

[†] The Contribution of Ploidy to Evolutionary Divergence of Gene Expression in Yeasts

Eric Delgado Regev Group Summer Research Program in Genomics

Ploidy Varies Among Organisms



Haploid



Triploid









The Evolutionary Contribution of Ploidy is Unclear



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An Evolutionary Advantage of Haploidy in Large Yeast Populations

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Antisense Transcription Controls Cell Fate in Saccharomyces cerevisiae

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To determine the set of ploidy regulated genes.

To determine whether differences in gene expression are the caused by species-specific traits, environmental factors or conserved ploidy dependent differences.

Unique Regulation of Ploidy-Dependent Genes May Be Due To Multiple Factors



Ascomycota Fungi Provide an Ideal System for Studying Ploidy



These species span over 300 million years of evolution

Experimental Approach

- Create stable haploids from diploid species
- Create stable diploids from haploid species
- Compare expression profiles among strains



Ascomycota Fungi Provide an Ideal System for Studying Ploidy



Isogenic Strain Construction Allows For Ploidy-Based Gene Expression Comparison

Diploid To Haploid



Isogenic Strain Construction Allows For Ploidy-Based Gene Expression Comparison



Cells

Organisms Included In Gene Expression Analysis



- S. cerevisiae
 - S288c Haploid
 - S288c Diploid
 - Bb32 Diploid
- S. paradoxus
 - Wild Parental
 - Parental Haploid
 - Diploid Isogenic To Haploid
- Single Channel Data For LOG and PLAT

Haploid and Diploid Global Expression Patterns Are Similar



MET17 IME4 ASG7 STE18



Several Groups of Genes Are Differentially Expressed In Haploids And Diploids



Meiosis Genes Are Differentially Expressed In Haploids And Diploids



Differential Gene Expression By Gene Type

Plateau Log

Increased Expression in Meiosis Genes Provides Molecular Basis For Ploidy Advantage



[•] Conclusions

- Clear ploidy expression signatures were identified
- Experimental approach validated by expected regulation of previously identified ploidy genes
- Novel ploidy genes identified
- Gene expression data may provide a molecular basis for evolutionary observations
- Future Direction
 - Expand comparisons across phylogeny
 - Introduce stress conditions

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