

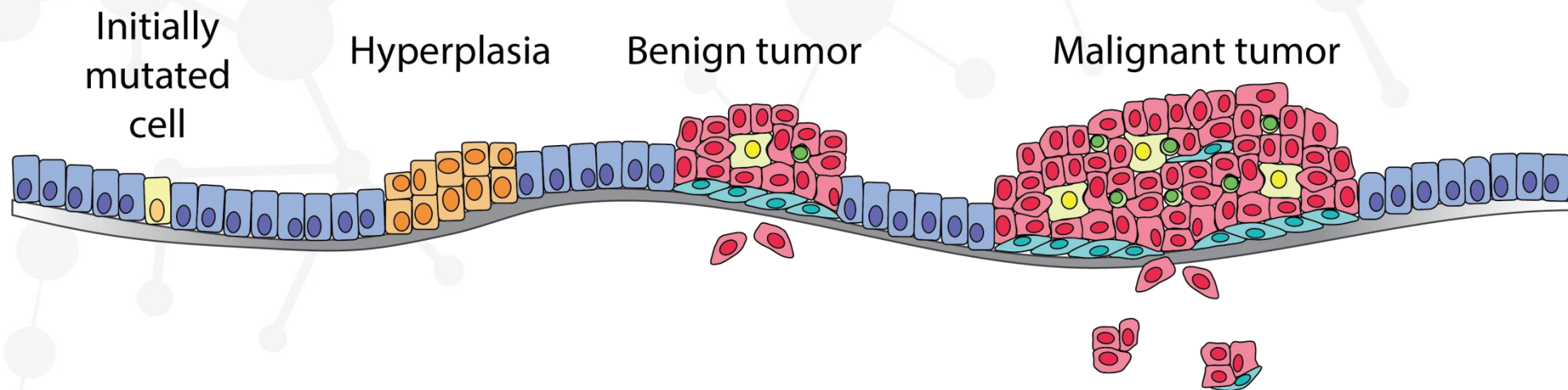
Disclosures

- Currently employed by AbbVie Inc.



Society for Immunotherapy of Cancer

Tumor development requires the acquisition of tumor-cell intrinsic and extrinsic phenotypes, both of which are driven by mutations and alterations in the tumor cell



Effect on tumor cell growth

Initial disruption of cell-cycle controls

Development of tolerance to anti-growth signals

Metabolic and proliferative adaptations. Possible early metastatic seeding. Initial immunoevasive strategies.

Acquisition of invasive and widespread metastatic potential. Fully established immunoevasive strategies.

Effect on tumor microenvironment

Establishment of growth-supportive microenvironment

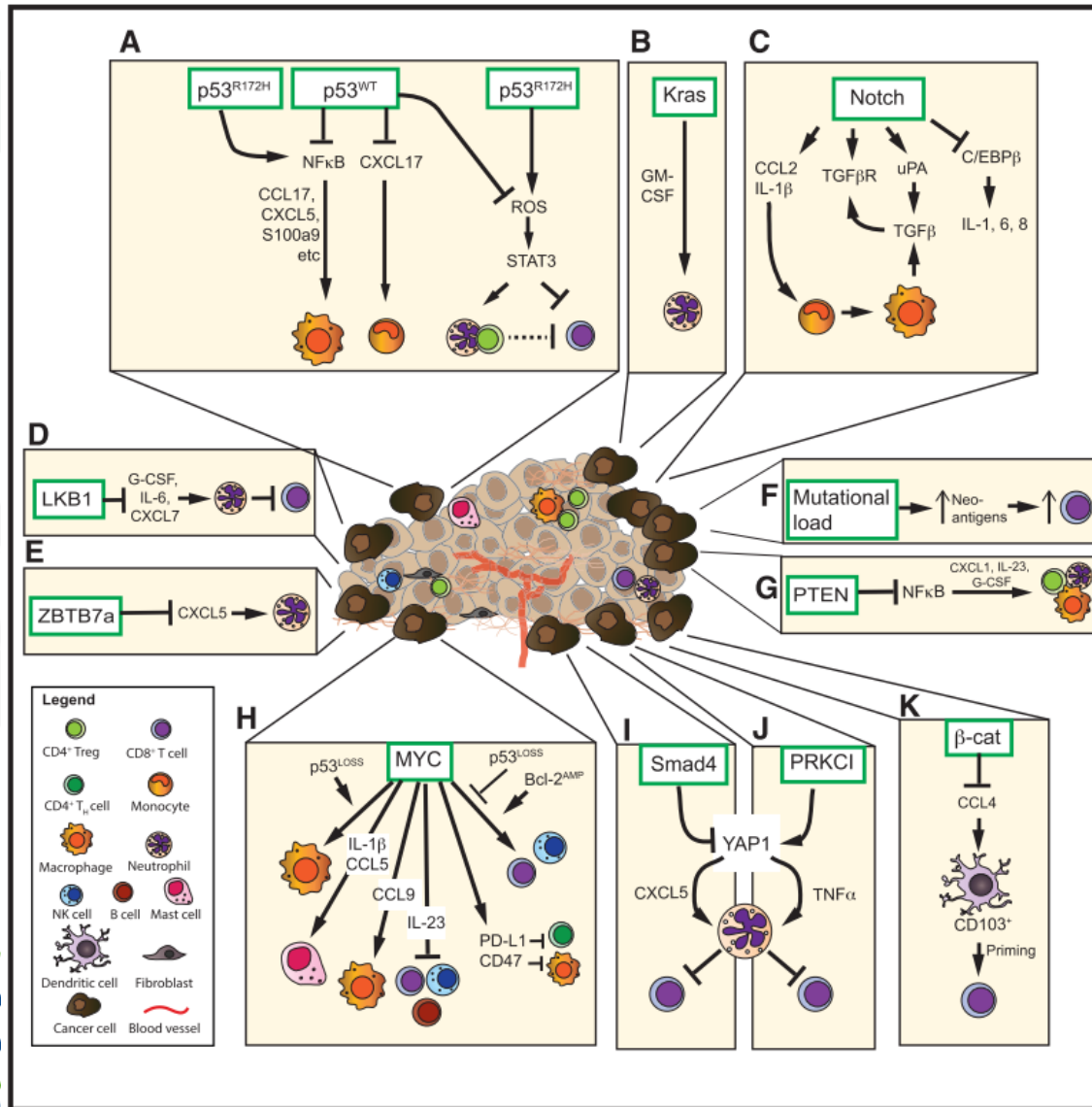
Engagement of tumor supportive and suppressive immunity. Adaptation of local non-tumor cells.

Erosion of local mechanical barriers. Conversion of distant sites into viable tumor soil. Incapacitation of local and distant anti-tumor immunity.



Society for Immunotherapy of Cancer

Specific tumor-cell alterations with clear effects on the tumor-immune context and IO response



Types of mutations interacting with the immune context:

Immunogenic drivers

Mutations driving the development of anti-tumor immunity. Includes mismatch repair deficiency, polymerase mutations.

Immuno-evasive mutations

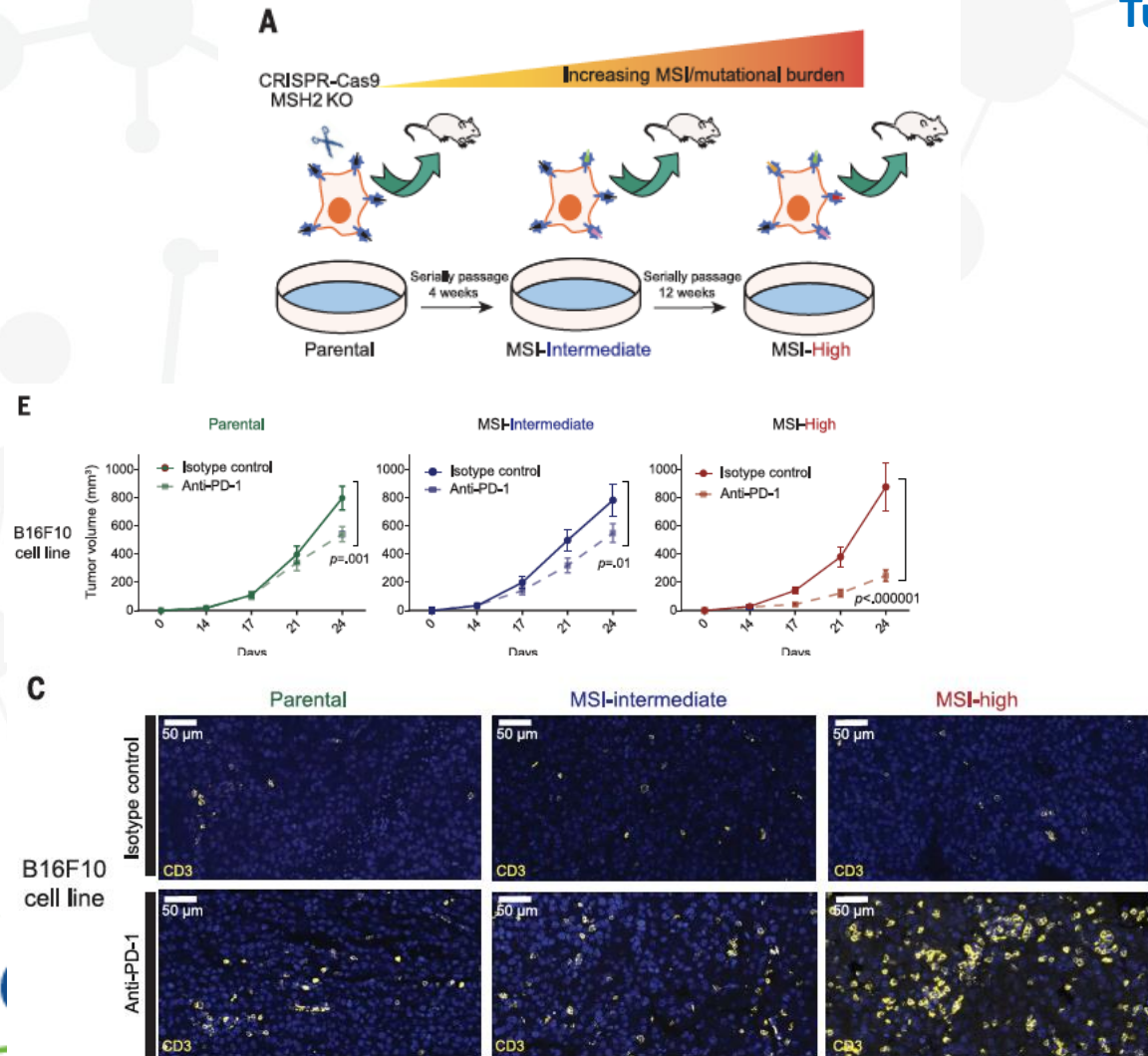
Mutations with direct effects on the ability of the immune system to eliminate tumor cells. Includes antigen presentation and processing mutations, PD-L1 amplification, and many others.

Context drivers

Mutations with clear effects on the immune context, but which may not directly impair the ability of the immune system to eliminate tumor cells. Examples include aneuploidy and LKB1/STK11, EGFR, BRAF.

Genetic diversity associated with MMR-d influences response to ICI

Tuning MSI severity with CRISPR-Cas9



- Study authors used CRISPR-Cas9 guide RNAs directed to Msh2 to knock out MSH2 in a poorly immunogenic mouse models B16F10, CT26
- Used serial passage of 1 & 4 months to model varying degrees of MSI, MSI-intermediate and MSI-high respectively
- As expected, MSI-high displayed higher non-synonymous SNVs and coding indel mutations
- MSI-high associated with increased anti-PD1 responsiveness and immune infiltration compared with parental and intermediate groups

Emerging questions from the breakout session

- How do we deal with the molecular subtypes that define intrinsic resistance to ICI and expand CIR?
- What is the relationship between primary and secondary resistance to ICI
 - What data is available to address this questions?
 - We need data from treated patients? PACT initiative.
- What would we do with this information? Could we leverage it for combinations with targeted therapies or just for exclusion criteria?
- There is a need to develop methodologies to more accurately identify key driver mutations and their connection to CIR.
- What technologies & biomarker development are needed to translate this knowledge into clinical practice?
 - How do we convince the clinicians to use the data?

