

Comparative Genomics and Core Fungal Genes

Tom Hsiang

Dept. Environmental Biology

University of Guelph

Guelph, Ontario, Canada

Genomic sequencing

- 1995: first prokaryote sequenced
 - *Haemophilus influenzae* (2 Mb)
- 1997: first eukaryote - fungus
 - *Saccharomyces cerevisiae* (12 Mb)
- 1998: first animal - nematode
 - *Caenorhabditis elegans* (97 Mb)
- 2000: first plant
 - *Arabidopsis thaliana* (115 Mb)
- 2002: *Oryza sativa* (400 Mb)
- 2004: *Populus trichocarpa* (5.6 Gb)

SIZE



Genomic sequencing - animals

- 2001, *Homo sapiens* (human, 2.9 Gb)
- Some insects, sea animals, mouse complete
- Rat, cow, chimpanzee, bee almost complete

Genomic sequencing - fungi

- Size of published genomes averages 30 Mb
- Over 20 species to date in publicly accessible databases
- other fungal genomes held by private companies

Objectives

- Find core fungal sequences by comparing each yeast gene to 13 other fungal genomes
- Find sequences which are unique to fungi by comparing these core fungal sequences to other organisms

Method of gene comparison

- Yeast genome 12 Mb -> 6,000 predicted genes
- Use Standalone TBLASTN to find homologs
 - download TBLASTN program
 - download genomes & create personal databases

yeast protein: MYYIMFLYNMLLIILIFYSI...

expect value $\leq 10^{-5}$

Predicted protein: MREIVHLQTLIIILIFYS.....

translate (6-frame)

fungus genome: gttcaccttcagaccggccagtgtgtaagtt.....

6356 Yeast predicted genes

3340 Yeast genes with homologs in 12 of 14 fungi

Ascomycetes

Aspergillus fumigatus

Magnaporthe grisea

Aspergillus nidulans

*Neurospora crassa**

Candida albicans

Podospora anserina

Gibberella zeae

Trichoderma reesei

(Saccharomyces cerevisiae)*

Basidiomycetes

Cryptococcus neoformans

Phakopsora pachyrhizi

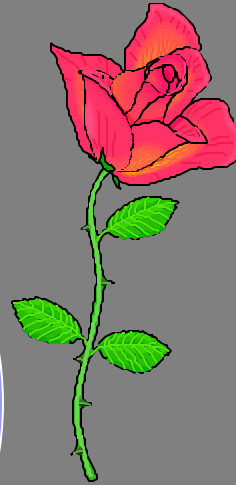
Phanerochaete chrysosporium

Coprinus cinerea

Ustilago maydis

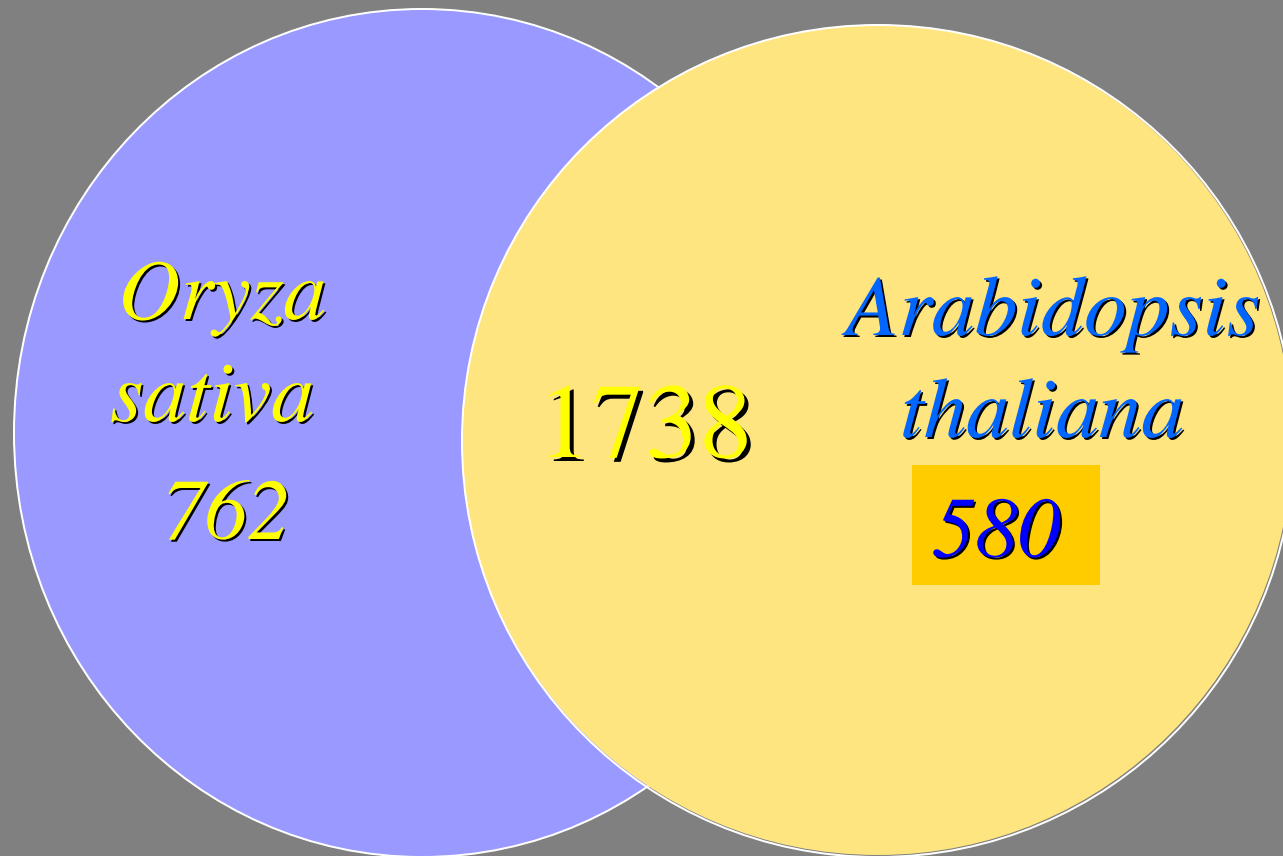
3340 genes common to 12 of 14 fungi compared to plant genomes

Oryza sativa
2500



Arabidopsis thaliana
2318

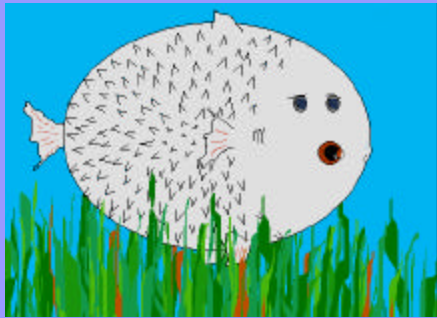
3340 genes common to 12 of 14
fungi compared to plant genomes



Total: 3080

3340 genes common to 12 of 14 fungi compared to animal genomes

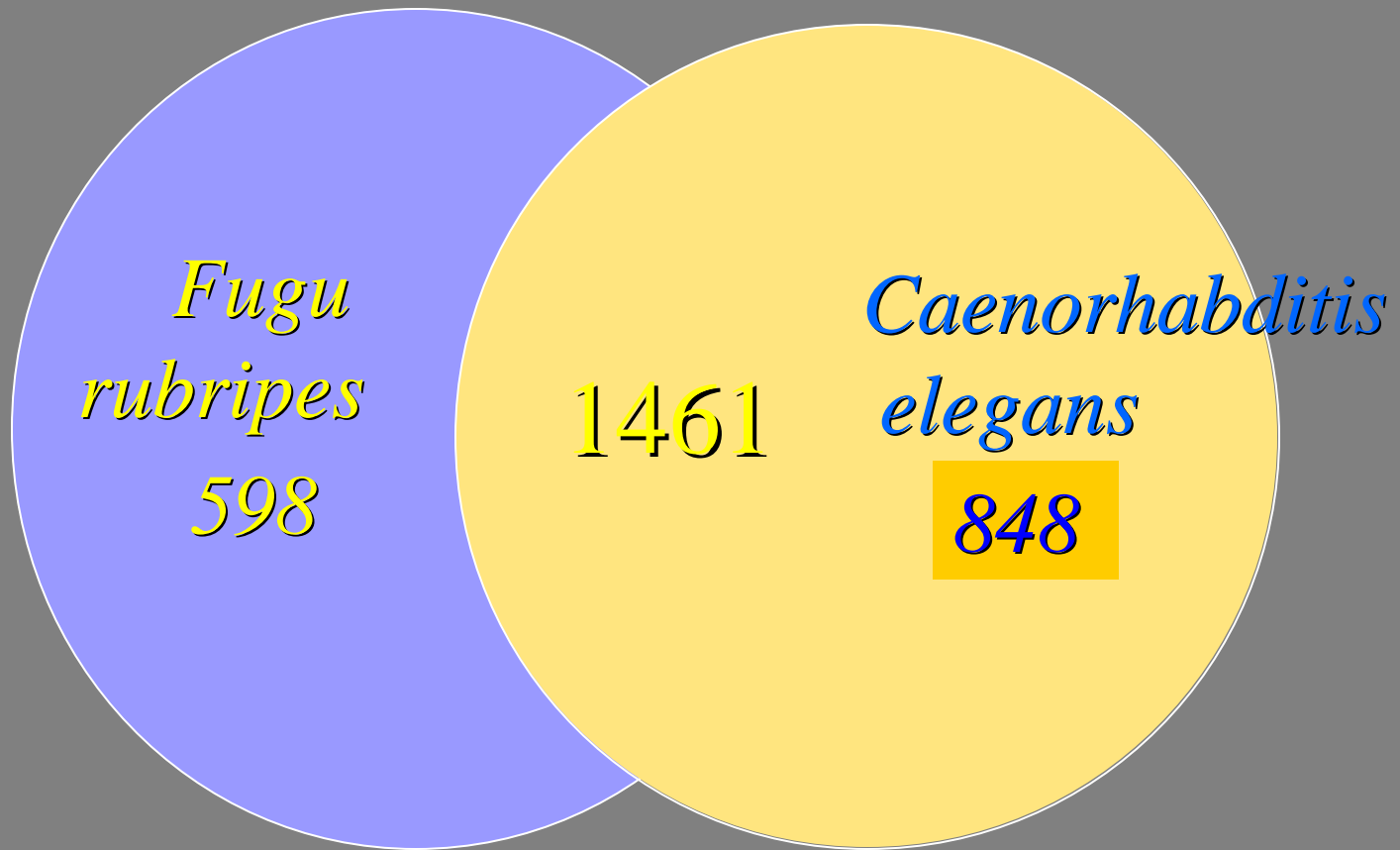
Fugu rubripes
2059



Caenorhabditis
elegans
2309



3340 genes common to 12 of 14
fungi compared to animal genomes



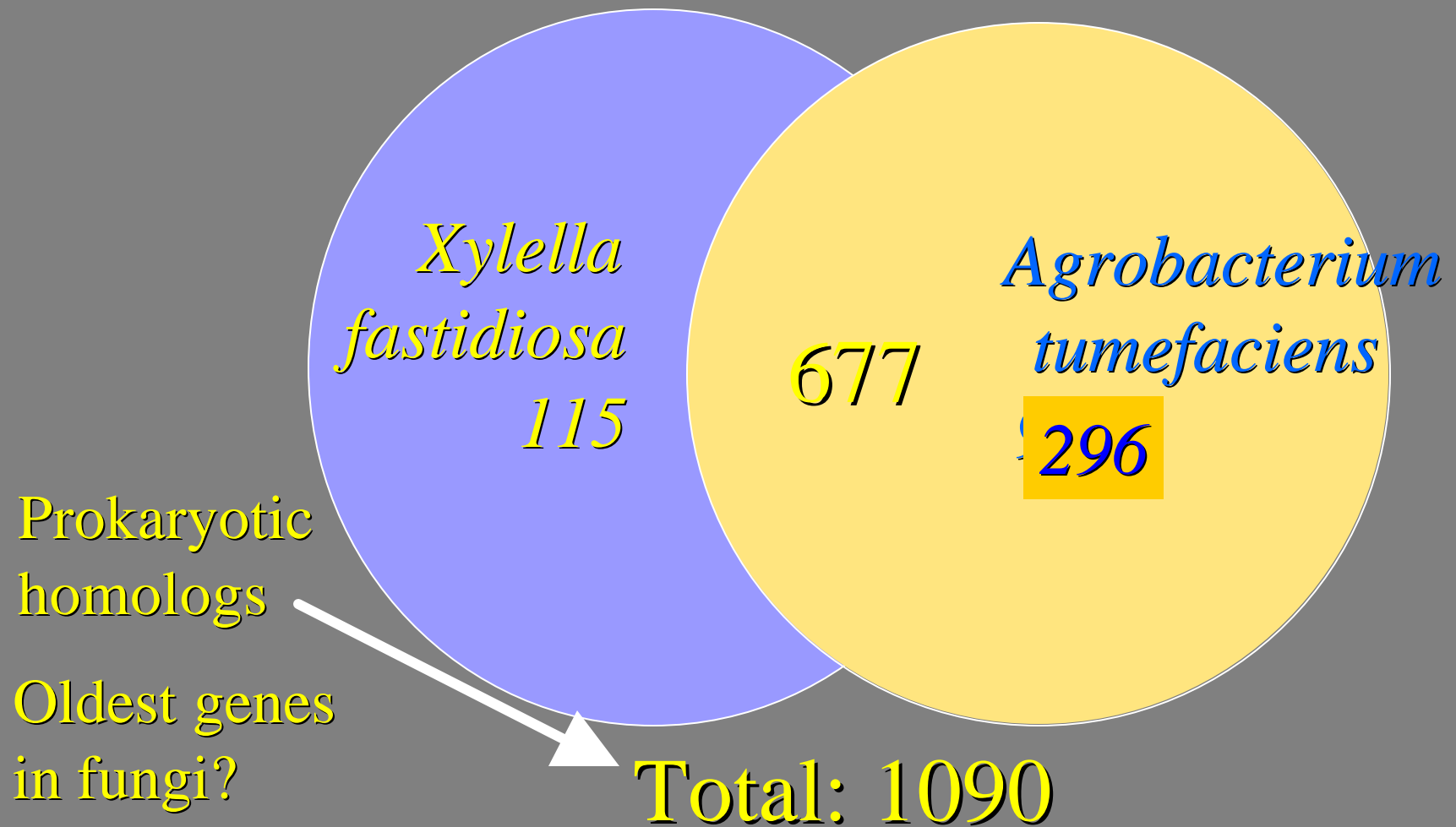
Total: 2907

3340 genes common to 12 of 14
fungi compared to bacterial genomes

Xylella fastidiosa
792

Agrobacterium
tumefaciens
973

3340 genes common to 12 of 14 fungi compared to bacterial genomes



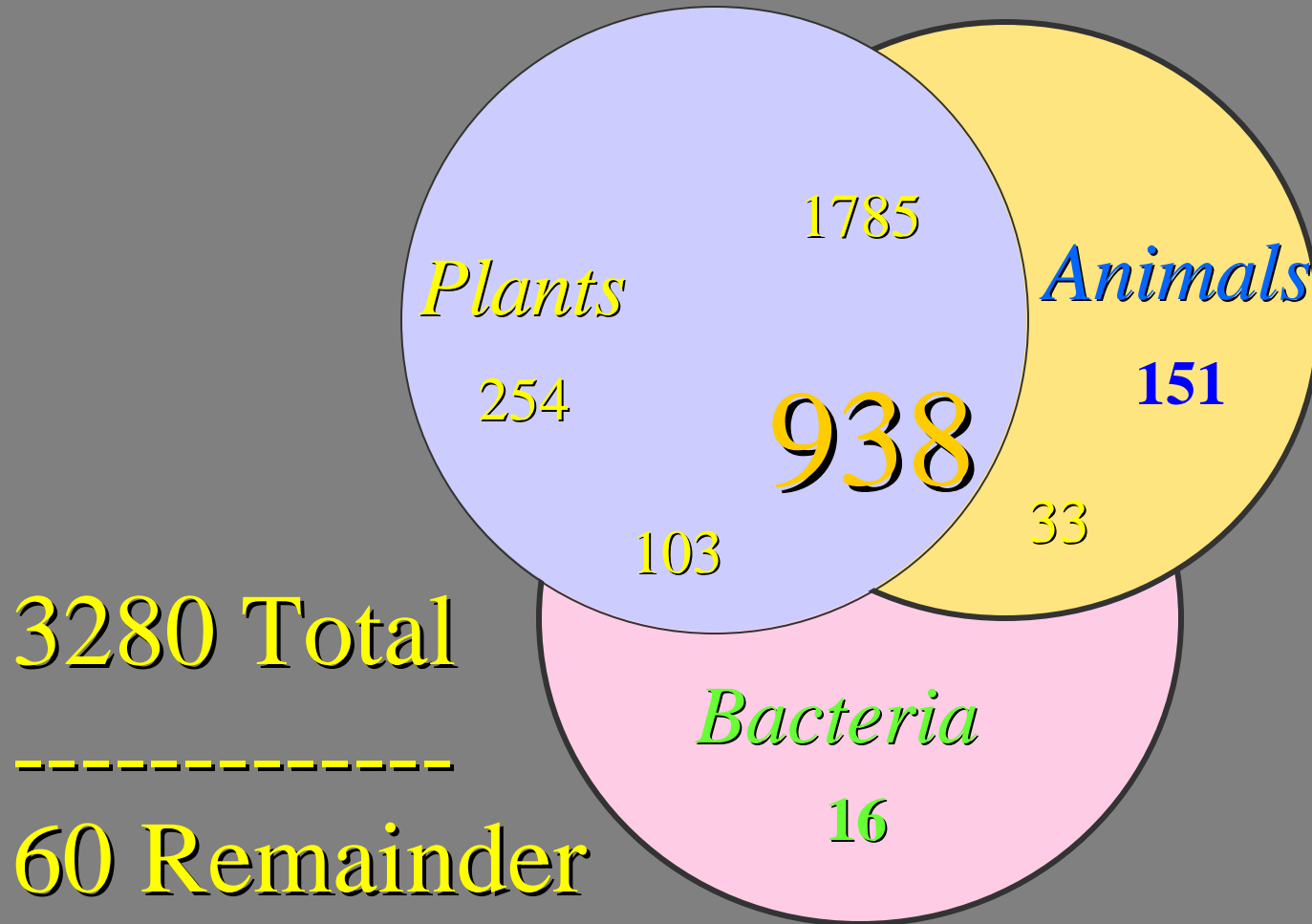
3340 genes common to 12 of 14 fungi compared to other genomes

Plants
3080

Animals
2907

Bacteria
1090

3340 genes common to 12 of 14 fungi compared to other genomes



What are these 60 fungal genes?

- are they really found only in fungi?
 - sent to BLAST against GenBank NR , EST, GSS, and HTGS databases
- 43 have a non-fungal match at $E () \leq 10^{-5}$ so these are not exclusively fungal (many of these have homologs in the human or mouse genomes)
- this leaves 17 genes found only in fungi

What are these 17 fungal genes?

- 5 have unknown function
- 2 involved in protein biosynthesis
- 2 involved in transport
- 7 have miscellaneous functions
- 1 involved in sporulation (clearly fungal)

Note: functional annotation from Yeast Genome Database

Methods

- set up Linux Operating System (free)
 - also works in Windows, but slower
- download genomic data from internet (free)
 - see my web copy for details
- set up Standalone Blast program (free)
 - from www.ncbi.nlm.nih.gov/BLAST/
- learn PERL script programming (free but takes much time for biologists)