Sekwencja1: GATGCCTACACAGATTATCAAGG
F: TTCGGATGCCTACACAGATTATCA
R:AAACTGATAATCTGTGTAGGCAT
czy specyficzna:

Details

| Bulge Type | DNA bulge | RNA bulge | Mismatch | Filter | Download filtered result |

Bulge Type	Target	Chromosome	Position	Direction	Mismatches	Bulge Size
X	crRNA: GATGCCTACACAGATTATCANGG DNA: GATGCCTACACAGATTATCAAGG	chr5	18428304	-	0	0

Sekwencja2: GAAGCGAATCCACCATACCAAGG
F:TTCGGAAGCGAATCCACCATACCA
R:AAACTGGTATGGTGGATTCGCTT
czy specyficzna:

Details

| Bulge Type | DNA bulge | RNA bulge | Mismatch | Filter | Download filtered result |

Bulge Type	Target	Chromosome	Position	Direction	Mismatches	Bulge Size
X	crRNA: GAAGCGAATCCACCATACCANGG DNA: GAAGCGAATCCACCATACCAAGG	chr5	18427419	-	0	0

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   1 ccaaccagaa taattcaaac cttcacacac aacaaaaagc ctcaaaagca aaaaaactca
  61 tagtaaaacc agaaaaagaa agaagtcaac ctctttagat gtatctgtgt tagatctcat
 121 aaaccatcag ttcttgtctg tttcagcaaa tcagtttcaa gatggtgaac gcaatcccaa
 181 agcctatcaa ggacatttgg gacgaatgga gcattcgaag taccttaatc ttcagtctca
 241 gccttcaaac atttctcatt ttctttgcac ctcaaagaaa acgctcatcc agaaaagttt
 301 tactctcttt catttggtca gcttaccttc ttgctgattg gtctgccaat ttcgccgccg
 361 gacaaatctc agatagtcaa ggagatgatc cagaacccgg agaaccgaaa aagagcgccg
 421 agttgtttgc tttctgggtt cctttcttgc tcttgcatct cggtggtcca gatacaatca
 481 ctgcccttgc ccttgaagat aacgagctct ggctcaggca cttgttgggt cttttctttc
 541 aatcagttgc aaccgtttac gtgcttttgc agtcgttacc caatgctctt tggaaaccta
 601 ttttgctggt gtttgcgacc ggggtgatca agtatgtcga gaggacatta gctttgtacc
 661 tggcgagtct tgacaagttc aaagattcca tgattcagcg gcctgatcct ggaccaaact
 721 acgcaaagct catggaggaa tacgcagcca agaaagatat gaagatgcct acacagatta
 781 tcaaggttgg cgagcctgag aaagatccaa gagacgatgc tccggtaaag cctcctgacg
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1381 ctcatgaagt gcccaacacg agtactaaga aaacaccatt tcttaaaagg attcttaaca
1441 ggattctcag ggtgagggag ttgaagacag agaagtctca tgaagttctc gacaagagta
1501 ccagcaaaat accaggtcag gaagtgcccg acaagagtgc caagacaata ccatgtcata
1561 aagtgctcga cacgagtttc atgtatcgcc ggtggtctga gtatgtacat gcacacaacc
1621 tgatcgaata ctgcctaggt ttaaaaccga agcgaatcca ccataccaag ggtttcatac
1681 acatcgcctt tgacaaggtt atcaacatac tttacattgg ccccgcattc acaaaacttg
1741 gaagtgtcat tgaatcctgt tttagagtca caaaacagag aattcatcag actttcaagt
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1801 ggattgacgg taaagtaagt cgtttatgca agaaataccc gaagttgaat gaagagtaca
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2941 taactcatgt ccttcttctg tccctgataa attcatttat gtgagct
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