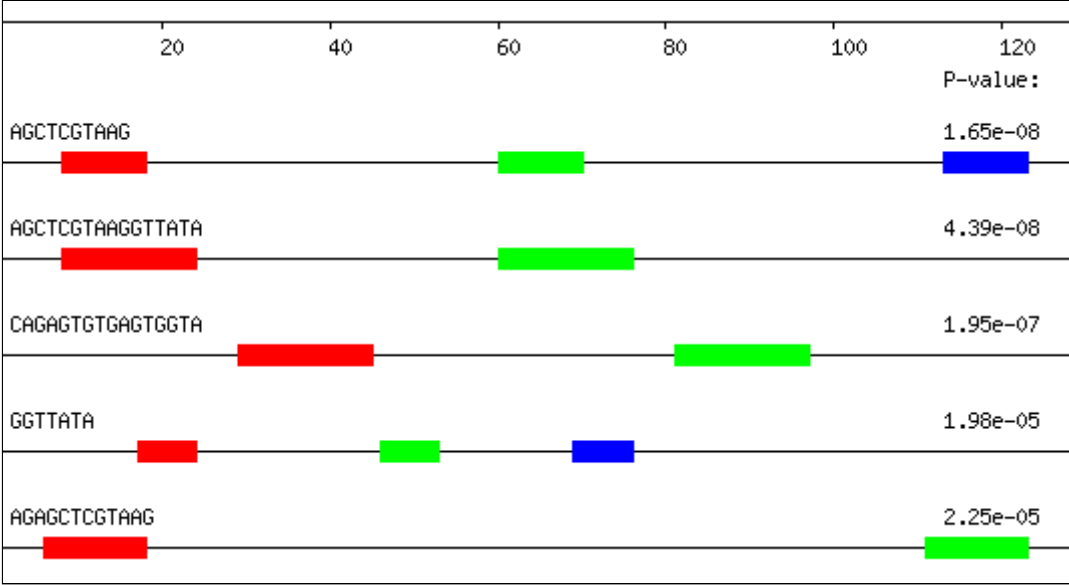# REPFIND

Results for your sequence

Nucleotides 1 to 127
P-value cutoff: 0.0001
Minimum repeat length: 6
Maximum repeat length: infinity
Low complexity filter: on
Statistical background: query sequence
Order of background Markov model: 1

REPFIND home

Word: **AGCTCGTAAG**
Word locations: 8 60 113
Most significant cluster: 8 to 122
P-value: 1.64589e-08

Word: **AGCTCGTAAGGTTATA**
Word locations: 8 60
Most significant cluster: 8 to 75
P-value: 4.39185e-08

Word: **CAGAGTGTGAGTGGTA**
Word locations: 29 81
Most significant cluster: 29 to 96
P-value: 1.94576e-07

Word: **GGTTATA**
Word locations: 17 46 69
Most significant cluster: 17 to 75
P-value: 1.97892e-05

Word: **AGAGCTCGTAAG**
Word locations: 6 111
Most significant cluster: 6 to 122
P-value: 2.24601e-05