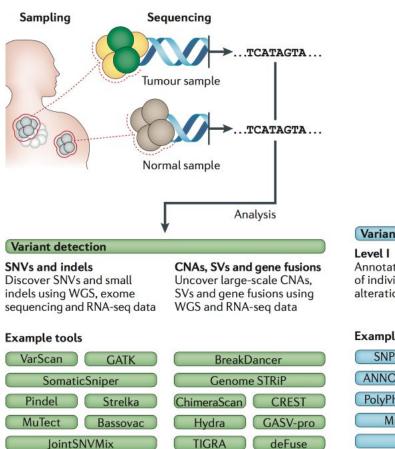
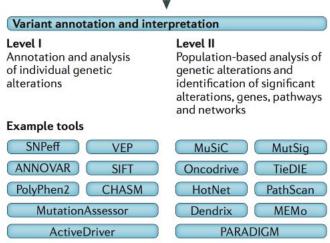
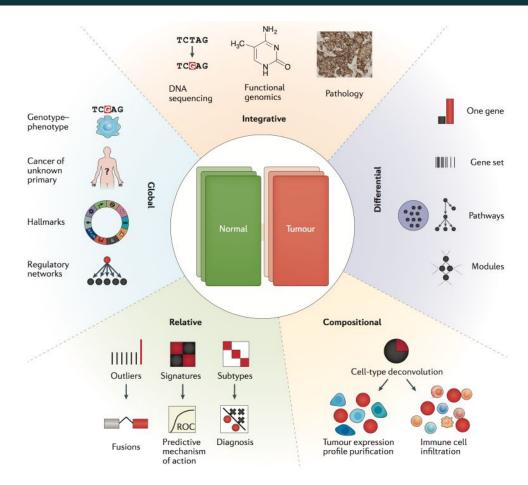
Lecture 27: Cancer genomics

Overview

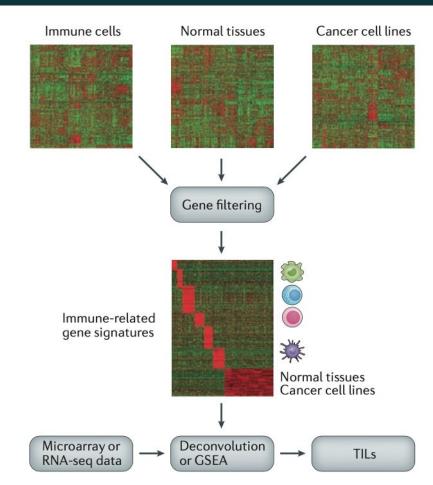




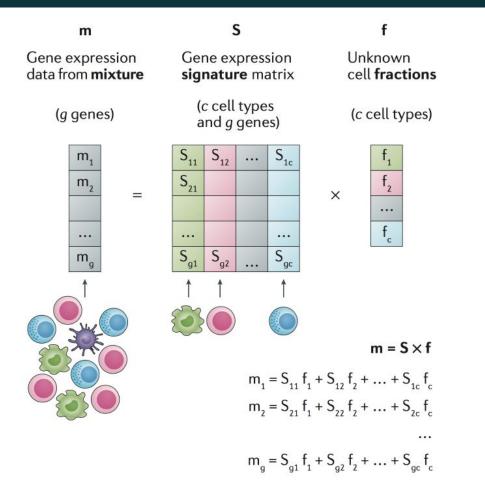
Uses of gene-expression assays in cancer genomics



Deriving cell-type-specific gene signatures

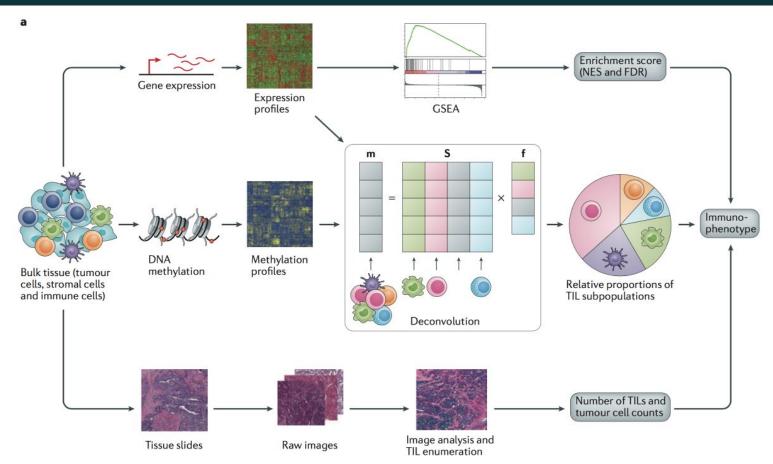


Cell-type deconvolution from bulk tissue data

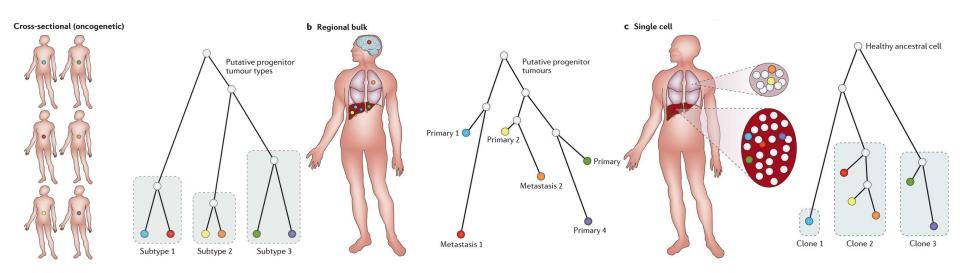


Hackl (2016) Nat. Rev. Genet.

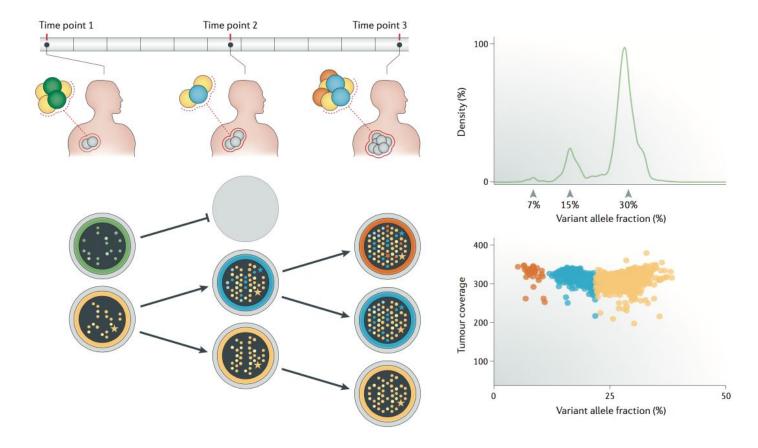
Combining these analyses



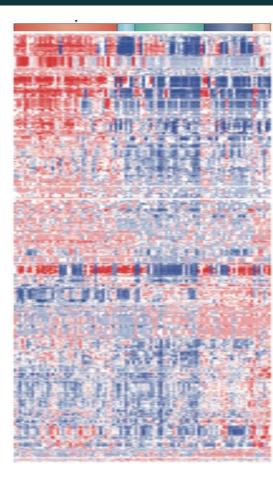
Cancer evolution



Cancer evolution



Tumor subtyping



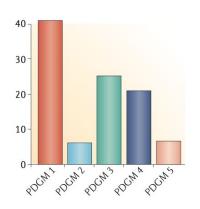
IL4 signalling Thromboxane A2 signalling IL23 signalling IL12 signalling TCR signalling

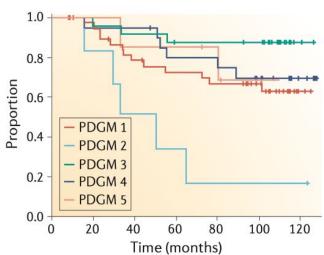
NFAT-calcineurin

transcription

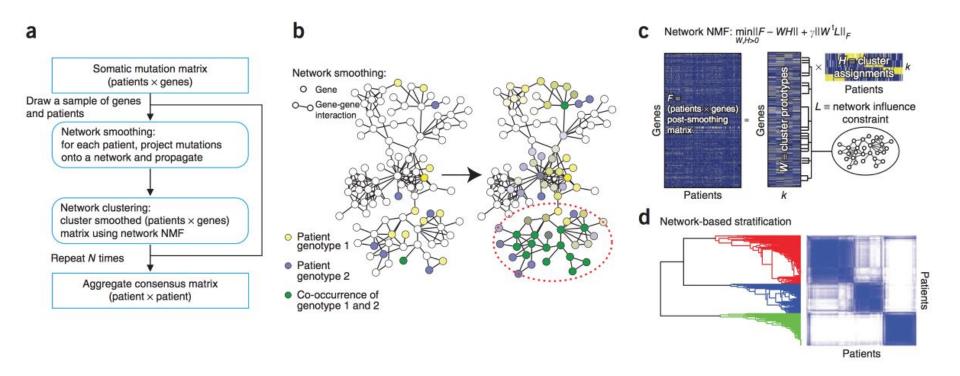
FOXM1 transcription ERBB4

Endothelins Angiopoietin receptor TIE2-mediated signalling

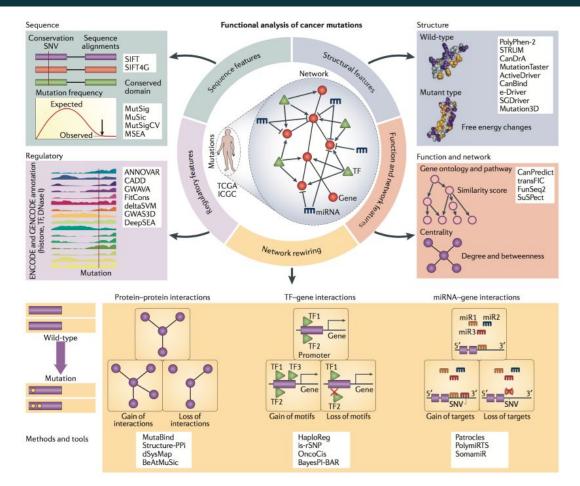




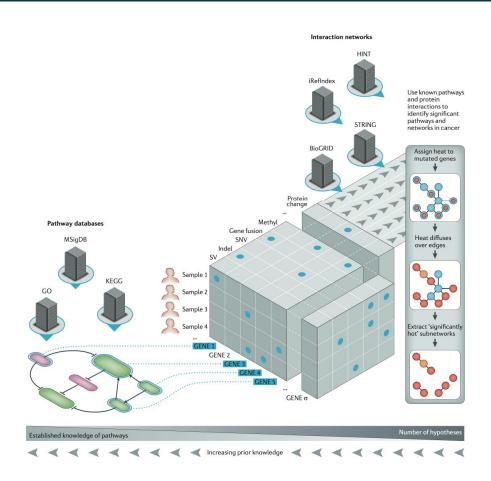
Tumor subtyping



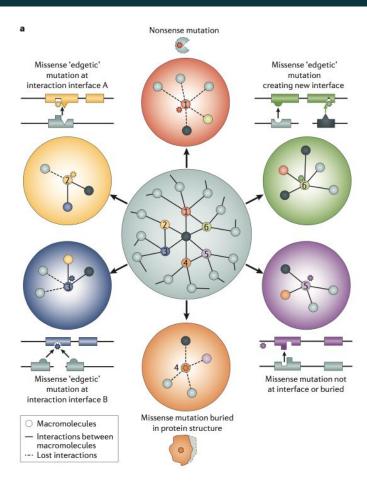
Functional analysis of cancer mutations



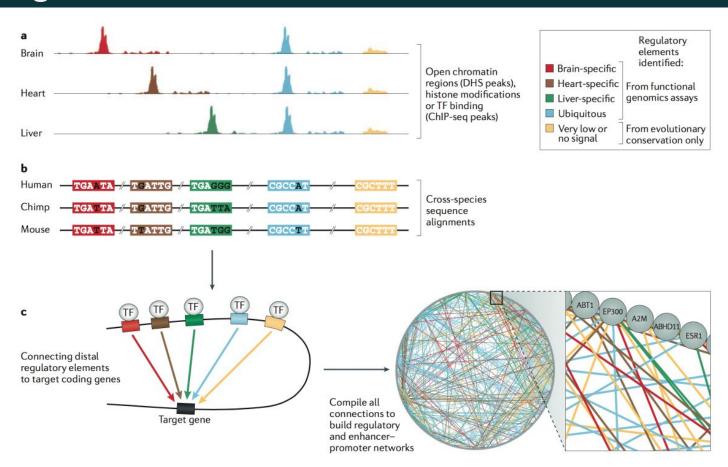
Genes, pathways, & networks



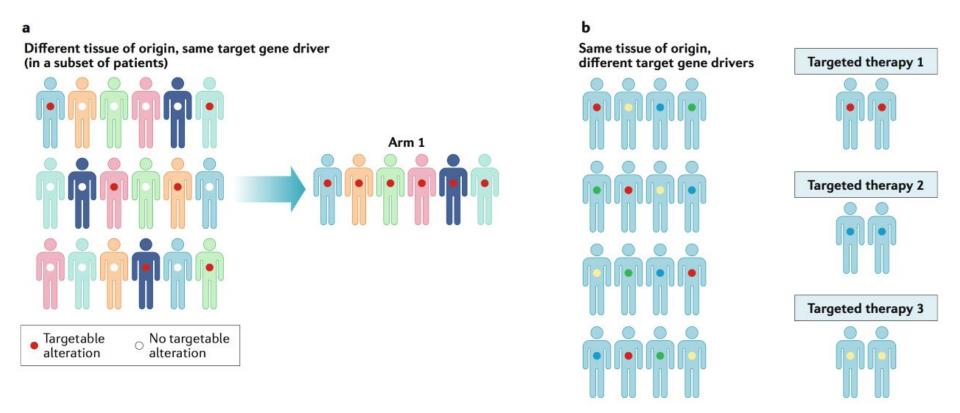
Genes, pathways, & networks



Non-coding variants



Clinical trial designs informed by genomics



Integrative data generation & analysis of breast cancer

