





Mechanisms of epigenetic dynamics

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Progression in induced by mutations in *BTK* or *PLCG2* genes



Does MRD genetically evolve?

Clonal evolution is not responible for MRD survival



MRD under ibrutinib lacks BTK and PLCG2 mutations



Does MRD epigenetically evolve?

MRD is in a predominantly closed chromatin state



Transcription factor map of MRD is rewired: NF-kB, JUN, NFAT targets decrease their accessibility



BCR signaling is downregulated under ibrutinib



What is the pathway that supports MRD?

Transcription factor map of MRD is rewired: ERK targets increase their accessibility





Is RAS-RAF-MEK-ERK active in the tissue compartment?

In vivo modeling of MRD in the tissue



TCL1 leukemia cells persisting in the spleen under ibrutinib have active RAF-RAF-MEK-ERK



Ibrutinib mice were sacrificed on days 25, 26, 26 and 29

Pre-treatment MRD

Does BCR^{low}/ERK^{high} cells pre-exist before ibrutinib start?

Single cell RNA-seq reveals a pre- existing BCR^{low}/ERK^{high}



What does activate of RAS-RAF-MEK-ERK?

MRD cells adapt their surface machinery



MRD maintains signalling via RAS-RAF-MEK-ERK





NA



Is RAS-RAF-MEK-ERK a vulnerability of MRD?

Opportunities for pharmacologicaly manipulatethe RAS-RAF-MEK-ERK pathway



ERK inhibition synergizes with ibrutinib





CDI <1 indicates synergism CDI <0.7 indicates a significantly synergistic effect

Ulixertinib in combination with ibrutinib prolongs survival



- BTK inhibition results into a large chromatin rewiring
- RAS-RAF-MEK-ERK remains active
- BCR engagement triggers RAS-RAF-MEK-ERK to sustain MRD viability
- Clones BCR^{low}/ERK^{high} pre-exist ibrutinib treatment
- RAS-RAF-MEK-ERK is a vulnerability of MRD