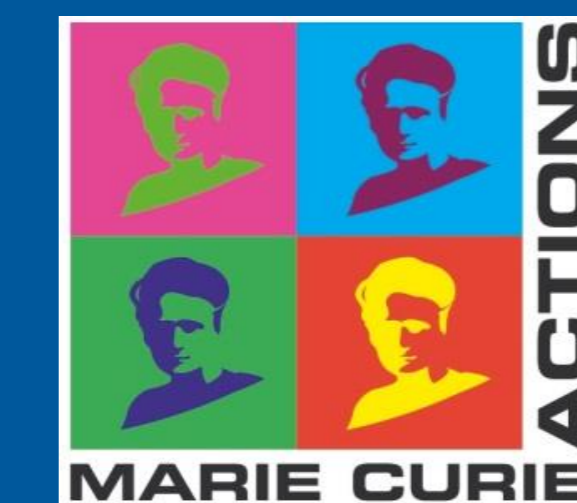


Glucocorticoid deficiency due to disruption of mitochondrial steroidogenesis leads to dysregulation of antioxidant pathways and nucleotide biosynthesis

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Aim

The role of glucocorticoids (GCs) as regulators of systemic homeostasis has been mainly studied by using synthetic GCs or in states of GC excess. Thus, the pathophysiologic consequences of cortisol deficiency on metabolic and biosynthesis pathways remain largely elusive.

Here we make use of a recently published ferredoxin (*fdx1b*) null-allele zebrafish line with massively decreased cortisol concentrations and a severely impaired stress response in order to define the global pathophysiologic response *in vivo* to glucocorticoid deficiency.

Summary

- Systemic profiling of the *fdx1b* null-allele zebrafish line was performed by a combination of RNA-sequencing and metabolomics analysis.
- An enrichment of genes in the *fdx1b* null-allele zebrafish line linked with pathways altered in metabolic disease was observed.
- This includes significant alteration in expression of genes and metabolites acting in pathways of energy and biomolecule synthesis (e.g., amino acids), and antioxidant pathways.

Conclusion

We provide *in vivo* evidence on the global pathophysiologic effects of GC deficiency, which can be vital for improving the understanding of the pathophysiology of adrenal insufficiency in humans.

Introduction

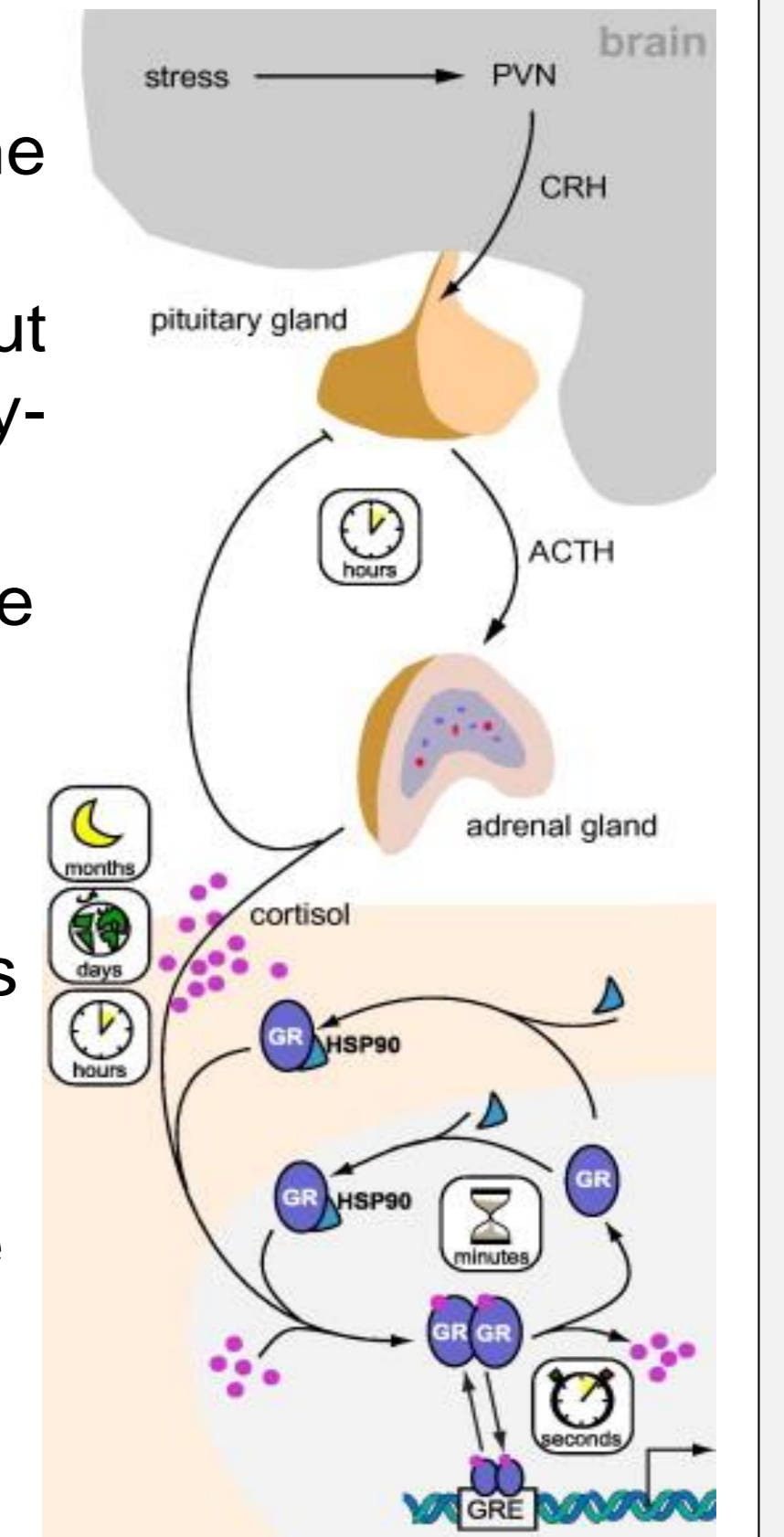
GCs are important steroid hormones for the regulation of physiology such as metabolism. Their release occurs in a circadian fashion, but is also regulated by the Hypothalamus-Pituitary-Adrenal gland (HPA) axis upon stress.

Zebrafish is a well-established vertebrate model for studying whole organism biology.

The zebrafish is day active and has a conserved endocrine system similar to human.

The main GC in both human and fish is cortisol.

A functional stress response in zebrafish is developed 4-5 days post-fertilization (dpf)

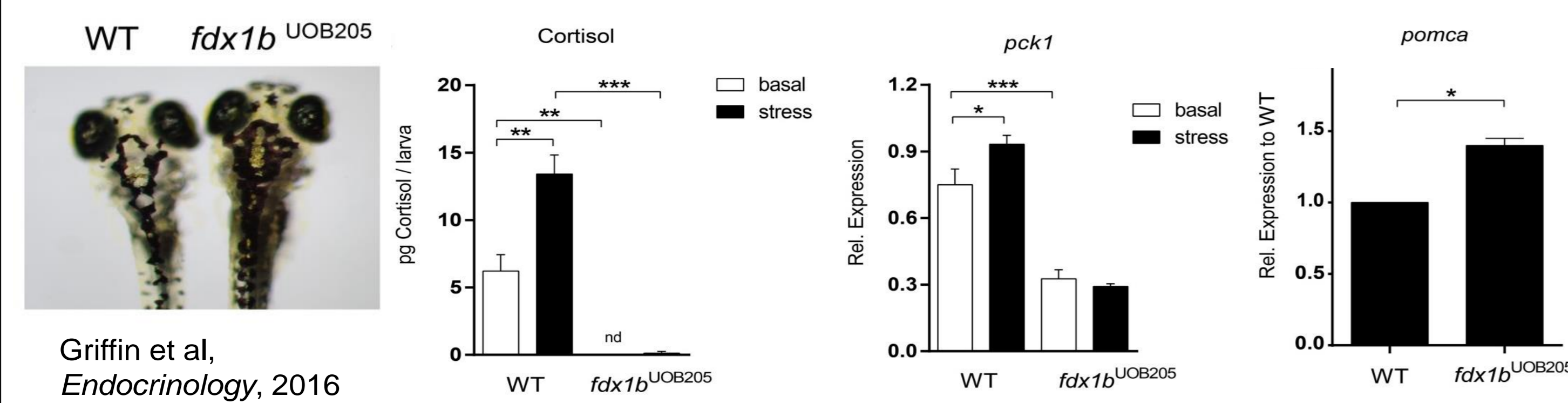


Weger BD, Weger M et al, ACS Chem Biol, 2012

Dickmeis T, Weger BD, Weger M, Mol Cell Endocrinol, 2013

Results

fdx1b null-allele zebrafish larvae are impaired in cortisol synthesis, cortisol regulated gene expression and in their stress response

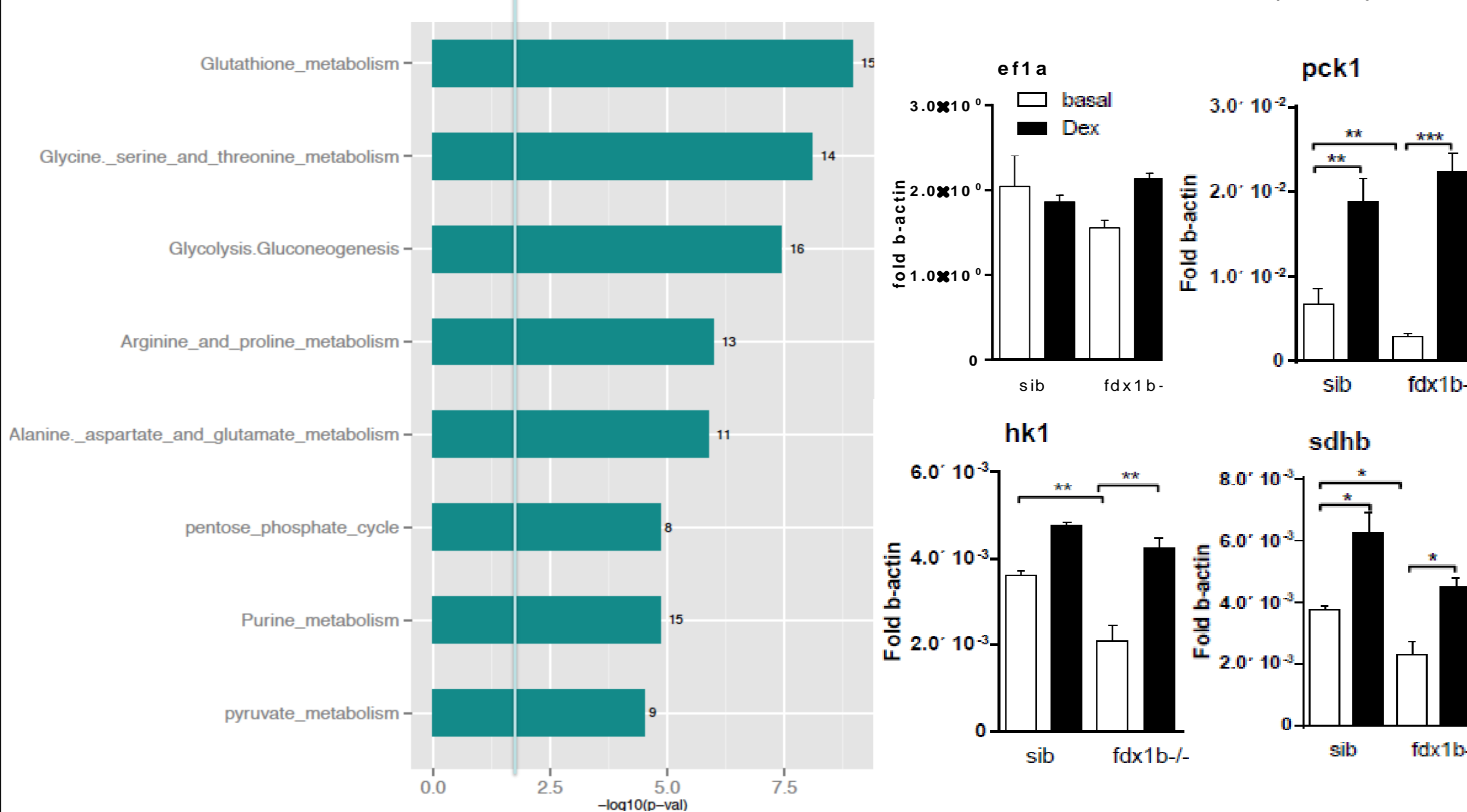


Enrichment of genes linked to metabolism and metabolic disease

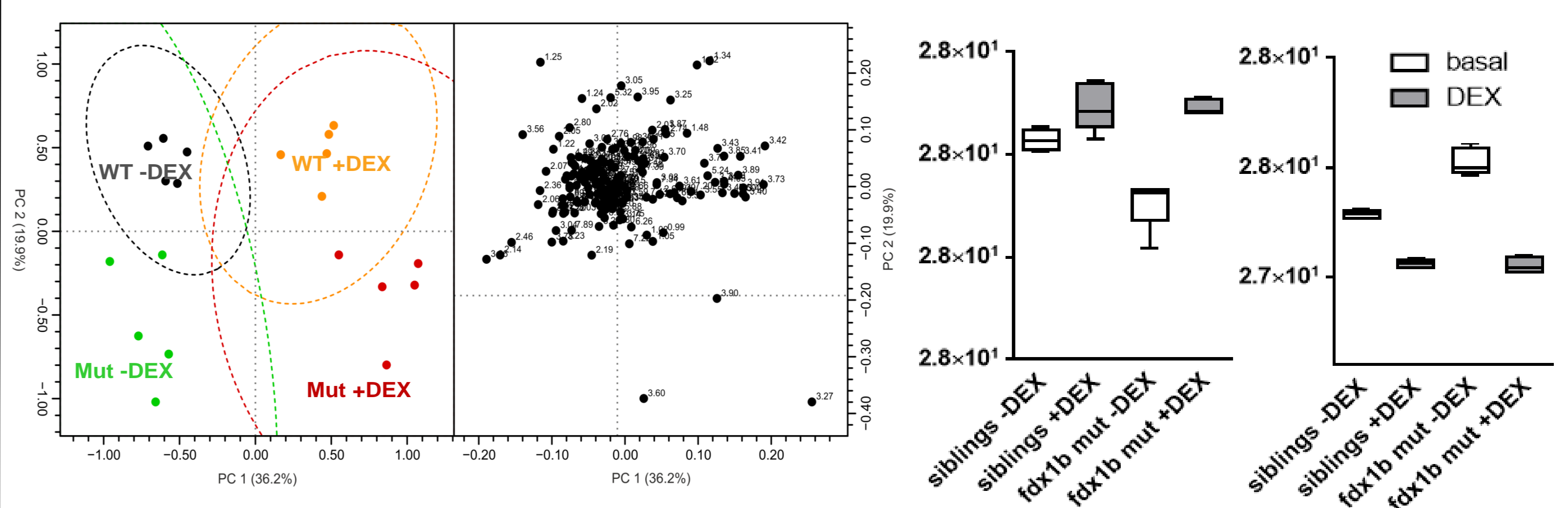
Pathway analysis reveals genes linked with metabolic disease

Top Diseases and Bio Functions	p-value	#Molecules
Metabolic Disease	1.17E-03 - 1.77E-11	166
Neurological Disease	1.26E-03 - 3.14E-11	228
Psychological Disorders	1.01E-03 - 3.14E-11	180
Cardiovascular Disease	1.23E-03 - 5.10E-09	89
Inflammatory Response	1.13E-03 - 2.57E-08	144

Genes involved in energy and biomolecule synthesis are altered in cortisol deficient *fdx1b* mutants, but can be rescued with dexamethasone (DEX).

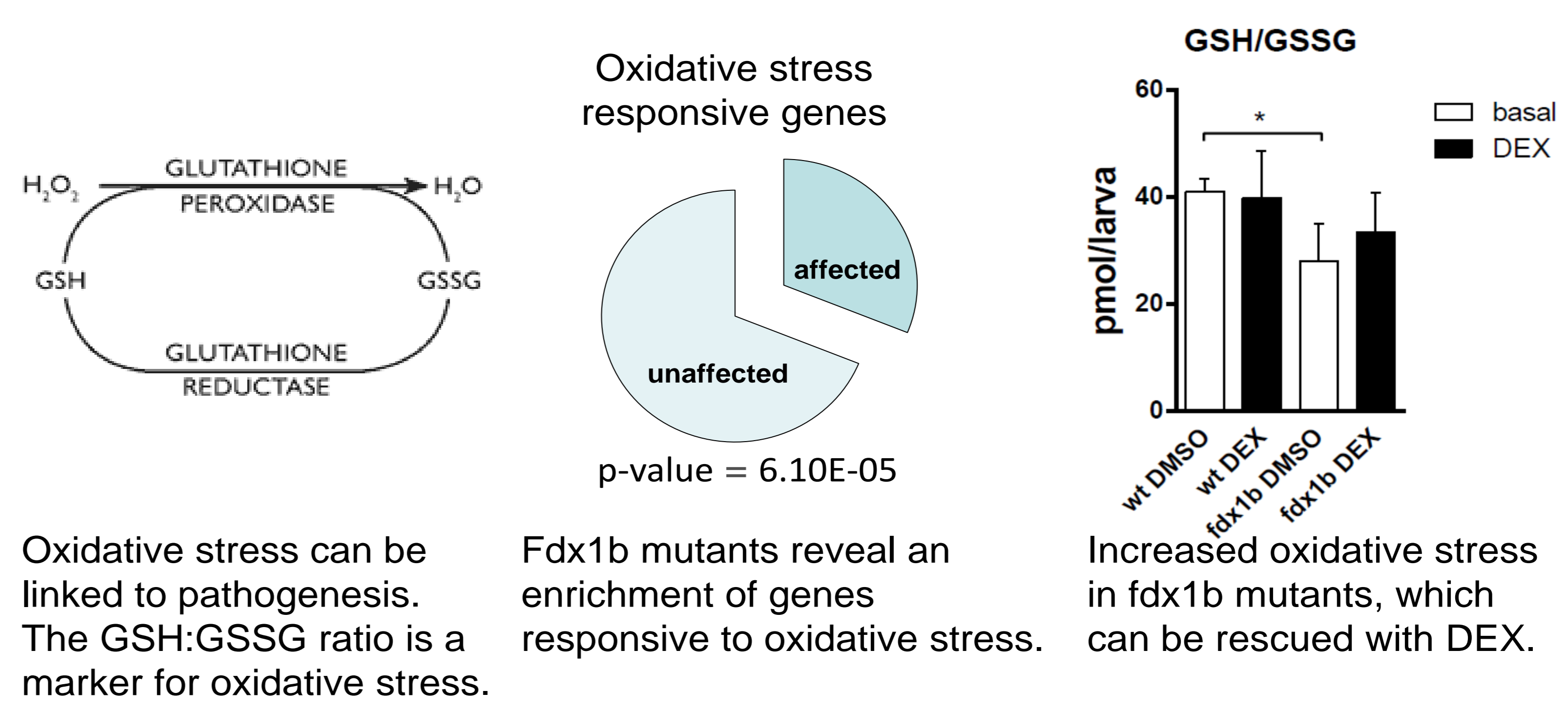


An enrichment of oxidative stress responsive genes can be observed in the cortisol deficient *fdx1b* mutant



Untargeted NMR analysis shows differences in the metabolomes between *fdx1b* mutants and wild-type siblings. Some of the altered metabolites can be rescued in the *fdx1b* mutants with DEX.

Alterations in oxidative stress levels in cortisol deficient *fdx1b* mutants



Material and Methods

Establishing a *fdx1* null-allele zebrafish mutant line using Transcription Activator-like Effector Nucleases (TALENs)

