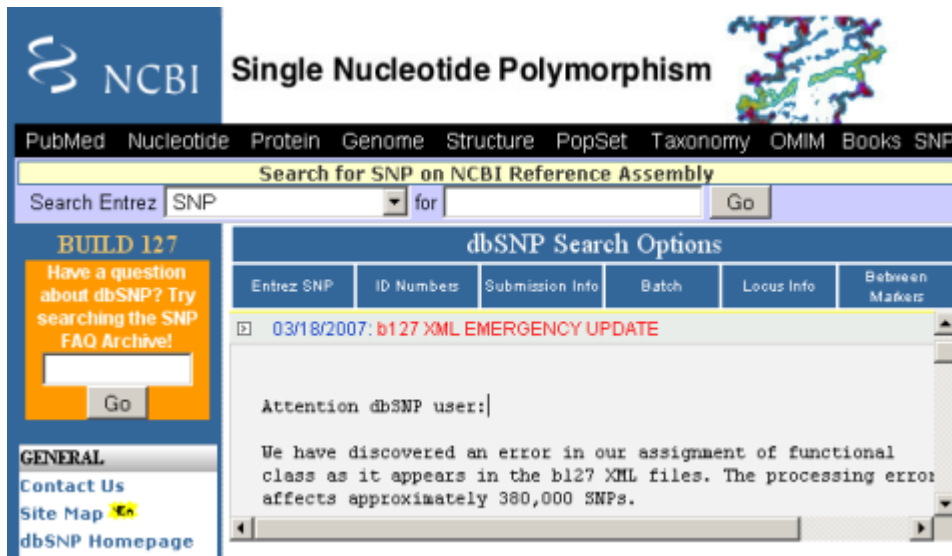


GENETICS

# DBSNP USERS ATTENTION PLEASE

14.05.2007

dbSNP has a warning that the assignment of functional class might be wrong in many SNPs:



The screenshot shows the NCBI dbSNP homepage. At the top left is the NCBI logo and the text "Single Nucleotide Polymorphism". Below this is a navigation bar with links: PubMed, Nucleotide, Protein, Genome, Structure, PopSet, Taxonomy, OMIM, Books, SNP. A search bar is present with the text "Search for SNP on NCBI Reference Assembly" and a "Go" button. Below the search bar is a section titled "dbSNP Search Options" with tabs for Entrez SNP, ID Numbers, Submission Info, Batch, Locus Info, and Between Markers. A message box is displayed with the following text:

**03/18/2007: b127 XML EMERGENCY UPDATE**

Attention dbSNP user:|

We have discovered an error in our assignment of functional class as it appears in the b127 XML files. The processing error affects approximately 380,000 SNPs.

On the left side of the page, there is a "BUILD 127" section with a "Go" button and a "GENERAL" section with links for "Contact Us", "Site Map", and "dbSNP Homepage".

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