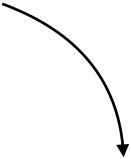
Functional analysis and allelic association



Functional analysis









Missense mutations

- Polyphen2
 - Aligns homologues and identifies conserved AAs
 - 8 sequence based,
 3 structure based features
 - Sens: 88 %
 - Spec: 62 %

- SIFT
 - Aligns homologues and identifies conserved AAs

- Sens: 85 %
- Spec: 69 %

Functional analysis and allelic association

