

Output results of CLIME (CLustering by Inferred Models of Evolution)

Dataset:

Num of genes in input gene set: 7
Total number of genes: 20834
Prediction LLR threshold: 0

The CLIME PDF output two sections:

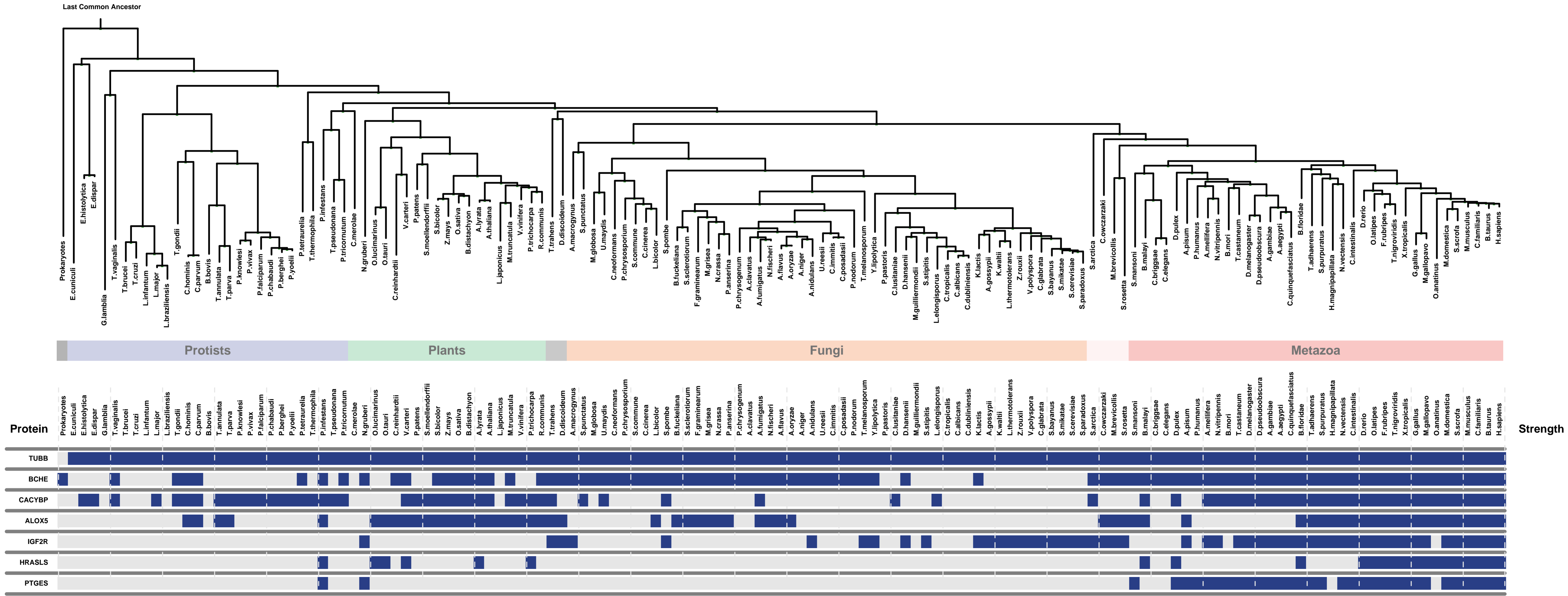
1) Overview of Evolutionarily Conserved Modules (ECMs)

- Top panel shows the predefined species tree.
- Bottom panel shows the partition of input genes into Evolutionary Conserved Modules (ECMs), ordered by ECM strength (shown at right), and separated by horizontal lines.
- Each row show one gene, where the phylogenetic profile indicates presence (blue) or absence (gray) of homologs in each species (column).
- Gene symbols are shown at left. Gray color indicates that the gene is a paralog to a higher scoring gene within the same ECM (based on BLASTP $E < 1e-3$).

2) Details of each ECM and its expansion ECM+

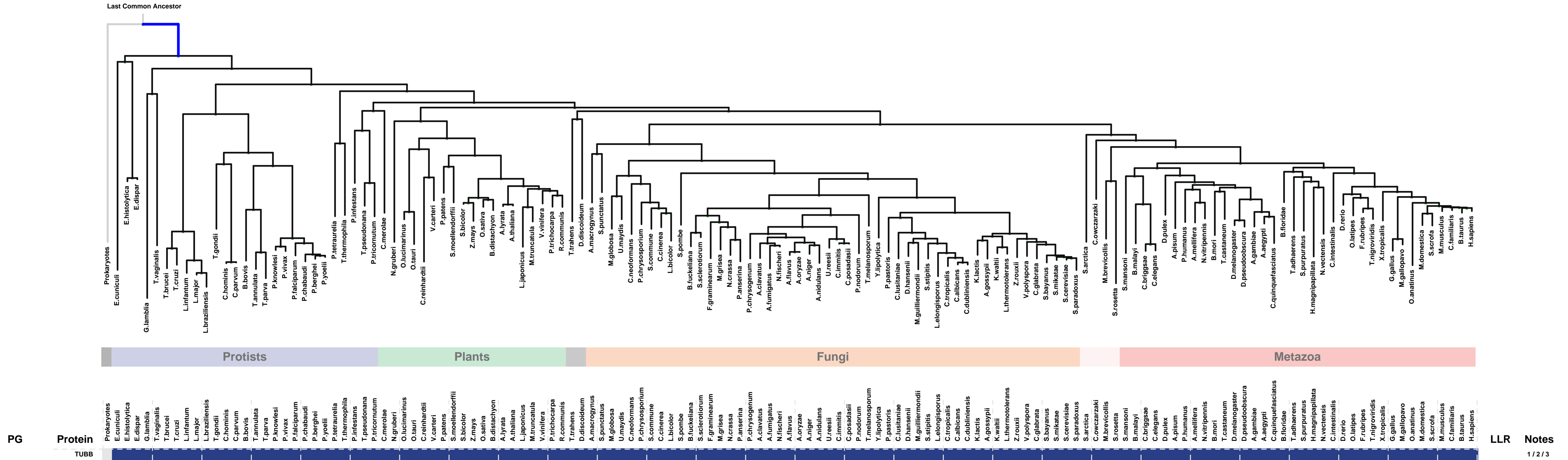
- Top panel shows the inferred evolutionary history on the predefined species tree. Branch color shows the gain event (blue) and loss events (red color, with brighter color indicating higher confidence in loss). Branches before the gain or after a loss are shown in gray.
- Bottom panel shows the input genes that are within the ECM (blue/white rows) as well as all genes in the expanded ECM+ (green/gray rows). The ECM+ includes genes likely to have arisen under the inferred model of evolution relative to a background model, and scored using a log likelihood ratio (LLR).
- PG indicates "paralog group" and are labeled alphabetically (i.e., A, B). The first gene within each paralog group is shown in black color. All other genes sharing sequence similarity (BLAST $E < 1e-3$) are assigned to the same PG label and displayed in gray.

Overview of Evolutionarily Conserved Modules (ECMs)



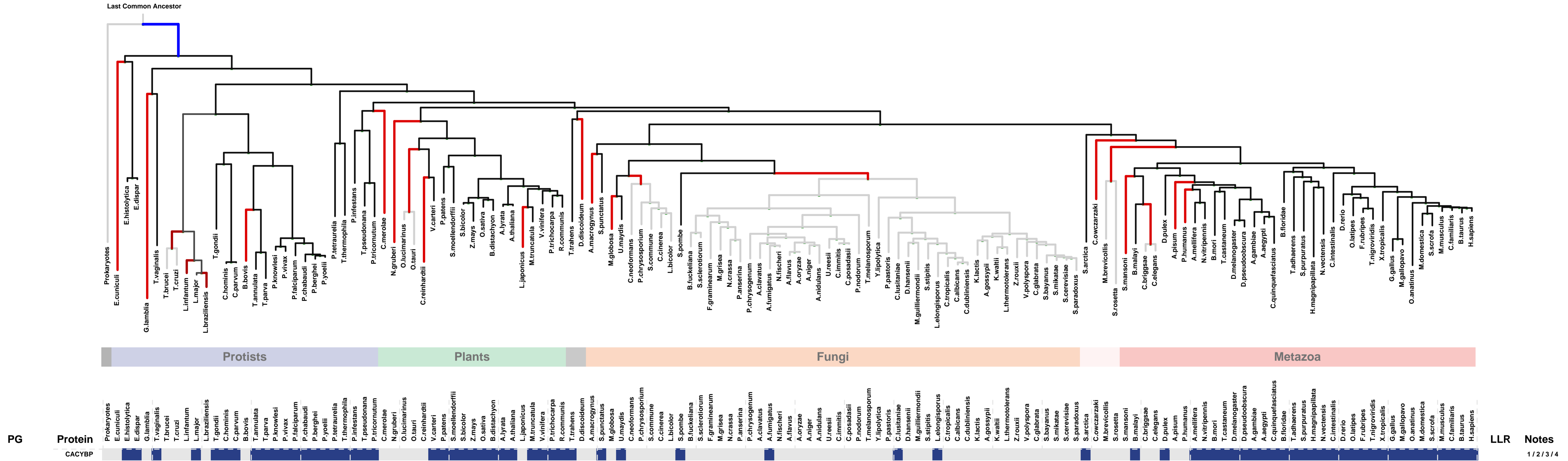
ECM 1, Gene set "nuclear envelope lumen", Page 1

Num of ECM Genes: 1. Num of Predicted Genes: 0



ECM 3, Gene set "nuclear envelope lumen", Page 1

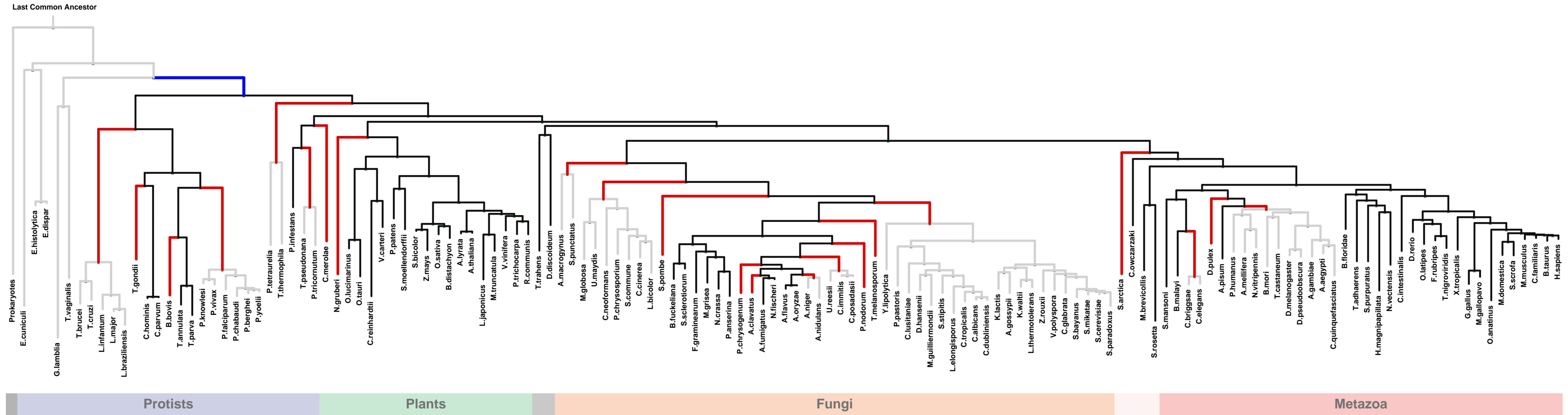
Num of ECM Genes: 1. Num of Predicted Genes: 0



1: beta-catenin destruction complex || 2: cell body || 3: neuron projection || 4: nuclear envelope lumen

ECM 4, Gene set "nuclear envelope lumen", Page 1

Num of ECM Genes: 1. Num of Predicted Genes: 14

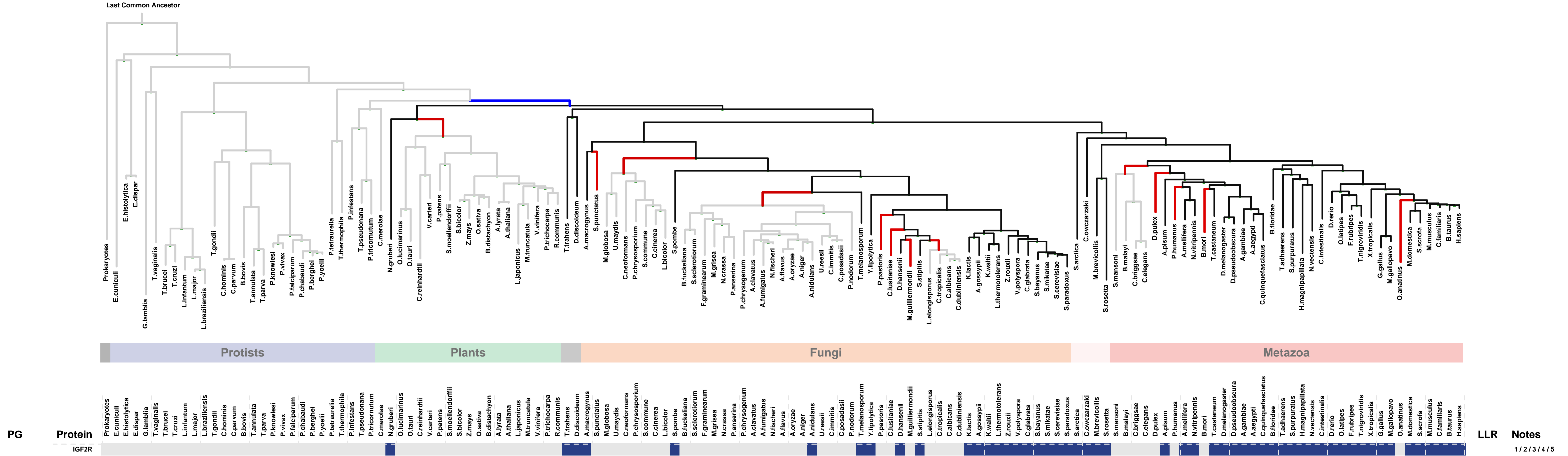


PG	Protein	Prokaryotes	Protists	Plants	Fungi	Metazoa	LLR	Notes
A	ALOX5							1/2/3/4/5/6
A	ALOX12B							
A	ALOXE3							
A	ALOX12							6
A	ALOX15							
A	ALOX15B							
B	SLC17A9							17.4
B	SLC8A1							12.2
B	SLC17A1							7.9
	MTA1							4.7
	FKRP							1.1
	NAGK							1.0
	PROZ							1.0
	NECAB2							0.7
	NID1							0.2

1: dendrite || 2: nuclear envelope || 3: nuclear envelope lumen || 4: nuclear matrix || 5: nuclear membrane || 6: sarcolemma || 7: dendritic spine || 8: intercalated disc || 9: T-tubule || 10: Z disc || 11: NuRD complex || 12: dystrophin-associated glycoprotein complex || 13: rough endoplasmic reticulum || 14: endoplasmic reticulum lumen || 15: Golgi lumen || 16: basal lamina || 17: cell periphery || 18: extracellular matrix

ECM 5, Gene set "nuclear envelope lumen", Page 1

Num of ECM Genes: 1. Num of Predicted Genes: 0



1: endocytic vesicle || 2: endosome || 3: lysosomal membrane || 4: nuclear envelope lumen || 5: trans-Golgi network transport vesicle

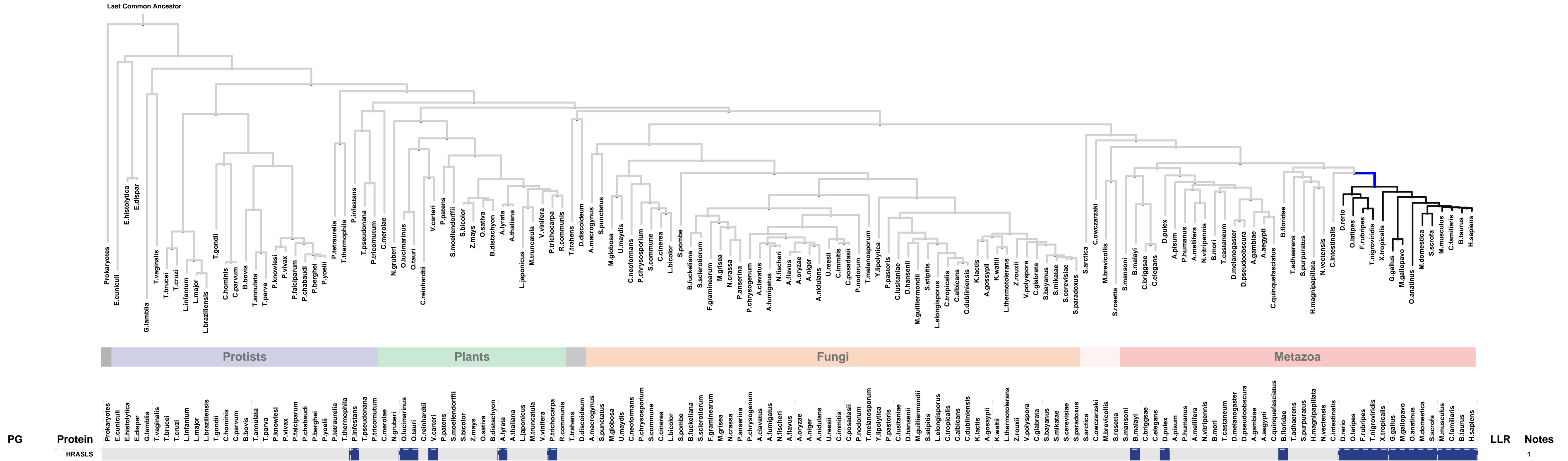
ECM 6, Gene set "nuclear envelope lumen", Page 1

Num of ECM Genes: 1. Num of Predicted Genes: 0

PRESENCE ——— ABSENCE ———
GAIN ——— LOSS ———

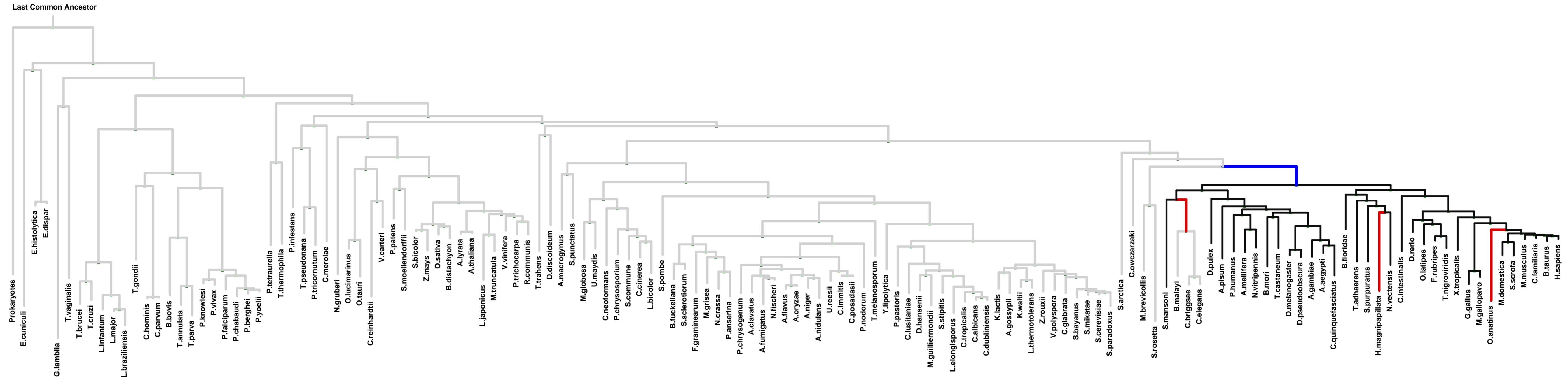
Log-likelihood Ratio Scale

0 10 20 30 40 50 60



ECM 7, Gene set "nuclear envelope lumen", Page 1

Num of ECM Genes: 1. Num of Predicted Genes: 23



PG	Protein	Prokaryotes	Protists	Plants	Fungi	Metazoa	LLR	Notes
A	PTGES							1
A	MGST1						15.5	2/3/4
B	DAXX						8.4	5/6/7/8/9/10
B	OR4C3						6.2	
C	ZMYM2						6.2	10
D	NOG						4.9	
D	BOD1L1						4.3	
D	BOD1						4.3	11/12
B	OR2A14						3.7	
B	OR4B1						3.7	
B	OR52R1						3.3	
E	SCAF1						3.2	
E	OAZ2						3.2	
B	OAZ1						2.8	
B	OR4F5						2.6	
B	LOC650293						1.3	
B	OR6K2						0.9	
F	CCDC28A						0.7	
F	CCDC28B						0.7	
C	ZMYM4						0.6	
C	HEY1						0.0	
C	ZMYM3						0.0	
C	TCF24						0.0	
C	WFS1						0.0	13/14

1: nuclear envelope lumen || 2: apical part of cell || 3: mitochondrial outer membrane || 4: peroxisomal membrane || 5: cell body || 6: cell cortex || 7: chromosome, centromeric region || 8: heterochromatin || 9: neuron projection || 10: PML body || 11: condensed chromosome kinetochore || 12: microtubule organizing center || 13: dendrite || 14: integral to endoplasmic reticulum membrane