

Massachusetts Institute of Technology



Cancer as an evolutionary process

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Theory



Impact of deleterious passenger mutations on cancer progression

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Tug-of-war between driver and passenger mutations in cancer and other adaptive processes

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Experiments

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Molecular and Cellular Pathobiology

The Damaging Effect of Passenger Mutations on Cancer Progression №

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https://cancerres.aacrjournals.org/content/77/18/4763.long

Main points

1. Cancer is an evolutionary process

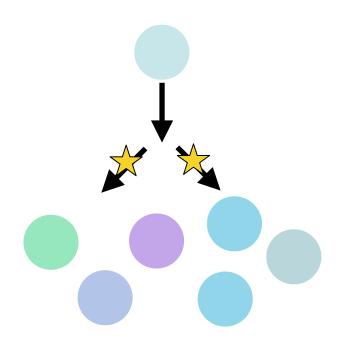
Cancer genomics allows to look under the hood of this process

Treating cancer using its own evolutionary mechanisms

Evolution

mutations

diversity



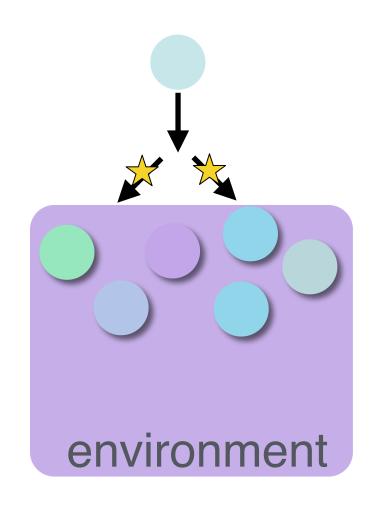
selection

Evolution

mutations

diversity

selection

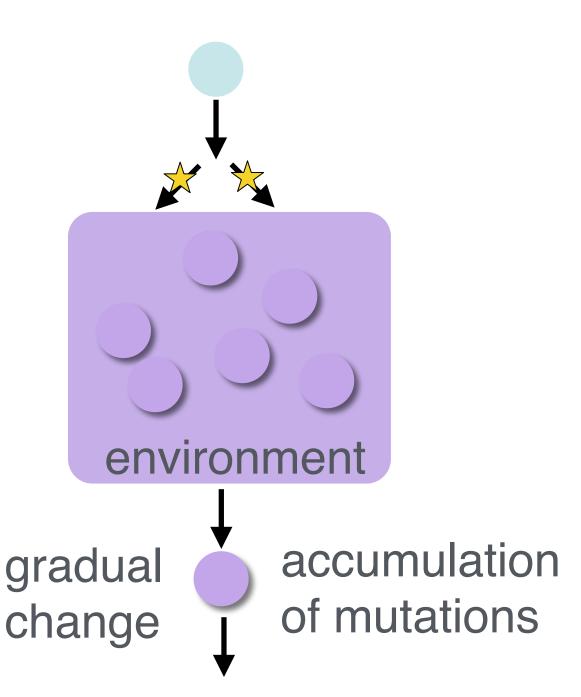


Evolution

mutations

diversity

selection



Mutant is new normal

mutations

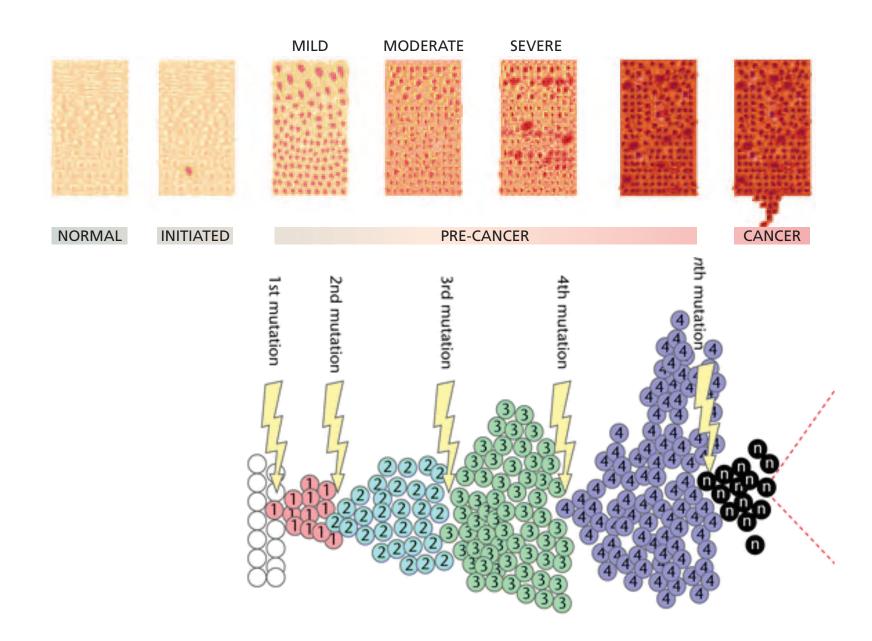
new phenotype is acquired

diversity

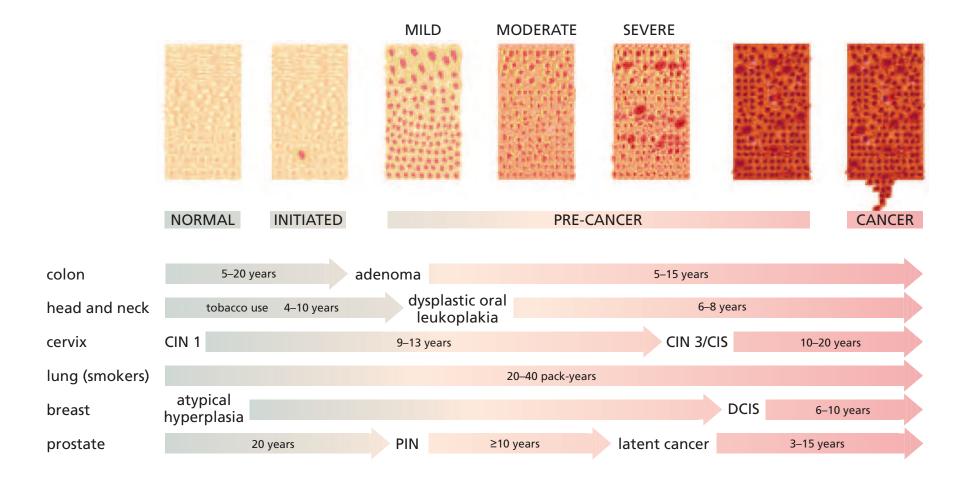
selection

gradual change accumulation of mutations

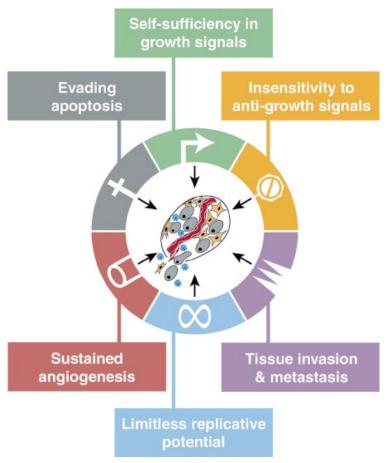
Cancer = evolution



Cancer = evolution



Acquired phenotypes of cancer



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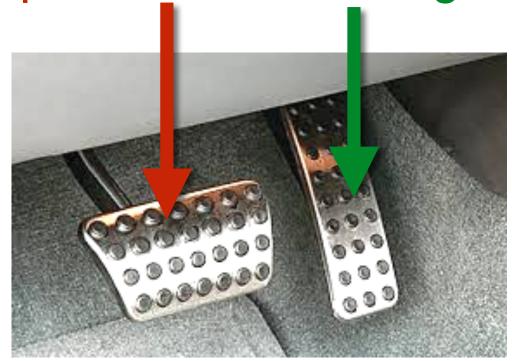
The Hallmarks of Cancer

Acquired phenotypes of cancer acquired *mutations*

Mutation targets tumor suppressors and oncogenes

Acquired phenotypes of cancer acquired *mutations*

Mutation targets tumor suppressors and oncogenes



Acquired phenotypes of cancer acquired *mutations*

Mutation targets tumor suppressors and oncogenes



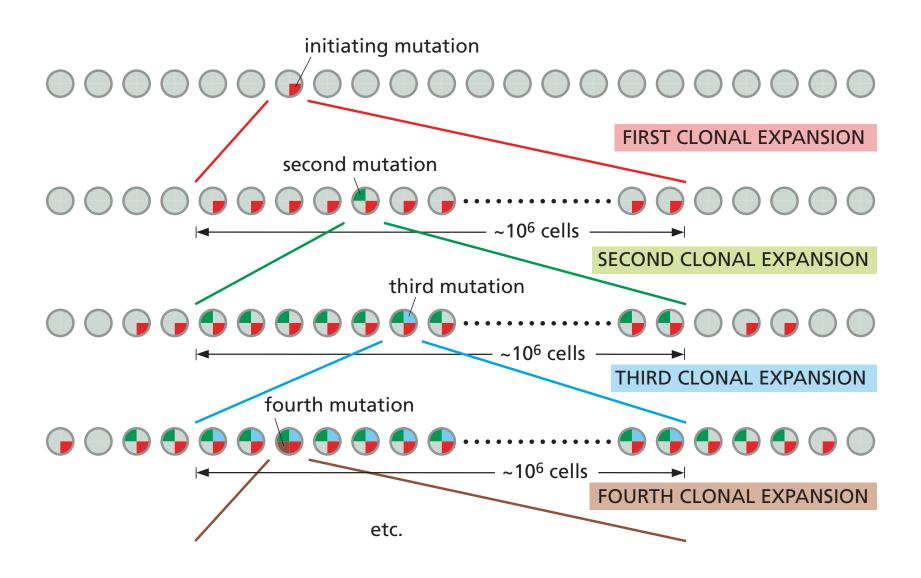
Mutation targets tumor suppressors and oncogenes [drivers]

Oncogenes and tumor suppressors

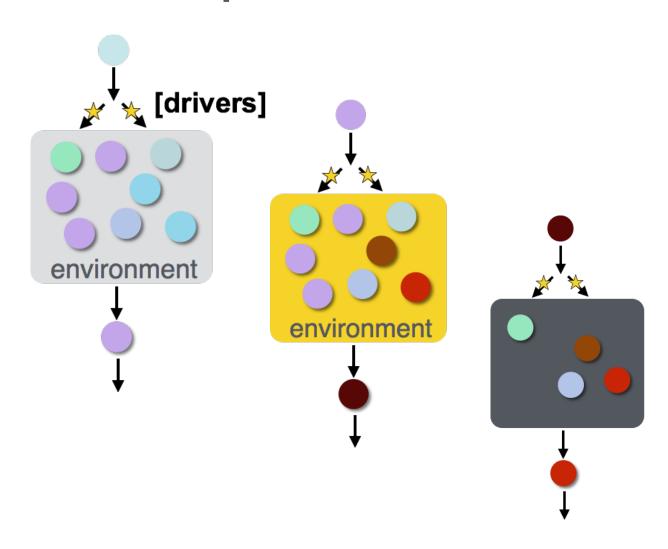
- Oncogenes -- need to be activated
 - -by mutations (within a gene or regulatory regions)
 - by chromosomal alterations
 - overexpression/modifications
- Tumor suppressors -- need to be inactivated
 - mutations, chromosomal loss, modifications

[drivers]

Cancer: series of driver mutations



Cancer: is hard to stop because it's an evolutionary process



Main points

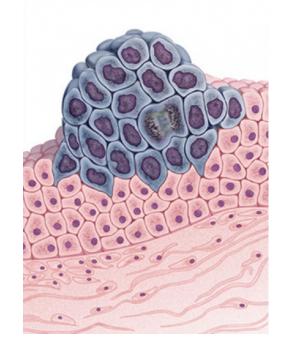
1. Cancer is an evolutionary process

2. Cancer genomics allows to look under the hood of this process

Treating cancer using its own evolutionary mechanisms

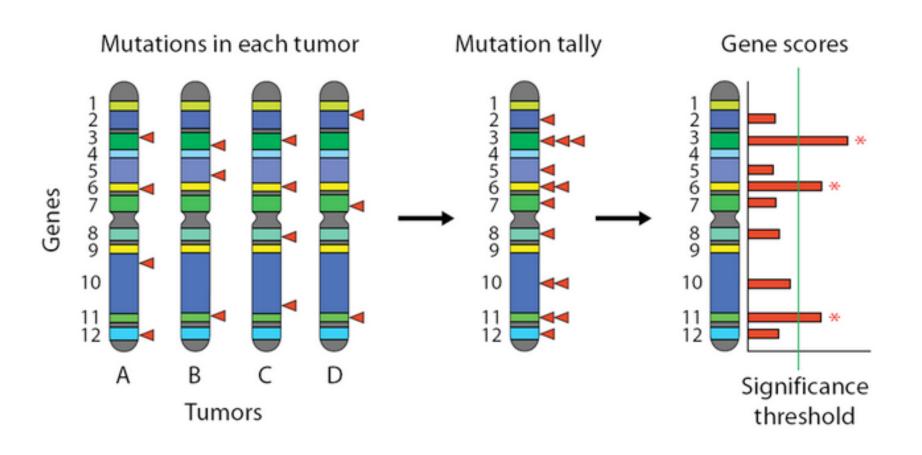
Cancer genomics

- Get a sample of cancer => sequence
- Get a sample of normal tissue
 (from the same patient) =>

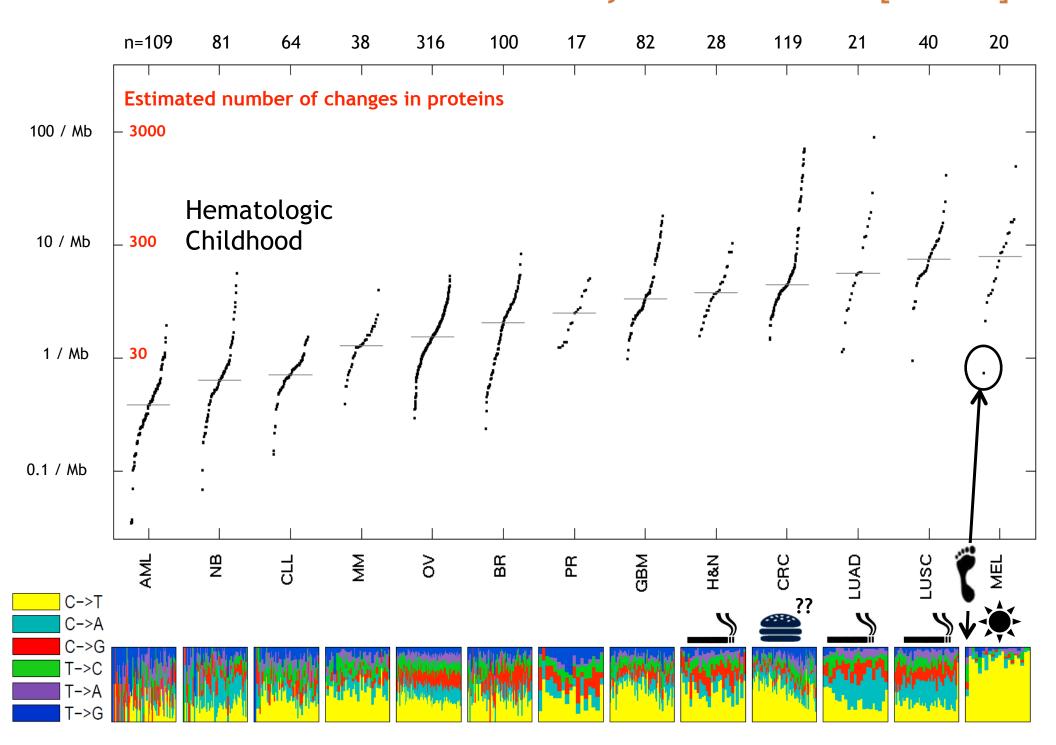


GATGTTTTATCAGTATCTTTTGACTTTTTAACATTCAAAACACTCCCTACTAATTTCTGCTTTTGGTAACAGTACATGCCATGTTAACCT TTTGCCCCTCCTTCCACCCCCATGATTCATAATGTCTGCAGACATTTTTCACTGTCACAACTGGGGGATGCTGCTAAACATCCTACAAA AGGACAGTTCTCTAGGTCAAAGCTGTTCTTAGGCAAAAAAGTCAAGTGCCAAAGGTGAGAAATCCTAATATAGAGGAATTTACTGTCTC TGAAAATTTTTCTCAAGCAATTTCATGATTTAAATAATTTCCCAGTCATAGGGTTGAATCCATGAGGTAATGCTAGCAATATGAAACA(GCAGGATTATTAATTATCACTAATTCTTCCAAGGCTACCTAACAGAATATCTCTGCTCTCCACAGGCCCATCAATTTGAAAACTCAAGT TAAGAGTAAAAAAGTAGATAATGGCTTTGAAGTTTATAAGAAAATTATGCAGCAAAGCTTTTTGTTTTACATAAGCTCATGTAAGAATAA AATTTCCTAAATCTGCATAAAACAGTTGTTATTTGGATCCACTTTTACATGTTAAGTTAGAATCTGGCAAATTCTGTCTAAATAGTCC TTTCACCAGCCCTACAAGATTATTCATGGGAGAGACTATATTAACGAATTTTGTTTCTAAAAAATTAAACCTCTCTTTTTCCCTACAATAT AGATGGAGGCACTGTCTGACCATTTTACTGAAAGCATTGTAAACGTGGTCAAACCAAACATACACAGACTGTGGCATTTCTCTGCACT(ATTTAAAGACAAAAGGAAAAAAAGCCTAAGCCATTGTCATATGTTAACAAAGGGCTGCCAACATTGTAATCTTGCCTCGAAATGTCCAC TATTTAAAATTACCCGAACGGAAACATGTAAGTGATATGAGCACACAATTCACAAAGATCAAGGTGCCAATGGTTAGCAGATACAAAA GTTCAACCTCATCAGTATTCCAGAATATGCACAAAGATTCTATCTTTAAACCTGCAAAATTATCAGAGTACAAGATATACTCTCAAAAT GCTAGTGCAAACATGTCATAAACTCCTACTACTGTCTGCATAGAATTGTTCATCCAAATGGATTTTTTCAAAGGAAATTTAAAACTCAG AAATGGACAAATGTGGTTTTTTTTTAATAGCAAGCAACATGACAATGAAGAATTGTGTCCTGGTATCTATGTCCTGGTAGGTGGGCCAA GCAAGAGTGCTCTGCTGATCTGACTTAAATGTGTTTTCTTCAGTGAATCCCTCTGTAGAGGTTTAATTTGGTAGACGTTCTATAGAGAA AAAAAATAAGATATCATCTTGATCAATTATAAAATGTGTACTTCAATTTCTTGGTTTCTATCATTGCAAATAGCAGTTCATGTTATACA AAACCCAGGTGTGGTCAAATTTCATTGTCAAGGAAAAGGGAACATTTTGGTGCTTCTTGAGATTATCATCATGAAAACACAATAAAAGG CTTAACTTTTCTTGGTAGAGAGGTTATGTGTGCCAATTCATGCACTGGTACATTAATGTCTAGCTCACATCAAATAAAAAGCAACATC GATACTGCTATGAATAAAAGACTGTTCTCTACACTTTCCTGTACTGTTTTGTAATTTCTGAAGGGAAAAAAGAAGAAATGAATTAGAGAA AGCTAGAAAGGTAAAAGTATATGAACAACACTTTTCTATTTAGTTCCCTCATTTGTTTCATAGTGCTTTAACTGCCATCATTTCATTAC AAAAAAGGTTAAATCTAACAATATATGCTAAAAACTCAATTTCACTGCAACAAAAGAATGAAAGTCCCAGGCTGGGCGTGGTGGCTCAC CCTGTAATCCCAGCACTTTGGGAGGCCAAAGCAGGCGGATCACCTGAGATCAGGAGTTCGAGACCAGCCCAGCCAACATGGTAAAACC 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CCTGTAATCCCAGCACTTTGGGAGGCCAAAGCAGGCGGATCACCTGAGATCAGGAGTTCGAGACCAGCCCAGCCAACATGGTAAAACC AAAAGAAAAAAAAGAAAGTCTCTTGCATTAGTGTCAAAAGTATAATATAGATATTTCAAGTTCCCCAGATTAATAATATTACCTTAACT GATGTTTTATCAGTATCTTTTGACTTTTTAACATTCAAAACACTCCCTACTAATTTCTGCTTTTGGTAACAGTACATGCCATGTTAACCT TTTGCCCCTCCTTCCACCCCCATGATTCATAATGTCTGCAGACATTTTTCACTGTCACAACTGGGGGATGCTGCTAAACATCCTACAAA AGGACAGTTCTCTAGGTCAAAGCTGTTCTTAGGCAAAAAAGTCAAGTGCCAAAGGTGAGAAATCCTAATATAGAGGAATTTACTGTCTC TGAAAATTTTTCTCAAGCAATTTCATGATTTAAATAATTTCCCAGTCATAGGGTTGAATCCATGAGGTAATGCTAGCAATATGAAACA(GCAGGATTATTAATTATCACTAATTCTTCCAAGGCTACCTAACAGAATATCTCTGCTCTCCACAGGCCCATCAATTTGAAAACTCAAGT TAAGAGTAAAAAAGTAGATAATGGCTTTGAAGTTTATAAGAAAATTATGCAGCAAAGCTTTTTGTTTTACATAAGCTCATGTAAGAATAA AATTTCCTAAATCTGCATAAAACAGTTGTTATTTGGATCCACTTTTACATGTTAAGTTAGAATCTGGCAAATTCTGTCTAAATAGTCC TTTCACCAGCCCTACAAGATTATTCATGGGAGAGACTATATTAACGAATTTTGTTTCTAAAAAATTAAACCTCTCTTTTTCCCTACAATAT AGATGGAGGCACTGTCTGACCATTTTACTGAAAGCATTGTAAACGTGGTCAAACCAAACATACACAGACTGTGGCATTTCTCTGCACT(ATTTAAAGACAAAAGGAAAAAAAGCCTAAGCCATTGTCATATGTTAACAAAGGGCTGCCAACATTGTAATCTTGCCTCGAAATGTCCAC TATTTAAAATTACCCGAACGGAAACATGTAAGTGATATGAGCACACAATTCACAAAGATCAAGGTGCCAATGGTTAGCAGATACAAAA GTTCAACCTCATCAGTATTCCAGAATATGCACAAAGATTCTATCTTTAAACCTGCAAAATTATCAGAGTACAAGATATACTCTCAAAAT GCTAGTGCAAACATGTCATAAACTCCTACTACTGTCTGCATAGAATTGTTCATCCAAATGGATTTTTTCAAAGGAAATTTAAAACTCAG AAATGGACAAATGTGGTTTTTTTTTAATAGCAAGCAACATGACAATGAAGAATTGTGTCCTGGTATCTATGTCCTGGTAGGTGGGCCAA GCAAGAGTGCTCTGCTGATCTGACTTAAATGTGTTTTCTTCAGTGAATCCCTCTGTAGAGGTTTAATTTGGTAGACGTTCTATAGAGAA AAAAAATAAGATATCATCTTGATCAATTATAAAATGTGTACTTCAATTTCTTGGTTTCTATCATTGCAAATAGCAGTTCATGTTATACA AAACCCAGGTGTGGTCAAATTTCATTGTCAAGGAAAAGGGAACATTTTGGTGCTTCTTGAGATTATCATCATGAAAACACAATAAAAGG CTTAACTTTTCTTGGTAGAGAGGTTATGTGTGCCAATTCATGCACTGGTACATTAATGTCTAGCTCACATCAAATAAAAAGCAACATC GATACTGCTATGAATAAAAGACTGTTCTCTACACTTTCCTGTACTGTTTTGTAATTTCTGAAGGGAAAAAAGAAGAAATGAATTAGAGAA 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Finding driver events



Rates of somatic mutation vary across cancers: [G.Getz]



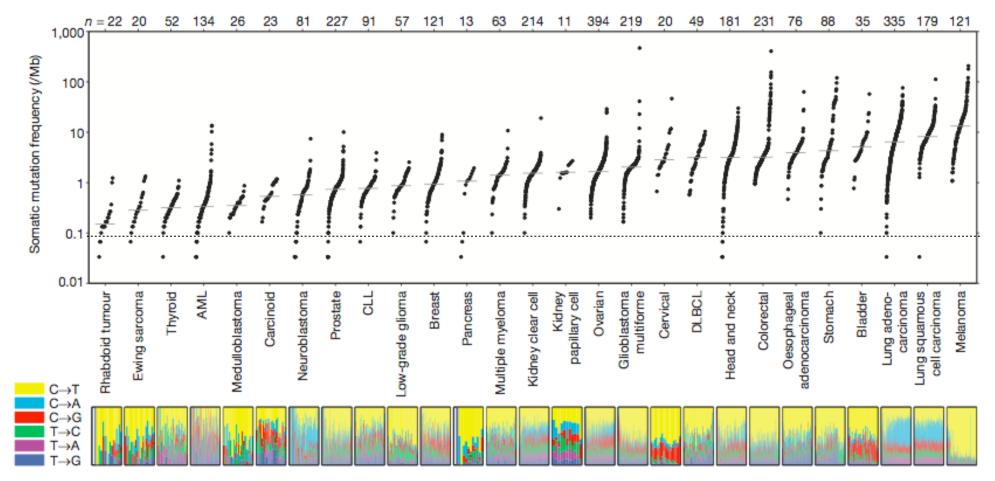


Figure 1 | Somatic mutation frequencies observed in exomes from 3,083 tumour-normal pairs. Each dot corresponds to a tumour-normal pair, with vertical position indicating the total frequency of somatic mutations in the exome. Tumour types are ordered by their median somatic mutation frequency, with the lowest frequencies (left) found in haematological and paediatric tumours, and the highest (right) in tumours induced by carcinogens

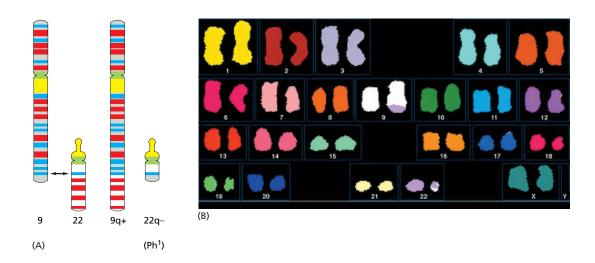
such as tobacco smoke and ultraviolet light. Mutation frequencies vary more than 1,000-fold between lowest and highest across different cancers and also within several tumour types. The bottom panel shows the relative proportions of the six different possible base-pair substitutions, as indicated in the legend on the left. See also Supplementary Table 2.

M. Lawrence et al Nature 2013

Cancer genomics

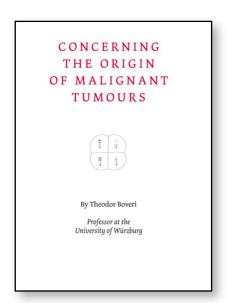
- Whole-genome sequences (cancer vs normal)
- Whole-exome sequences (cancer vs normal)

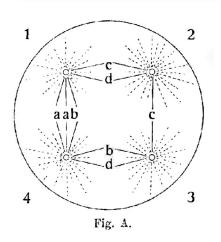
Chromosomal alterations

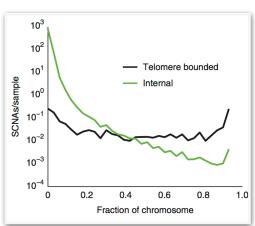


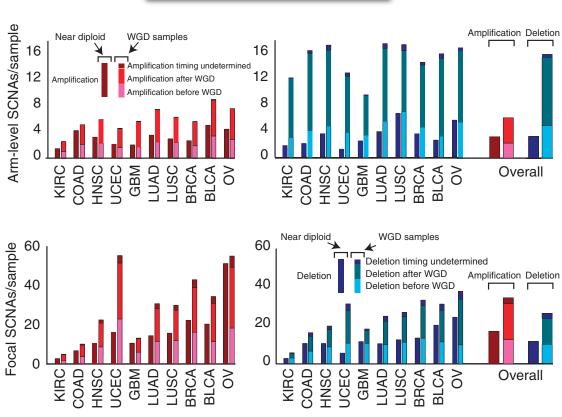
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Somatic Copy Number Alterations (SCNAs)

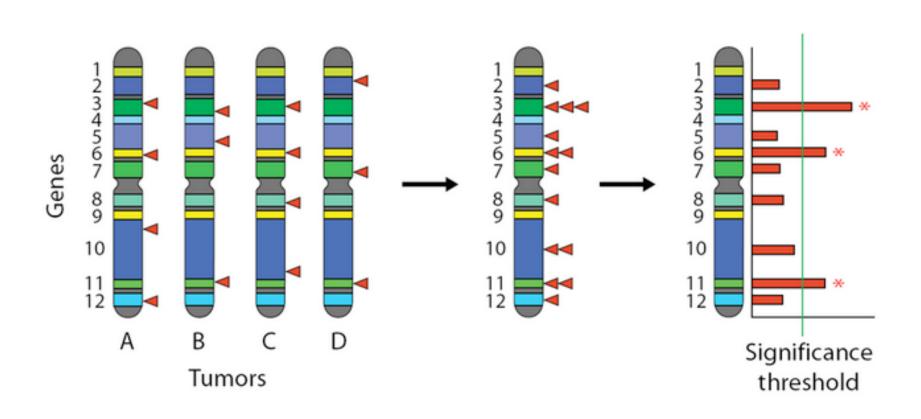




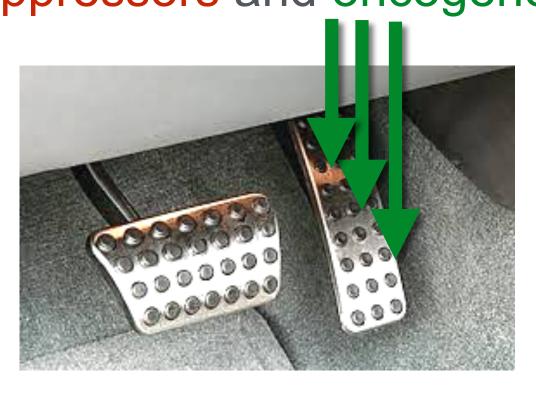




Finding oncogenes and tumor suppressors



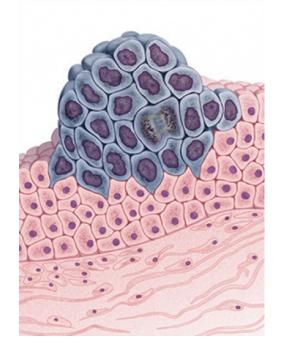
Deletions Amplifications tumor suppressors and oncogenes



Cancer genomics

Finding new oncogene and tumor suppressors Whole mutational landscape of cancer Precision medicine:

mutations in each patient



Main points

1. Cancer is an evolutionary process

2. Cancer genomics allows to look under the hood of this process

Treating cancer using its own evolutionary mechanisms

Cancer genomics

100-400 amino acid substitutions

10-40 chromosomal alterations

2-5 drivers

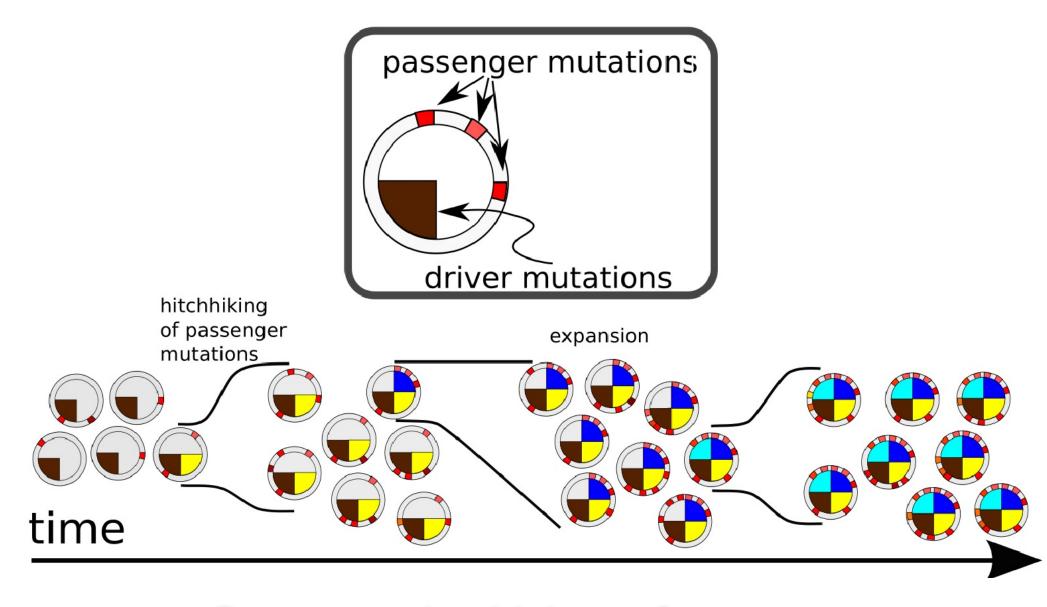
the rest are passengers

Can some passengers

... be deleterious to cancer cells?

... affect progression?

Passengers hitchhike of drivers



Passengers hitchhike to fixation

Theory



Impact of deleterious passenger mutations on cancer progression

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Molecular and Cellular Pathobiology

The Damaging Effect of Passenger Mutations on Cancer Progression №

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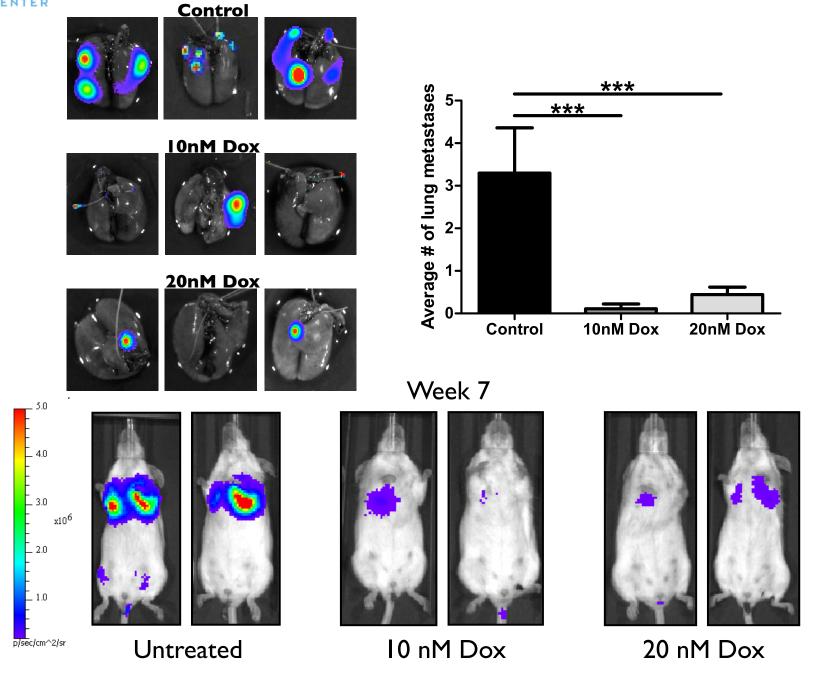




https://cancerres.aacrjournals.org/content/77/18/4763.long



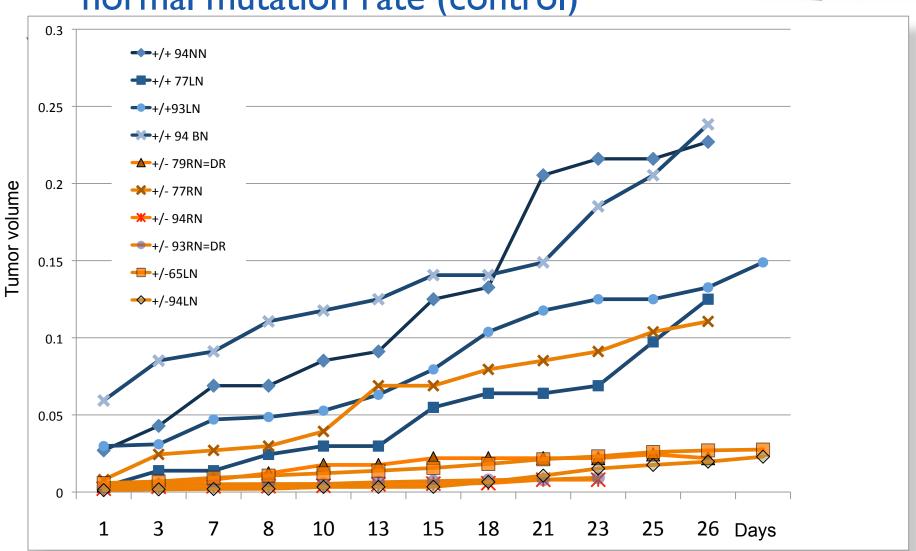
Passanger load negatively correlates with metastasis



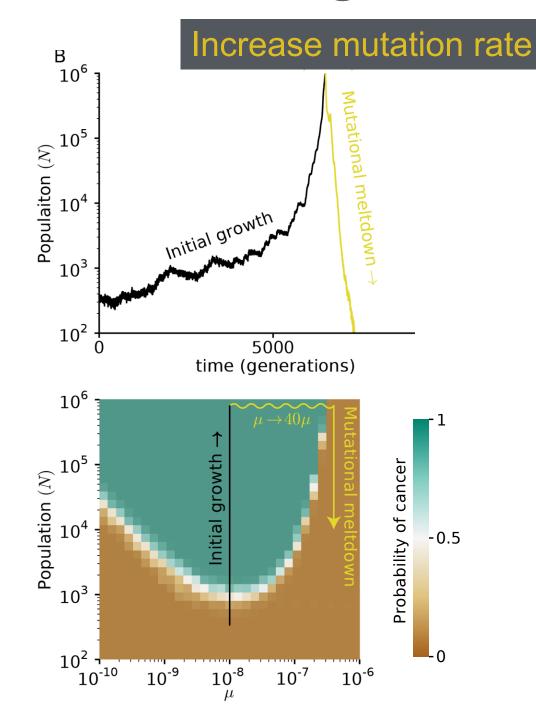
Passengers slowdown cancer

New Experiment: Her2+ breast cancer mouse model:

mildly elevated mutation rate (H2AX+/-) normal mutation rate (control)



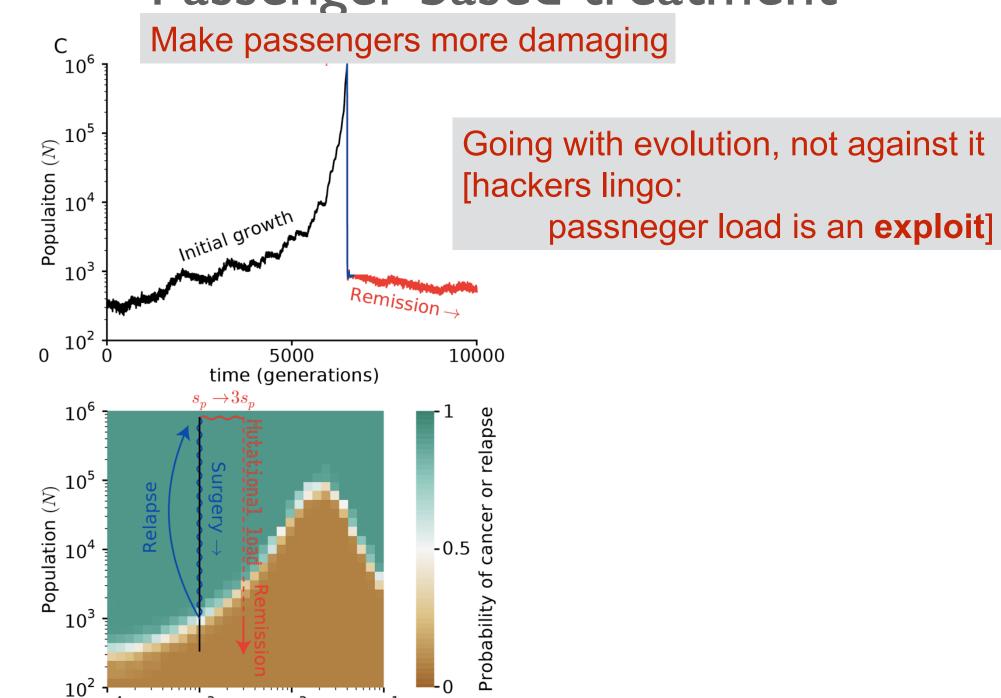
Passenger-based treatment



Mutagenic chemo

- requires very high mutation rate
- likely relapse

Passenger-based treatment



10⁻³

10⁻⁴

10⁻²

10⁻¹

Main points

1. Cancer is an evolutionary process

Cancer genomics allows to look under the hood of this process

Treating cancer using its own evolutionary mechanisms

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Experiments

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Metastatic potential



Genomics

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