

Supplementary Material

March 17, 2019

| Name | Base Value | Single Patient | Survival Curve | Vary MIE | Vary MGA |
|----------------------|------------|----------------|----------------|----------|----------|
| RP Cancer Line | 0.5 | | | | |
| MIE | 0.6 | 0.9 | 0.4-0.9 | 0.7 | 0.5 |
| MGA | 0.2 | | 0.1-0.4 | 0 | 0.2 |
| INFL High Duration | 30 | | | | 60 |
| INFL Low Duration | 30 | | | | |
| Mes Threshold | 0.7 | | | | |
| p | 0.28 | | | | |
| d | 0.14 | | | | |
| N_{00} | 8 | | | | |
| N_0 | 80 | | | | |
| NK Efficacy Low | 0.2 | 10 | | 10 | 10 |
| Treg EC50 Low | 5 | | | | |
| CTL Efficacy Low | 4 | 200 | | 200 | 200 |
| NK Efficacy Up | 2 | | | | 1.2 |
| CTL Efficacy Up | 3 | | | | 3 |
| Treg Efficacy Up | 200 | | | | 10 |
| NK Source Low | 1.3 | | | | |
| CTL Recruitment Low | 100 | | | | |
| Treg Recruitment Low | 200 | | | | |
| NK Recruitment Up | 1 | | | | |
| CTL Recruitment Up | 1 | | | | |
| Treg Recruitment Up | 1 | | | | |
| $p_{mutation}$ Start | 1e-2 | | | | |
| $p_{mutation}$ | 1e-4 | | | | |
| Apoptosis Down | 0.3 | | | | |
| Immune Evasion | 0.48 | | | | |
| Proliferation Up | 0.36 | | | | |
| k_{EMT} | .01 | | | | |
| c | 6 | | | | |

| | | | | | |
|--------------------|------|-----|--|--|--|
| TGFB Received EC50 | 200 | | | | |
| TGFB Max | 500 | 700 | | | |
| Mutant TGFB | 5e-2 | | | | |
| Treg TGFB | 5e-1 | | | | |

| Name | Description | Value |
|----------------------|---|-------|
| RP Cancer Line | When proportion of mutant cells reaches this number, then cancer is declared | 0.5 |
| MIE | (mesenchymal immune evasion) proportional decrease of probability immune cells clear mutated mesenchymal cells | 0.6 |
| MGA | (mesenchymal growth arrest) proportional decrease of probability mesenchymal cells proliferate; this decreased probability is accompanied and an equal and opposite increase in probability of rest | 0.2 |
| INFL High Duration | Number of consecutive cell cycles a patient will experience high inflammation before returning to a low inflammatory state | 30 |
| INFL Low Duration | Number of consecutive cell cycles a patient will experience low inflammation before returning to a high inflammatory state | 30 |
| Mes Threshold | the threshold EMT score above which a cell is labeled mesenchymal and below which it is labeled epithelial | 0.7 |
| p | weight of a cell proliferating during a cell cycle | 0.28 |
| d | weight of a cell dying during a cell cycle | 0.14 |
| N_{00} | EC50 term for Hill functions describing likelihood of cytotoxic immune cells locating mutated cells | 8 |
| N_0 | EC50 term for Hill function describing negative feedback of total cell population on cell proliferation rates | 80 |
| NK Efficacy Low | weight of NK cell clearing mutated cells during low inflammation | 0.2 |
| Treg EC50 Low | EC50 term for Hill functions describing Treg-mediated inhibition of efficacy of cytotoxic immune cells during low inflammation | 5 |
| CTL Efficacy Low | weight of CTL cell clearing mutated cells during low inflammation | 4 |
| NK Efficacy Up | proportional increase of NK Efficacy Low during high inflammation | 2 |
| CTL Efficacy Up | proportional increase of CTL Efficacy Low during high inflammation | 3 |
| Treg Efficacy Up | during high inflammation, Treg EC50 is Treg EC50 Low / Treg Efficacy Up | 200 |
| NK Source Low | rate at which NK cells enter TME during low inflammation | 1.3 |
| CTL Recruitment Low | rate at which CTL cells are recruited after mutated cells are lysed during low inflammation | 100 |
| Treg Recruitment Low | rate at which Treg cells are recruited after mutated cells are lysed during low inflammation | 200 |

| | | |
|----------------------|---|------|
| NK Recruitment Up | proportional increase of NK source rate during high inflammation | 1 |
| CTL Recruitment Up | proportional increase of CTL recruitment rate during high inflammation | 1 |
| Treg Recruitment Up | proportional increase of Treg recruitment rate during high inflammation | 1 |
| $p_{mutation}$ Start | maximum probability of mutating after the warmup period ends | 1e-2 |
| $p_{mutation}$ | increase in probability of a cell mutating after it proliferates but does not mutate | 1e-4 |
| Apoptosis Down | proportional decrease in apoptosis weight for cells with apoptosis pathway mutated | 0.3 |
| Immune Evasion | proportional decrease in immune clearance weight for cells with immune pathway mutated | 0.48 |
| Proliferation Up | proportional increase in proliferation weight for cells with proliferation pathway mutation | 0.36 |
| k_{EMT} | rate parameter controlling speed of EMT and MET | .01 |
| c | standard deviation of noise on TGF- β received by each cell | 6 |
| TGFB Received EC50 | EC50 term for Hill function determining how much TGF- β in TME enters a cell | 200 |
| TGFB Max | maximum amount of TGF- β that can enter a cell from the TME during a cell cycle | 500 |
| Mutant TGFB | amount of TGF- β produced by each mutated cell each cycle | 5e-2 |
| Treg TGFB | amount of TGF- β produced by each Treg cell each cycle | 5e-1 |