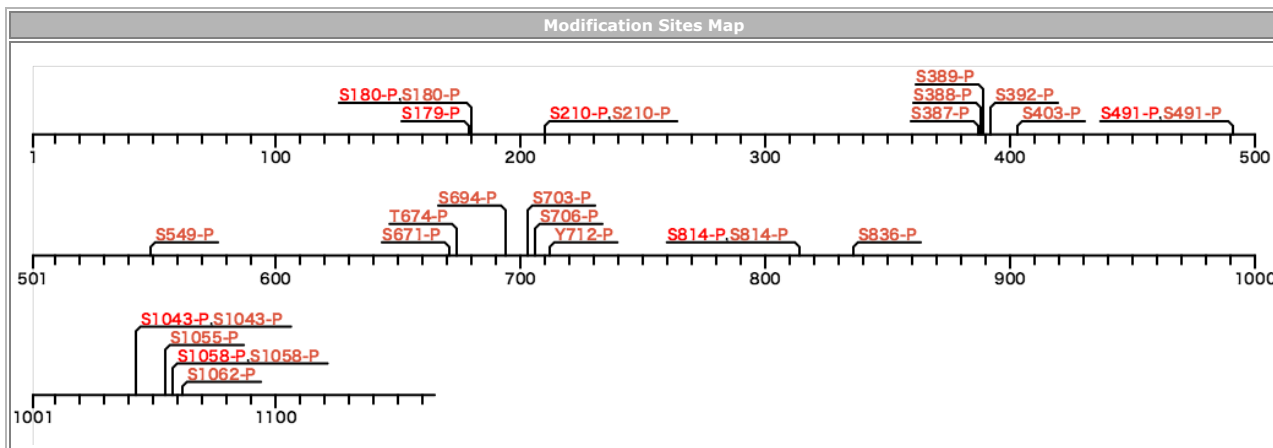


| ID | Accession | GeneName | Chr.No. | | Description |
|------------|-----------|----------|---------|----------------------|--------------------------|
| AFF4_HUMAN | Q9UHB7 | AFF4 | 5q31.1 | 132211071..132299326 | AF4/FMR2 family member 4 |



Click a modification site to display information in detail.

| Site no | Amino acid | Type | Division | Detail |
|---------|------------|------|----------|--------|
|---------|------------|------|----------|--------|

| Protein Sequence | |
|------------------|--|
| MNREDRNVL | RKERERRNQE IQQGEDAFPP SSPLFAEPYK VTSKEDKLS |
| W | DSPSRVFSS GQHSTQSFPP SLMSKSNMQL QKPTAYVRPM DGQESMEPKL SSEHYSSQSH GNSMTELKPS SKAHLTKLKI PSQPLDASAS GDVSCVDEIL KEMTHSWPP |
| P | LTAIHTPCKT EPSKFPFPTK ESQSNFGTG EQKRYNPSKT SNGHQSKSML KDDLKLSSE DSDGEQDCDK TMRSTPGSN SEPSHHNSEG ADNSRDDSSS HSGSESSSG |
| S | DSESESSSD SEANEPSQSA SPEPEPPTN KWQLDNWLNK VNPCHKVSPAS SVDSNIPSSQ GYKKEGREQG TGNSYDTSG PKETSSATPG RDSKTIQKGS ESGRGRQKS |
| P | AQSDSTTQRR TVGKKQPKKA EKAAAEPRG GLKIEETPV DLASSMPSSR HKAATKGSRK PNIKESKSS PRPTAEKKKY KSTKSSQKS REIETDTSS SDSDESESLP P |
| SSQTPKYPE | SNRTPVKPS VEEEDSFRQ RMFSPEEKE LLSPLSEPDD RYPLIVKIDL NLLTRIPGKP YKETEPKGE KKNVPEKTR EAQKQASEKV SNKGRKRHKH EDDN |
| RASESK | KPKTEDKNSA GHKPSNRES SKQSAAKEKD LLSPAGPVP SKDPKTEHGS RKRTISQSSS LKSSNSNKE TSGSSKNSSS TSKQKTEGK TSSSSKEVKE KAPSS |
| SSNCP | PSAPTLDSK PRRTKLVFDD RNYADHYLQ EAKKLKHNAD ALSDRFEKAV YYLDVAVSFI ECGNALEKNA QESKSPFPMY SETVDLIKYT MKLKNYLAPD ATAADKRL |
| TV | LCLRCESLY LRLFLKKN ALKYSKLTLE HLKNSYNSQ APSPLGSKA VGMPSPVSPK LSPGNSGNYSGASSASASG SSVTIPQKIH QMAASYVQVT SNFLYATEI |
| DQAEQLSKEQ | KEFFAELDKV MGPLIFNASI MTDLVRYTRQ GLHWLRQDAK LI |

Backcolor of amino acid : Yellow -> site of modification, gray -> in front of processing