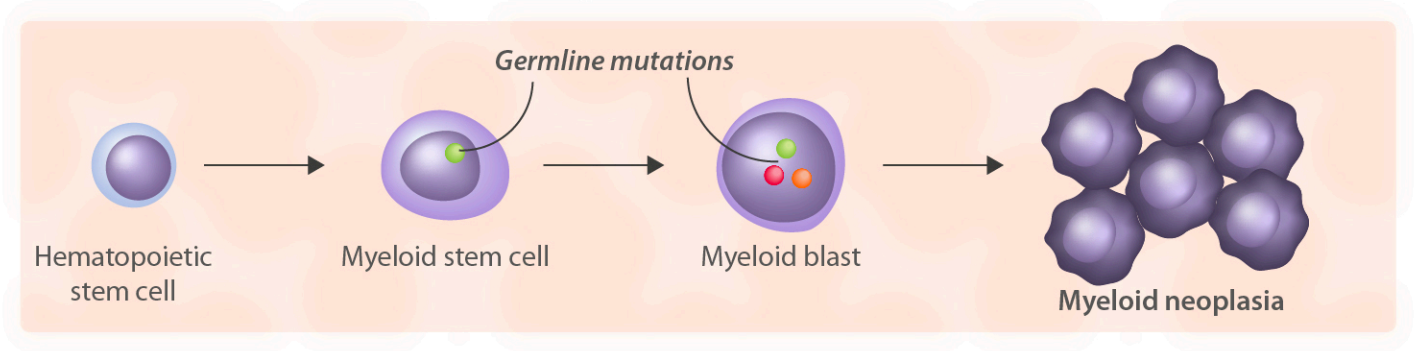


Classification of germline RUNX1 variants predisposing to myeloid neoplasms

Myeloid differentiation



Detection of germline variants

- Next generation sequencing (NGS)
- Microarrays



The 2015 American College of Medical Genetics and Genomics (ACMG) and Association for Molecular Pathology (AMP) guidelines for sequence variant interpretation

Benign (BEN)

Likely Benign (LBEN)

Variant of Uncertain Significance (VUS)

Likely Pathogenic (LPATH)

Pathogenic (PATH)