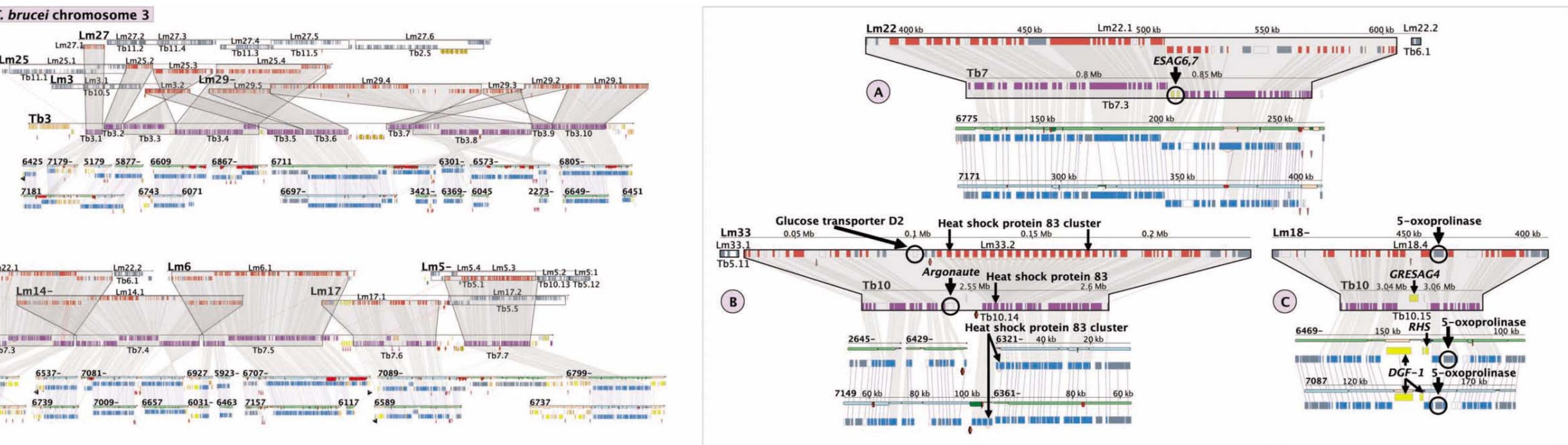
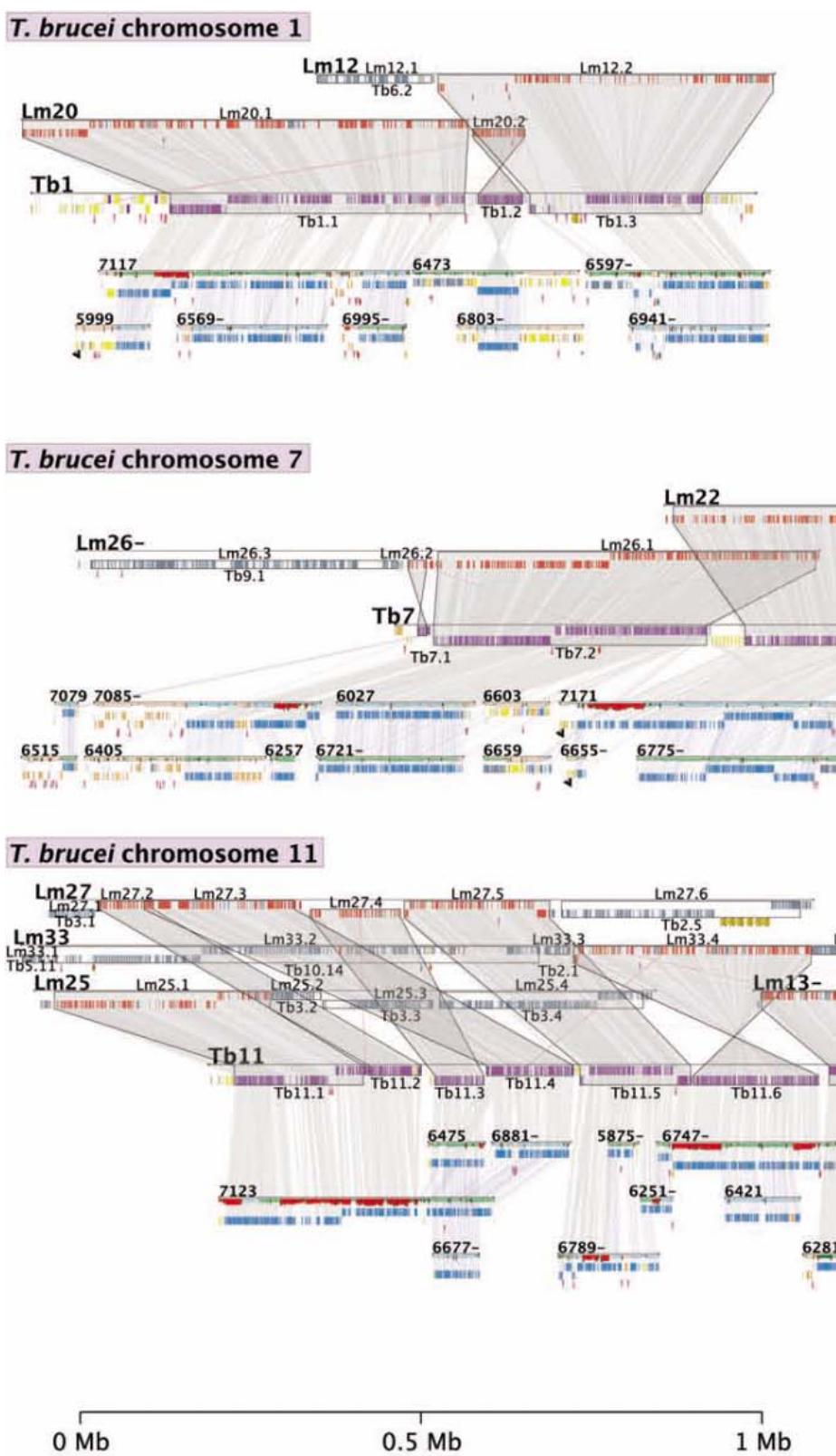


## PLATE 1



Protein-coding genes	
L. major gene with a COG match (match shown)	Gene with no COG matches
T. brucei gene with a COG match (match shown)	Gene with a COG match (match not shown)
T. cruzi gene with a COG match (match shown)	

Highlighted protein families	
Putative surface proteins: mucin, MASP, trans-sialidase, GP63, VSG RHS, ESAG/ESAG-like protein, DGF-1	

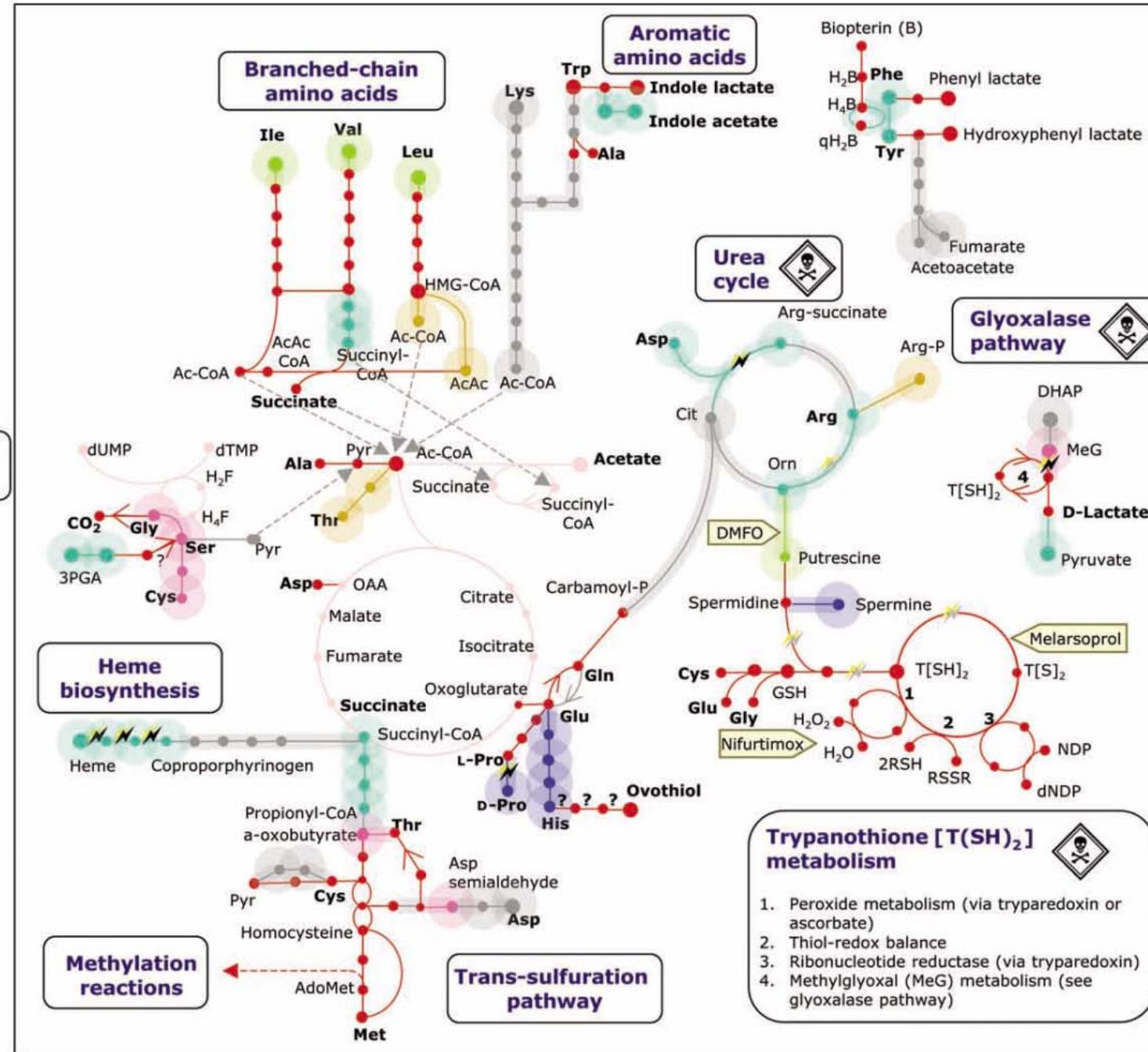
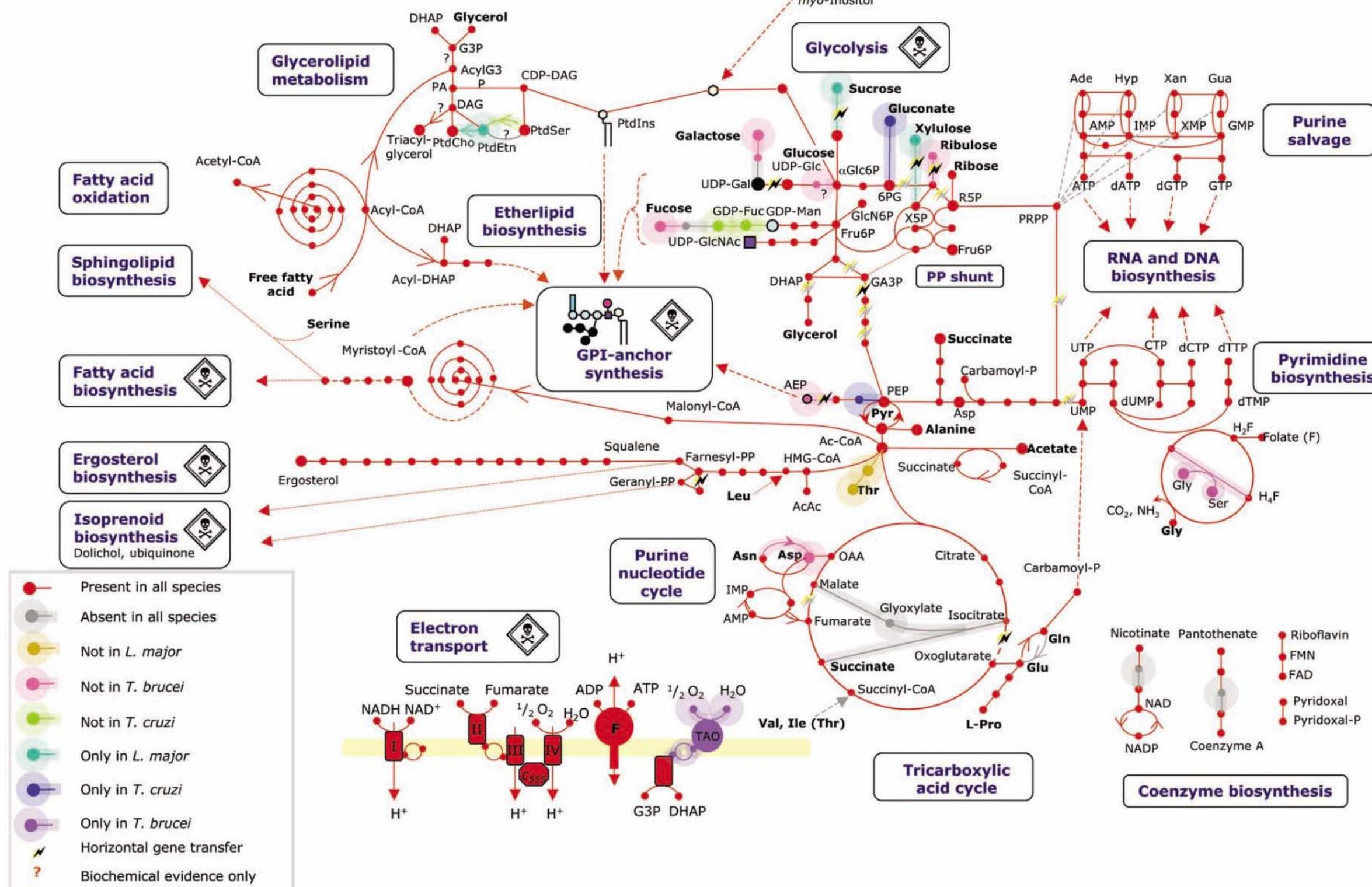
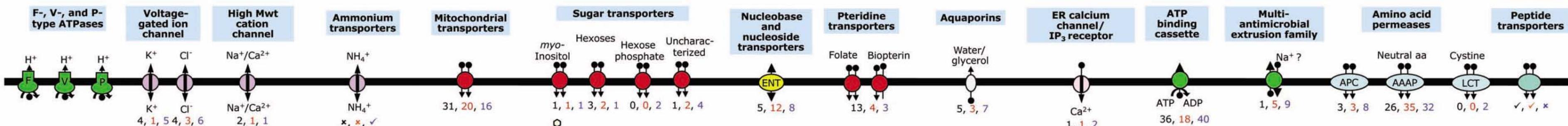
Clusters of orthologous genes (COGs)	
Genes in the same COG (syntenic match)	Genes in the same COG (nonsyntenic match)
Syntenic block	

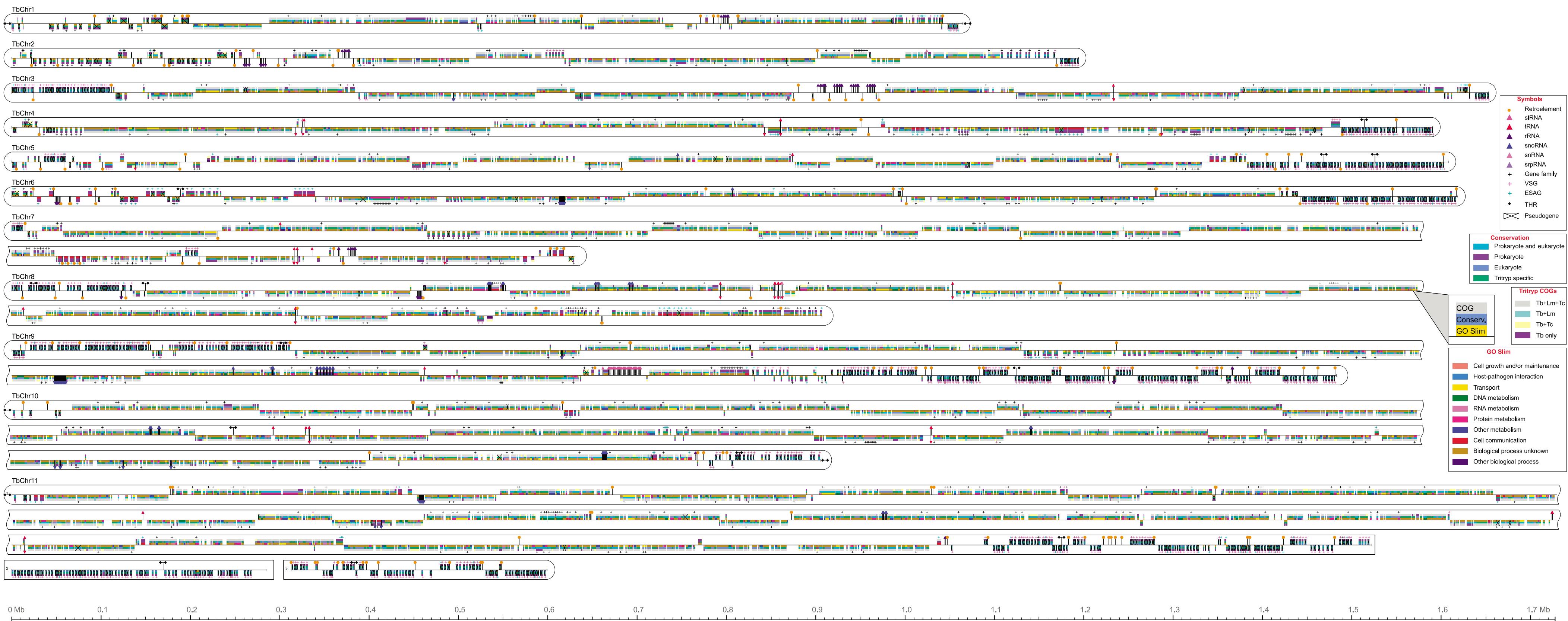
RNA genes	
tRNA	sRNA
snoRNA	rRNA
snRNA	

Repeats	
Subtelomeric repeat	
Telomeric repeat	
Retrotransposons: ingi, DIRE, L1Tc, VIPER	Tandemly repeated genes
Repetitive region	

T. cruzi-specific features	
Heterozygous Esmeraldo-like haplotype	Heterozygous non-Esmeraldo-like haplotype
Haplod Esmeraldo-like haplotype	Haplod non-Esmeraldo-like haplotype
Region of combined assembly	T. cruzi haplotypes

PLATE 2



*Trypanosoma brucei* genome organization

*Leishmania major* genome organization