

Next-Generation Sequencing for the Detection of Somatic Mutations in Myeloid Neoplasms

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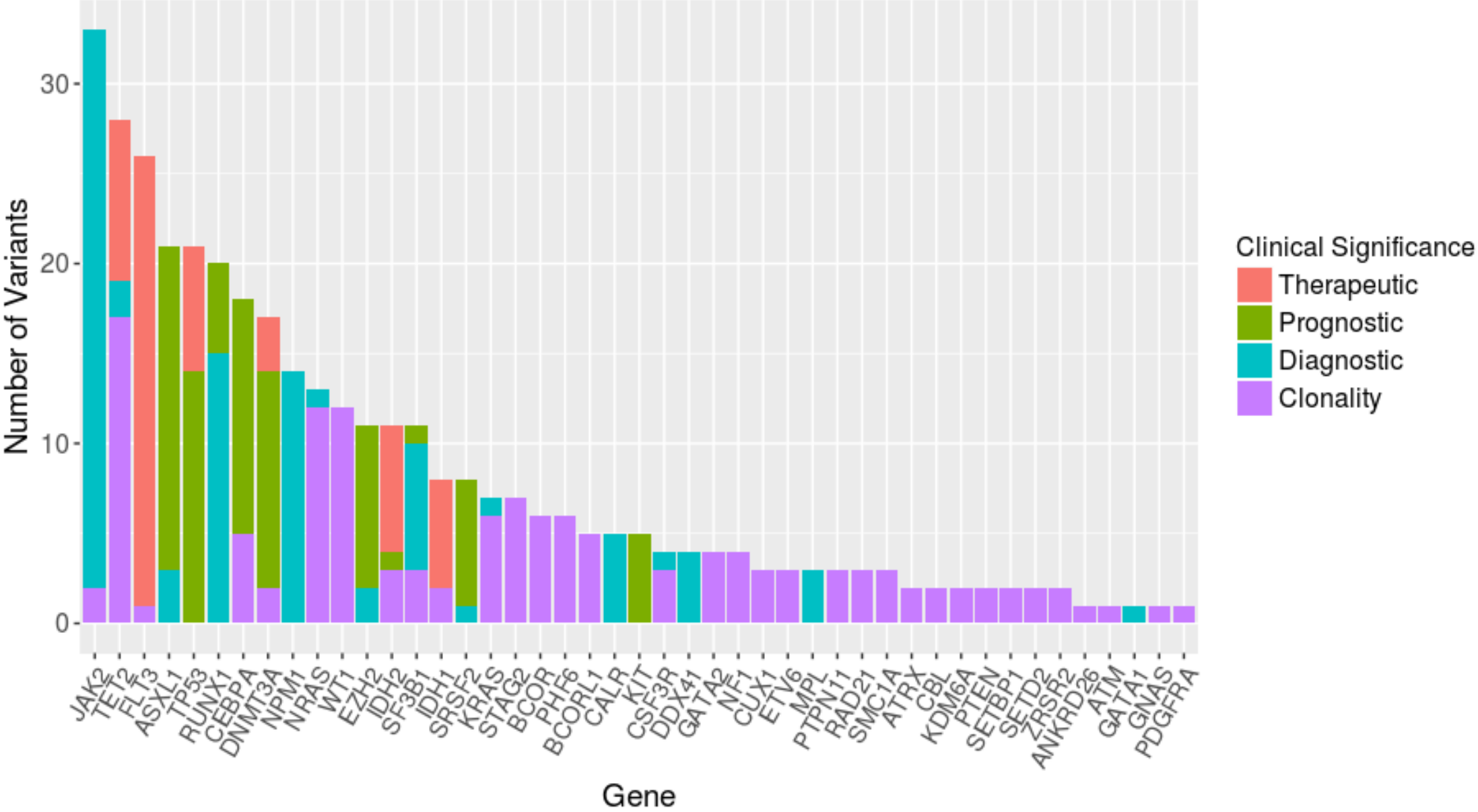
Genomic Testing for Haematological Malignancies

Myeloid Panel in Queen Mary Hospital

- 67 genes implicated to be of **diagnostic, prognostic and therapeutic significance** in myeloid neoplasms
 - Acute myeloid leukaemia (AML)
 - Myelodysplastic syndrome (MDS)
 - Myeloproliferative neoplasms (MPN)
 - Myelodysplastic/myeloproliferative neoplasms (MDS/MPN)
- **Hybridisation-based capture** approach
- **Paired-end sequencing** by Illumina MiniSeq sequencer
- Tumour sample: Peripheral blood or bone marrow aspirate
- Matched normal sample: Buccal swab



Clinical Significance of Variants by Genes in Myeloid Neoplasms



Clinical Roles of NGS Myeloid Panel

Myeloid Neoplasms with Germline Predispositions

Myeloid neoplasm classification

Myeloid neoplasms with germ line predisposition without a preexisting disorder or organ dysfunction

AML with germ line *CEBPA* mutation

Myeloid neoplasms with germ line *DDX41* mutation*

Myeloid neoplasms with germ line predisposition and preexisting platelet disorders

Myeloid neoplasms with germ line *RUNX1* mutation*

Myeloid neoplasms with germ line *ANKRD26* mutation*

Myeloid neoplasms with germ line *ETV6* mutation*

Myeloid neoplasms with germ line predisposition and other organ dysfunction

Myeloid neoplasms with germ line *GATA2* mutation

Myeloid neoplasms associated with BM failure syndromes

Myeloid neoplasms associated with telomere biology disorders

JMML associated with neurofibromatosis, Noonan syndrome or

Noonan syndrome-like disorders

Myeloid neoplasms associated with Down syndrome*

- M/75
 - AML with minimal differentiation
 - No family history
 - Normal karyotype
 - *DDX41* germline p.Gln306*
 - *DDX41* somatic p.Arg525His
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- M/57
 - MDS-EB-1
 - Father died of acute leukaemia at 60 y.o.
 - Normal karyotype
 - *DDX41* germline p.Tyr259Cys
 - *DDX41* somatic p.Arg525His
 - Referred for HSCT
 - 12 siblings. 3 HLA-matched.

Clinical Roles of NGS Myeloid Panel

Summary

- **High positive rate** (91%) for variants in myeloid neoplasms
- Most detected variants have some forms of **clinical significance**
- Facilitate detection of variants in **multiple genes simultaneously**
- **Cost reduction** per gene tested