



上海交通大学
SHANGHAI JIAO TONG UNIVERSITY



Chapter 10

Pan-genomics

Spring 2019



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Reading materials

1. Tettelin, Hervé, et al. “Genome analysis of multiple pathogenic isolates of *Streptococcus agalactiae*: implications for the microbial “pan-genome”. *PNAS*, 2005, 102(39): 13950-13955.
2. Computational Pan-Genomics Consortium, “Computational pan-genomics: status, promises and challenges”, *Brief Bioinform*, 2016, Oct 21.
3. Hu, Z., Sun, C., Zhao, Y., Wei, C.* , “EUPAN enables pan-genome studies of a large number of eukaryotic genomes”, *Bioinformatics*, 2017, 23(15):2408-2409
4. Sun, C., Hu, Z., Lu, K., Zhao, Y., Lu, J., Zheng, T., Wang, W., Shi, J., Zhang, D., Li, Z., Wei, C., “RPAN: Rice Pan-genome Browser for ~3,000 rice genomes”, *Nucleic Acids Research*, 2017, 45(2): 597-605.
5. Wang, W., et al., “Genomic variation in 3,010 diverse accessions of Asian cultivated rice”, *Nature*, 2018

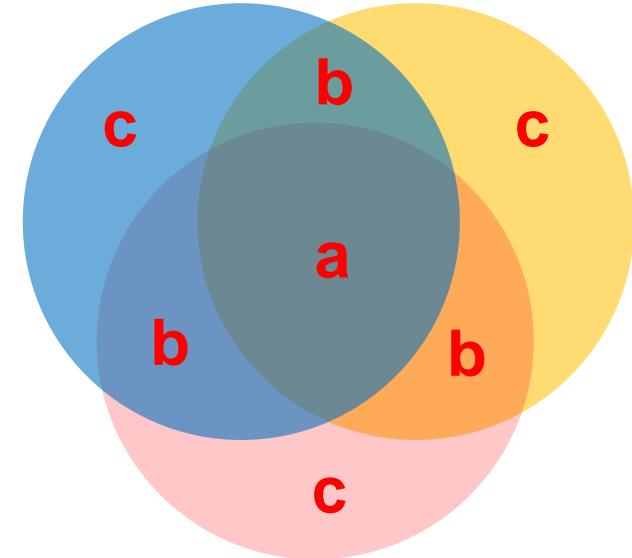


What is pan-genome?

Individual genome

Pan-genome*

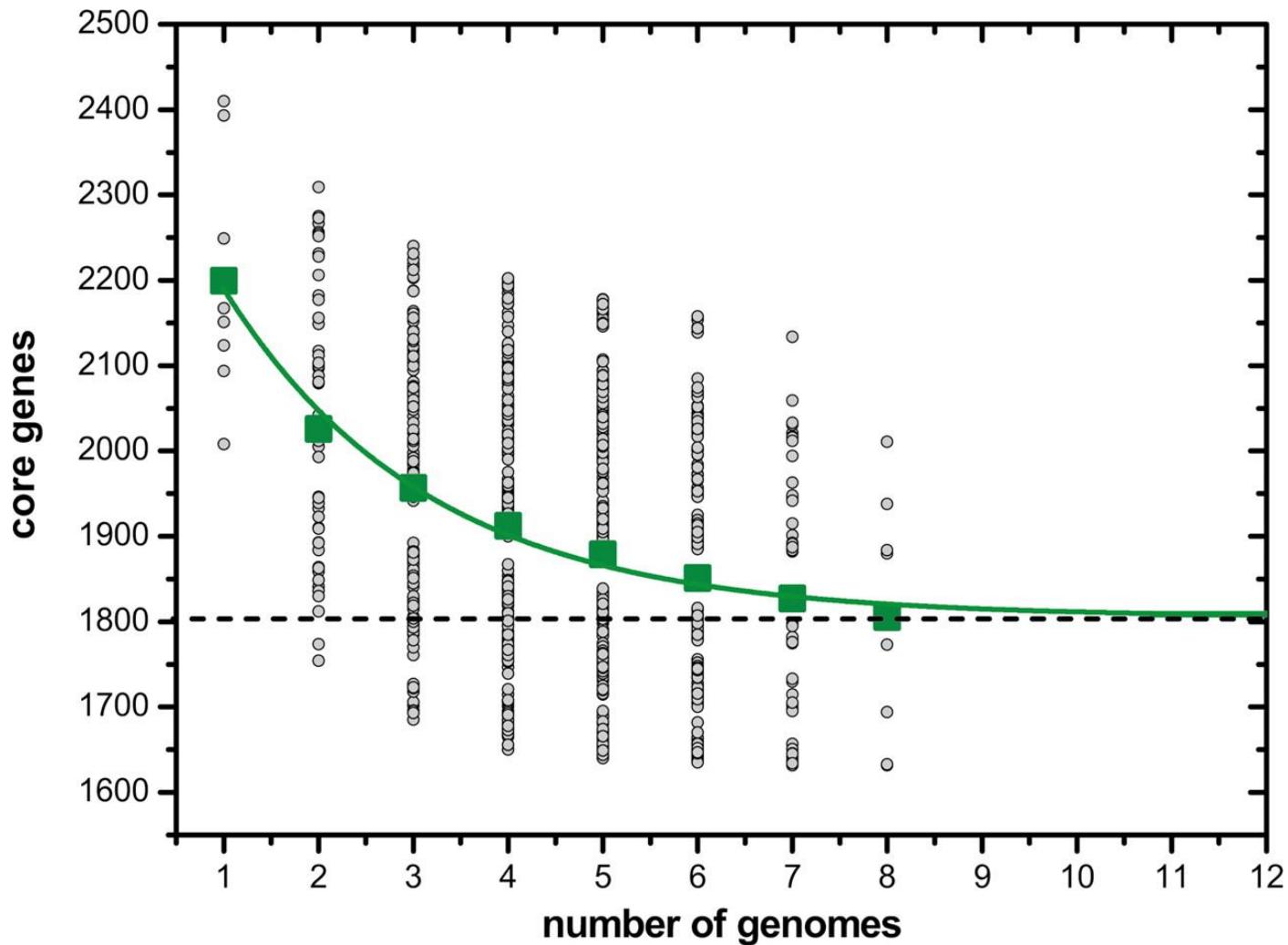
- pan, from Greek πᾶν, means all, everything
- Core genes [a]
- Dispensable genes [b]
- Unique genes [c]



* Tettelin, Hervé, et al. "Genome analysis of multiple pathogenic isolates of *Streptococcus agalactiae*: implications for the microbial “pan-genome”." *PNAS* 102.39 (2005): 13950-13955.



GBS core genome.

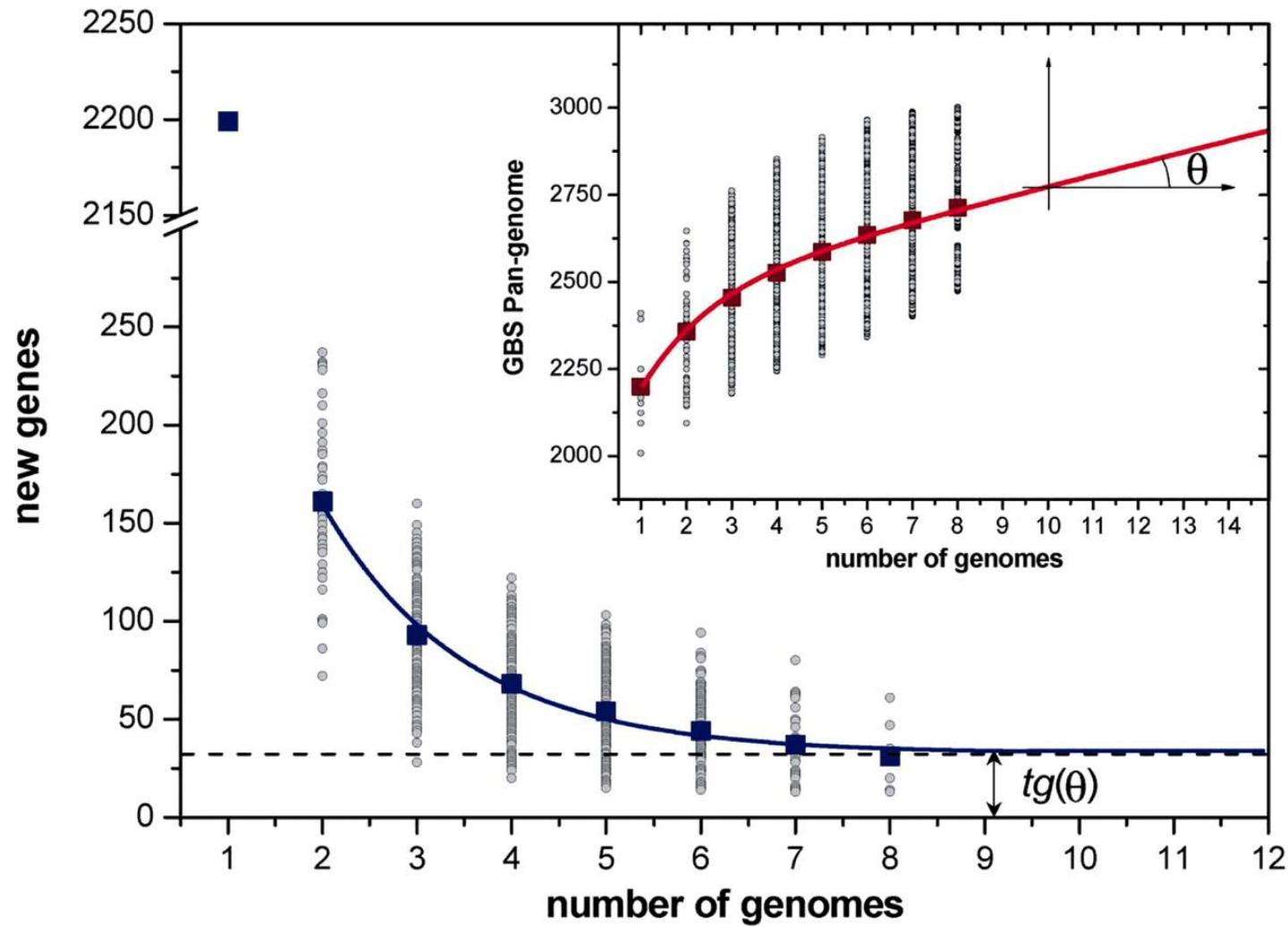


Hervé Tettelin et al. PNAS 2005;102:13950-13955

PNAS



GBS pan-genome (An open pan-genome!)

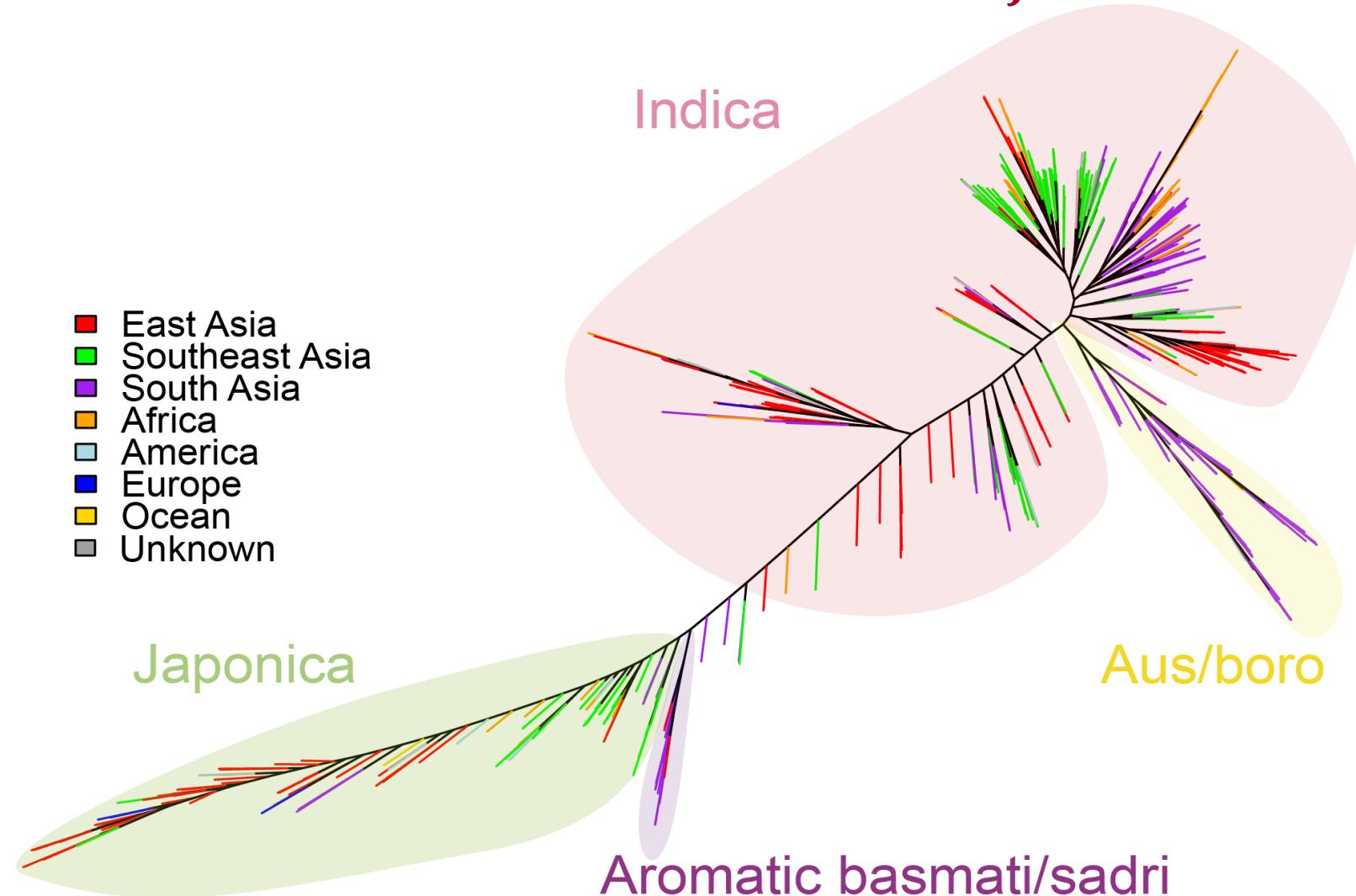


Hervé Tettelin et al. PNAS 2005;102:13950-13955

PNAS



The 3000 Rice Genome Project



Sun, C. et al., "RPAN: Rice Pan-genome Browser for ~3,000 rice genomes",
Nucleic Acids Research, 2017, 45(2): 597-605.



EUPAN: Eukaryote Pan-genome ANalysis

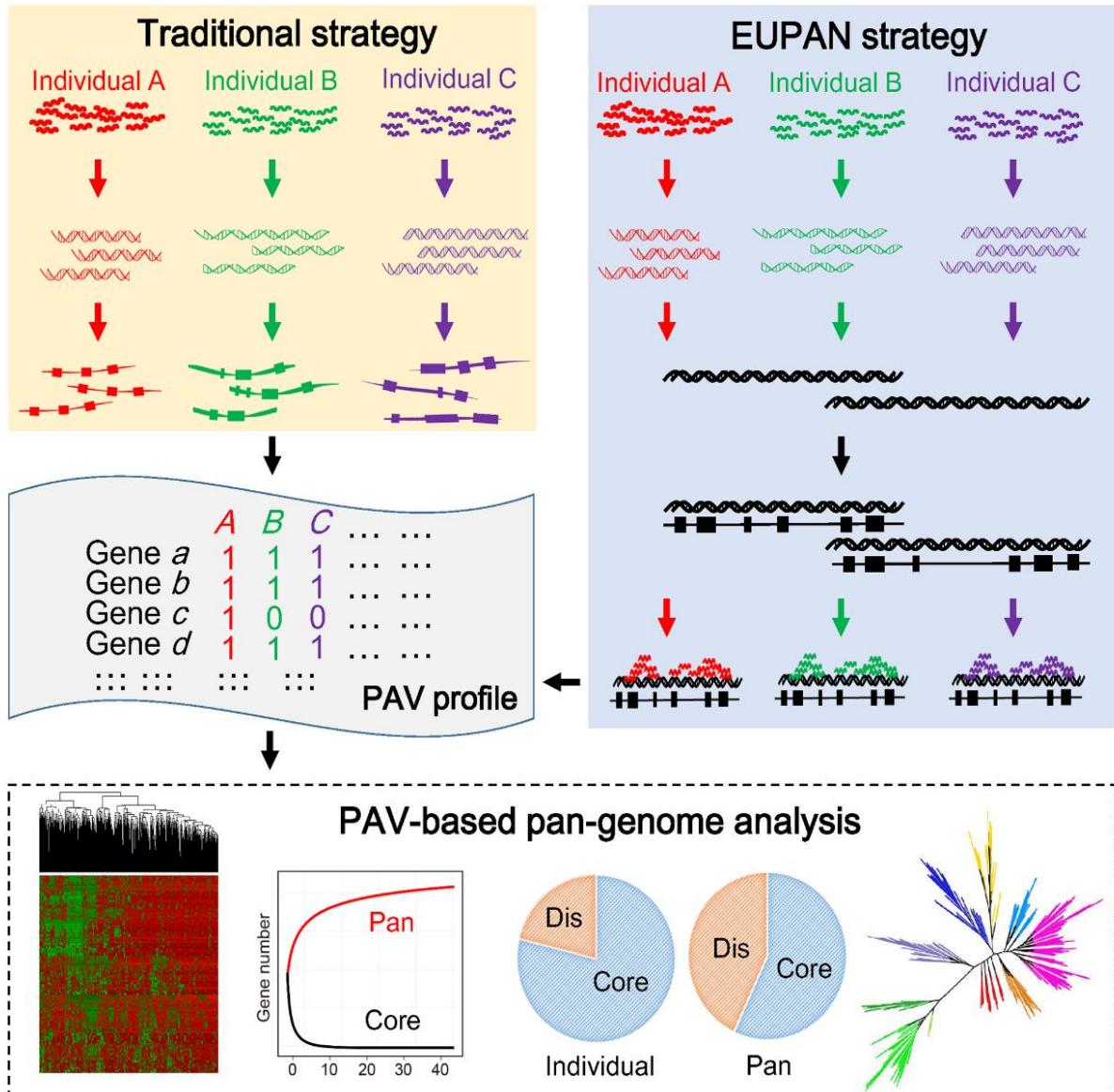
Sequencing data



PAV profile

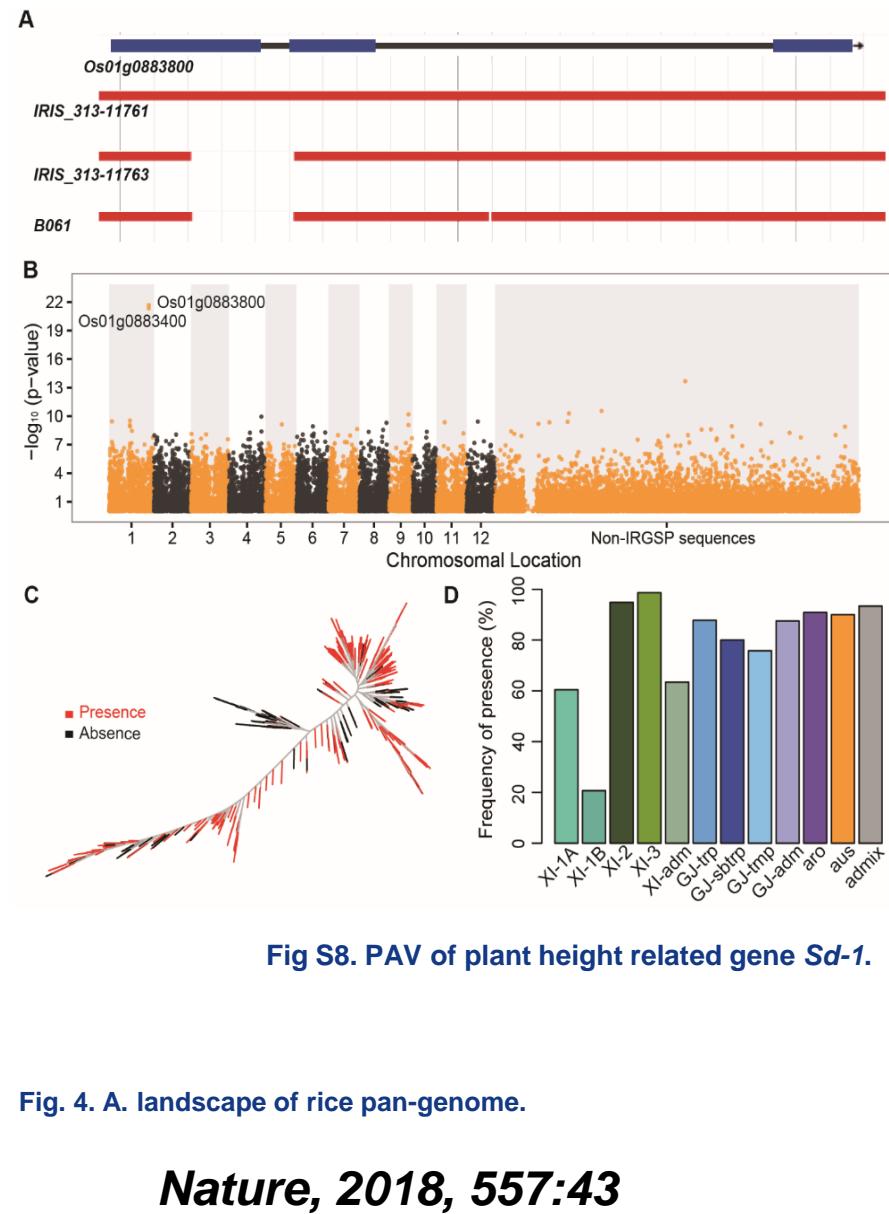
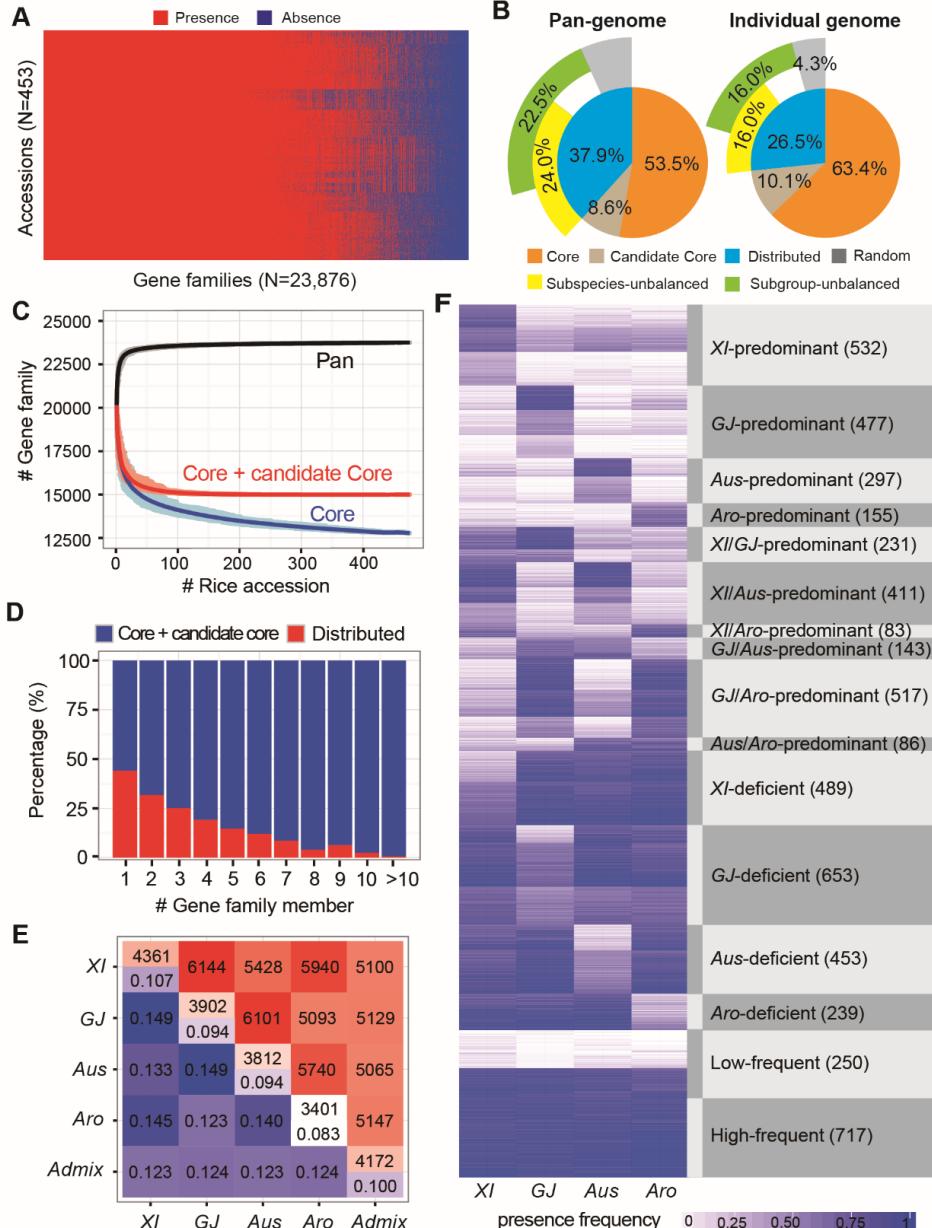


Pan-genome analysis





The rice pan-genome (3k rice genomes)





RPAN:

Rice PAN-genome browser for >3000 genomes

[Home](#)[Search](#)[Visualization](#)[Rice table](#)[Gene table](#)[Expression](#)[Download](#)[About & Manual](#)

3K Rice Pan-genome Browser

Lab of Computational Genomics and Metagenomics

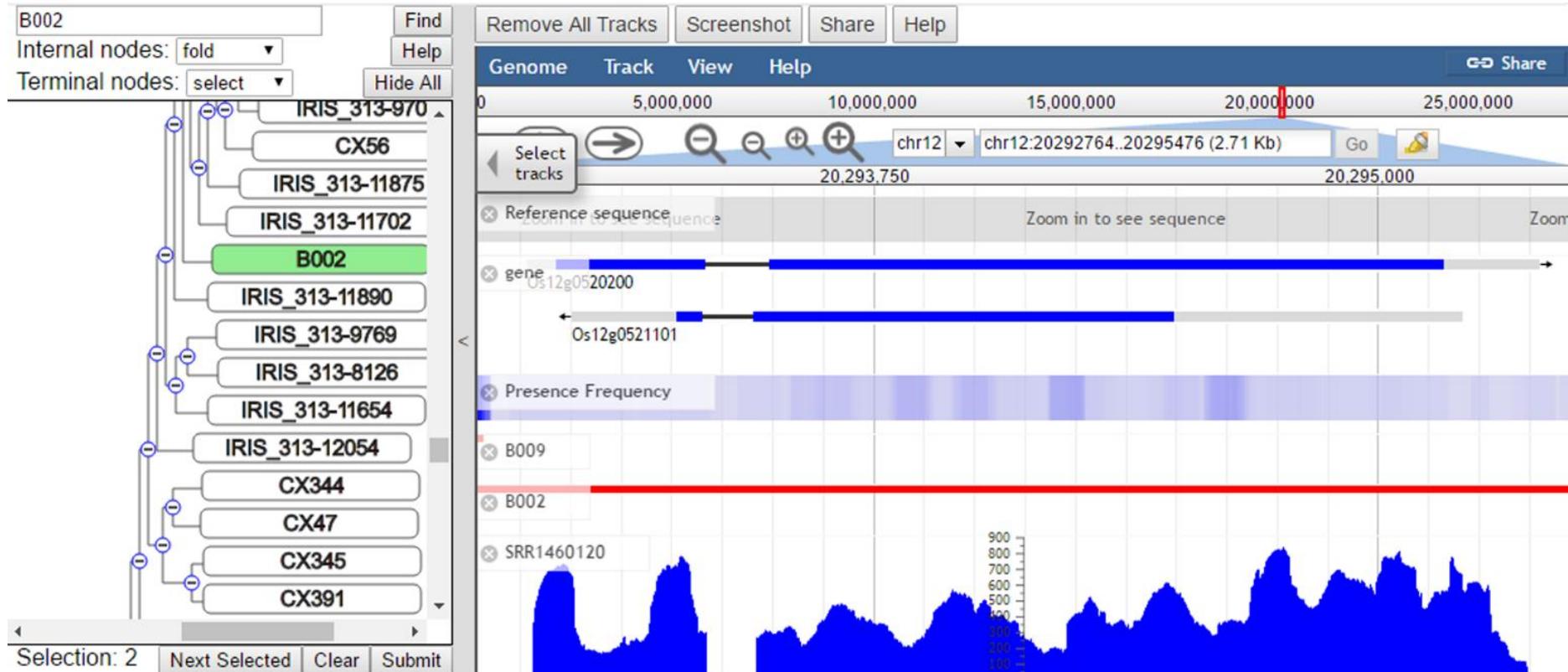
Shanghai Jiao Tong University

● <http://cgm.sjtu.edu.cn/3kricedb/>

Sun, C. et al., “RPAN: Rice Pan-genome Browser for ~3,000 rice genomes”,
Nucleic Acids Research, 2017, 45(2): 597-605



RPAN: Rice PAN-genome Browser

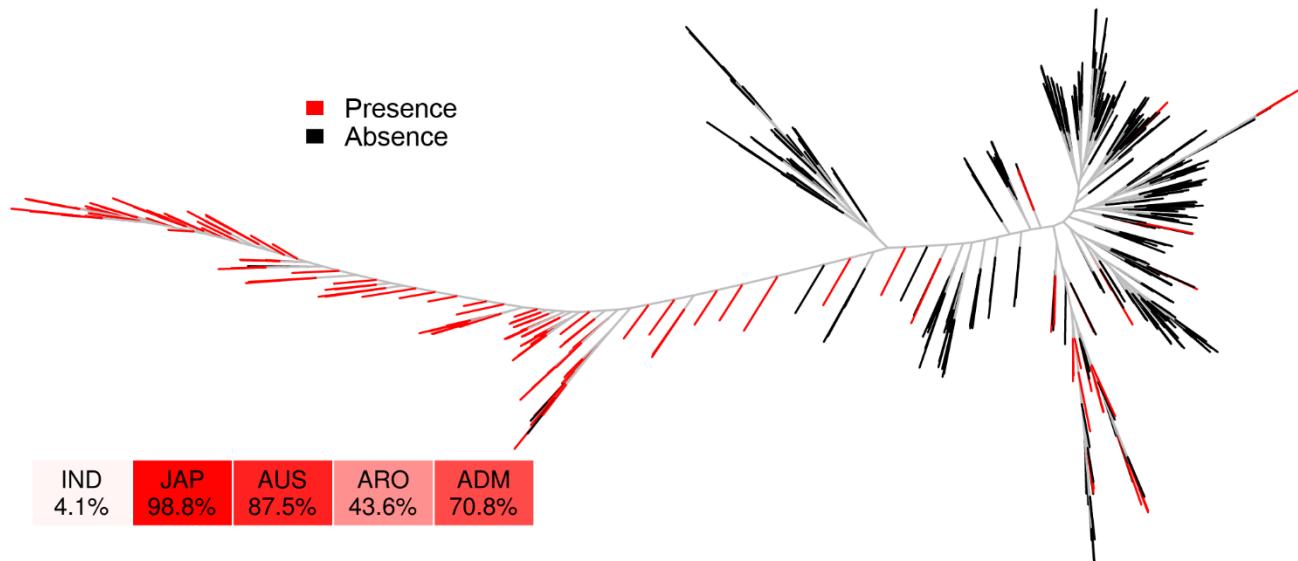


Sun, C. et al., "RPAN: Rice Pan-genome Browser for ~3,000 rice genomes",
Nucleic Acids Research, 2017, 45(2): 597-605



RPAN

A

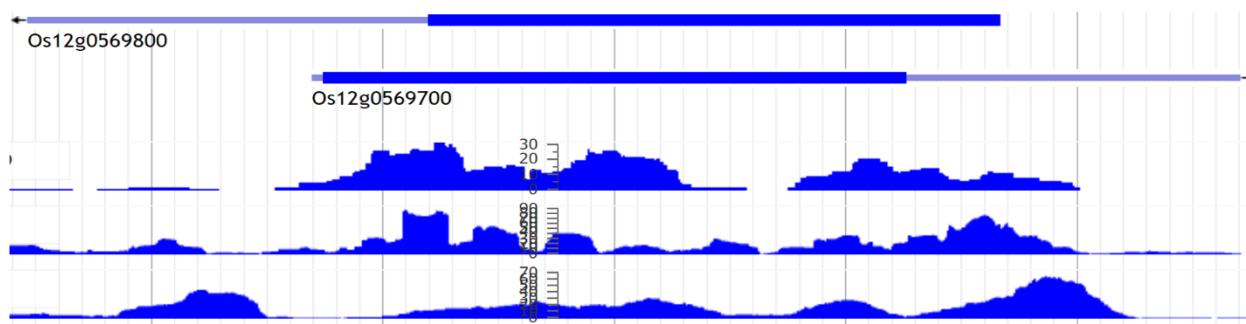


B

IND	4.1%	JAP	98.8%	AUS	87.5%	ARO	43.6%	ADM	70.8%
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IG1	9.8%	IG2	0%	IG3	0%	IG4	7.9%	IG5	1.8%	AUSG6	43.6%	JG7	96.6%	JG8	100%	JG9	100%	JG10	100%	AROG11	87.5%	ADM	70.8%
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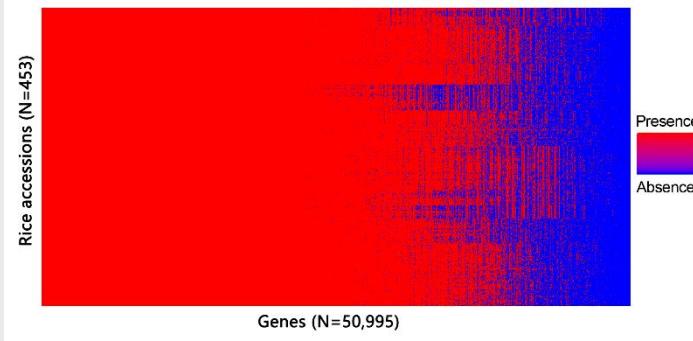
C



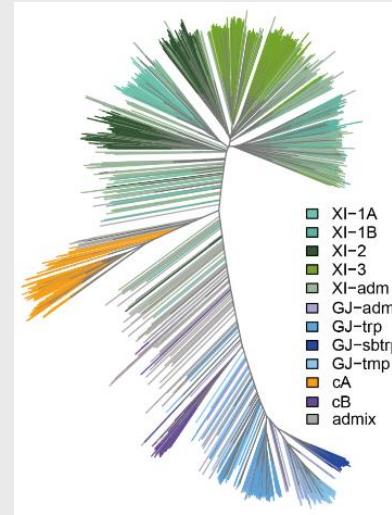
Sun, C. et al., "RPAN: Rice Pan-genome Browser for ~3,000 rice genomes", *Nucleic Acids Research*, 2017, 45(2): 597-605.



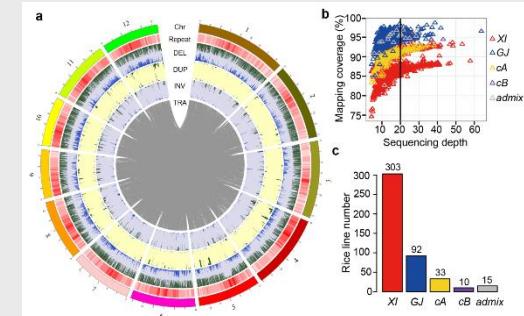
Pan-genomics based genomics analysis



Pan-genome
analysis
PAV



SNP



SV

Biomarker identification

Phenotype



Computational pan-genomics: status, promises and challenges

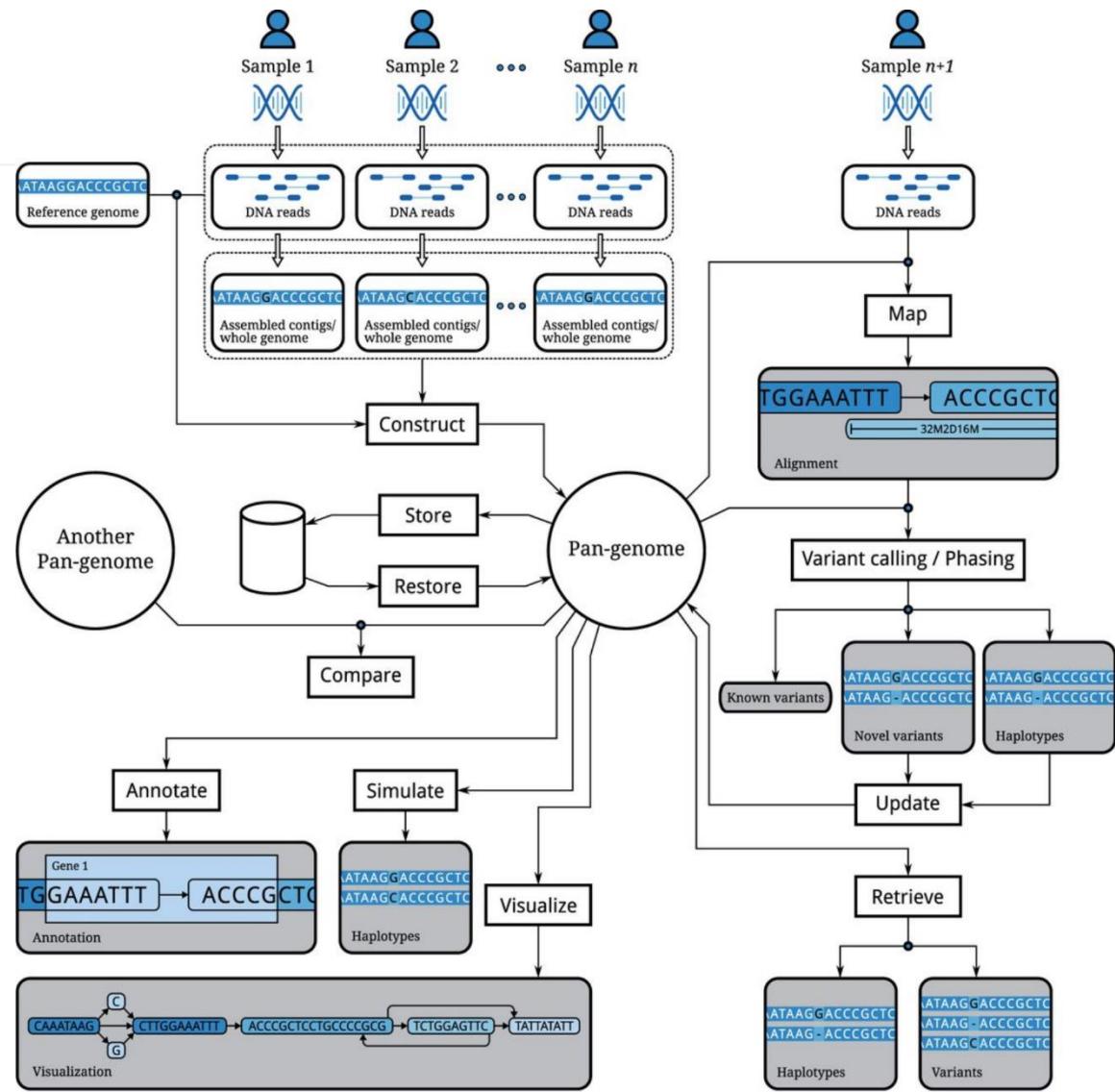


Illustration of operations to be supported by a pan-genome data structure.



Summary

It's pan-genome era now !

Crop pan-genomes

wheat pan-genomes

Human pan-genomes

Cancer genomes

Human body single cell genomes