

Block annotations

Block info

| | |
|----------------------|---------------------------------------|
| genomic range | chr17:34,366,820-34,375,832 <i>e!</i> |
| block size | 9,013 bp |
| variant count | 4 variants |

Basic features

| Conservation/deleteriousness | | Linked genes | |
|------------------------------|--------------------------------|--------------------------------------|---|
| phyloP | $\mu = 0.167$ [-0.156 – 0.434] | gene(s) hit or close-by | - |
| phastCons | $\mu = 0.208$ [0.013 – 0.709] | eQTL gene(s) | - |
| GERP++ | $\mu = -0.199$ [-0.794 – 0] | potentially regulated gene(s) | - |
| CADD score | $\mu = 2.542$ [0.796 – 5.671] | disease gene(s) | - |

