→ Primer-BLAST Primer-Blast results

```
NCBI/ Primer-BLAST : results: Job id=kJpP1N2V0D33A8oGx2buNL19_waQbuQbkQ more...

•
Input PCR template
    none
Specificity of primers
    Target templates were found in selected database: Nucleotide collection (nt)
    (Organism limited to Homo sapiens)
Other reports
    Search Summary
```

Detailed primer reports

Primer pair 1

| | Sequence (5'->3') | Length | Tm | GC% | Self complementarity | Self 3' complementarity |
|-------------------|--------------------------------|--------|-------|-------|----------------------|-------------------------|
| Forward primer | CGCAGAAGTAAGTAGCCGGG | 20 | 60.53 | 60.00 | 4.00 | 3.00 |
| Reverse primer | ACTGGATGAGCGTGAAGTGG | 20 | 60.04 | 55.00 | 3.00 | 0.00 |
| Products | on intended target | | | | | |
| Products | on allowed transcript variants | | | | | |

Products on potentially unintended templates

Products on target templates

>AY615271.1 Homo sapiens growth arrest and DNA-damage-inducible beta (GADD45B) mRNA, partial cds, alternatively spliced

>AY615270.1 Homo sapiens growth arrest and DNA-damage-inducible beta (GADD45B) mRNA, partial cds, alternatively spliced

>AF533019.1 Homo sapiens growth arrest and DNA-damage-inducible, beta (GADD45B) gene, complete cds

>AC005624.1 Homo sapiens chromosome 19, cosmid R30017, complete sequence

 product length = 725
 CGCAGAAGTAAGTAGCCGGG
 20

 Forward primer 1
 CGCAGAAGTAAGTAGCCGGG
 20

 Template
 27862
 27881

 Reverse primer 1
 ACTGGATGAGCGTGAAGTGG
 20

 Template
 28586
 28567