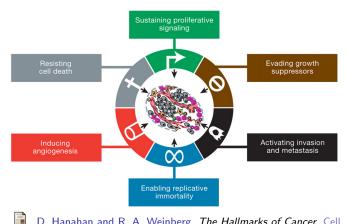
# Hallmarks of cancer

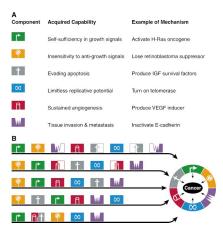


D. Hanahan and R. A. Weinberg. *The Hallmarks of Cancer*, Cell, vol. 100, no. 1, pp. 57-70, 2000.

D. Hanahan and R. A. Weinberg. *Hallmarks of Cancer: The Next Generation*, Cell, vol. 144, no. 5, pp. 646-674, 2011.

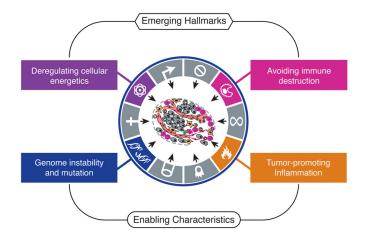
### Tumor progression

#### Only certain paths are available to tumors as they acquire hallmarks.



# Emerging hallmarks

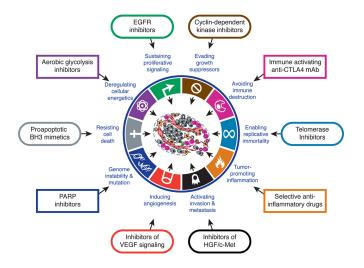
New hallmarks are being proposed by different researchers.



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## Therapeutic agents

Different hallmarks are associated with different therapeutic agents.



# More therapeutic agents

| Target        | Addiction   | Hallmarks   | Potential mechanisms  | References   |
|---------------|---|---|---|--|
| HSP90         | NOA   | Ì   | A geldanamycin analog that binds to the<br>ATP-binding pocket of HSP90 and inhibits its<br>catalytic activity   | Whitesell and<br>Lindquist, 2005   |
| IDO           | NOA   | <b>Y</b>  | Inhibits tryptophan catabolism in tumor mi-<br>croenvironment to allow T cell proliferation   | Muller and Scherle, 2006   |
| DNA           | NOA   | X   | Inhibits pyrimidine metabolism, incorporation in to DNA and RNA causes cell-cycle arrest  | Longley et al., 2003   |
| BCL-XL, BCL-2 | OA  | +   | Bind to the BH3 pocket of BcI-XL and inhibit its<br>antiapoptotic function  | Stauffer, 2007   |
| CDKs          | OA  | r   | Inhibit CDKs and induce cell-cycle arrest   | Lee and Sicinski,<br>2006  |
| TGFβ 2        | NOA   |   | Inhibits tumor autocrine and paracrine signal-<br>ing, reverses immune suppression in the tumor<br>microenvironment   | Muller and Scherle, 2006   |
| PARP1         | NOA   | X   | Inhibit base excision repair in homologous<br>recombination repair-deficient cancer cells   | Bryant et al., 2005;<br>Farmer et al., 2005  |
| VEGF          | NOA   | P   | Inhibits endothelial cell recruitment and tumor<br>vasculature  | Folkman, 2007  |
| PI3K          | OA  | r 🖻   | Causes cell-cycle arrest in tumor cells and<br>inhibits tumor angiogenesis  | Maira et al., 2008   |
| Proteasome    | NOA   | )   | Inhibits the catalytic activity of 26S proteasome<br>and induces apoptosis  | Roccaro et al.,<br>2006  |
| COX2          | NOA   | M <b>t</b> M A  | Reverses immune suppression in the tumor<br>microenvironment, inhibits tumor autocrine and<br>paracrine signaling   | Muller and Scherle,<br>2006  |
| DNA           | NOA   | 2   | Induces DNA crosslinks  | Siddik, 2003   |
| EGFR          | OA  | 100 B 10  | Inhibit EGFR tyrosine kinase by competing with  | Sharma et al., 2007  |
|               | HSP90<br>DNA<br>BCL-XL, BCL-2<br>CDKs<br>TGFp 2<br>PARP1<br>VEGF<br>PI3K<br>Proteasome<br>COX2<br>DNA | HSP90      NOA        DO      NOA        DDA      NOA        DNA      NOA        BCL-XL, BCL-2      OA        CDKs      OA        TGF/p 2      NOA        PPARP1      NOA        PI3K      OA        Proteasome      NOA        COX2      NOA | HSP90      NOA      Image: Constraint of the sector o | HSP90  NOA  Application and inhibits to the<br>ATP-binding pocket of HSP90 and inhibits its<br>catalytic activity    DO  NOA  Inhibits typophan catabolism in turnor mi-<br>crosivorment to allow T cell proliferation    DNA  NOA  Inhibits typophan catabolism, incorporation in<br>to DNA and RNA causes cell-cycle arrest    BCL-XL, BCL-2  OA  Inhibits typophan catabolism, incorporation in<br>to DNA and RNA causes cell-cycle arrest    BCL-XL, BCL-2  OA  Inhibits DNA and RNA causes cell-cycle arrest    BCL-XL, BCL-2  OA  Inhibits to DNA and RNA causes cell-cycle arrest    TGFβ 2  NOA  Inhibits turnor autocrine and paracrine signal-<br>ing, reverses immuse suppression in the turnor<br>ing, reverses immuse suppression in the turnor<br>recombination repain-definit in homologous    VEGF  NOA  Inhibits turnor autocrine and paracrine signal-<br>ing, reverses immuse suppression in the turnor<br>recombination repain-definit in homologous    VEGF  NOA  Inhibits turnor autocrine and paracrine signal-<br>ing reverses immuse suppression in the turnor<br>recombination regain-definit or acore cells and<br>inhibits turnor angle-definit activity of 26S proteasome<br>and induces apoptosis    COX2  NOA  Inhibits Immor autocrine and paracrine and<br>and induces apoptosis    DNA  Inhibits turnor autocrine and<br>paracrine signal-<br>crine signal earlier activity of 26S proteasome<br>and induces apoptosis |

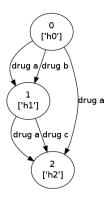
### (the list goes on)

J. Luo, N. L. Solimini, and S. J. Elledge. Principles of Cancer Therapy: Oncogene and Non-oncogene Addiction, Cell, vol. 136, no. 5, pp. 823-837, Mar. 2009.

# "Hallmark automata"

- With increasing numbers of very specifically targeted therapeutic agents being identified, combining them optimally into cocktails and in temporal succession becomes complex
- We propose a framework to automatically generate therapeutic regimens
- Represent progression models as Kripke structure / finite automaton
- Personalize model to specific cancer type and stage of patient
- Use model to automatically generate therapeutic regimens:
  - Specify therapeutic objective using
    - Temporal logic
    - Cost function to be minimized
    - Combination of the two
  - Generate supervisory controller to achieve therapeutic objective:
    - Model checking
    - Reachability analysis
    - Cost optimization

## Simple example

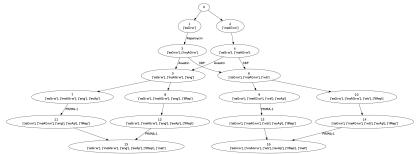


- E.g.,  $AG \neg h2$  will yield controllers that
  - give drug a plus drug b at state 0, or
  - give drug a at state 0 and drug a plus drug c at state 1,

depending, e.g., on the costs of drug b vs drug c

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# More complex model



E.g.,  $AG \neg met$  will yield controllers that

- give Rapamycin, or Avastin and 3BP, if the patient comes at early stage
- give Avastin at stage 3 and 4 and PRIMA-1 at stage 9 and 14 if 3BP has high toxicity
- give 3BP at stage 3 and 4 and PRIMA-1 at stage 7 and 12 if the patient's genome indicates adverse reaction to Avastin
- give PRIMA-1 if the disease status is advanced but unknown

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# Ideas / extensions

- (Multi-dimensional) costs for drugs, states, observations (biopsies), (violated) properties
- Timing and probabilities for transitions
- Include edges that can be enabled by drugs
- Include indistinguishabilities between states
- Represent model symbolically
  symbolic model checking
- Generate model automatically from data (GOALIE)