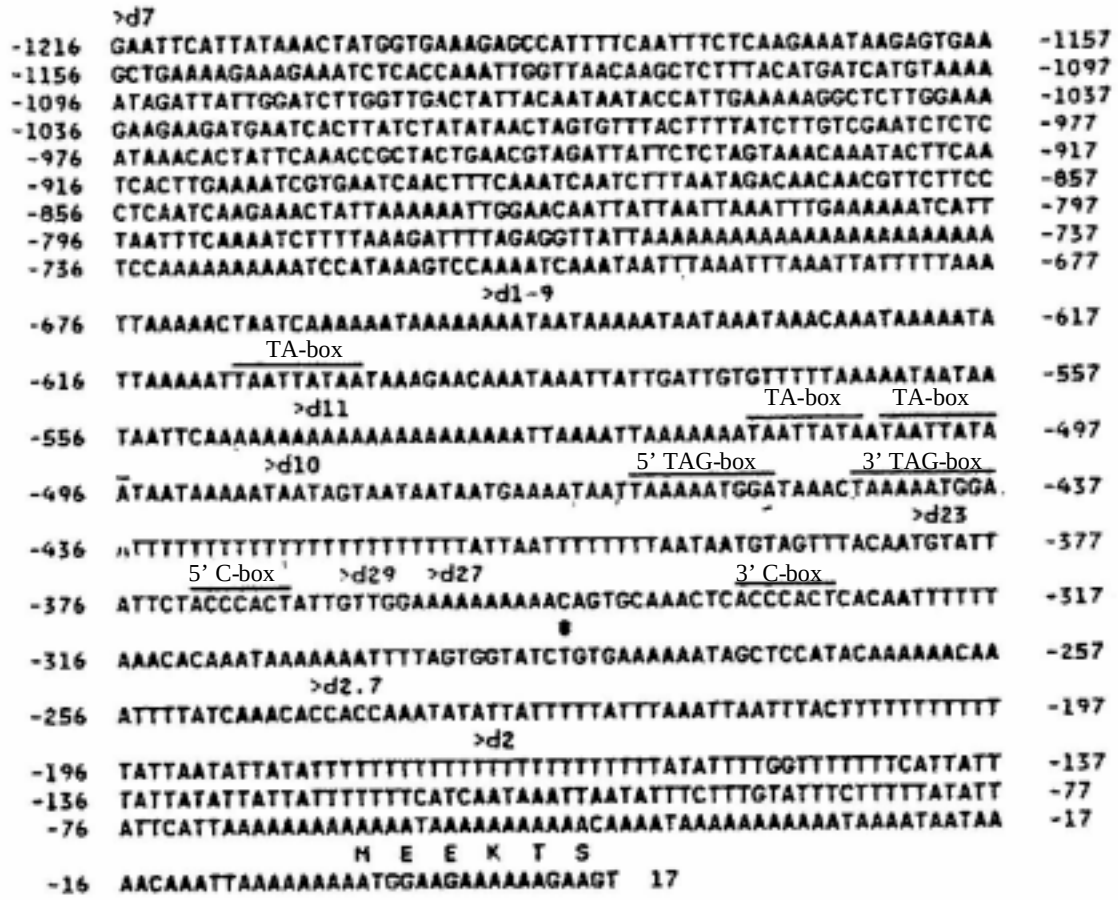
>d7
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-1156 GCTGAAAAGAAAGAAATCTCACCAAATTGGTTAACAAGCTCTTTACATGATCATGTAAAA -1097
-1096 ATAGATTATTGGATCTTGGTTGACTATTACAATAATACCATTGAAAAAGGCTCTTGGAAA -1037
-1036 GAAGAAGATGAATCACTTATCTATATAACTAGTGTTTACTTTTATCTTGTCGAATCTCTC -977
-976 ATAAACACTATTCAAACCGCTACTGAACGTAGATTATTCTCTAGTAAACAAATACTTCAA -917
-916 TCACTTGAAAATCGTGAATCAACTTTCAAATCAATCTTTAATAGACAACAACGTTCTTCC -857
-856 CTCAATCAAGAAACTATTAAAAAATTGGAACAATTATTAATTAAATTTGAAAAAATCATT -797
-796 TAATTTCAAAATCTTTTAAAGATTTTAGAGGTTATTAAAAAAAAAAAAAAAAAAAAAAA -737
-736 TCCAAAAAAAAAATCCATAAAGTCCAAAATCAAATAATTTAAATTTAAATTATTTTTAAA -677
>d1-9
-676 TTAAAAACTAATCAAAAAATAAAAAAAATAATAAAAATAATAAATAAACAAATAAAAATA -617
TA-box
-616 TTAAAAATTAATTATAATAAAGAACAAATAAATTATTGATTGTGTTTTTAAAAATAATAA -557
>d11
TA-box TA-box
-556 TAATTCAAAAAAAAAAAAAAAAAAAAAAATTAAAATTAAAAAAATAATTATAATAATTATA -497
>d10
5' TAG-box 3' TAG-box
-496 ATAATAAAAATAATAGTAATAATAATGAAAATAATTAAAAATGGATAAACTAAAAATGGA -437
>d23
-436 TTTTTTTTTTTTTTTTTTTTTTTTATTAATTTTTTTTAATAATGTAGTTTACAATGTATT -377
5' C-box >d29 >d27 3' C-box
-376 ATTCTACCCACTATTGTTGGAAAAAAAAAACAGTGCAAACTCACCCACTCACAATTTTTT -317
-316 AAACACAAATAAAAAAATTTTAGTGGTATCTGTGAAAAAATAGCTCCATACAAAAAACAA -257
>d2.7
-256 ATTTTATCAAACACCACCAAATATATTATTTTTATTTAAATTAATTTACTTTTTTTTTTT -197
>d2
-196 TATTAATATTATATTTTTTTTTTTTTTTTTTTTTTTTATATTTTGGTTTTTTTCATTATT -137
-136 TATTATATTATTATTTTTTTTCATCAATAAATTAATATTTCTTTGTATTTCTTTTTATATT -77
-76 ATTCATTAAAAAAAAAAAAAATAAAAAAAAAAAACAAAATAAAAAAAAAAATAAAATAATAA -17
M E E K T S
-16 AACAAATTAAAAAAAAATGGAAGAAAAAAGAAGT 17

Figure 2. Nucleotide Sequence of the *gp2* Promoter

The 5' non coding sequence of the *gp2* promoter has been presented here. Repeat sequences called TA-, TAG- and C-boxes have been overlined. The transcription start site has been indicated with a # symbol. The positions of the nucleotides have been calculated based on the translation start site (ATG). Numbers above the sequence preceded by >d refer to the names of fragments used in the deletion analysis [2].