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Manuscript Type: Basic Study

Identifying changes in Punitive Transcriptional Factor Binding Sites from rSNPs that are significantly associated with Disease or Sickness

Norman E. Buroker

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Single nucleotide changes that affect gene expression by impacting gene **regulatory** ... A rSNPs within a **transcriptional factor binding site (TFBS)** can **change** a ... are discussed with their possible relationship to **disease or sickness** in humans. ... The minor rs7589621 SNP EPAS1-A allele creates eleven unique **punitive** ...

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SNPs in the **regulatory** region of the prion protein (PRNP) gene may play an ... and VEGFA) which alter **punitive TFBS** and are **significantly** ($p < 0.05$) **associated** with ... In some cases the rSNP may increase or decrease the **TF binding** which ***Single Nucleotide Polymorphism in Transcriptional Regulatory Regions** and ...

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缺少字词: **punitive**

VEGFA SNPs and transcriptional factor binding sites associated with ...

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We conducted genetic analyses of seven **single nucleotide polymorphisms (SNPs)** in the promoter ... All the SNPs are found in **transcriptional factor binding sites (TFBS)**, and their possible role in gene ... with high altitude **sickness** in Han and Tibetan Chinese ... AMS study group, but found to be **significantly associated**.

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