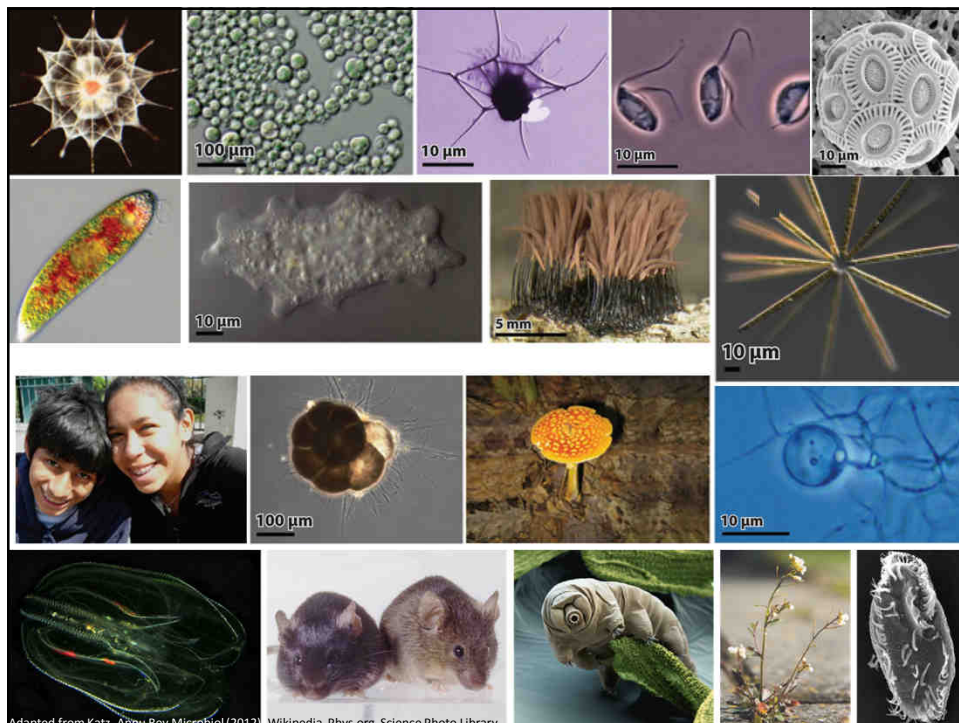


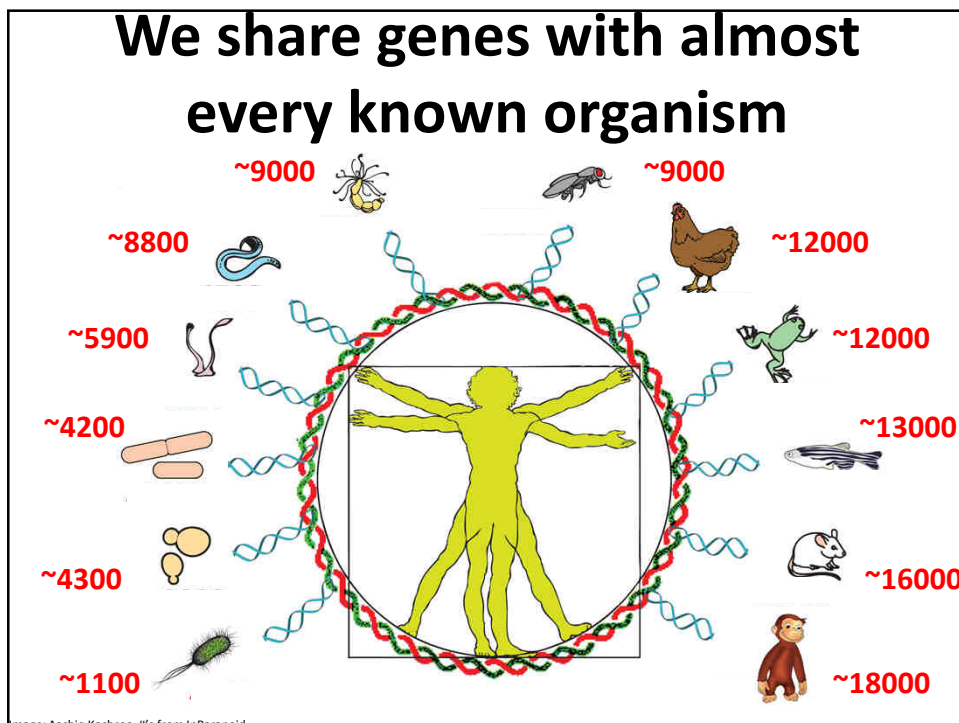
Phenologs

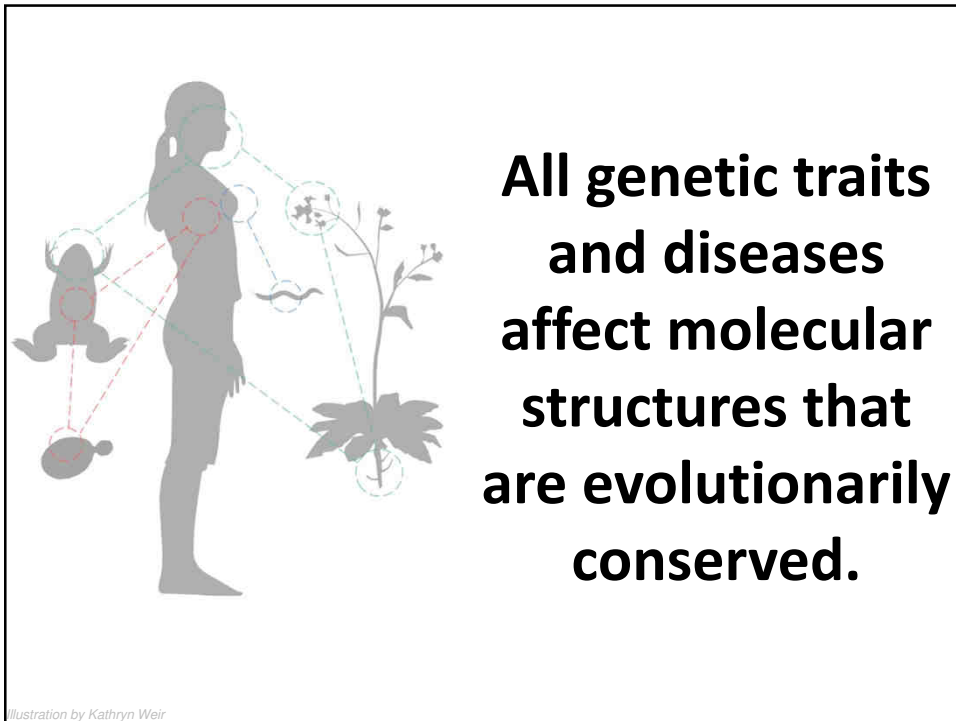
A case study of using bioinformatics to find new genes for genetic traits

BCH339N Systems Biology / Bioinformatics – Spring 2016

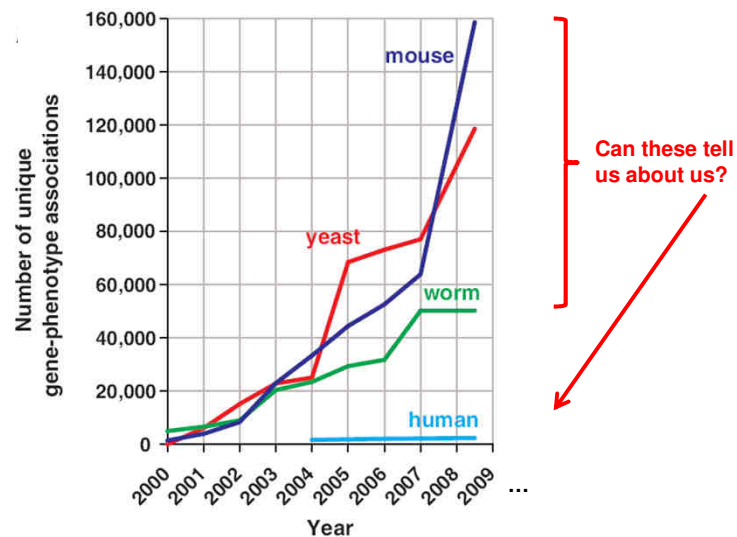
Edward Marcotte, Univ of Texas at Austin







We know far more about genes & traits in lower organisms than in us. Can we transfer that knowledge to us?



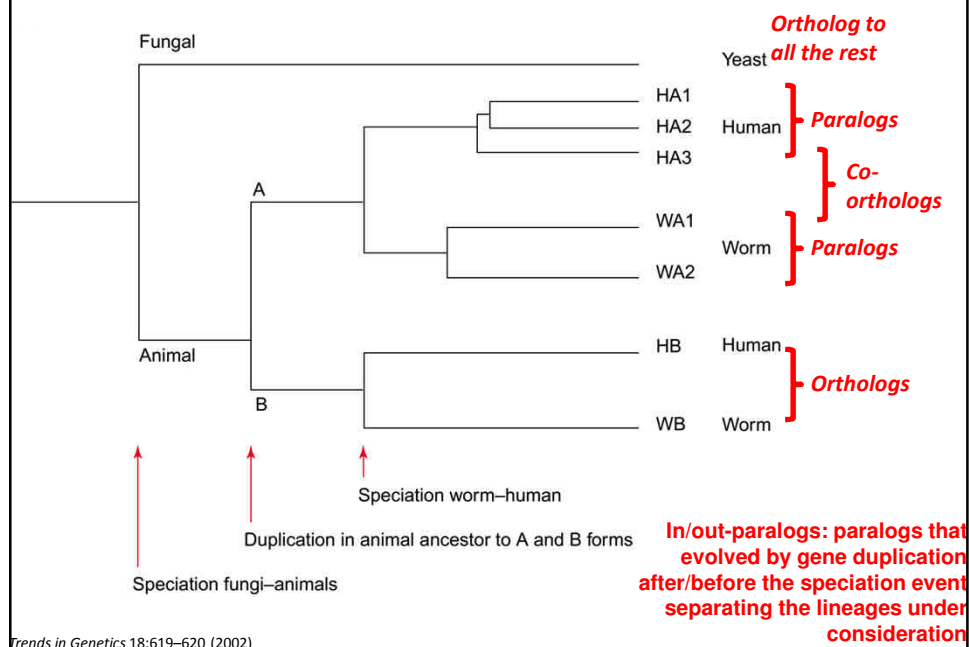
McGary, Park et al. PNAS 107:6544-9 (2010)

Comparative evolution studies rely on finding orthologs

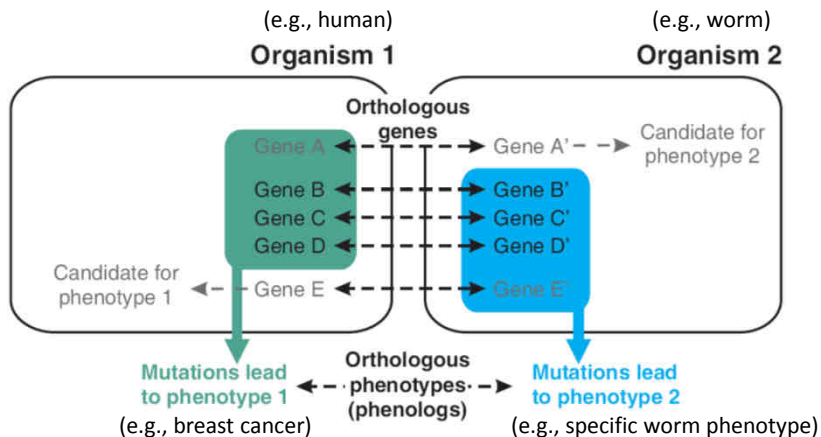
Orthologs = genes from different species that derive from a single gene in the last common ancestor of the species

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

Comparative evolution studies rely on finding orthologs

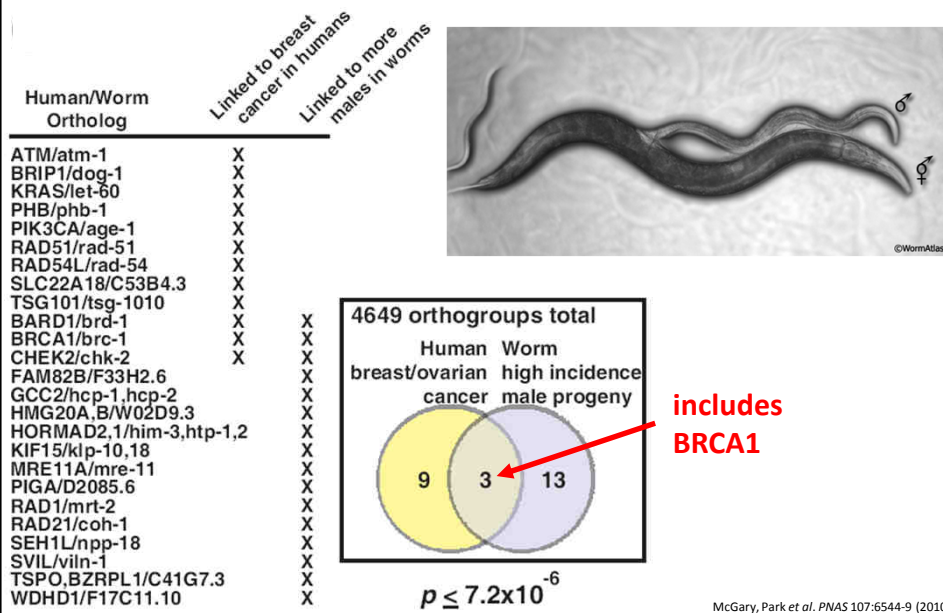


Phenologs = significantly overlapping sets of orthologous genes, such that each gene in a given set gives rise to the same phenotype in that organism



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

E.g., 'high incidence of male' *C. elegans* genes predict human breast/ovarian cancer genes



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

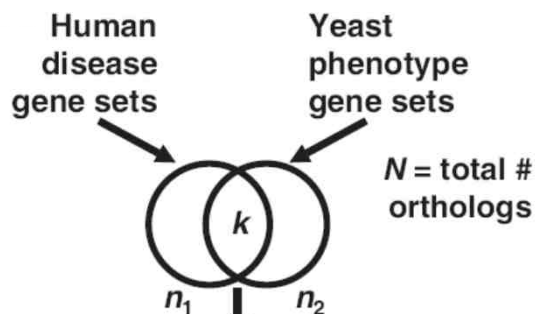
Spanning ~300 human diseases,
>7,000 model organism mutational phenotypes



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

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

**Computationally, we find many genes shared
between human diseases and
mouse, yeast, worm, and even plant traits**

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



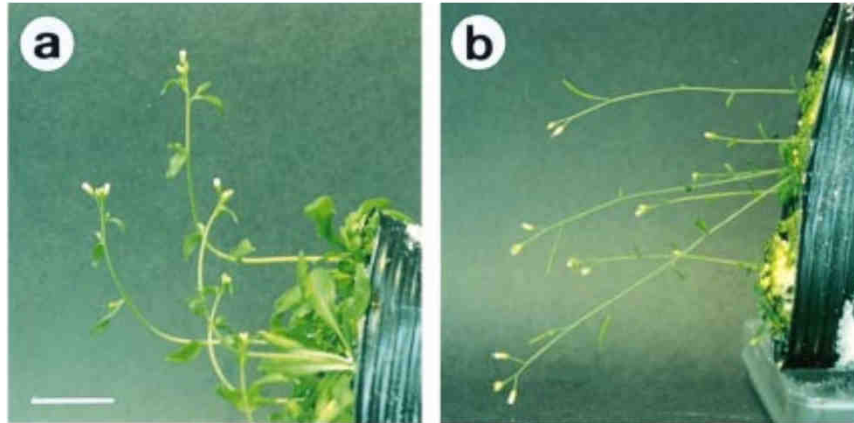
Michael Murphy, M.D.

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



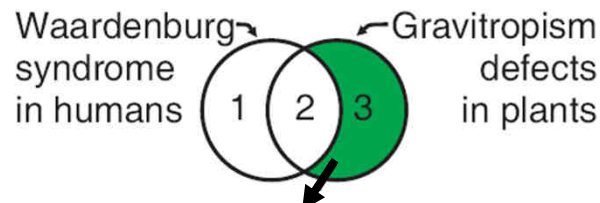
Associated websites: <http://www.varywell.com/waardenburg-syndrome-1045692>, <http://stephaniematesanchoz.blogspot.com/>

Plants sense and respond to gravity → gravitropism



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

Plant gravitropism predicts genes for Waardenburg syndrome, a human congenital deafness syndrome

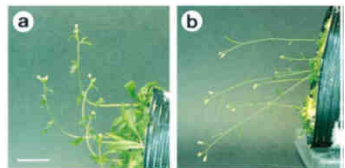


The human versions of these
plant genes are candidate
Waardenburg genes



Waardenburg syndrome

≈

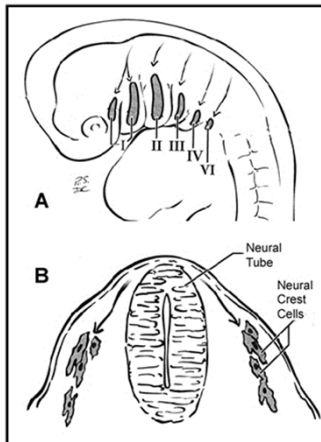


Gravitropism defects

Fukaki 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)



www.petplanet.co.uk

Variations in the Blenheim spot
Cavalier King Charles Spaniels

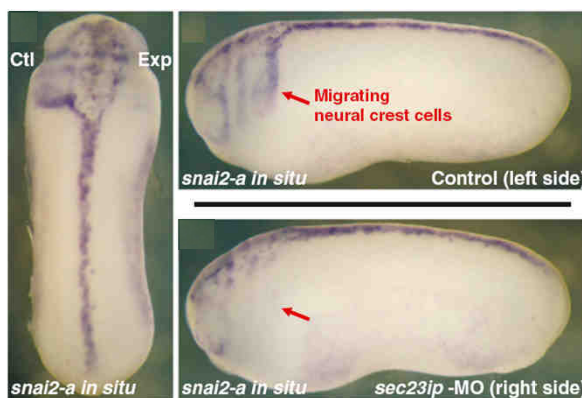
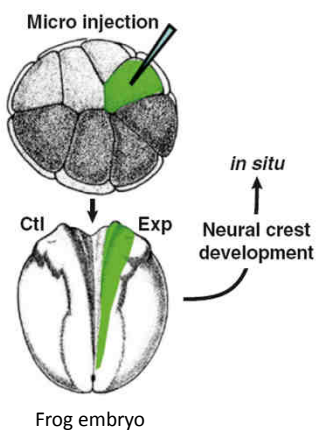


www.silvarcea.co.uk

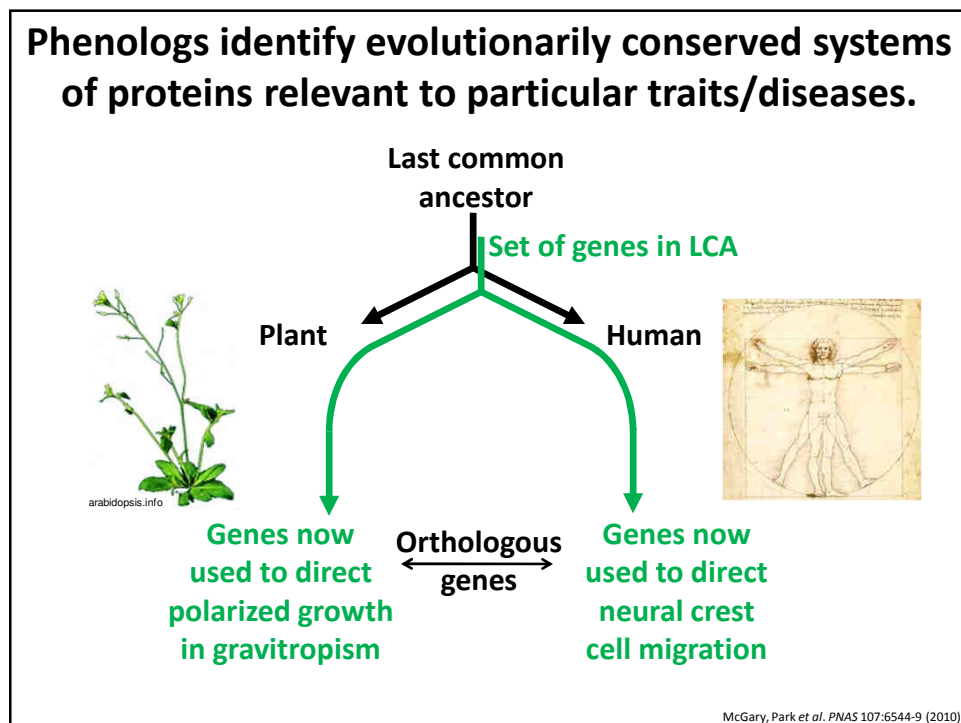
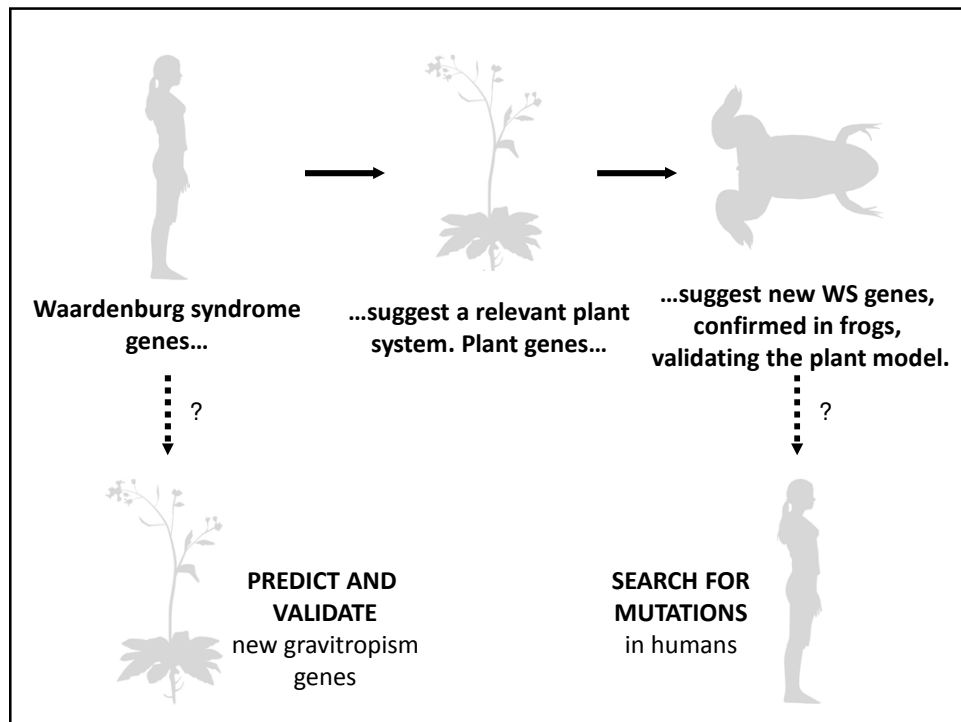
Association between white blue-eyed cats and deafness (noted by Darwin in 1859)

White forelock and deafness/bowel blockage in foals
& many more...

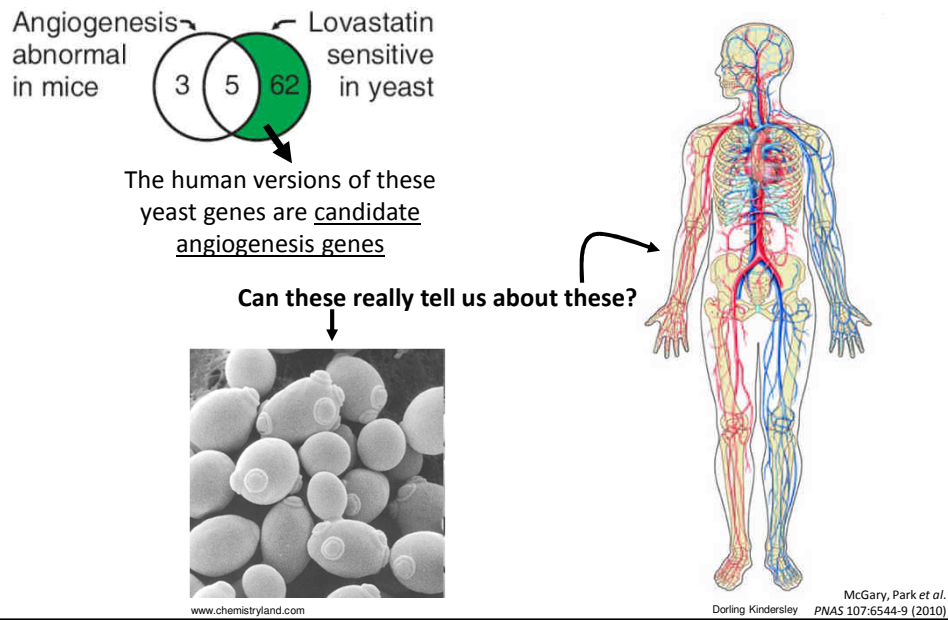
**Sure enough, inactivating one of the genes—
predicted from plants—in a tadpole disrupts neural
crest cells, consistent with Waardenburg syndrome**



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



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

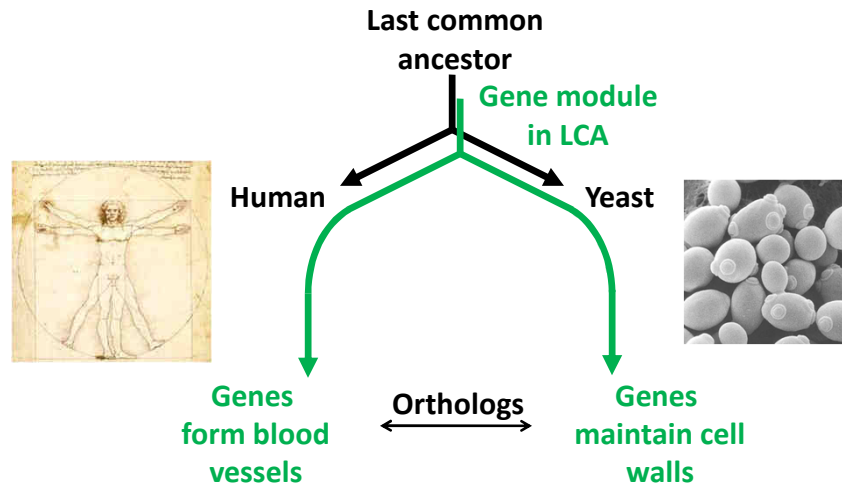


Disrupting the SOX13 gene causes strong blood vessel defects



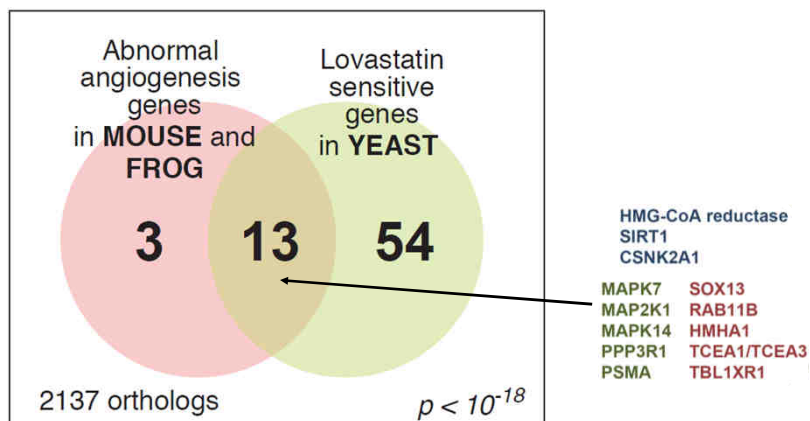
McGary, Park et al.
PNAS 107:6544-9 (2010)

A yeast model of angiogenesis = example of a deeply conserved, but “repurposed” gene module



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

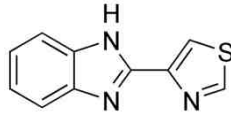
The yeast/angiogenesis gene module



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

TBZ = thiabendazole

FDA-approved antifungal drug with 40 years of safety data



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

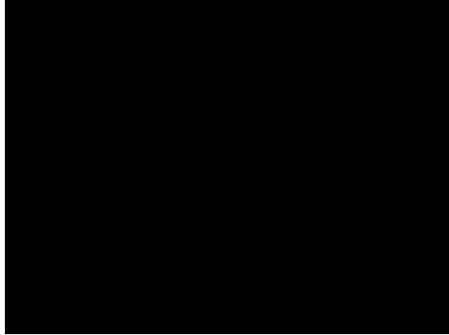
- Fungicide and parasiticide
- Not mutagenic or carcinogenic; 2 year dog safety trials
- Off-patent, marketed as a generic

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

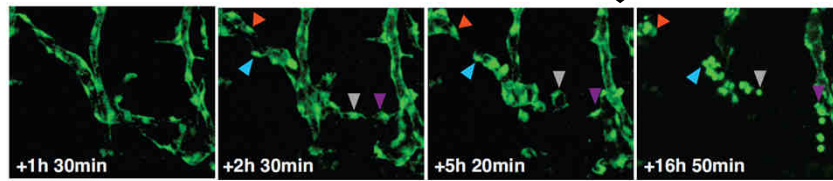


TBZ disrupts vascular integrity, making vascular endothelial cells retract & round up

Control (DMSO carrier)

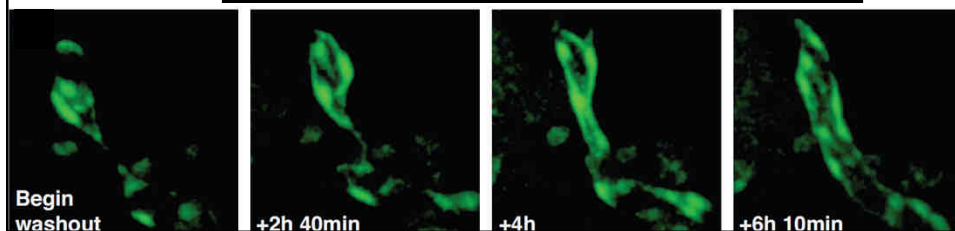
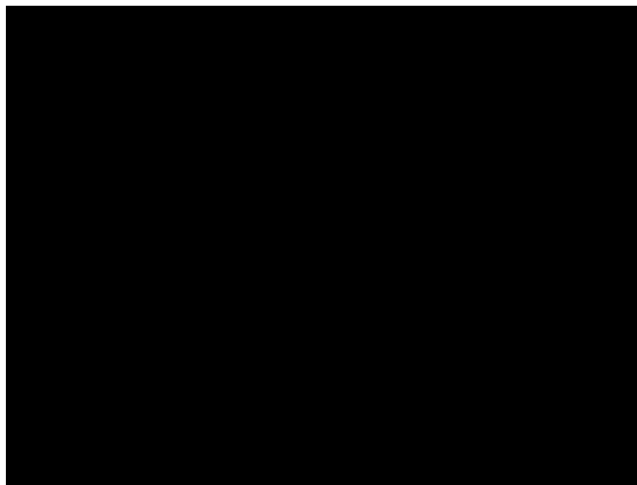


+ TBZ

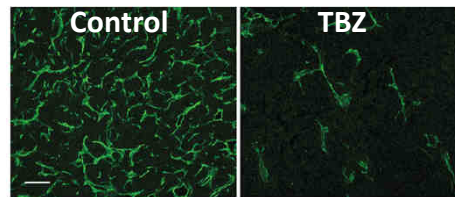
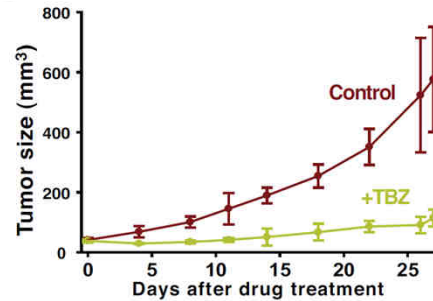
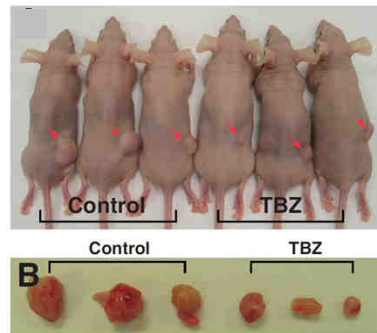


Cha et al., *PLoS Biology* (2012)

reversibly...



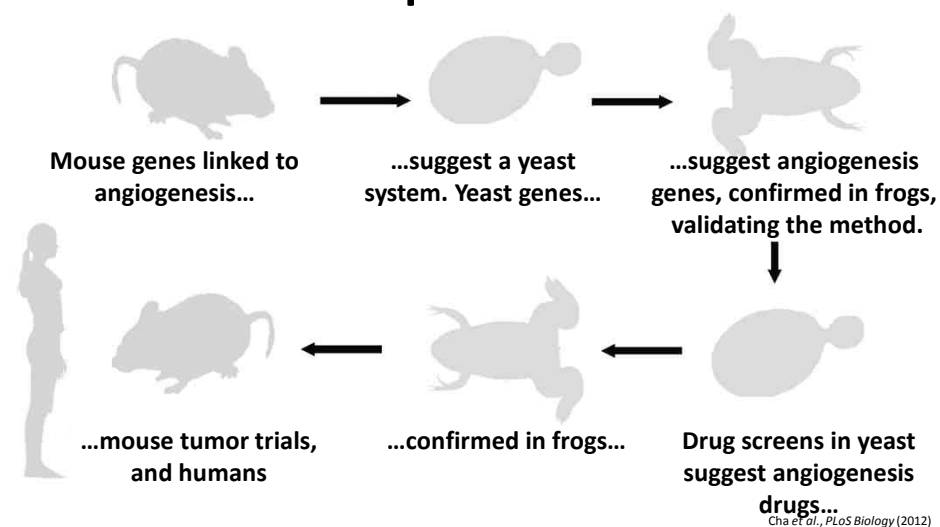
TBZ slows human fibrosarcoma tumors transplanted into immune-compromised mice



Vasculature in tumor sections

Cha et al., *PLoS Biology* (2012)

“Road map” to a new vascular disrupting agent, by mapping phenotypes across species



Cha et al., *PLoS Biology* (2012)