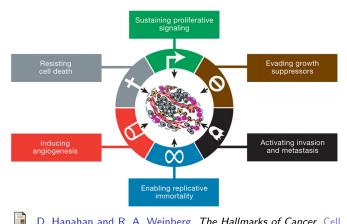
# Hallmarks of cancer

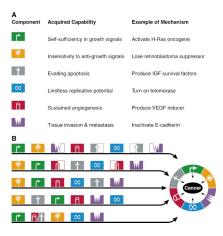


D. Hanahan and R. A. Weinberg. *The Hallmarks of Cancer*, Cell, vol. 100, no. 1, pp. 57-70, 2000.

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

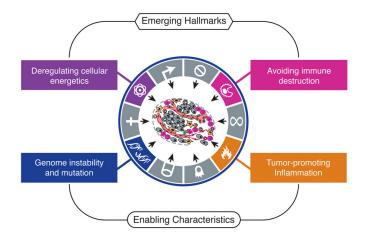
### Tumor progression

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



# Emerging hallmarks

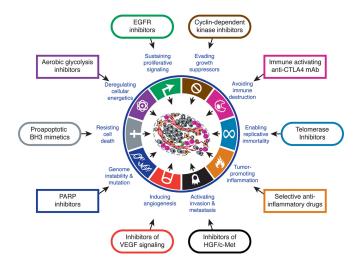
New hallmarks are being proposed by different researchers.



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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 ATP-binding pocket of HSP90 and inhibits its catalytic activity	Whitesell and Lindquist, 2005
IDO	NOA	<b>Y</b>	Inhibits tryptophan catabolism in tumor mi- 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 antiapoptotic function	Stauffer, 2007
CDKs	OA	r	Inhibit CDKs and induce cell-cycle arrest	Lee and Sicinski, 2006
TGFβ 2	NOA		Inhibits tumor autocrine and paracrine signal- ing, reverses immune suppression in the tumor microenvironment	Muller and Scherle, 2006
PARP1	NOA	X	Inhibit base excision repair in homologous recombination repair-deficient cancer cells	Bryant et al., 2005; Farmer et al., 2005
VEGF	NOA	P	Inhibits endothelial cell recruitment and tumor vasculature	Folkman, 2007
PI3K	OA	r 🖻	Causes cell-cycle arrest in tumor cells and inhibits tumor angiogenesis	Maira et al., 2008
Proteasome	NOA	)	Inhibits the catalytic activity of 26S proteasome and induces apoptosis	Roccaro et al., 2006
COX2	NOA	M <b>t</b> M A	Reverses immune suppression in the tumor microenvironment, inhibits tumor autocrine and paracrine signaling	Muller and Scherle, 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 DNA BCL-XL, BCL-2 CDKs TGFp 2 PARP1 VEGF PI3K Proteasome COX2 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 ATP-binding pocket of HSP90 and inhibits its catalytic activity    DO  NOA  Inhibits typophan catabolism in turnor mi- crosivorment to allow T cell proliferation    DNA  NOA  Inhibits typophan catabolism, incorporation in to DNA and RNA causes cell-cycle arrest    BCL-XL, BCL-2  OA  Inhibits typophan catabolism, incorporation in 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- ing, reverses immuse suppression in the turnor ing, reverses immuse suppression in the turnor recombination repain-definit in homologous    VEGF  NOA  Inhibits turnor autocrine and paracrine signal- ing, reverses immuse suppression in the turnor recombination repain-definit in homologous    VEGF  NOA  Inhibits turnor autocrine and paracrine signal- ing reverses immuse suppression in the turnor recombination regain-definit or acore cells and inhibits turnor angle-definit activity of 26S proteasome and induces apoptosis    COX2  NOA  Inhibits Immor autocrine and paracrine and and induces apoptosis    DNA  Inhibits turnor autocrine and paracrine signal- crine signal earlier activity of 26S proteasome and induces apoptosis

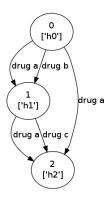
### (the list goes on)

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

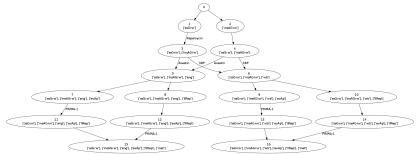


- E.g.,  $AG \neg h2$  will yield controllers that
  - give drug a plus drug b at state 0, or
  - give drug a at state 0 and drug a plus drug c at state 1,

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)