# Locus 395 (rs2075746)

SNiPAcard

## Block annotations

### Block info

| | |
|---|---|
| genomic range | chr17:34,325,532-34,325,532 |
| block size | 1 bp |
| variant count | 1 variants |

### Basic features

| Conservation/deleteriousness | | Linked genes | |
|---|---|---|---|
| phyloP | -0.706 | gene(s) hit or close-by | CCL15, CCL15-CCL14, RP11-104J23.1 |
| phastCons | 0 | eQTL gene(s) | – |
| GERP++ | 1.62 | potentially regulated gene(s) | – |
| CADD score | 5.599 | disease gene(s) | – |

## Putative effect on regulation

### Variation proximal to gene

| gene | variant type | min(distance) | transcript | RefSeq id | protein | variant(s) |
|---|---|---|---|---|---|---|
| CCL15 | upstream gene variant | 126 | ENST00000614368 | ? | ENSP00000484262 | 1 |
| RP11-104J23.1 | upstream gene variant | 2923 | ENST00000619334 | ? | ? | 1 |

## Putative effect on transcript

### Intron variant

| gene | affected transcript | RefSeq id | protein | variant(s) |
|---|---|---|---|---|
| CCL15 | ENST00000617897 | NM_032965.4 | ENSP00000484078 | 1 |
| CCL15-CCL14 | ENST00000616694 | ? | ENSP00000481402 | 1 |
| CCL15-CCL14 | ENST00000610751 | ? | ENSP00000481940 | 1 |