

Learning from Whales: Identifying key genes in genetic pathway responses to low oxygen

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Background

- Oxygen levels are declining in warming oceans, posing challenges for marine life
- Marine mammals tolerate oxygen scarcity (hypoxia) during their dives, making them a great model to understand the low oxygen response
- Responses to hypoxic conditions in early eukaryotes include physically moving away (cell motility) or reducing metabolic rate
- The stress response may also play a role in affecting the hypoxia response

How do cell motility, stress, and metabolism interact with the hypoxia response?



Methods

Data sourcing:

- Gene sets of hypoxia and glucocorticoid pulled from the Molecular Signature Database (MSigDB)

Analysis:

- Gene regulatory networks created using the Cytoscape framework with GeneMANIA application (Shannon et al., 2003)

RNA-sequencing:

- Human, dolphin, pilot whale, humpback whale, sperm whale, and beaked whale cells exposed to +/- hypoxia/cortisol
- Gene expression quantified with RNAseq

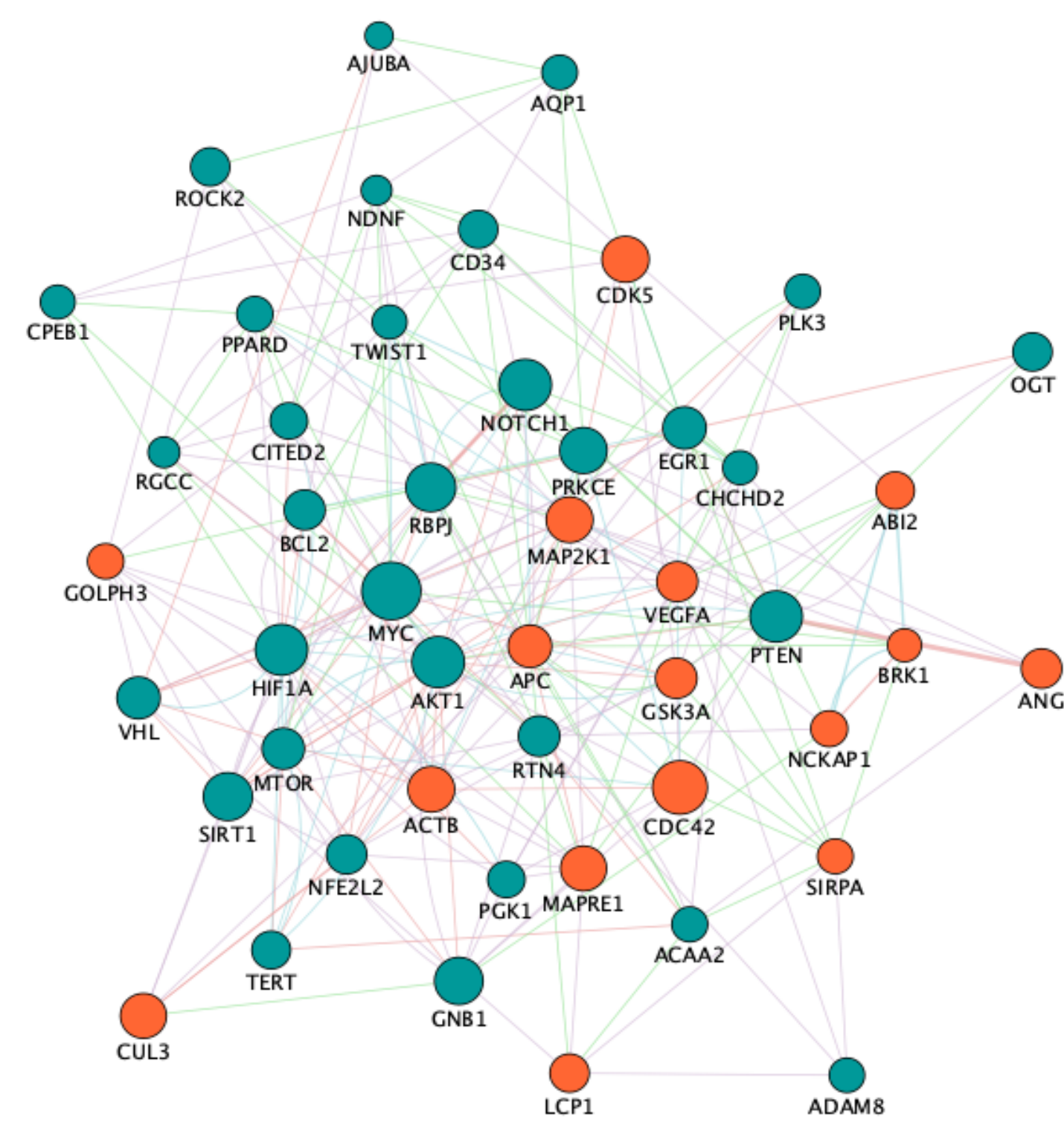
Results *Do these three genetic responses help tolerate low-oxygen conditions?*

Cell Motility

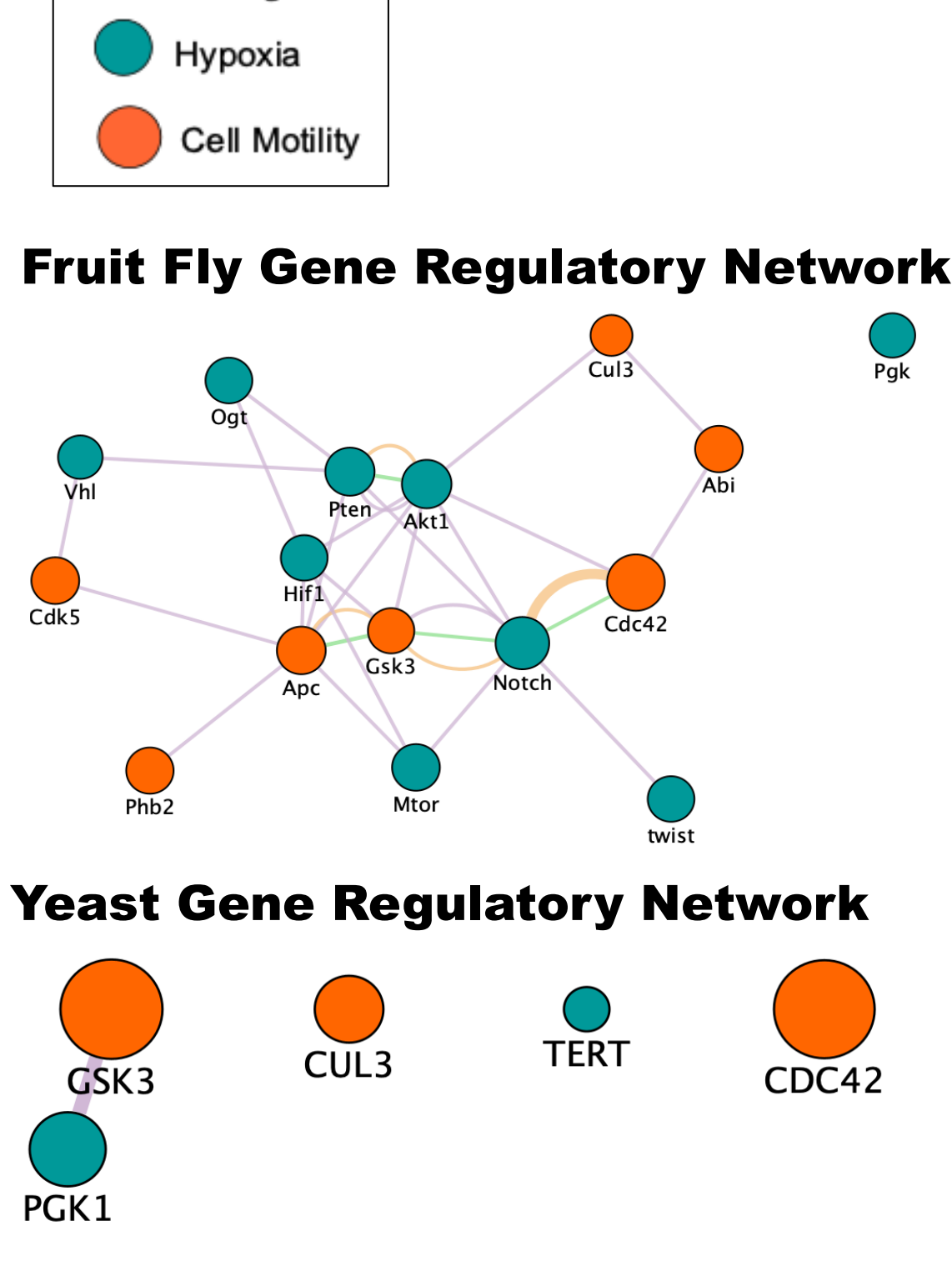
Evidence for coupling of the hypoxia and cell motility pathways in early eukaryotic evolution

- PGK1 and GSK3 are co-expressed even in non-motile yeast

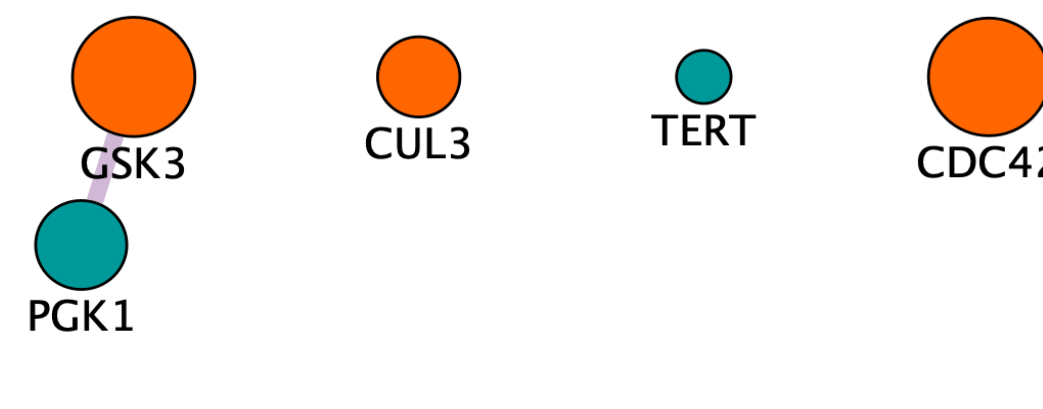
Human Gene Regulatory Network



Fruit Fly Gene Regulatory Network



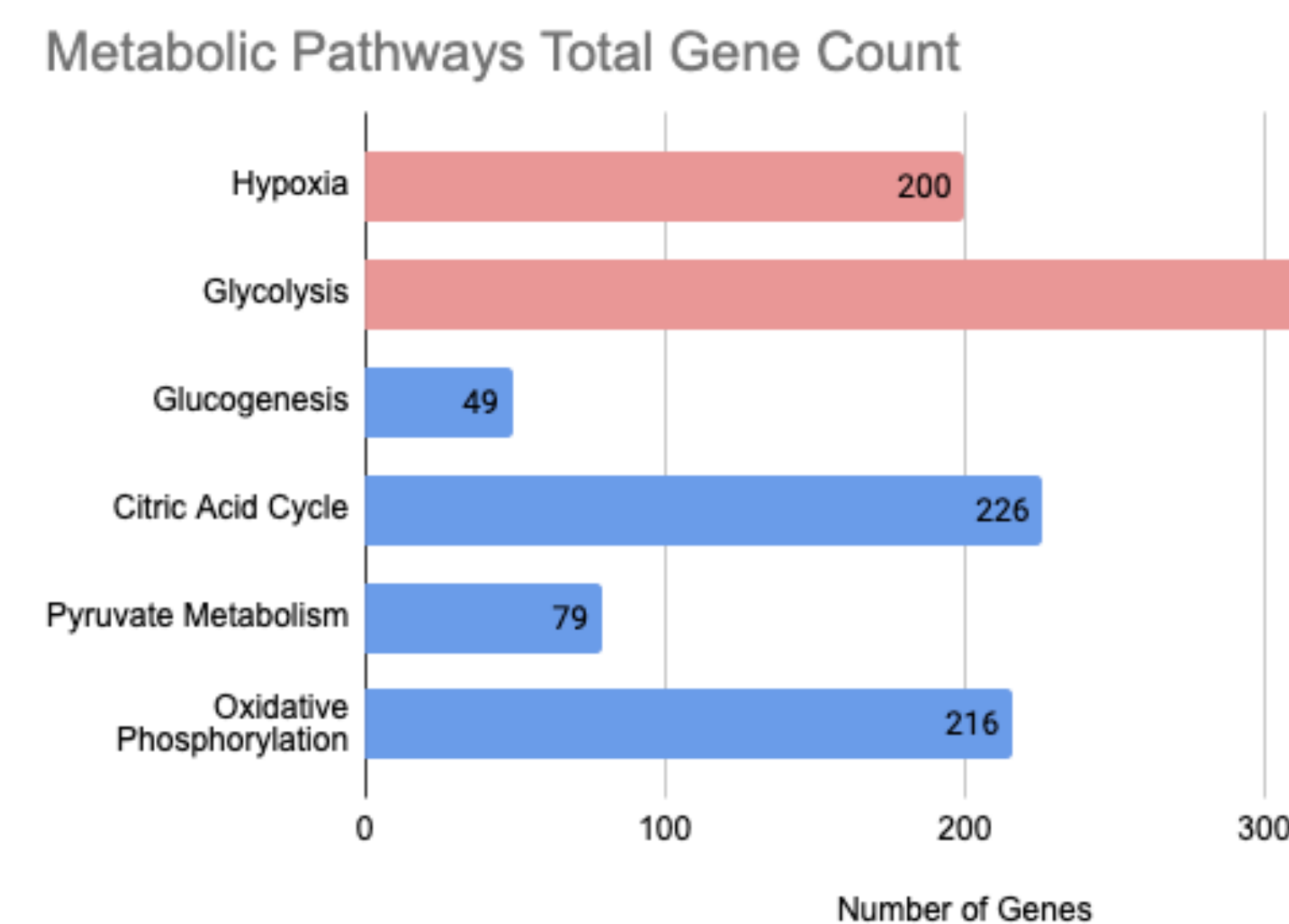
Yeast Gene Regulatory Network



Metabolism

The most genes present are used for anaerobic respiration

- Glycolysis is overrepresented, with 337 genes in the genome

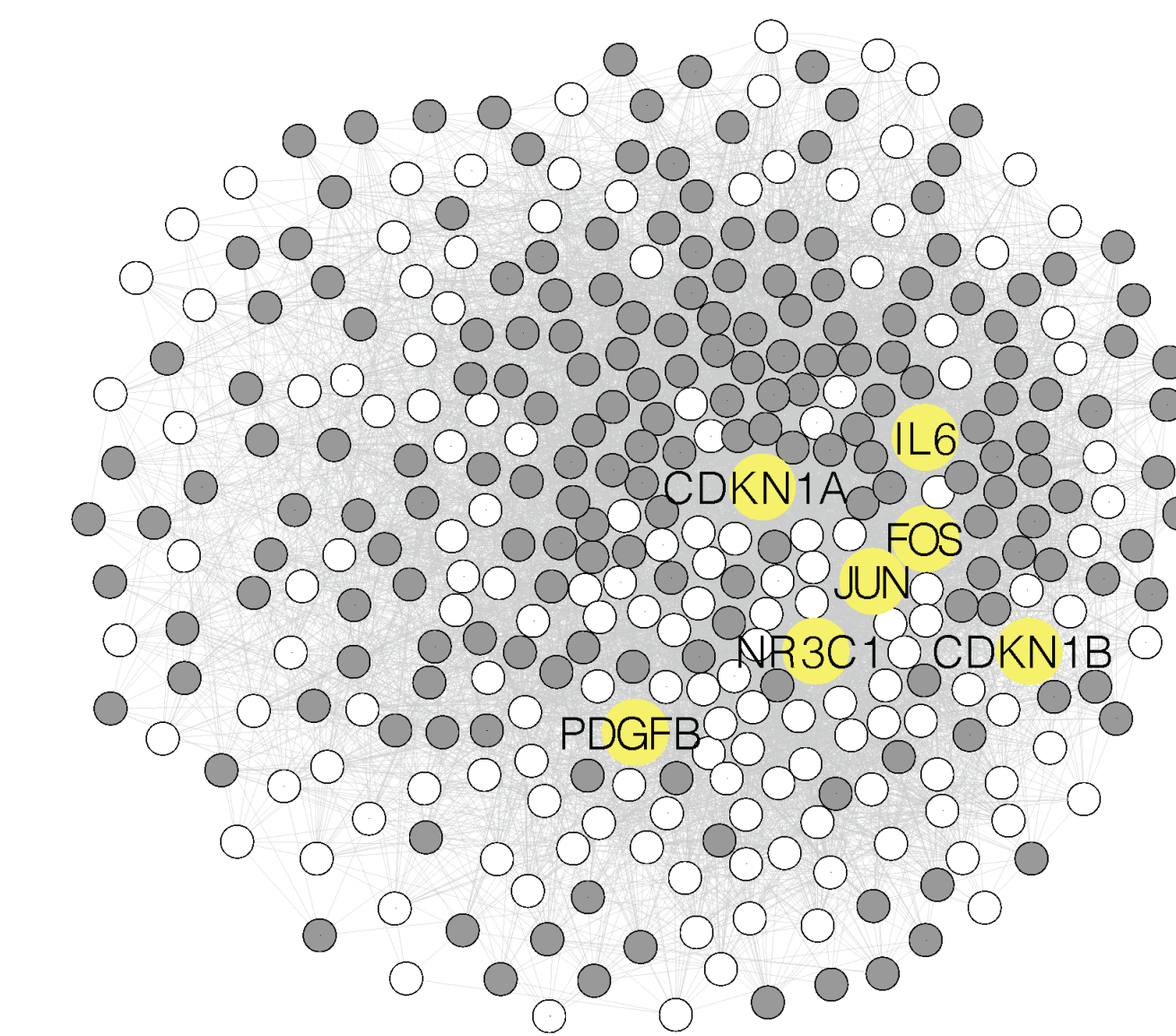


Takeaways and Applications

Marine Mammal Conservation

- Knowing how stress reduces low oxygen response in marine mammals can guide policy regulating anthropogenic stressors

Stress Hormones

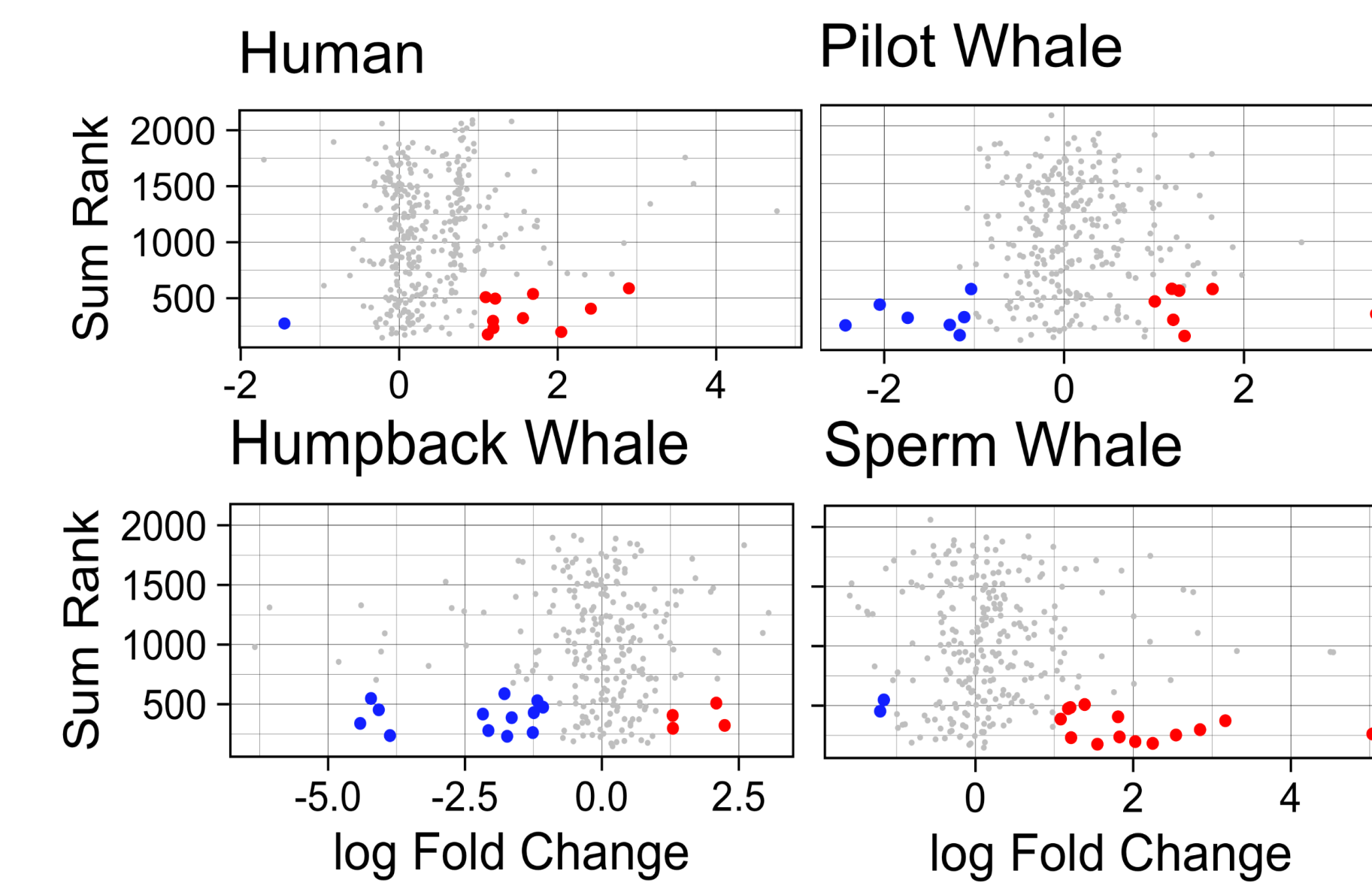
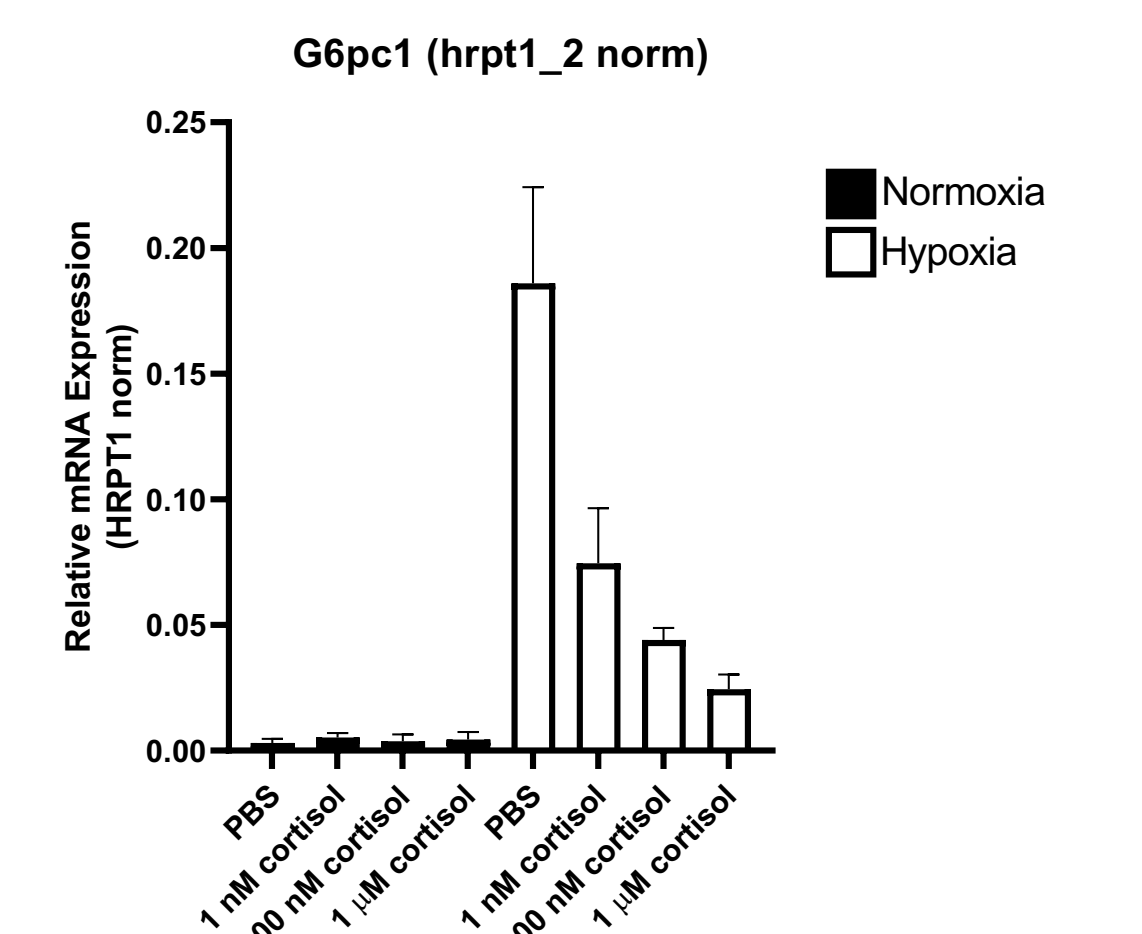


Hypoxia and GR pathways are highly interconnected

- GR: glucocorticoid – the molecular stress response
- NR4A2, ADM, and LOX are key genes mediating hypoxia GR interaction that were found only in the marine mammal species

Adding cortisol to beaked whale cell cultures suppressed the hypoxia response

- Tested in Pck1, G6pc, Per1, SGK1, and Igfbp1 (known GR downstream targets)



However, an abundance of shared genes were found between aerobic metabolism pathways

- Oxidative Phosphorylation and the Citric Acid Cycle share 98 genes

