

## Supplementary Data 2

### Examples of DNA sequencing profiles

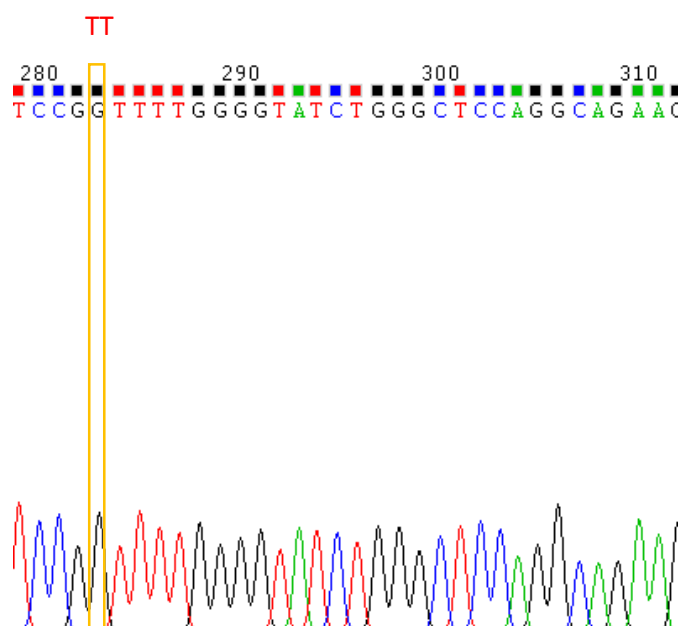
#### 1. *SOD2* c.47C>T (Ala16Val, rs4880)

*SOD2* Ala16Val rs4880 (Ala: GCT→Val: GTT) (NG\_008729.1:g.5482T>C)

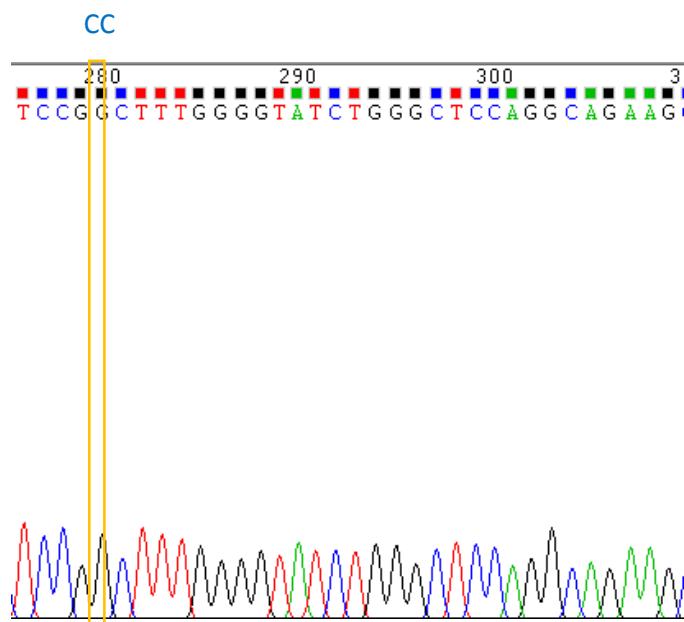
CAGCACCAGCAGGCAGCTGGCTCCGG[C/T]TTTGGGGTATCTGGGCTCCAGGCAG

Example: Sequencing results

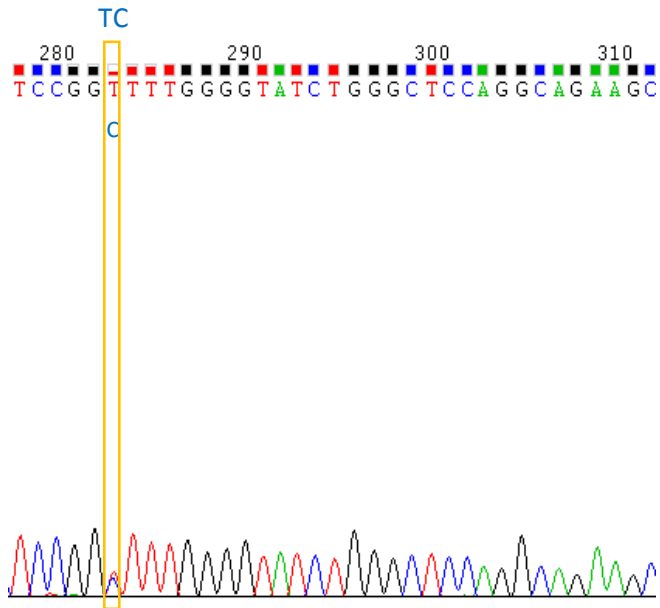
**CAD011: TT**



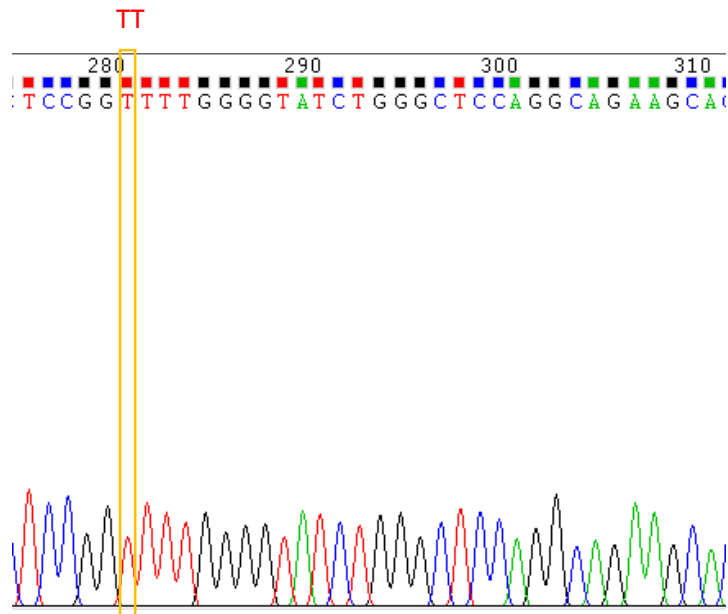
**CAD068: CC**

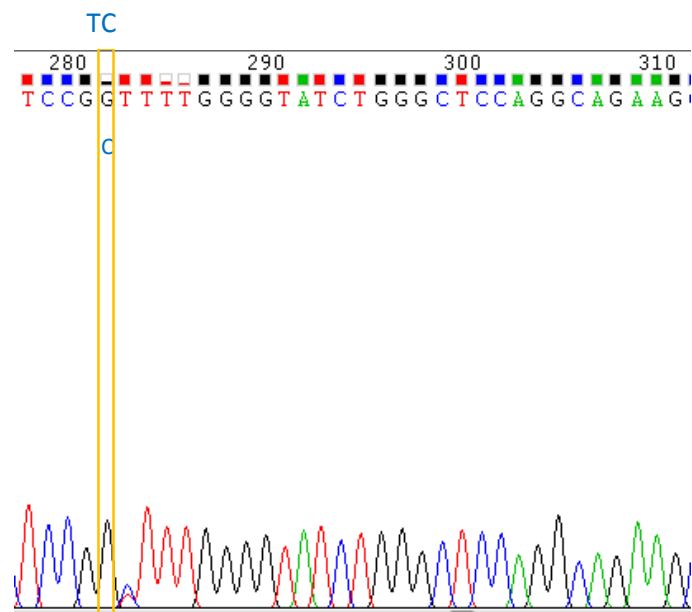
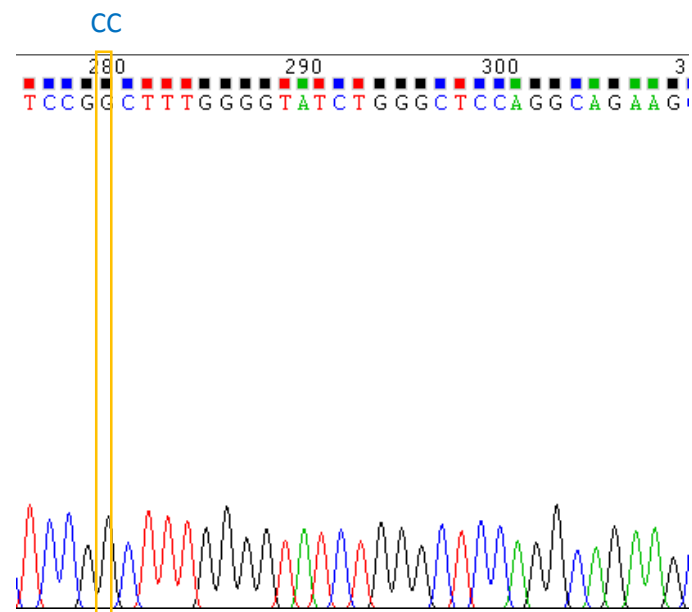


### CAD147: TC



### CADA030: TT



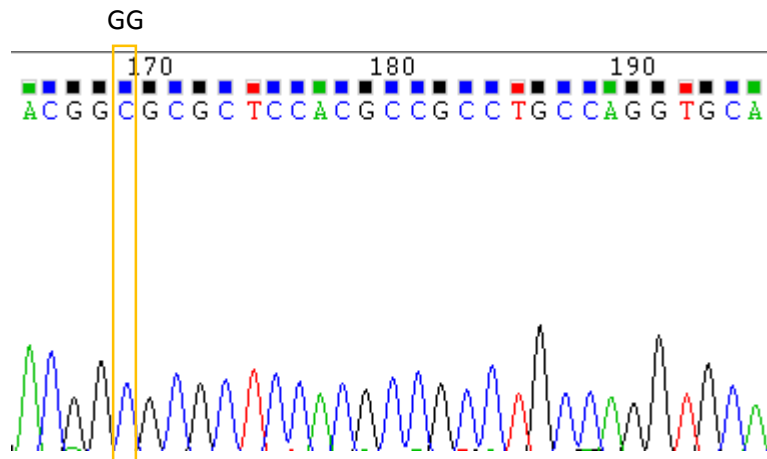
**CADA121: TC****CADA128: CC**

## 2. *SOD3* c.172G>A (Ala58Thr, rs2536512)

*SOD3* Ala58Thr rs2536512 (Ala: GCG → Thr: ACG)\_NG\_012213.1:g.9231G>A  
TCATGCAGCGGCGGGACGACGAGGC[A/G]CGCTCCACGCCGCCTGCCAGGTGCA

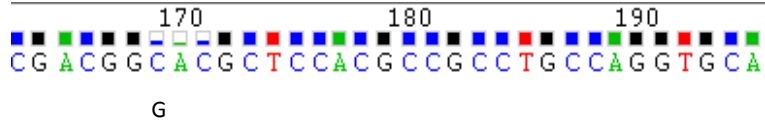
Example: Sequencing results

**CAD035: GG**

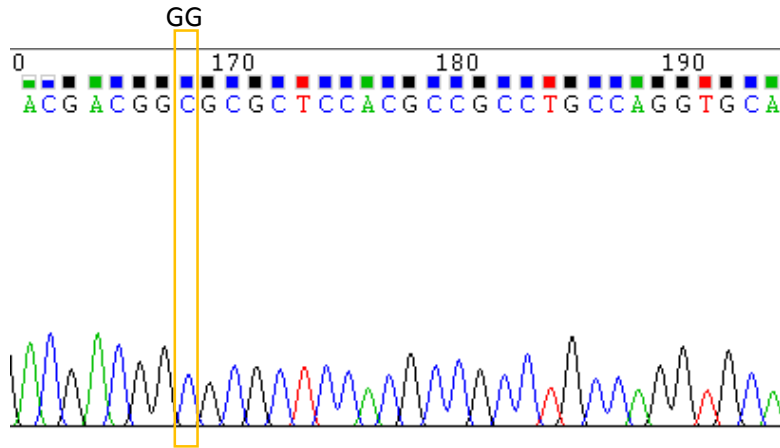


**CAD046: GA**

GA



**CADA017: GG**

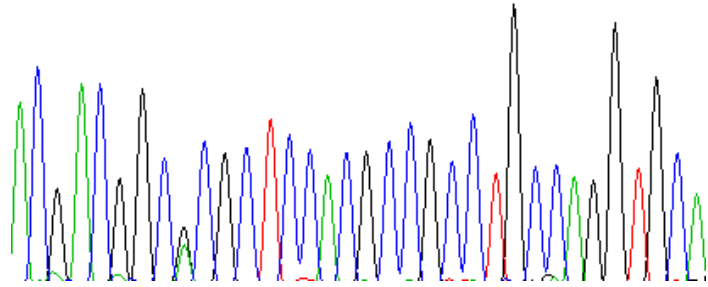


**CADA088: GA**

GA

Detailed description: This figure shows a vertical yellow box highlighting a region, with the label 'GA' above it. The box is positioned below the 'CADA088: GA' header.

170 180 190  
ACGACGGCGCGCTCCAGCCCGCCCTGCCAGGTGCA  
A

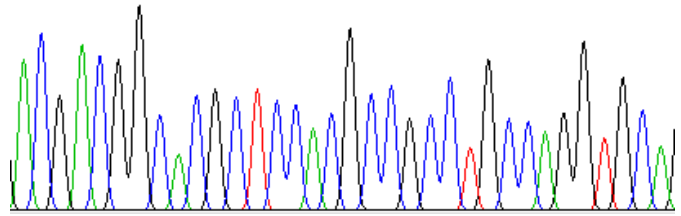


**CADA092: AA**

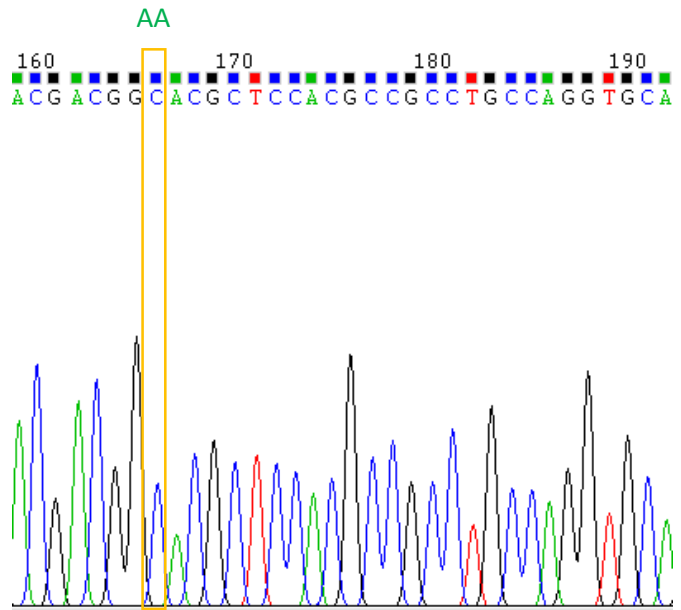
AA



160 170 180 190  
A C G A C G G C A C G C T C C A C G C C G C C T G C C A G G T G C A



**CADB025: AA**

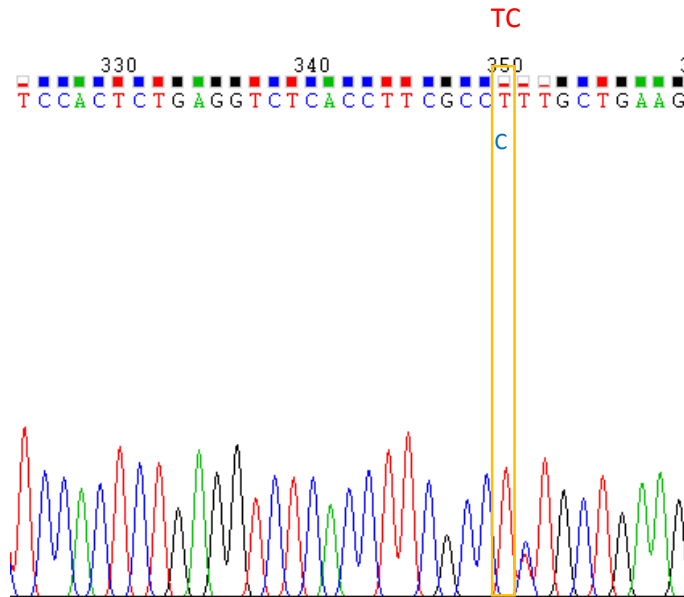


**3. SOD3 g.9892T>C (rs2855262)**

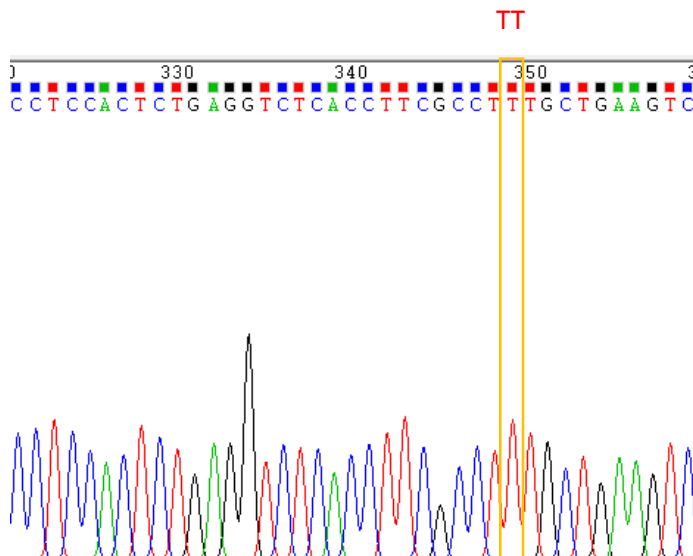
*SOD3* rs2855262 [NC\_000004.11:g.24801976T>C ] [NG\_012213.1:g.9892T>C ] [ NM\_003102.2:c.\*110T>C ] [NT\_006316.16:g.15983773T>C ] [NW\_001838900.1:g.15423814C>C] TCCACTCTGAGGTCTCACCTTCGCCT[C/T]TGCTGAAGTCTCCCCGCAGCCCTCT

Example: Sequencing results

### CAD148: TC

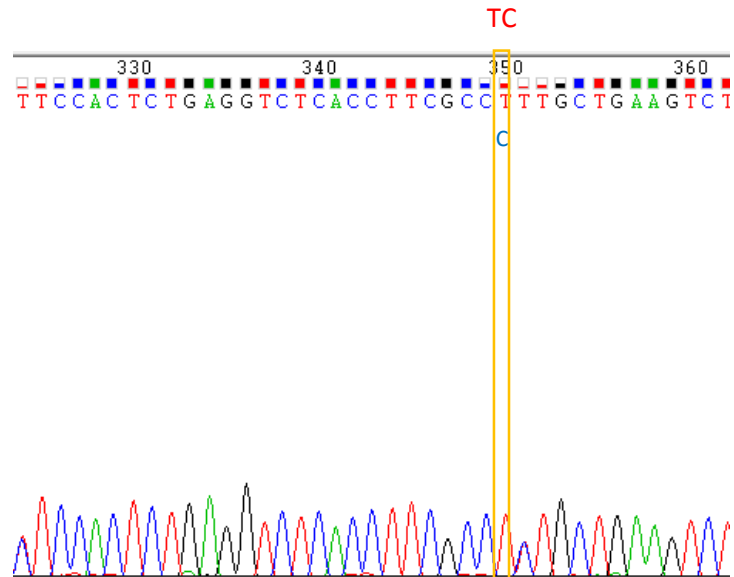


### CADA050: TT

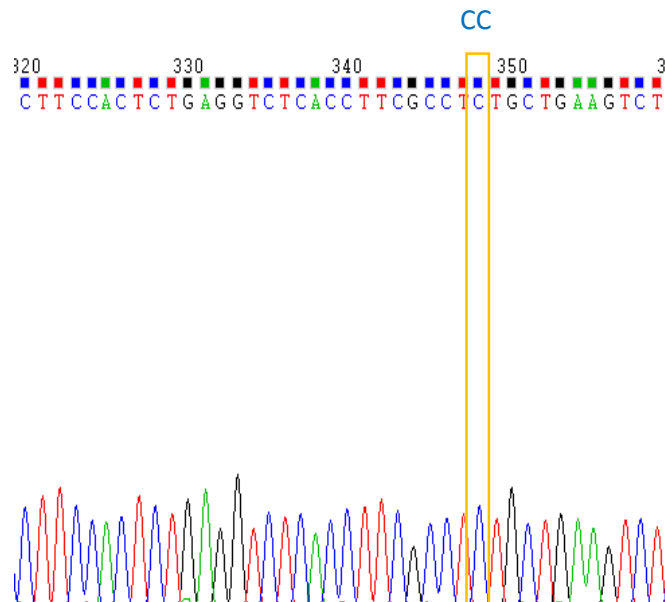




### CADA069: TC



### CADB016: CC



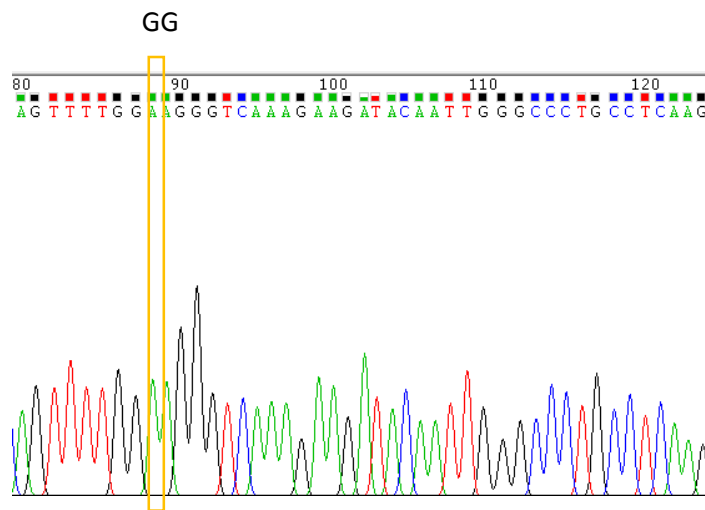
#### 4. *GPX3* c.87+1494A>G (rs3828599)

*GPX3* c.87+1494A>G [ NC\_000005.9:g.150401796A>G ] [ NM\_002084.3:c.87+1494A>G ]  
[NT\_029289.11:g.11564723A>G ]

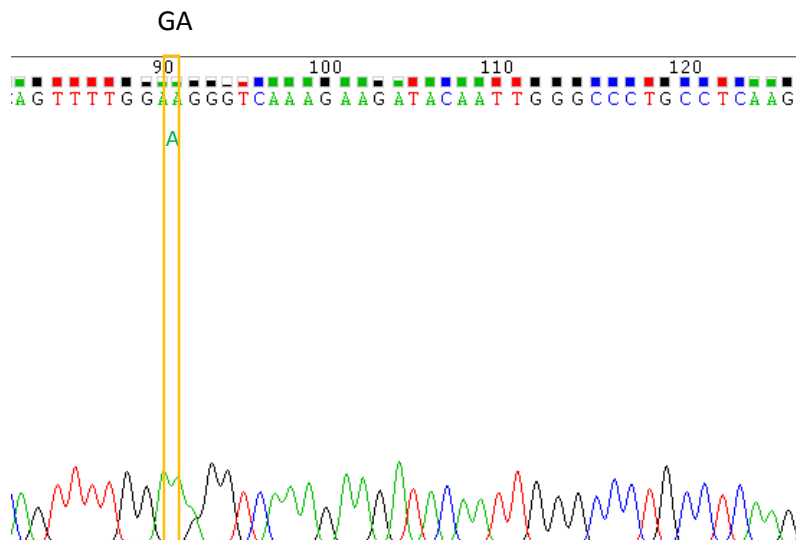
TCCCAACCTTCAGTTTTGGAA(G/A)GGTCAAAGAAGATAACAATTGGGCCCTGC

Example: Sequencing results

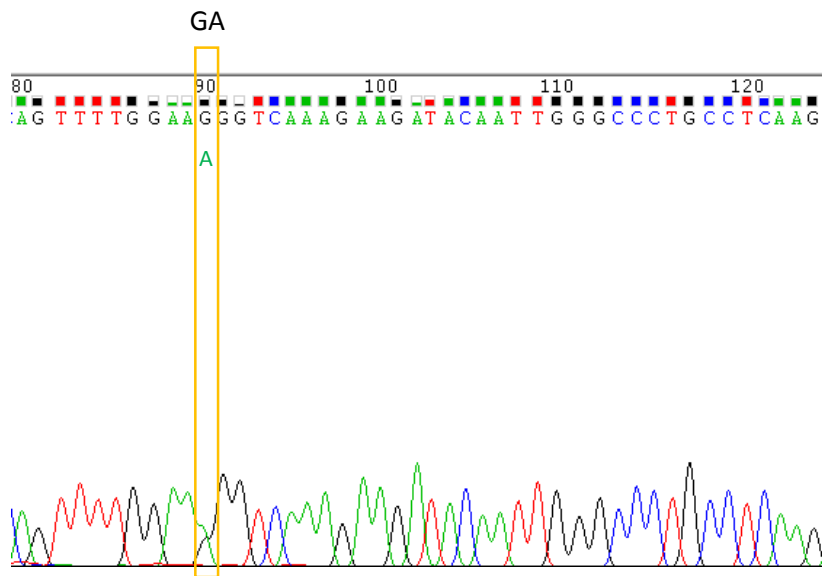
**CAD183: GG**



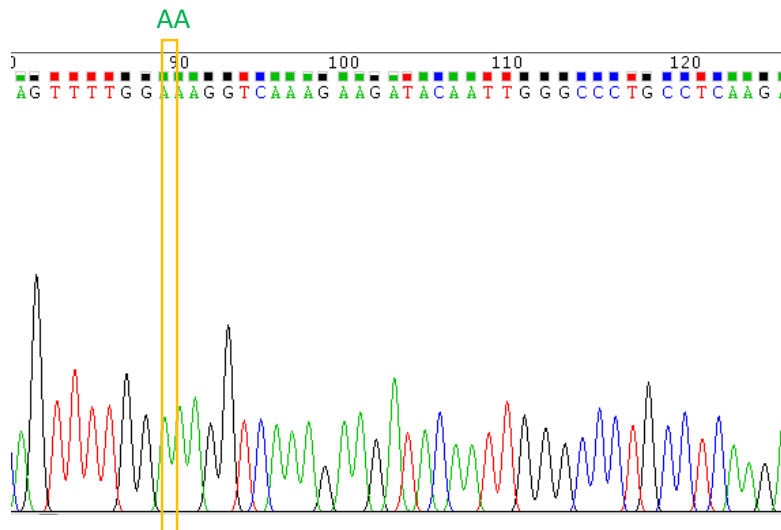
**CAD120: GA**



**CAD155: GA**



**CADA094: AA**



### 5. *GSTT1* deletion polymorphism

#### *GSTT1* + genotype

GenBank: AF240786.1 (minus strand)

```

85081 AAGCTGACCC ACACGGCCTG GGAAGGGGGT TGTCTTTTGC ATAGAGACCA TGACCAGGTC
85141 TGGGACAGAG GAAAGTCAAA TAAATCACAC ATTAGAGTTA GAAGCAGAGG CTCAGGCTGA GSTT1
85201 GCCCAGGTTT ATTATCCAAA ATCAAAATGA AATGCAGTGA TTAAAGGACA CAAGGCCTCA exon 5
85261 GTGTGCATCA TTCTCATTGT GGCTTTTCAGG CGGCTGTGGA AGACAGGGTG GGGATGGTGG (minus
85321 CTTCCGGGAGG TGAGGTGCTC TGGGACTTGG GCAAGTCTTA GGCAAGCCAT TCCTGCTTTC strand)
85381 TGGGCCTGGC TCCCATGGGC CATTAGAAAT GAAAATGCTT TGTGGACTGC TGAGGACGGT

```

GSTT\*A1 forward **CCAG CTCACCGGAT CATGGCCAG**

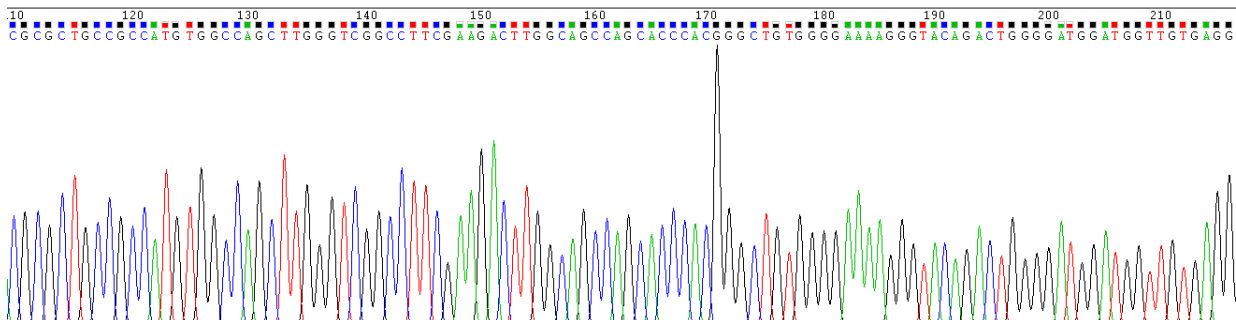
```

85441 GCAAGGGTGA GGTTTCCCAG CTCACCGGAT CATGGCCAGC ACCCAGGGCA TCAGCTTCTG
85501 CTTTATGGTG GGGTCTGCAG GTGGGAAGTC CTTGGCCTTC AGAATGACCT CATGGCCTC
85561 CTGGAAGAGG TCCTCCCCCA CTGCTGCCTC CACGCGCTGC CGCCATGTGG CCAGCTTGGG
85621 TCGGCCTTCG AAGACTTGGC AGCCAGCACC CACGGGCTGT GGGGAAAAGG GTACAGACTG
85681 GGGATGGATG GTTGTGAGGG CAGGGATGGG CAGCATCTGA TTTGGGGACC ACAGATCTCC
85741 AGGAGGTGTT TGCACACACA CTTAAGCACA GTGCCATAGC CCGGTGTGGC AGCATAAGCA
85801 GGA
```

#### Example: Sequencing results

CAD146: +/-

#### *GSTT1* + genotype



***GSTT1* - genotype**

## GSTT\*0 forward

```

6061 GGCCTCCCAA AGTACTGGGA TTACAGTTGT GAGCCACCGT ACCCGGCCTT CTGTGGGCCT
6121 TTTGACGCTG AGATTCTCTA GTTCAGAATG AAATGCCTCG AATGCTGTCC TGGGTGAGTC
6181 ACATCAGCCC CTCCCGTATG CCCTTCCCCT CGCCCTAATA AGACTCTTTC ATGCCCATTG
6241 TTTGAGTCCA CACTCCTGAC GGCTCTGTAA GGCGGGCAGG AGAAGGAGCT GAGGAAATGA
6301 GGCCAGAGA GGGAGGGAGA CTTGCTTGAG GTCGCCGCA GTCAGCAGGG CCAGAAGTGG
6361 CCTCCAGCCT TCCTGACTTC CCTGTGCCCG TGTCCCAAG CCCAGAAGT GGCCCTGCTC
6421 ACCTTGAGCC AGACTATCTT CTTTCATGGGC AGCTGAAAGG CCGTGTGTTG CCAAGCCACG
6481 AACTCATCCA CACGGGCACG TCGTGCAGGG TCTGGCGGGC ACCAGTGCGA TGGTGCCTG
6541 TACTTGCGGC ACAGGTAGTA AAGGATGGCC GCGCTGCAGA AGGGGCCGGT CAGGGGCACT
6601 GCCCTTGCCT TCCTGAGTGC CACTACATCA ACCACCCCGG TGTGGCCTGG GCCCAACTGC
6661 TGGGGCTTCC AGAGCAAAGA GGAGCCAAA CGGCCCCGAG AAAGACCTTC ACCAGAGCTG
6721 TCTGTCTGAC AGTCAGTAAG GGCTGGGAAG GAGCCCTGCG GGGTGTAGTAG GAGTTGGGGG
6781 CTGGTGGTAT AACAAAGAGT AGGCCAGCAG GGGGAACAAC ACGTGTGAA TTGGGATGCT
6841 GAGGTGGGAG GATCACTTGA TCCCAGGAAT TTGGGGCTAC TGTGAGCCAA GATCACACCA
6901 CTGCACTCCA GCTTGGGTGA AAGATCAAGA TCCTTTTTCA AAAACAAAAA CGGGGGGGCA
6961 CGATGGCTCA CACCTGTAAT CCCGGCACTT TGGGAGGCCA ATGGGGGCAG ATCCCTTGAG
7021 GCCAGGAGTT GGAGACCAGC CTGGCCAACA TGGTGAAACC CTGTCTCTAC TAAAATGAAA
7081 ATACAAAAAT TAGCTAGTTG TGGTGGCACA CACCTGTAAT CCCAGCTACT TGGGAAGCTG
7141 AGGCACGGGA GTCACCTGAA CCTGGGAGGC AGAGGTTGTA GTGAGCCAAG ATTGTGCCAC
7201 TGTACTCCAG CCTGGGCCAC AGAGCAAGAC TCTGTCTCAA AAAACCAACA AAGAAAAACA
7261 CATGCTGAAA TACGAGGGTA AAGGGAGCAA GGTAATCTG AAGAAAAGAG AGTAGGGGGT
7321 TGCAACTGGA AGAAGGGTGG GGGTGTATTG GGAGTGATGA GGCAGCCAGA GACTGTGG
7381 AGTCCACGGA GGGTAGCCCC TGGAGGTGCA GGGAGGTTAT GGAATTAATG CTTAAGATTA
7441 GGCATTATAT AAGCCAGGGC ATGAAAGGAT CCATCTCTCT GGTGCTGGAT GGAGGGTGAG
7501 CCCGAGGGGG CAGAATGGAC AATGAGGGGG CCAGCAACTA TCGGGAAGGT TGTGGTGTCT
CTCCCCC GGTCGTTGAT AGC GSTT*0 reverse

```

Example: Sequencing results

CADA033: -/-

***GSTT1* - genotype**

80 290 300 310 320 330 340 350 360 370 380 391  
G T G T C C C C A A G C C C C A G A A G T G G C C C T G C T C A C C T T G A G C C A G A C T A T C T T C A T G G C A G C T G A A A G G C C G T G T G T G C C A A G C C A C G A A C T C A T C C A C A C G G G C A C G

