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## Cancer–immune coevolution dictated by antigenic mutation accumulation

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We introduce a stochastic model of cancer–immune coevolution, where mutations trigger effector responses. Simulations show that interactions impact outcomes (e.g. suppression), with implications for immunotherapy. We find signatures of selection in sequencing statistics such as the single-cell mutational burden distribution. (based on the paper: <https://doi.org/10.7554/eLife.103970.3>)

### Pracovisko fakulty (katedra)/ Department of Faculty

KAMŠ

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