

Multiscale Models for Predicting Optimum Immune and Targeted Therapy Schedules

Goals

- We are combining multiscale mathematical approaches with novel cellular quantification experimental technologies in order to:
 - To gain a deeper, more robust understanding of tumor-immune dynamics
 - To optimize combination immunotherapy and receptor kinase targeted therapy

Aggressive Bladder Cancer Mutations

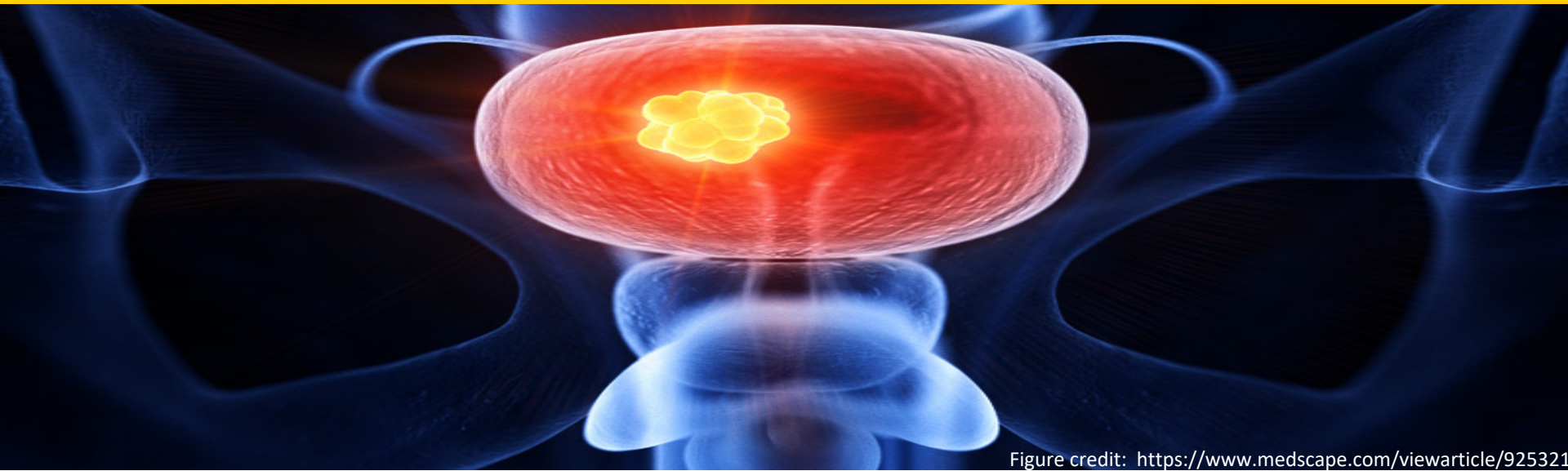
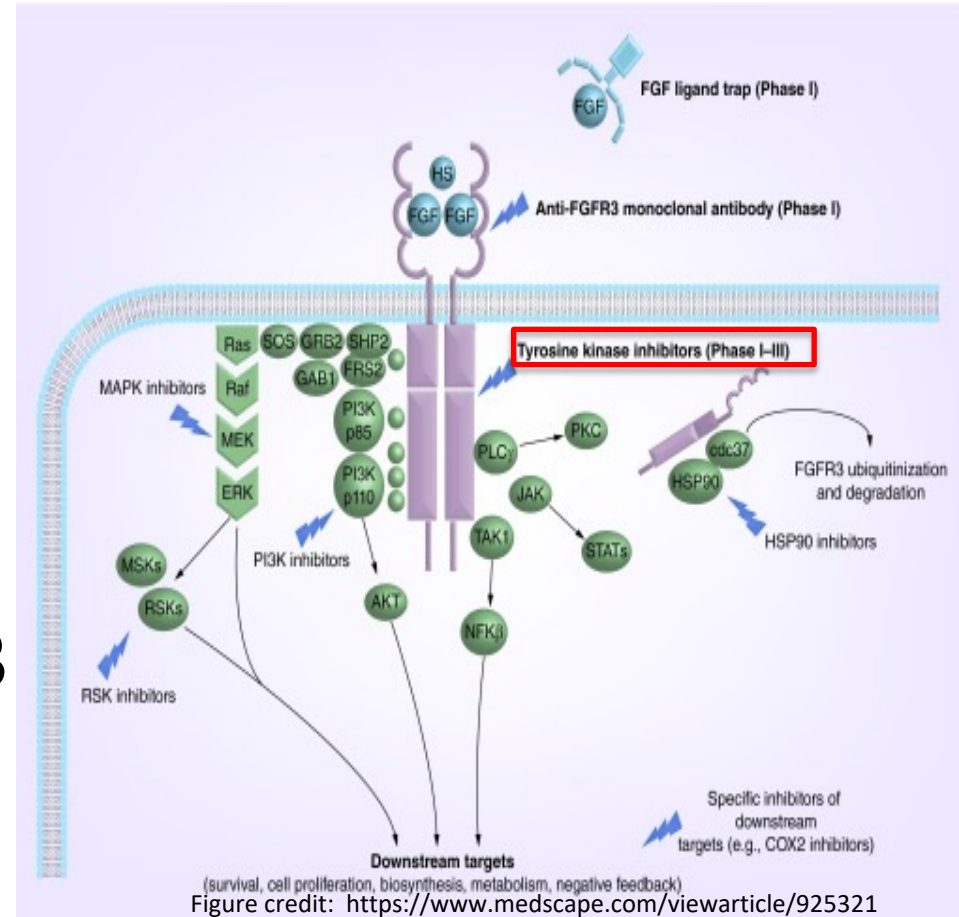


Figure credit: <https://www.medscape.com/viewarticle/925321>

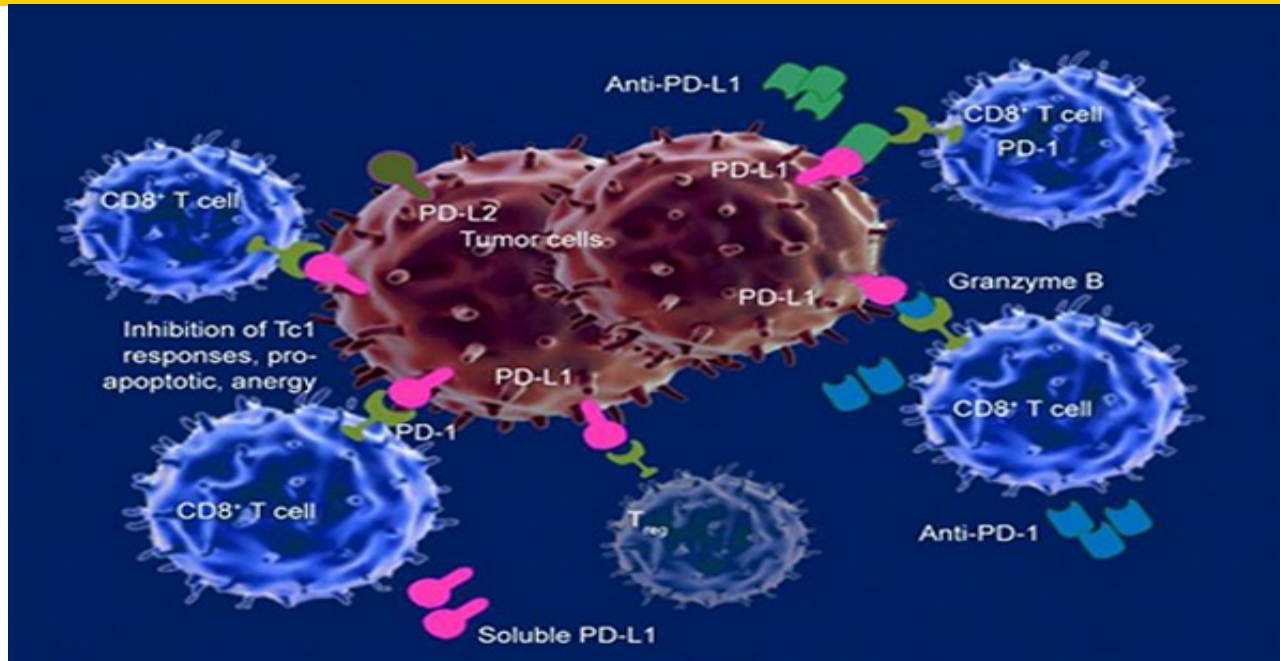
- Genomic analysis of bladder cancer has identified frequent alterations of FGFRs, including mutations of *FGFR3* that activate the receptor via ligand-independent dimerization → increased cell proliferation and survival.

Targeted Therapy

- Clinical trials using SMIs of FGFR3 are leading to promising clinical responses for patients with FGFR3 mutations.
- Last year, the FDA approved the first FGFR3 targeted therapy for bladder cancer.



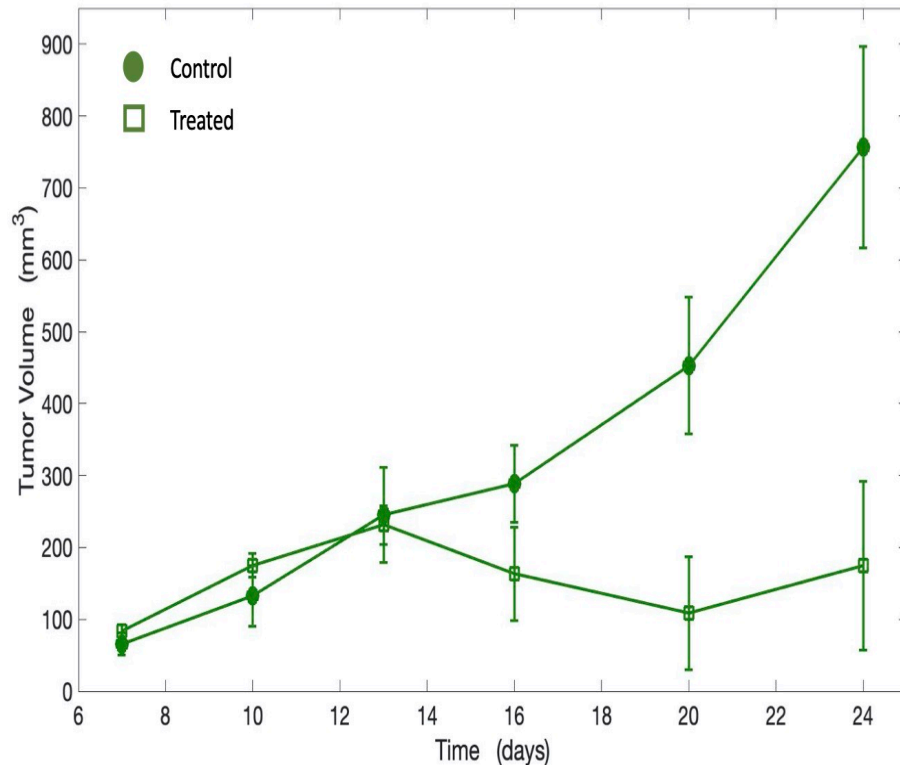
Immunotherapy



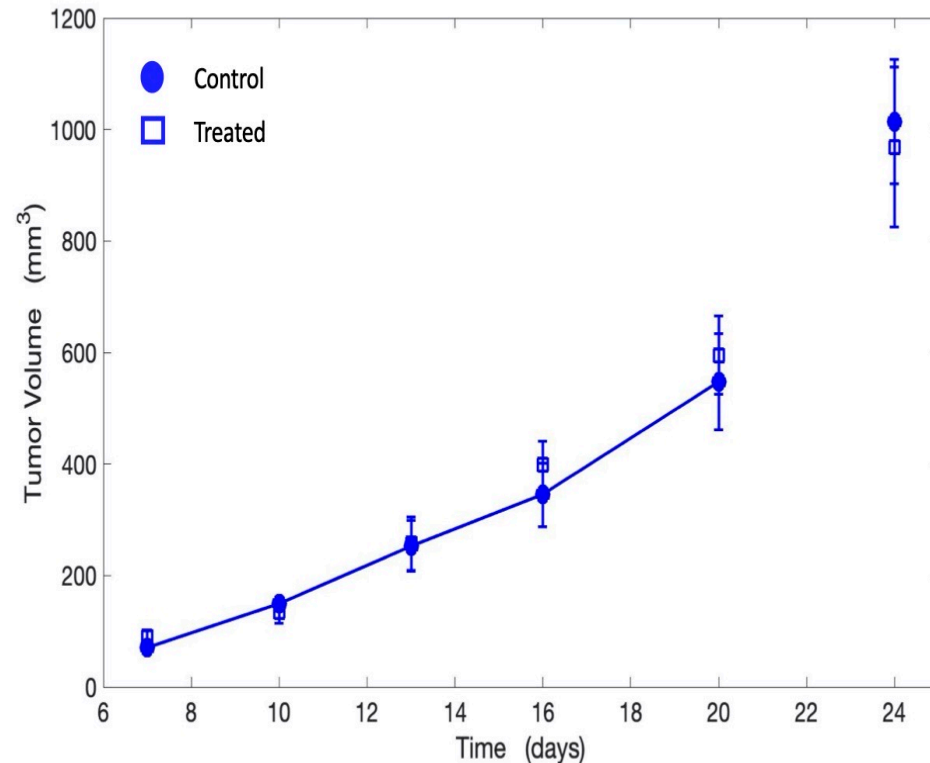
- MAbs targeting the PD-1/PD-L1 pathway have resulted in favorable outcomes in advanced bladder cancer.
- Despite the activity of these drugs in some patients, the objective response rate remains less than 25%.

Mutations Hinder Immunotherapy

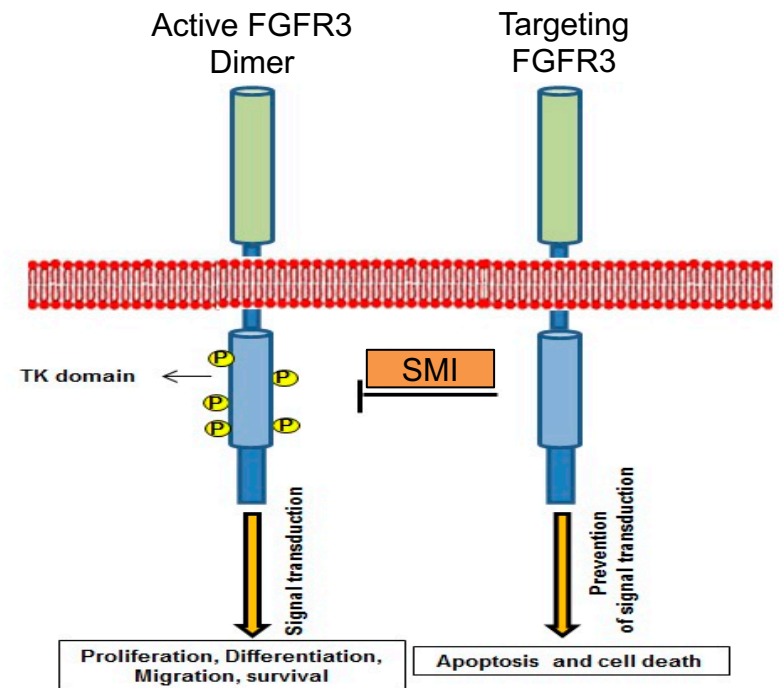
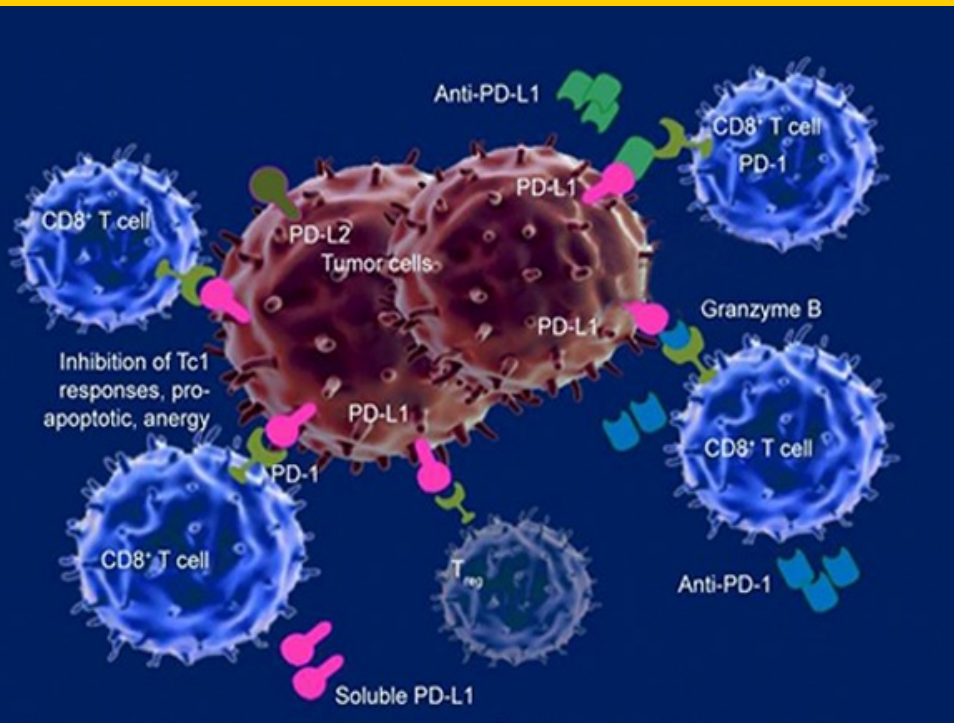
Wild Type Cells



Mutant Cells



Combination Therapy



Co-acting combination of potent immune checkpoint inhibitors and specific FGFR3 inhibitors potentially offers an advance in targeted therapeutics for cancer.

Optimizing Combination Therapy

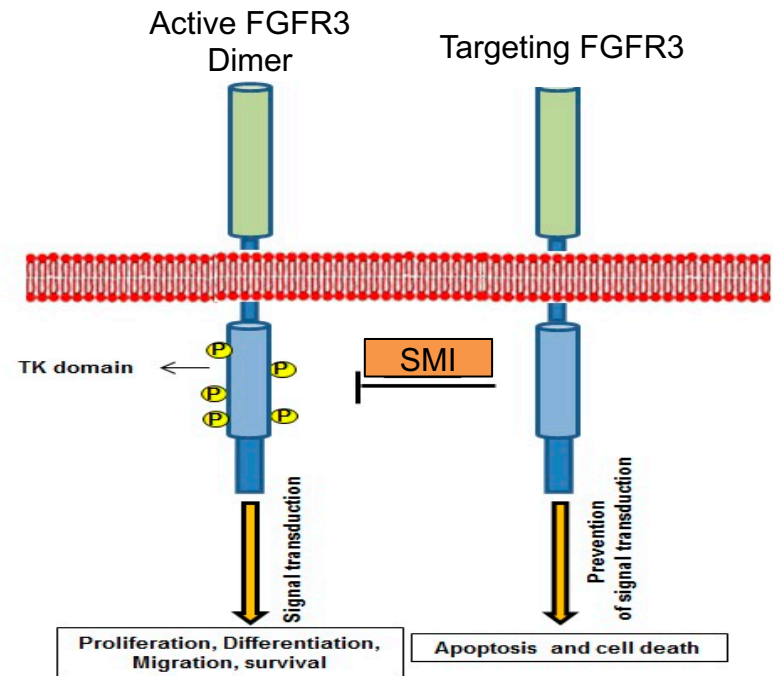
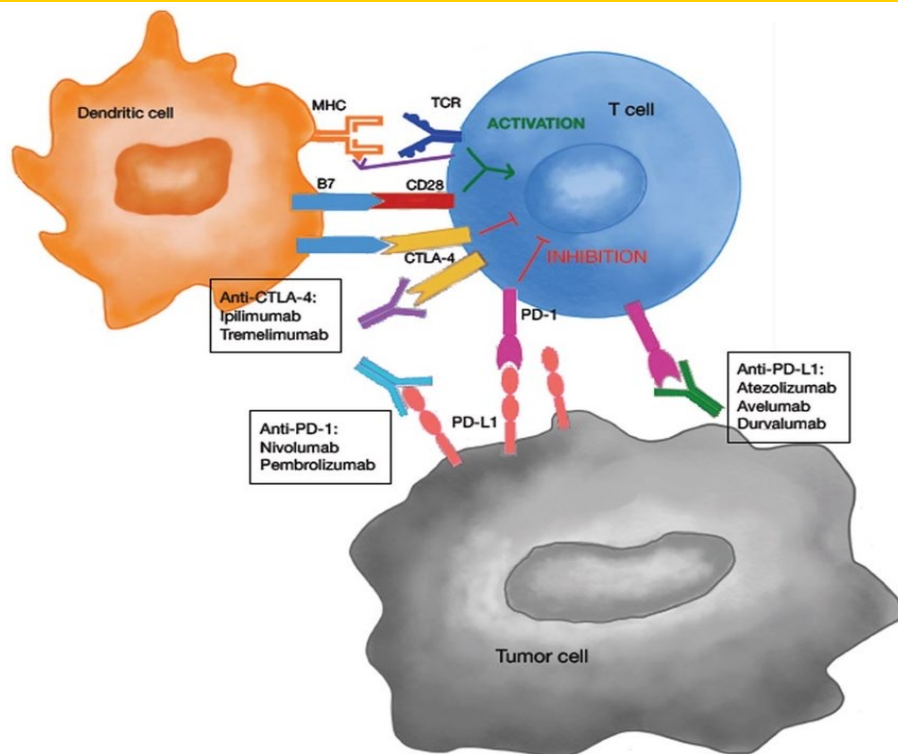
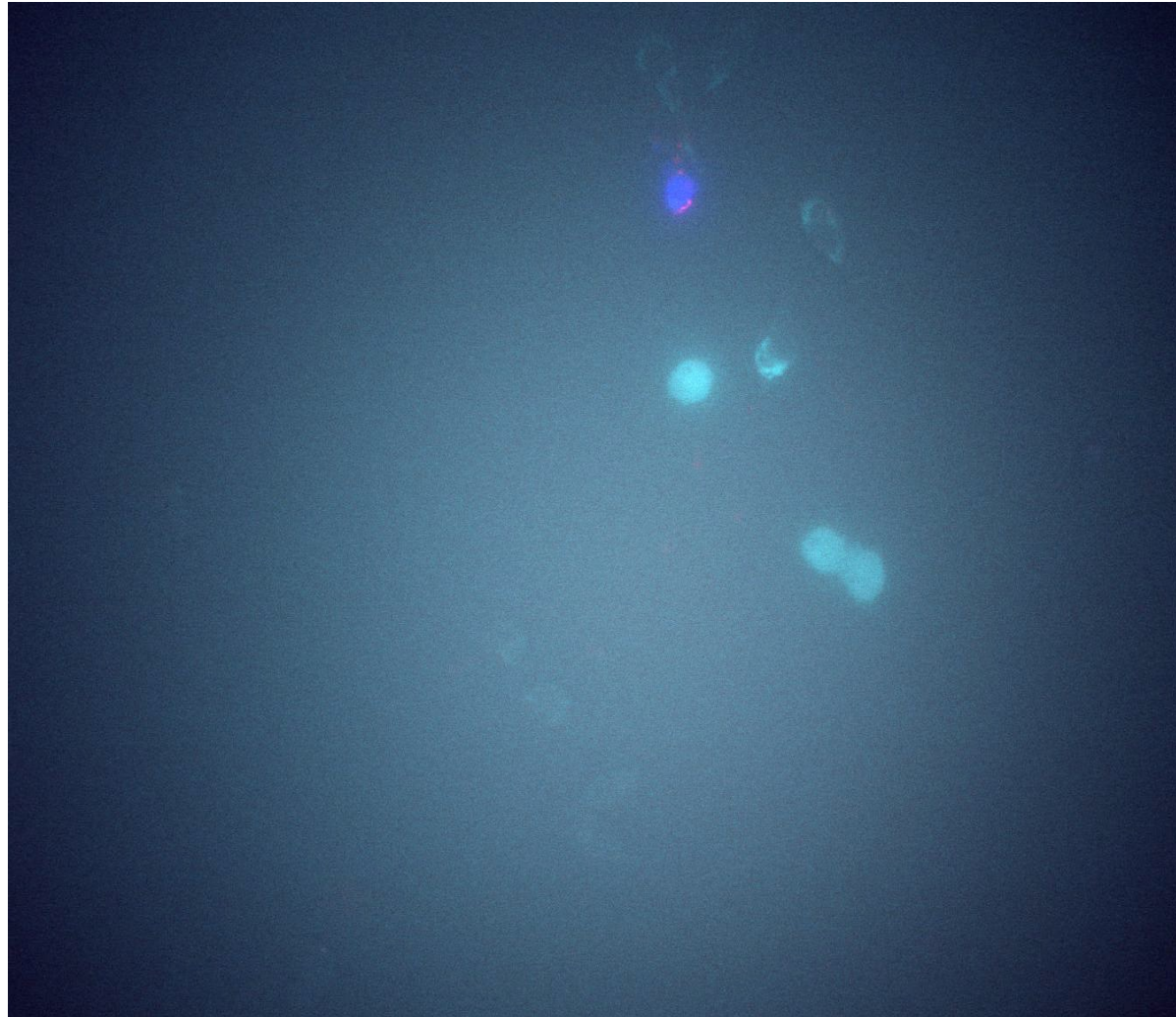


Figure Credit: Durvalumab in NSCLC: latest evidence and clinical potential. *Ther Adv Med Oncol*. 2018

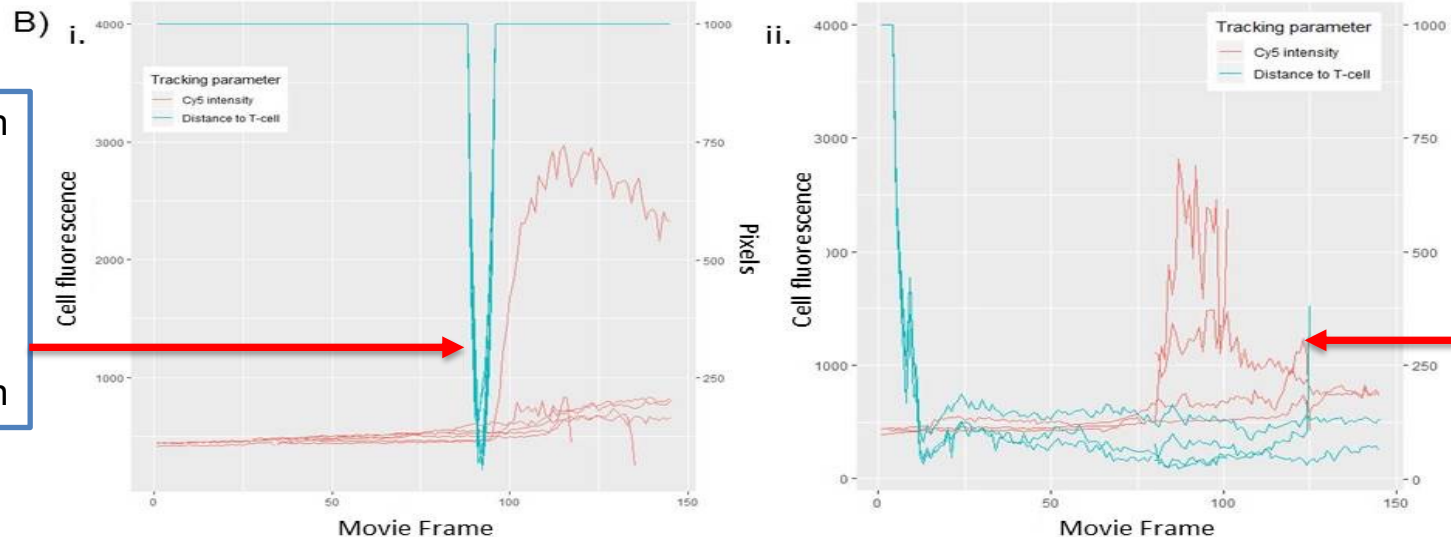
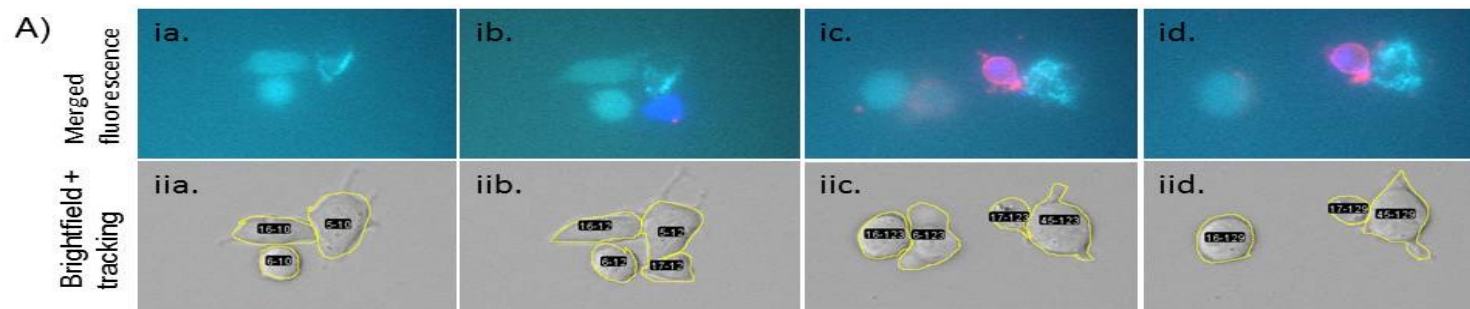
A powerful and practical way to optimize novel drug combinations for clinical cancer treatment is to use data-driven computational models.

Preliminary Data: Live Cell Tracking

- My collaborators developed a novel pipeline to track and quantify the interactions of living tumor cells and immune cells, including cell death.



Preliminary Data: Live Cell Tracking

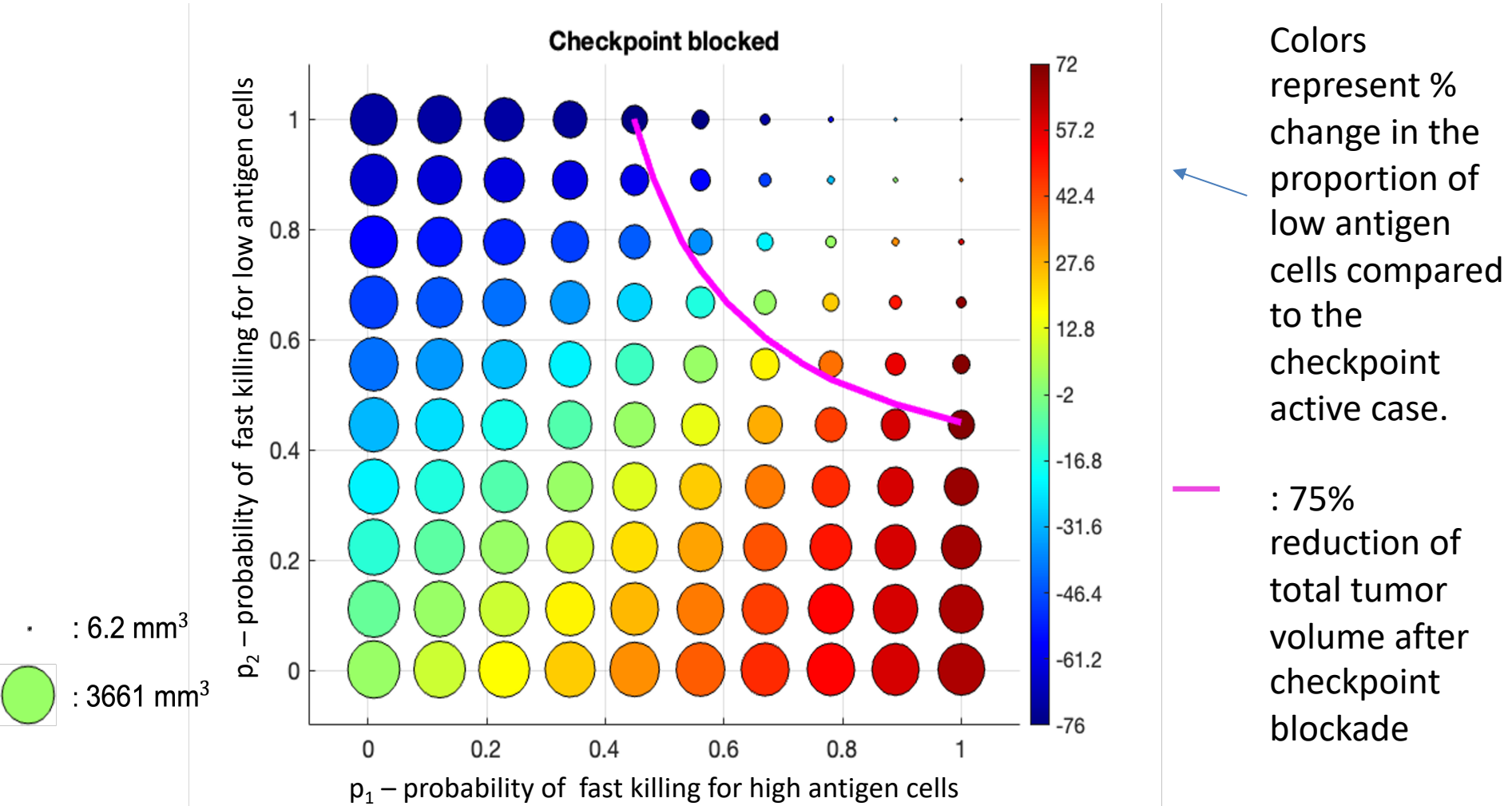


Cell death marker rapidly increases following minimal colocation

Extended cellular colocation of before cell death marker increases

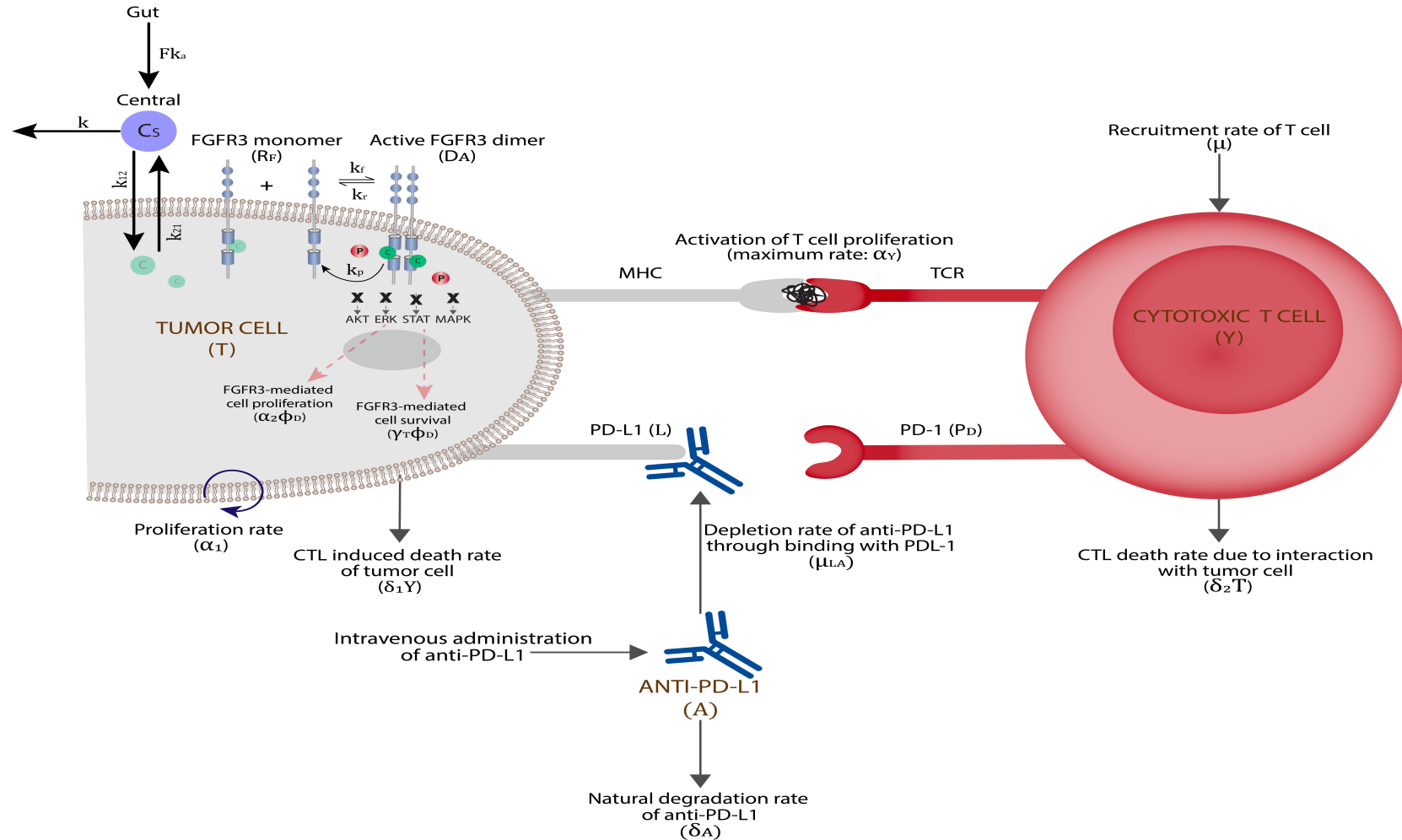
- Evidence of both rapid and slow killing during tumor-immune interactions.
- The proportion of slow and rapid killing within a solid tumor could have significant impact on immune mediated anti-cancer effects.

Impact of Fast/Slow Killing Probabilities



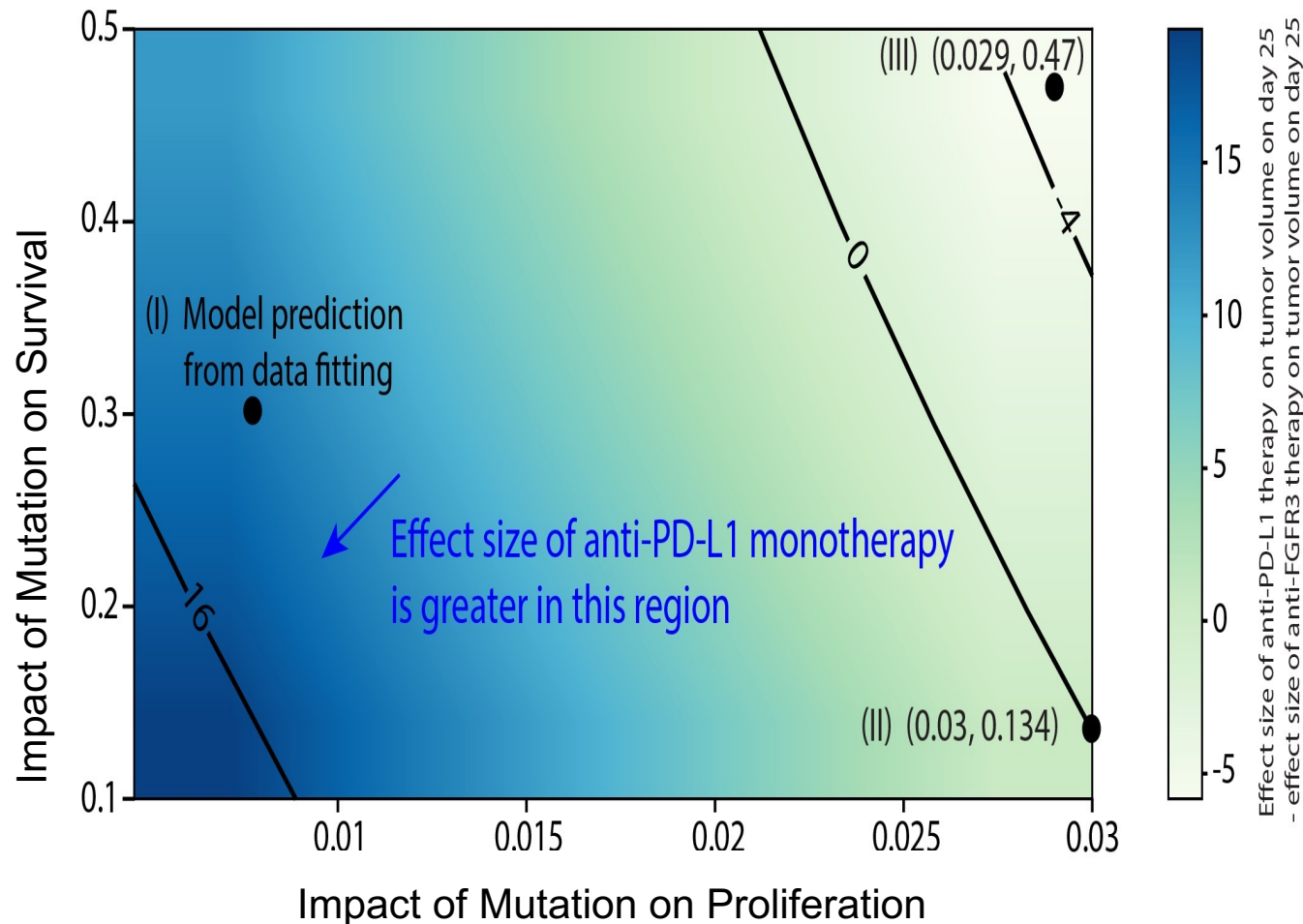
FGFR3 Mutation and Immune Dynamics

PHARMACOKINETICS OF ANTI-FGFR3



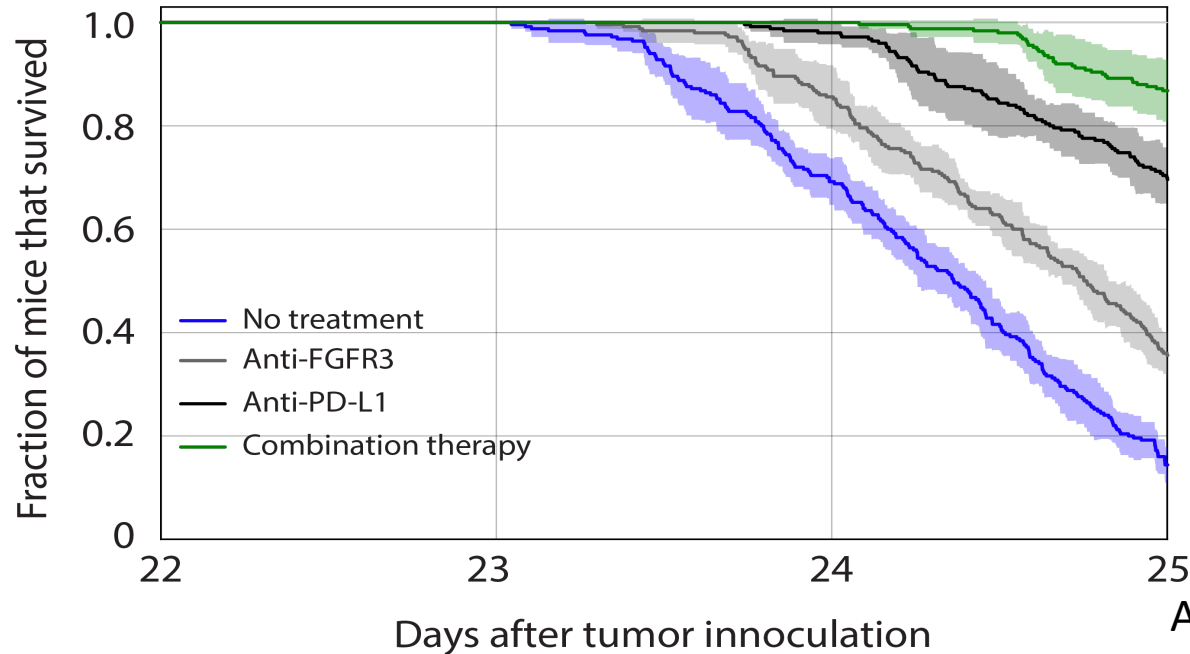
Model Predictions: Monotherapies

- We can predict when targeted therapy outperforms the immune therapy.
- The heatmap shows the difference between the immune and targeted therapies on day 25 as the impact of the mutation on proliferation and survival varies.



Model Predictions - Combo Therapy

| | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | 13 | 14 | 15 | 16 | 17 | 18 | 19 | 20 | 21 | 22 | 23 | 24 | 25 |
|------------|---|---|---|---|---|---|---|---|---|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|
| Anti-FGFR3 | | | | | | | | | | | | | | | | | | | | | | | | | |
| Anti-PDL1 | | | | | | | | | | | | | | | | | | | | | | | | | |



Mean number of mice at risk

No treatment

50

34

7

(14%)

Anti-FGFR3

50

42

18

(36%)

Anti-PD-L1

50

49

34

(70%)

Combination therapy

50

50

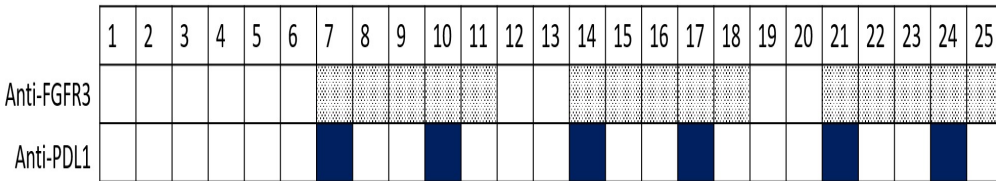
43

(86%)

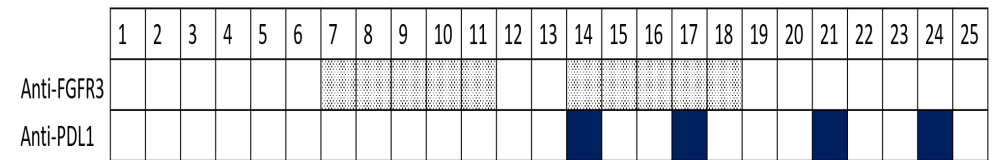
Comparing Dosing Strategies

Model prediction of tumor volume on day 25
(% reduction in tumor volume relative to no treatment)

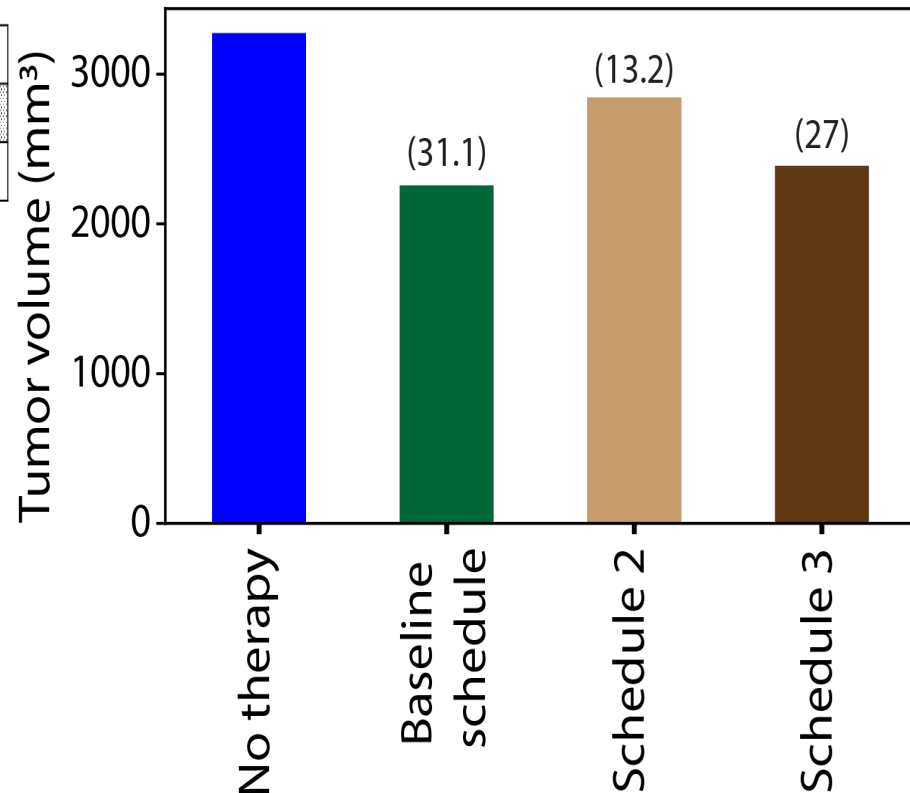
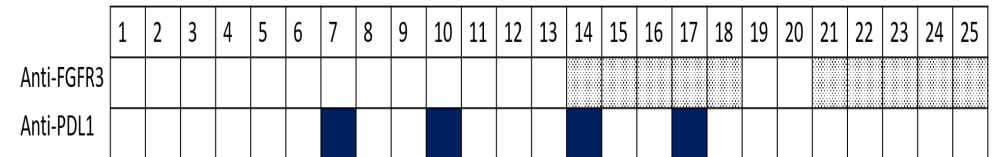
Baseline Schedule: Co-treatment



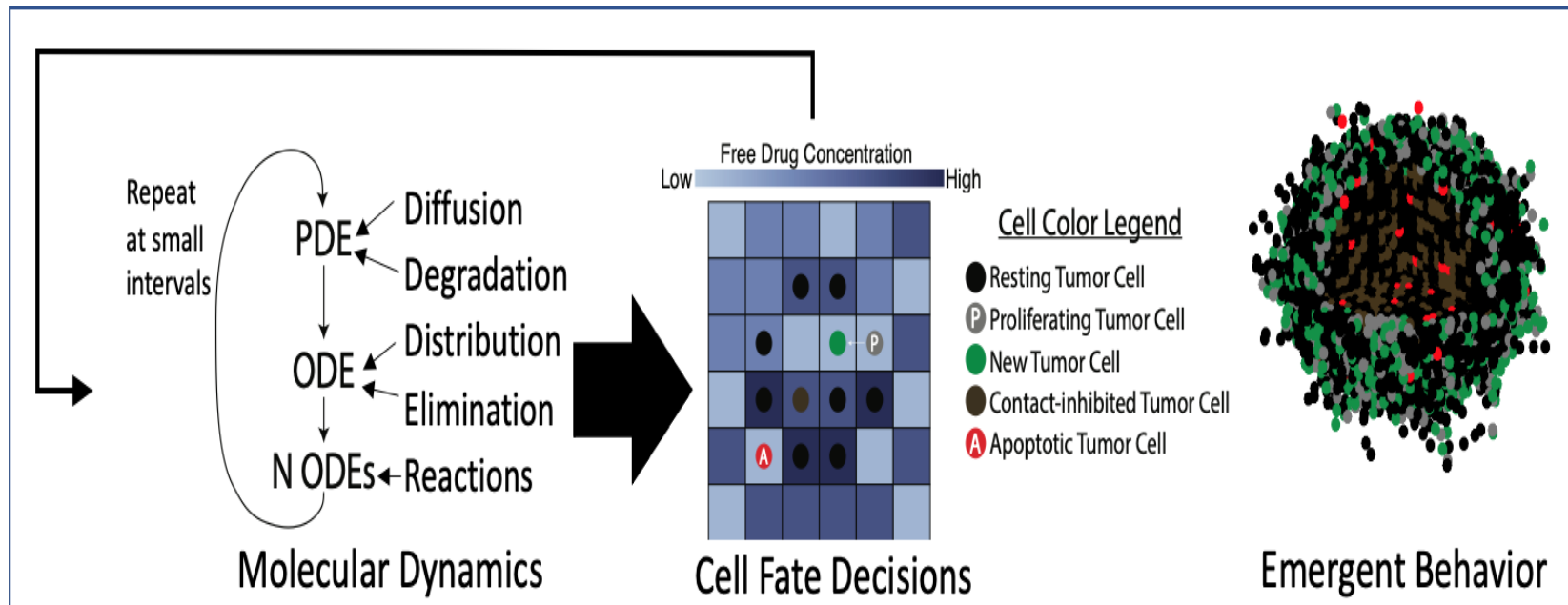
Schedule 2: Pretreatment with RTK Targeted Therapy



Schedule 3: Pretreatment with Immune Checkpoint Therapy

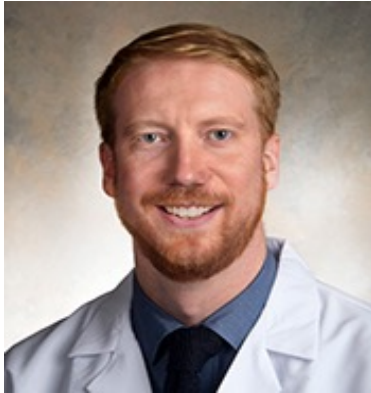


Next Steps: Agent-based Modeling



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