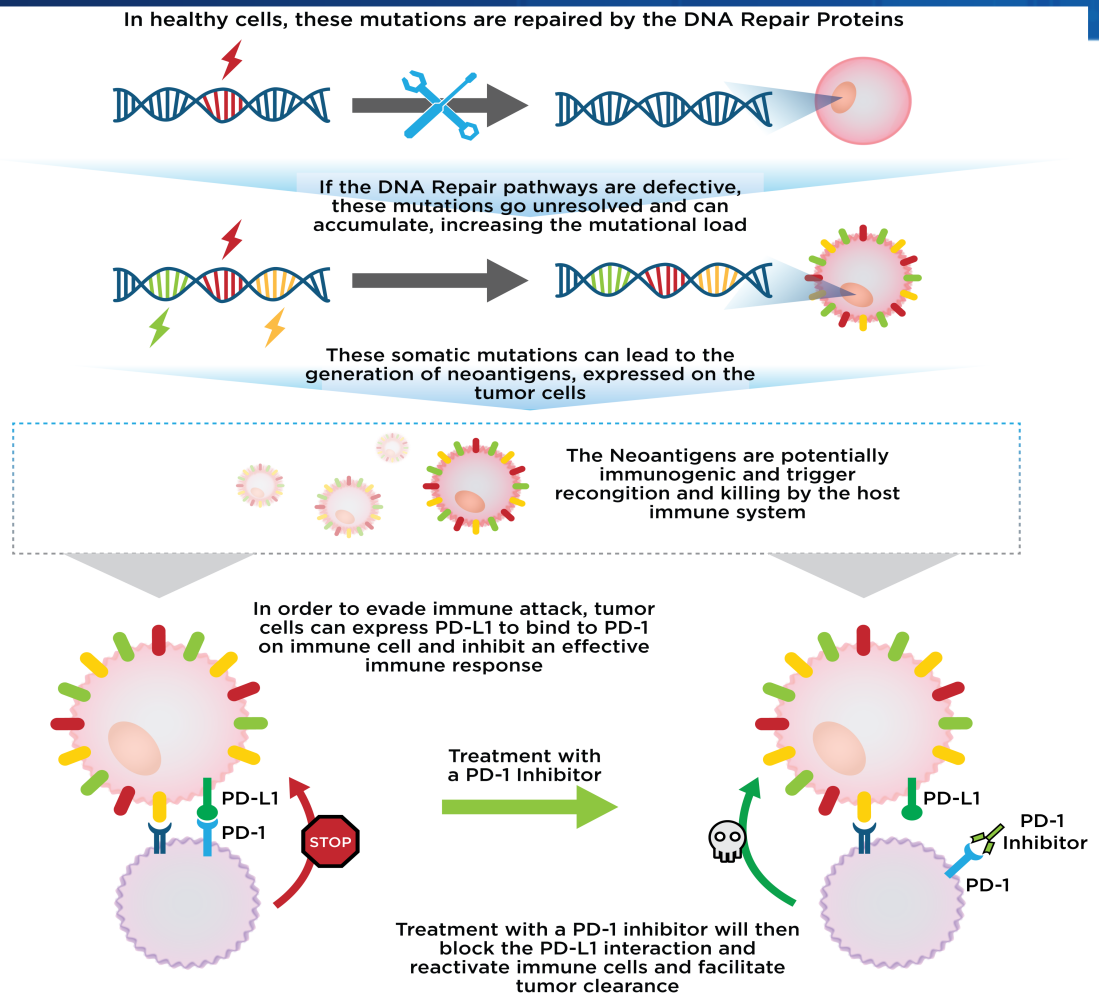
Supplemental Figure 2



Supplemental Figure 2. Tumor mutational burden as a surrogate measure of neoantigenic load. The accumulation of somatic mutations can trigger normal cells to become cancerous. These mutations are varied in nature, being single nucleotide variants or insertion-deletion events, and can be cumulatively measured and quantified as TMB. This emerging biomarker as been linked to predicting response to immune checkpoint inhibitor therapies in some tumor types.

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