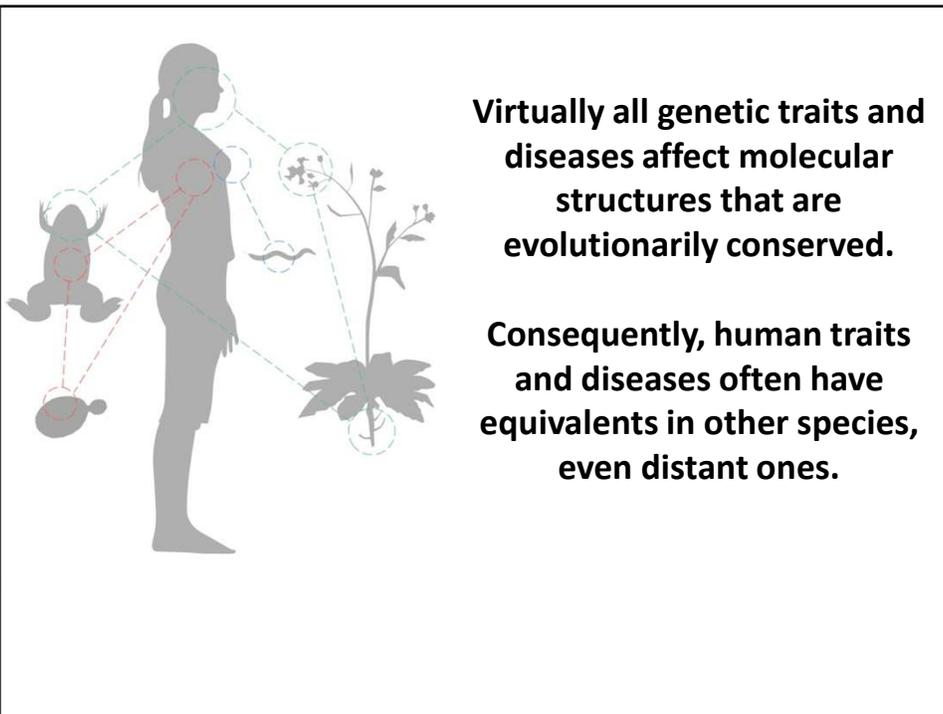


Phenologs

**An example of using bioinformatics to
find new genes for genetic traits**

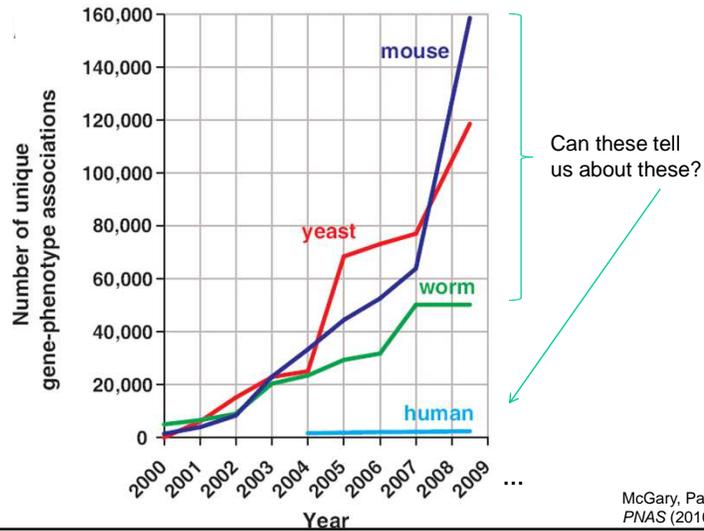
BCH364C/391L Systems Biology / Bioinformatics – Spring 2015

Edward Marcotte, Univ of Texas at Austin



We know far more about genes & traits in lower organisms than in us.

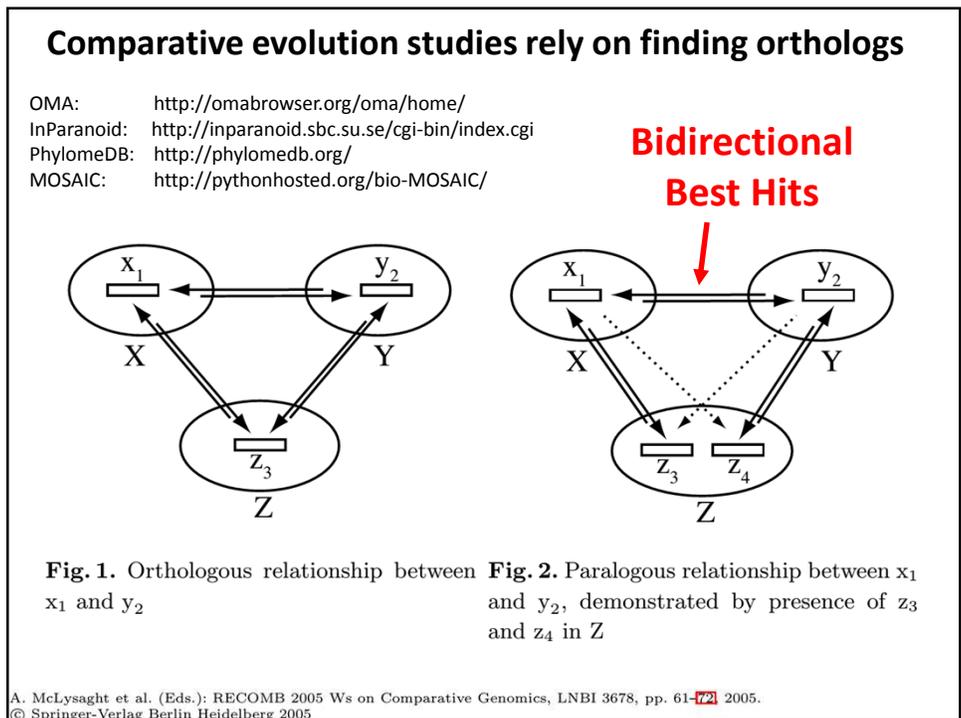
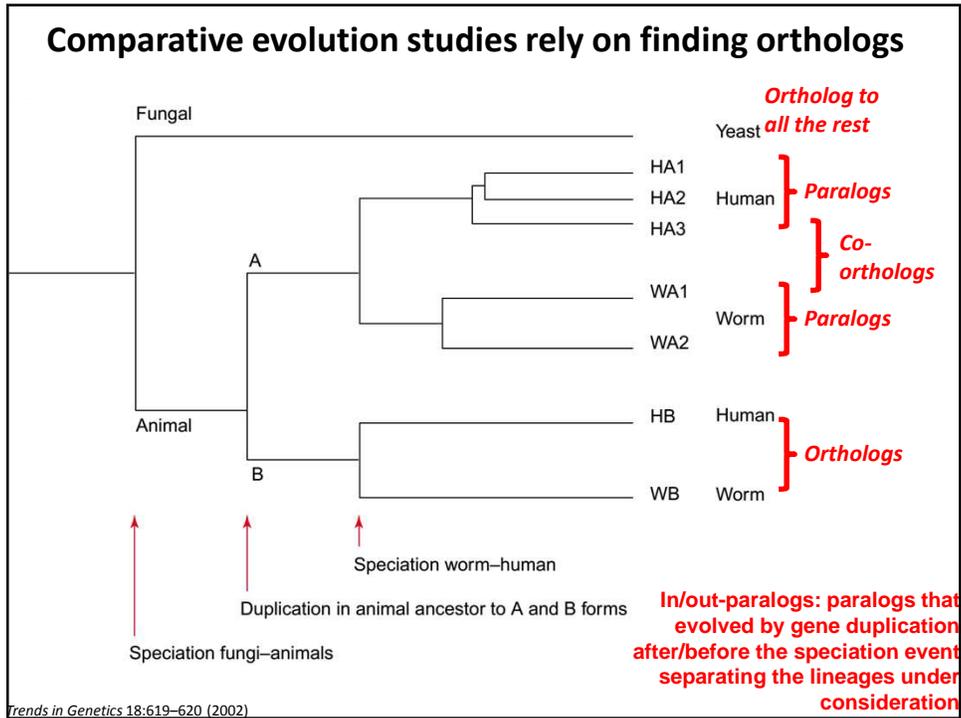
How do deeply conserved gene networks relate to traits, structures, and diseases in different organisms?



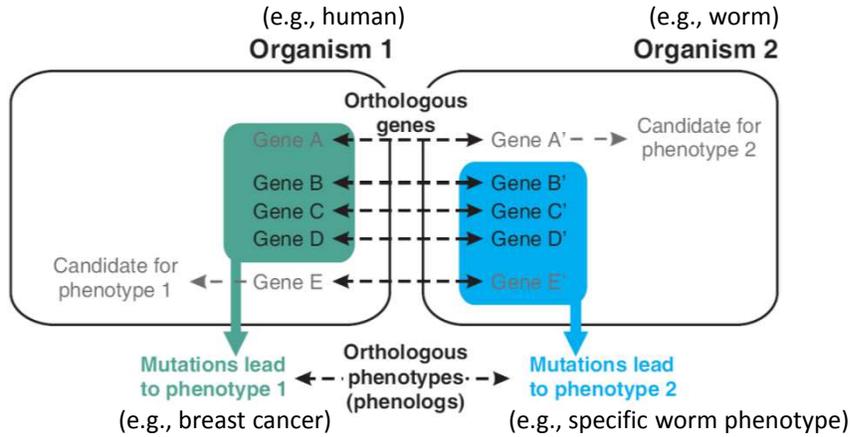
Comparative evolution studies rely on finding orthologs

Orthologs = genes from different species that derive from a single gene in the last common ancestor of the species

Paralogs = genes that derive from a single gene that was duplicated within a genome



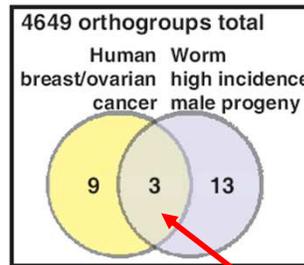
Phenologs = significantly overlapping sets of orthologous genes, such that each gene in a given set gives rise to the same phenotype in that organism



McGary, Park *et al.* PNAS 107:6544-9 (2010)

An example phenolog: a high incidence of male *C. elegans* maps to human breast/ovarian cancers

Human/Worm Ortholog	Linked to breast cancer in humans	Linked to more males in worms
ATM/atm-1	X	
BRIP1/dog-1	X	
KRAS/let-60	X	
PHB/phb-1	X	
PIK3CA/age-1	X	
RAD51/rad-51	X	
RAD54L/rad-54	X	
SLC22A18/C53B4.3	X	
TSG101/tsg-1010	X	
BARD1/brd-1	X	X
BRCA1/brc-1	X	X
CHEK2/chk-2	X	X
FAM82B/F33H2.6		X
GCC2/hcp-1, hcp-2		X
HMG20A, B/W02D9.3		X
HORMAD2, 1/him-3, htp-1, 2		X
KIF15/klp-10, 18		X
MRE11A/mre-11		X
PIGA/D2085.6		X
RAD1/mrt-2		X
RAD21/coh-1		X
SEH1L/npp-18		X
SVIL/viin-1		X
TSPO, BZRPL1/C41G7.3		X
WDHD1/F17C11.10		X



$p \leq 7.2 \times 10^{-6}$

includes
BRCA1

McGary, Park *et al.* PNAS 107:6544-9 (2010)

Building & searching a collection of phenotypes

Mining available databases +
manual collection from the primary literature



gene-phenotype
associations

<u>Organism</u>	<u># gene-phenotype associations</u>
human	1,923
mouse	74,250
worm	27,065
yeast	86,383
<i>Arabidopsis</i>	22,921

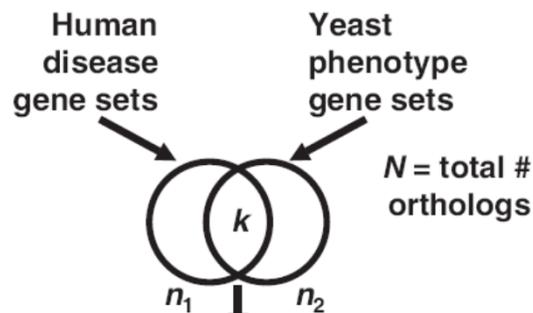
Spanning ~300 human diseases,
>7,000 model organism mutational phenotypes



**Computational scan phenotypes for novel models of a disease of interest,
identify significant phenologs using permutation tests**

McGary, Park *et al.*
PNAS (2010)

Discovering phenologs



Measure p (overlap $\geq k \mid n_1, n_2, N$) for each
disease-phenotype pair,
considering only human-yeast orthologs



**Identify all significant phenologs
by permutations or reciprocal best hits**

McGary, Park *et al.*
PNAS (2010)

There are 1,000's of phenologs between human diseases and mouse, yeast, worm, and even plant traits

Some cases we knew about already, serving as positive controls...

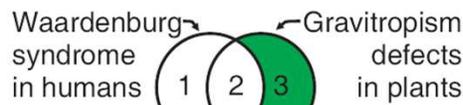
For example, genes for mouse cataracts suggest genes for human cataracts...

But many cases were surprising!

- | | |
|--|-------------------------------|
| <u>A defect in...</u> | <u>suggests genes for ...</u> |
| yeast lovastatin sensitivity | → angiogenesis defects |
| worm abnormal body wall muscle cell polarization | → gastrointestinal hemorrhage |
| yeast hydroxyurea sensitivity | → hemolytic anemia |
| plant cotyledon development defects | → mental retardation |
| <i>E. coli</i> chemical sensitivities | → chemically-induced seizures |

McGary, Park *et al.* *PNAS* 107:6544-9 (2010)
Woods, Blom *et al.* *BMC Bioinformatics*, 14:203 (2013)

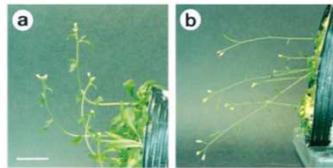
Example #2: plant *negative gravitropism* defects predict *Waardenburg syndrome*, a congenital disease with characteristic craniofacial, hearing, and pigmentation alterations



Vertebrate orthologs STX7/STX12, DDHD2/SEC23IP, and DNAJC13 are candidate Waardenburg genes



Waardenburg syndrome (accounts for ~2-5% of cases of deafness)



Plants failing to grow upwards

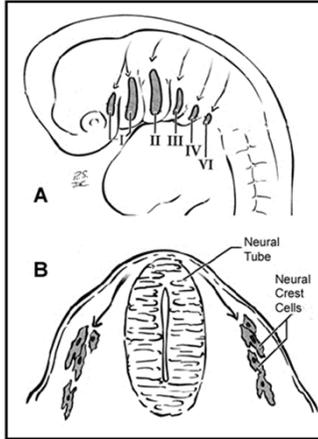
Michael Murphy, M.D.

Associated websites

Finkel *et al.*, *The Plant Journal* 14: 425-430 (1998)

Waardenburg syndrome is a defect of neural crest cells

Neural crest cells migrate during embryonic development



Heike & Hing, *Gene Reviews* (2009)

Some WS correlates in other animals:

Deafness in Dalmatian dogs (22% unilaterally deaf)



Variations in the Blenheim spot of Cavalier King Charles Spaniels



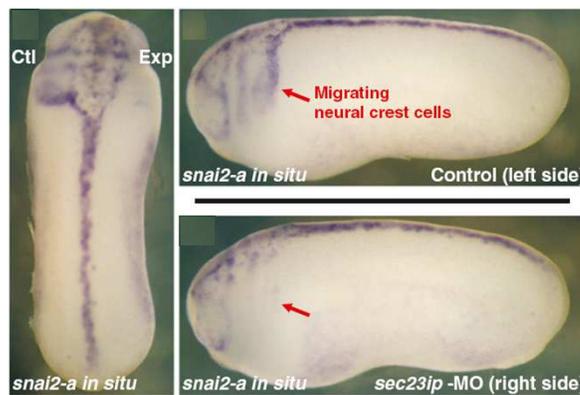
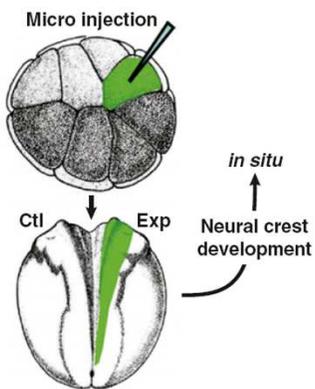
www.silversea.co.uk

Association between white blue-eyed cats and deafness (noted by Darwin in 1859)

White forelock and deafness/bowel blockage in foals

& many more...

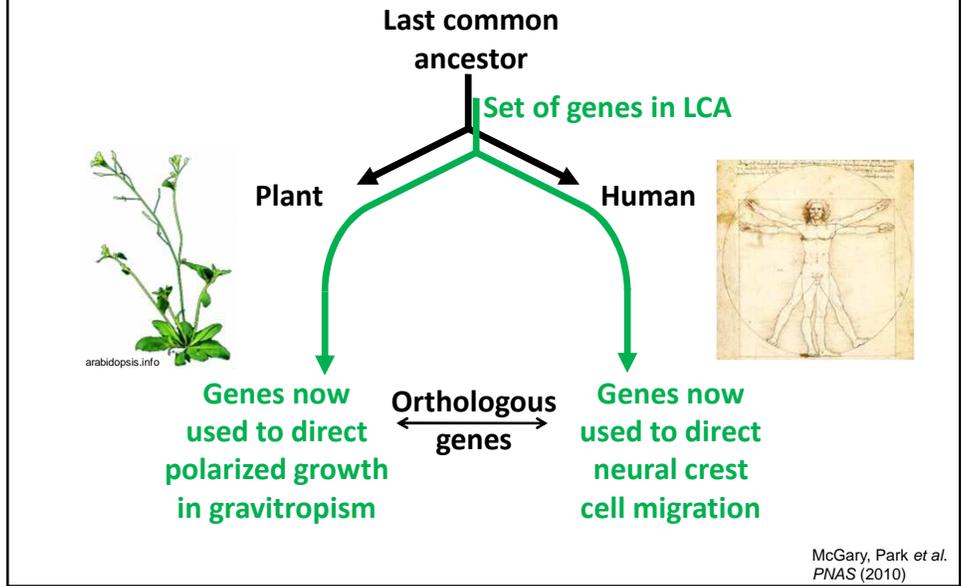
Inactivating SEC23IP—predicted from *Arabidopsis*—in a tadpole disrupts neural crest cells, consistent with Waardenburg syndrome



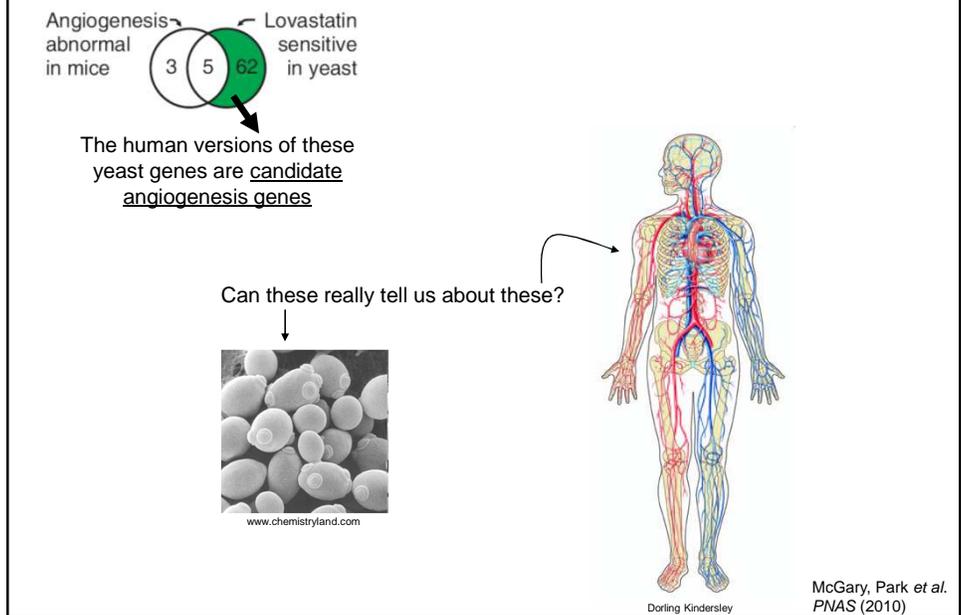
SEC23IP localizes to the neural crest cells & induces neural crest defects upon knockdown

McGary, Park *et al.* *PNAS* 107:6544-9 (2010)

Phenologs identify evolutionarily conserved systems of proteins relevant to particular traits/diseases.



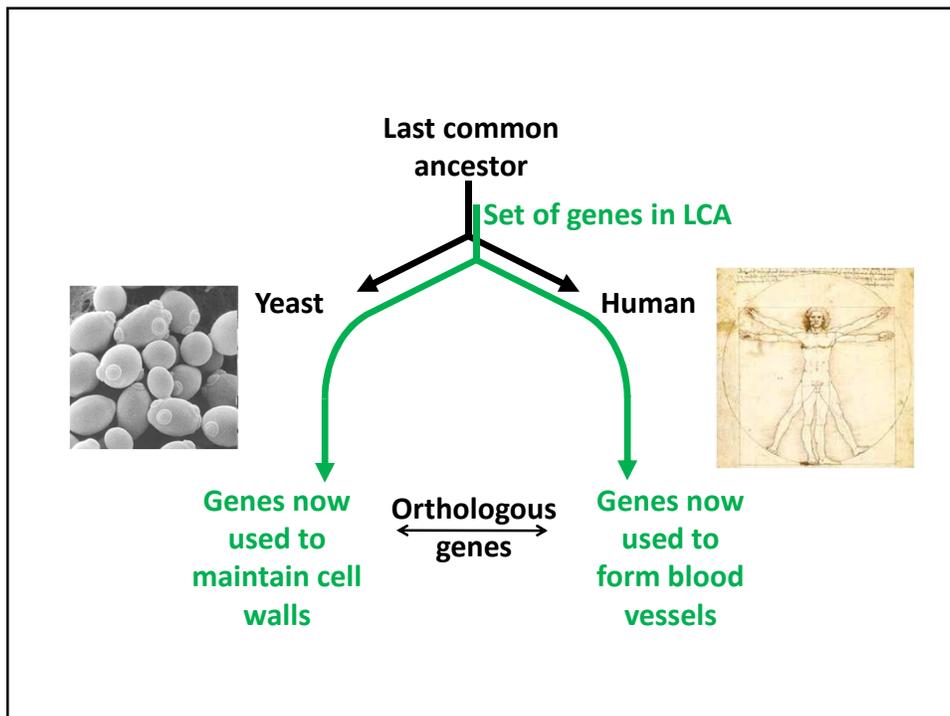
Example #3: Yeast genes linked to statin drug sensitivity predict mammalian blood vessel defects



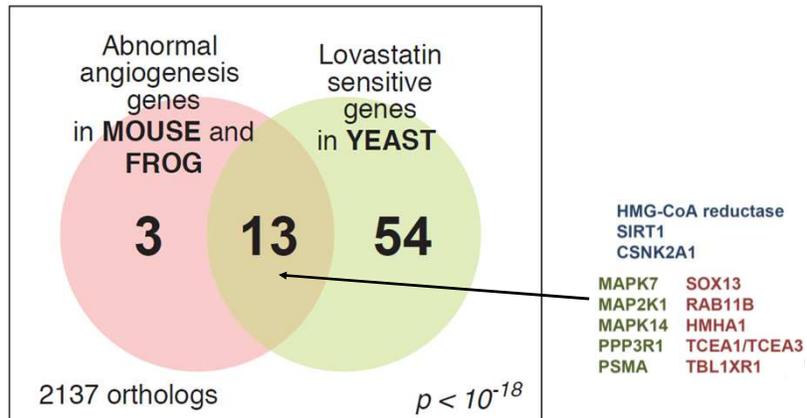
Disrupting the SOX13 gene causes strong blood vessel defects



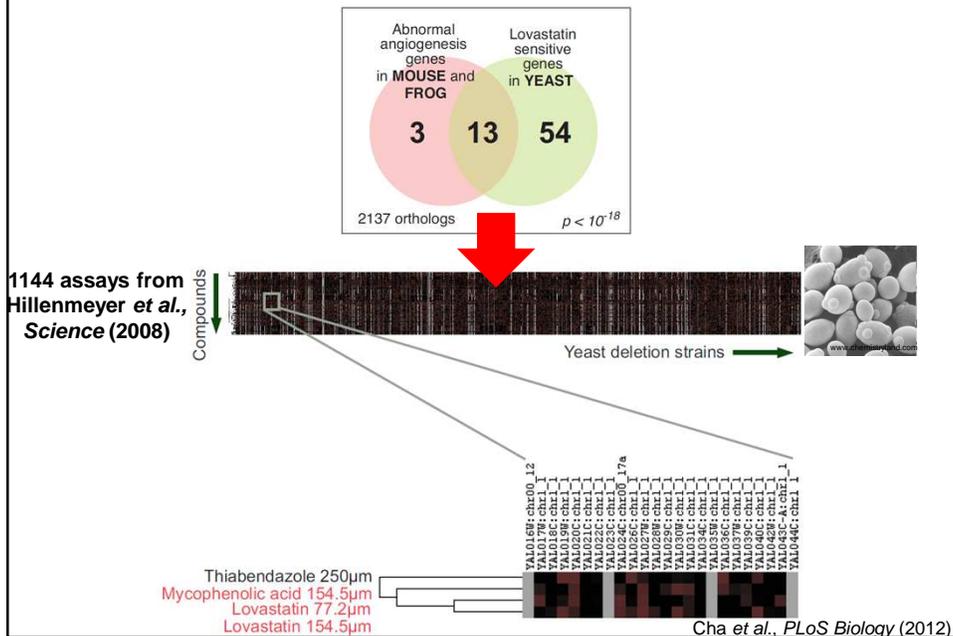
McGary, Park *et al.*
PNAS (2010)



The yeast/angiogenesis gene module



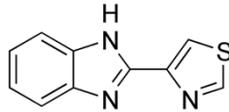
Chemicals that interact genetically with this module are candidate angiogenesis inhibitors



Screening for drugs that interact genetically with this yeast module led us to identify a new angiogenesis inhibitor

TBZ = thiabendazole

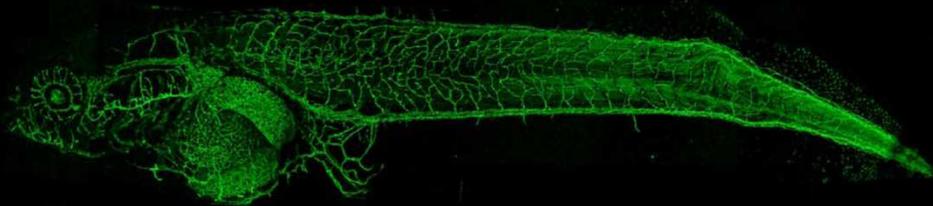
An FDA-approved antifungal drug with 40 years of safety data



- Approved by the U.S. Food and Drug Administration in 1967

- fungicide and parasiticide
- No mutagenic or carcinogenic effects
2 year safety trials in animals
- Off-patent, now marketed as a generic drug

Imaging the blood vessels of a living, transgenic tadpole in a dish of water



200 μm

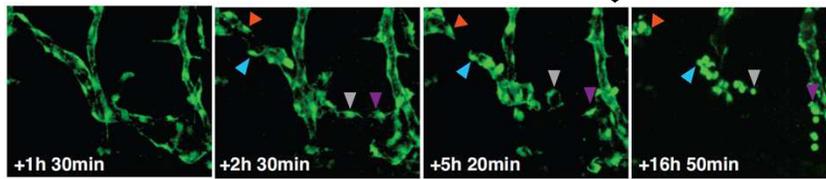
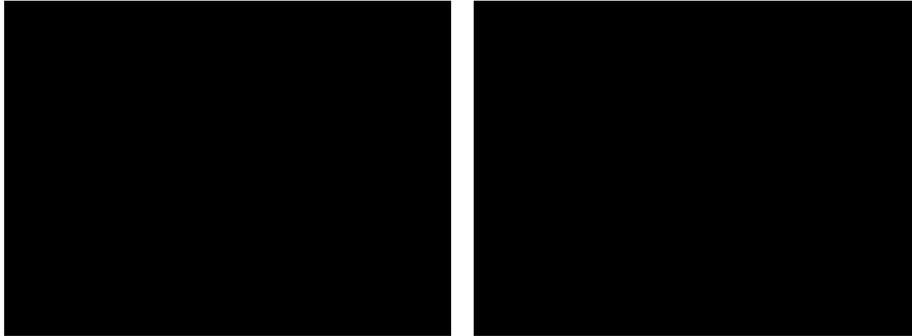
kdr:GFP transgenic *Xenopus laevis*

Image: Hve .li Cha

Thiabendazole disrupts vascular integrity, causing retraction and rounding of vascular endothelial cells

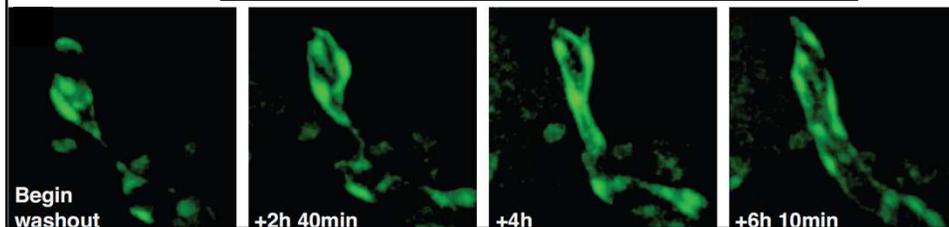
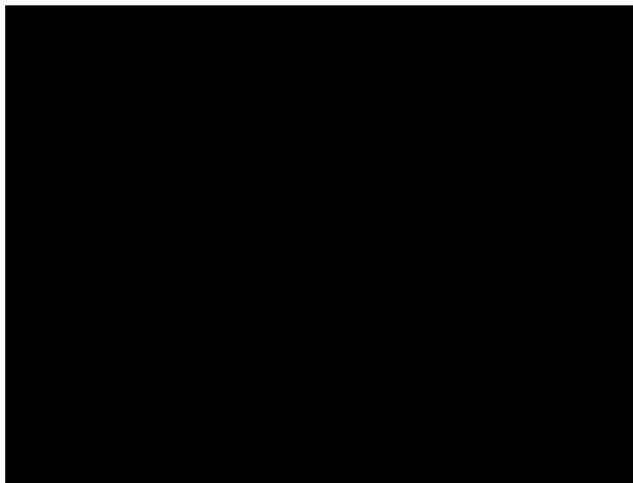
Control (DMSO carrier)

+ TBZ

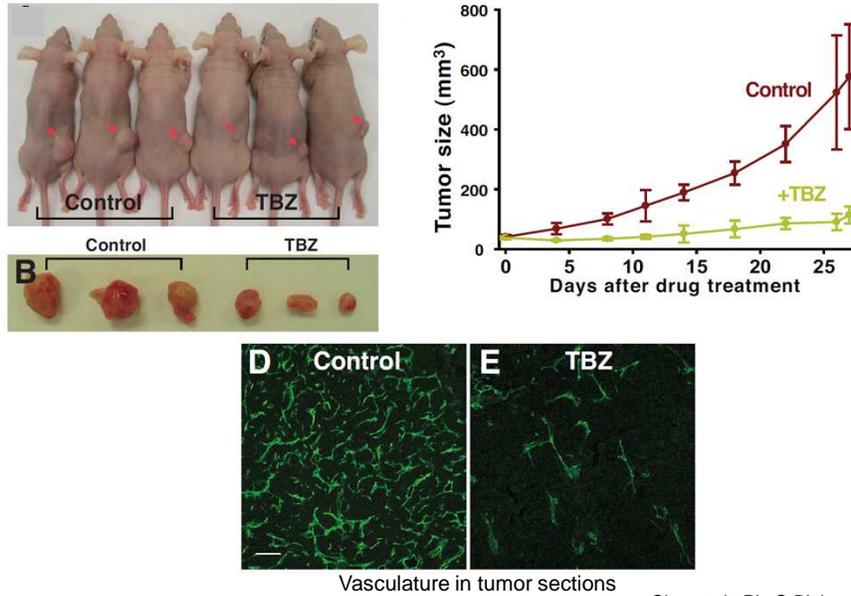


Cha et al., PLoS Biology (2012)

reversibly...

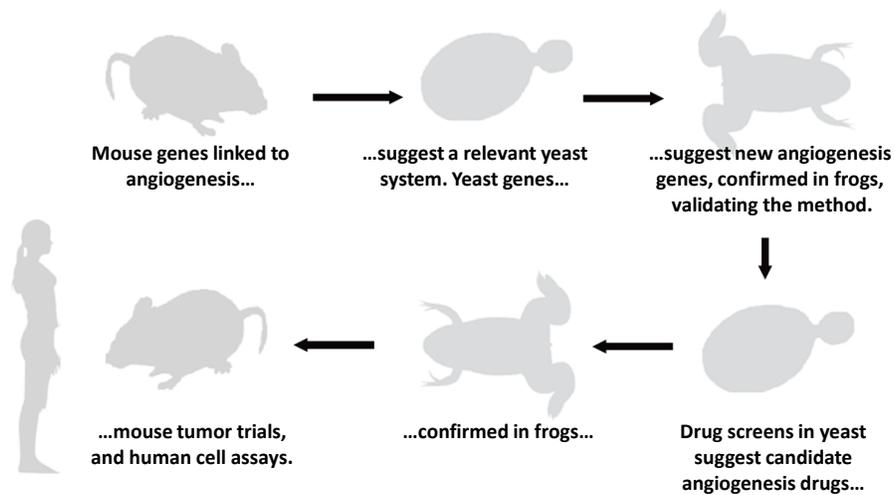


TBZ slows the growth of human fibrosarcoma tumors transplanted into immune-compromised mice



Cha et al., PLoS Biology (2012)

Summarizing the “road map” to a new vascular disrupting agent



Cha et al., PLoS Biology (2012)