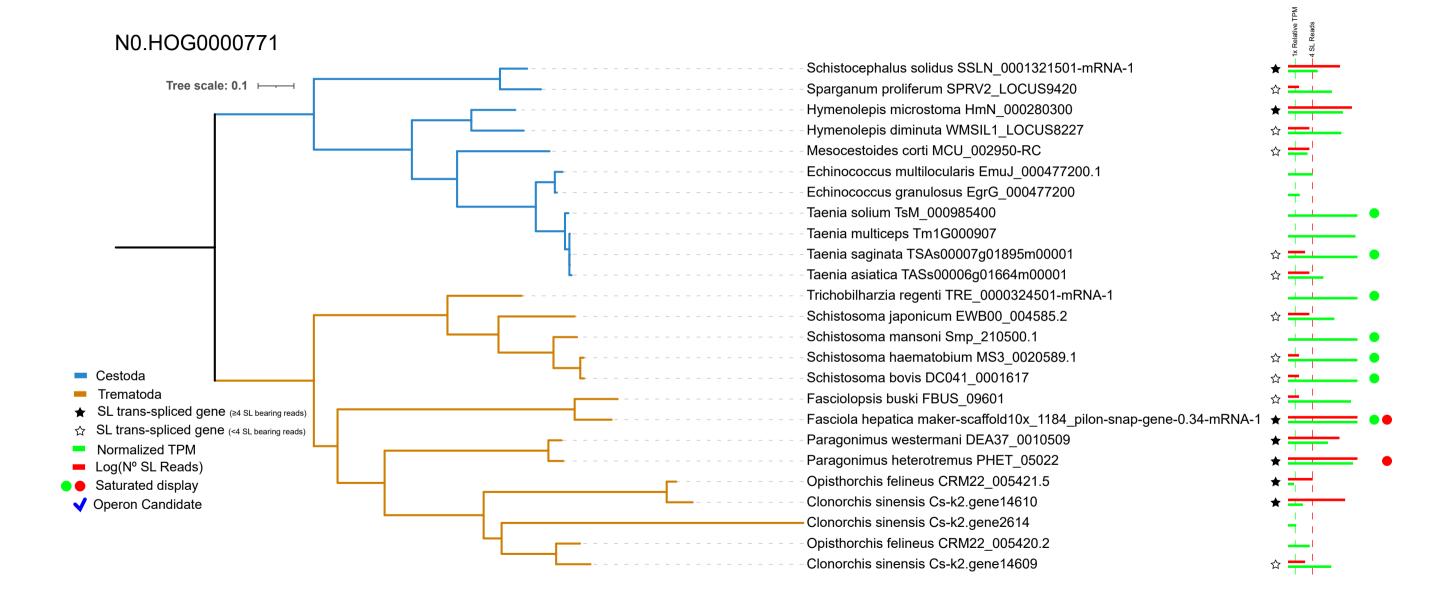
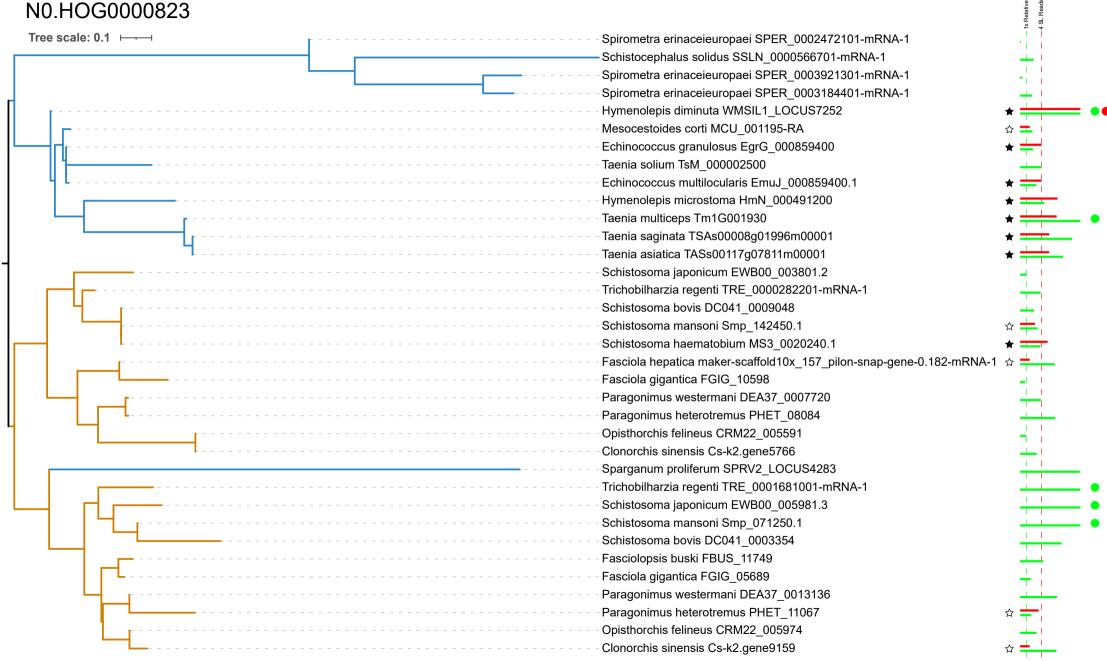
Supplementary File 5: Maximum Likelihood phylogenetic trees of the selected Phylogenetically Hierarchical Orthogroups (HOGs) based on their expression level and evidence of SL trans-splicing. For each gene it is indicated if it had evidence of SL trans-splicing above or below the 4 read threshold on the same acceptor site (Black Star or White star respectively). Followed by a Normalized TPM per species relative to the median TPM of SL trans-spliced genes and the total amount of SL bearing reads in the gene in logarithmic scale, displayed with green and red bars respectively. Both are capped (10 for the normalized TPM and 100 SL bearing reads), indicated with circles. Genes identified as members of operon candidates are indicated with a blue check mark.

Index:

Pag. 2: N0.HOG0000771	Pag. 23: N0.HOG0009450
Pag. 3: N0.HOG0000823	Pag. 24: N0.HOG0009887
Pag. 4: N0.HOG0000874	Pag. 25: N0.HOG0009968
Pag. 5: N0.HOG0000876	Pag. 26: N0.HOG0010306
Pag. 6: N0.HOG0000970	Pag. 27: N0.HOG0010318
Pag. 7: N0.HOG0001389	Pag. 28: N0.HOG0010377
Pag. 8: N0.H0G0002036	Pag. 29: N0.HOG0010496
Pag. 9: N0.HOG0002250	Pag. 30: N0.HOG0010643
Pag. 10: N0.H0G0002707	Pag. 31: N0.HOG0010669
Pag. 11: N0.H0G0003375	Pag. 32: N0.HOG0010755
Pag. 12: N0.H0G0004575	Pag. 33: N0.HOG0010762
Pag. 13: N0.H0G0005209	Pag. 34: N0.HOG0010786
Pag. 14: N0.HOG0006807	Pag. 35: N0.HOG0010915
Pag. 15: N0.HOG0007522	Pag. 36: N0.HOG0010953
Pag. 16: N0.H0G0007613	Pag. 37: N0.HOG0011542
Pag. 17: N0.H0G0007875	Pag. 38: N0.HOG0011745
Pag. 18: N0.H0G0007943	Pag. 39: N0.HOG0011763
Pag. 19: N0.H0G0008182	Pag. 40: N0.HOG0011930
Pag. 20: N0.HOG0008418	Pag. 41: N0.HOG0012348
Pag. 21: N0.H0G0008512	Pag. 42: N0.HOG0012854
Pag. 22: N0.HOG0008654	





Cestoda

Trematoda

Normalized TPM

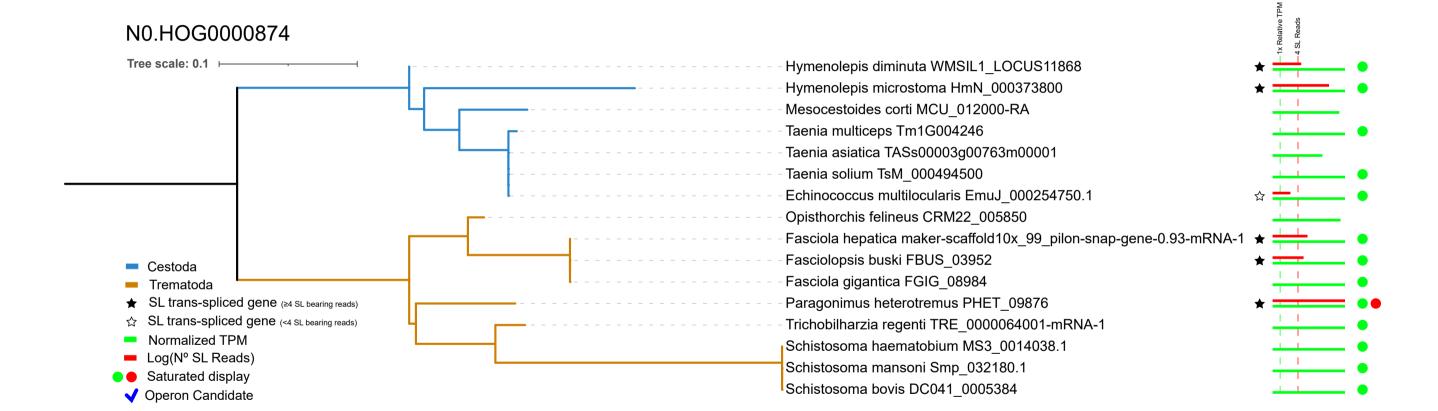
Log(N° SL Reads)

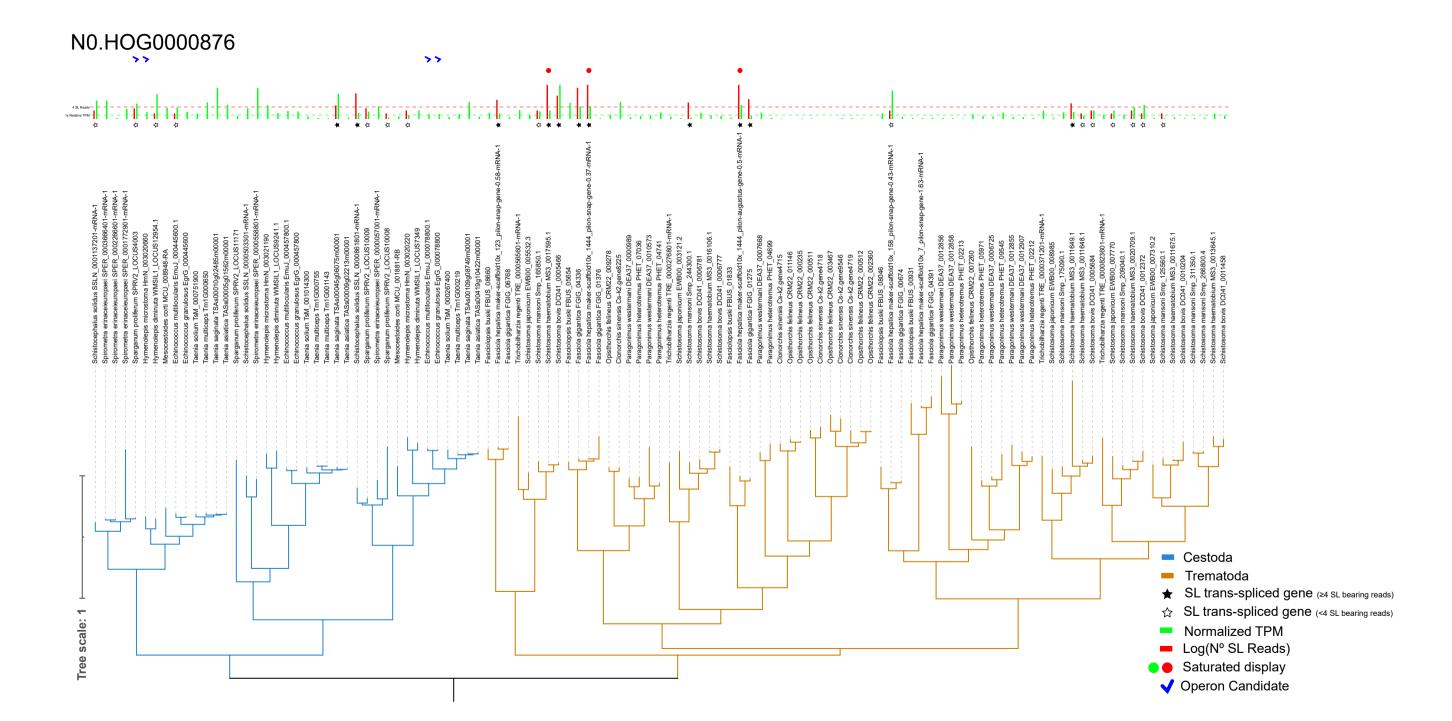
Operon Candidate

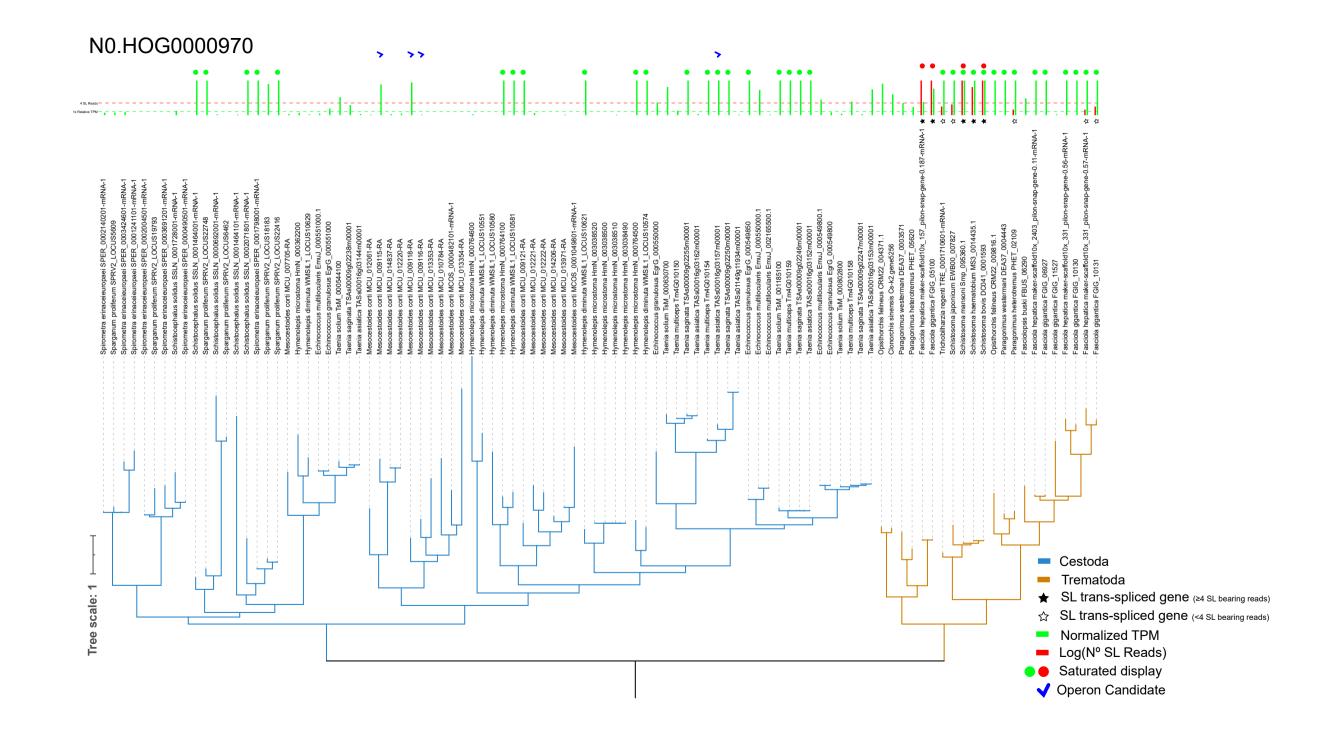
Saturated display

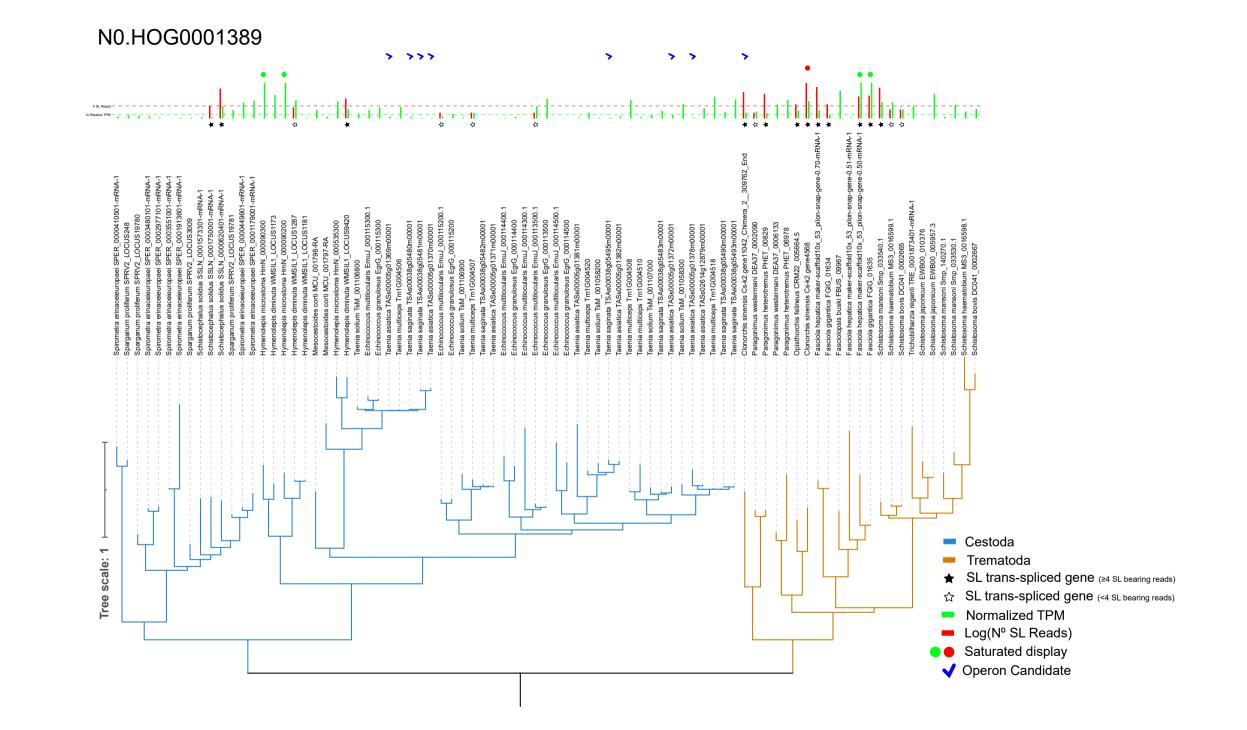
★ SL trans-spliced gene (≥4 SL bearing reads)

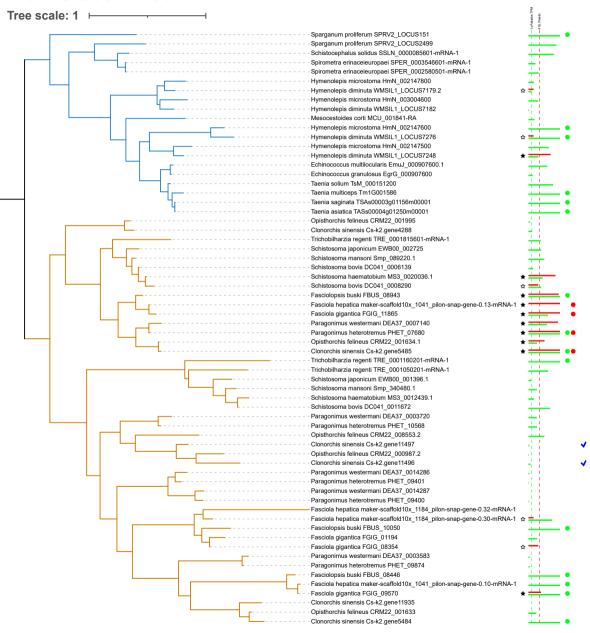
SL trans-spliced gene (<4 SL bearing reads)



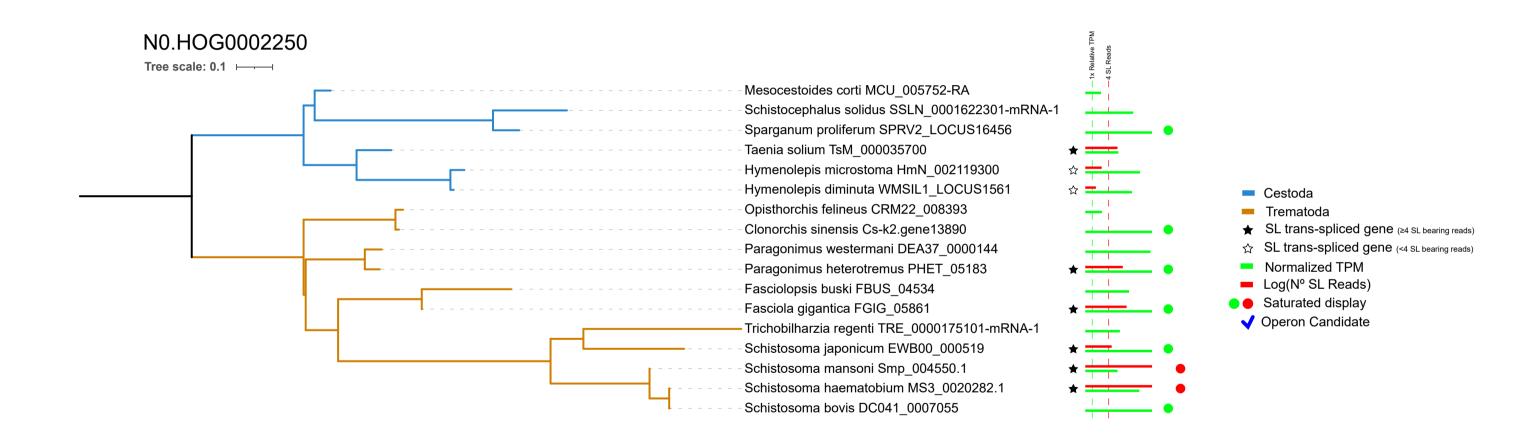


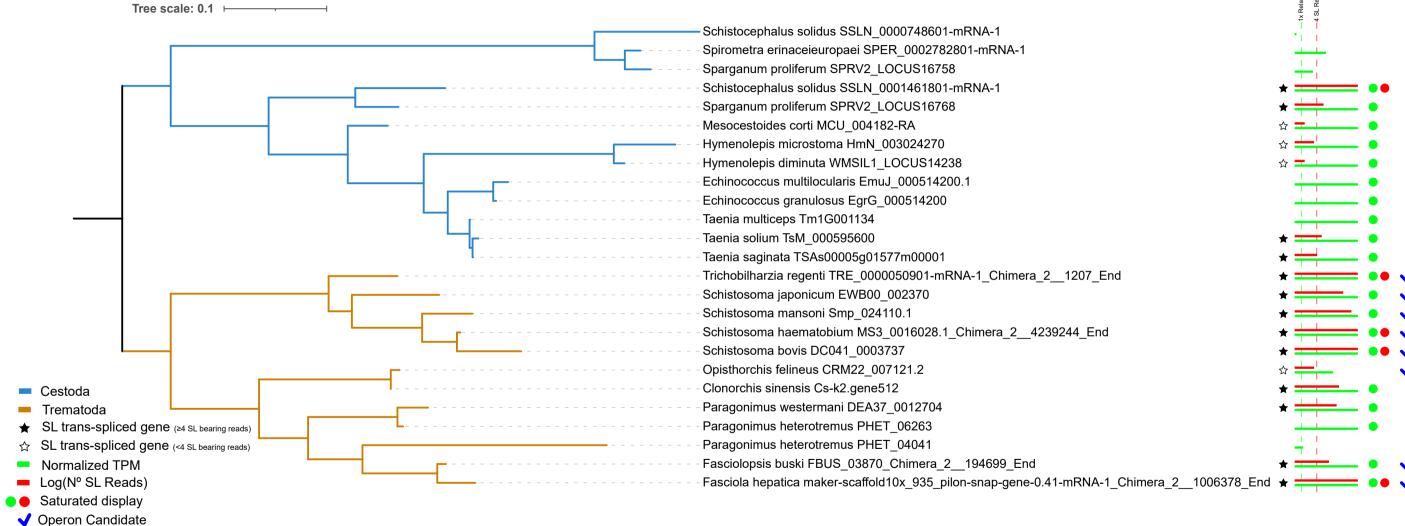


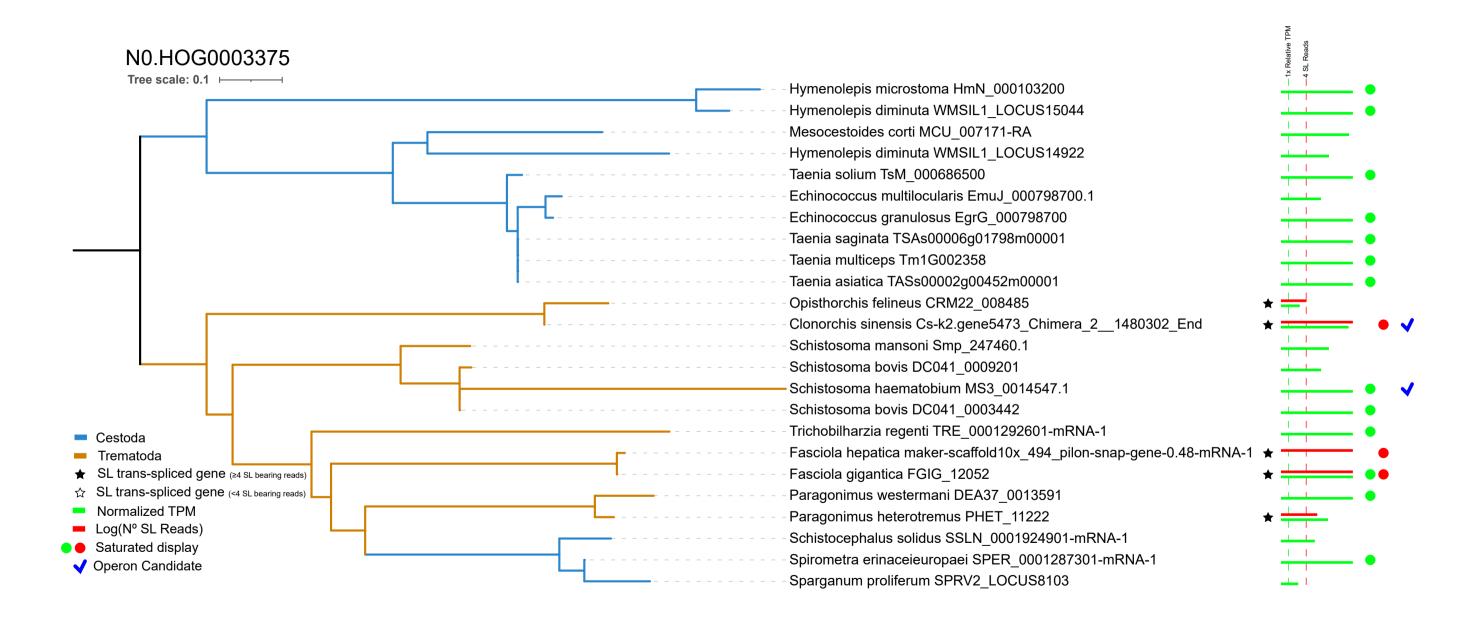


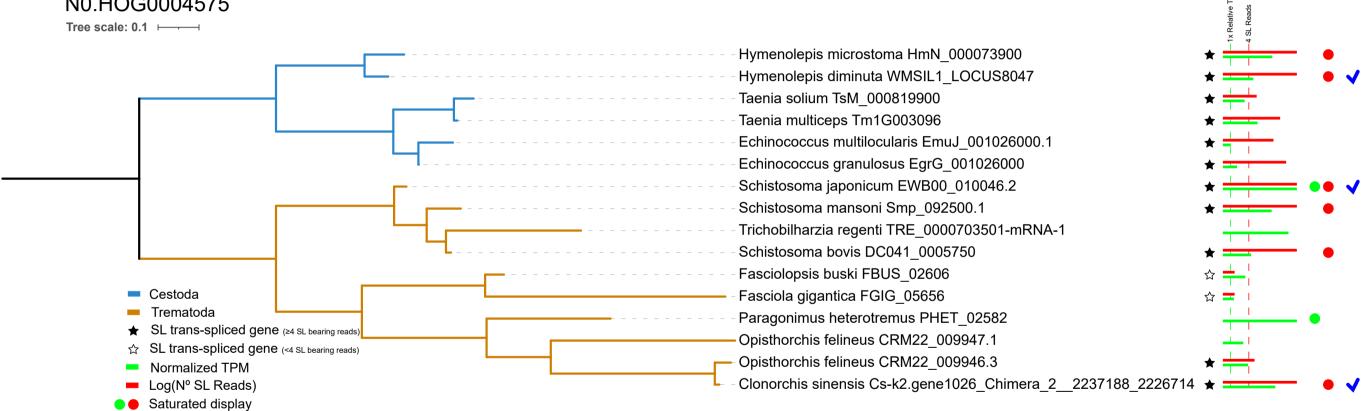


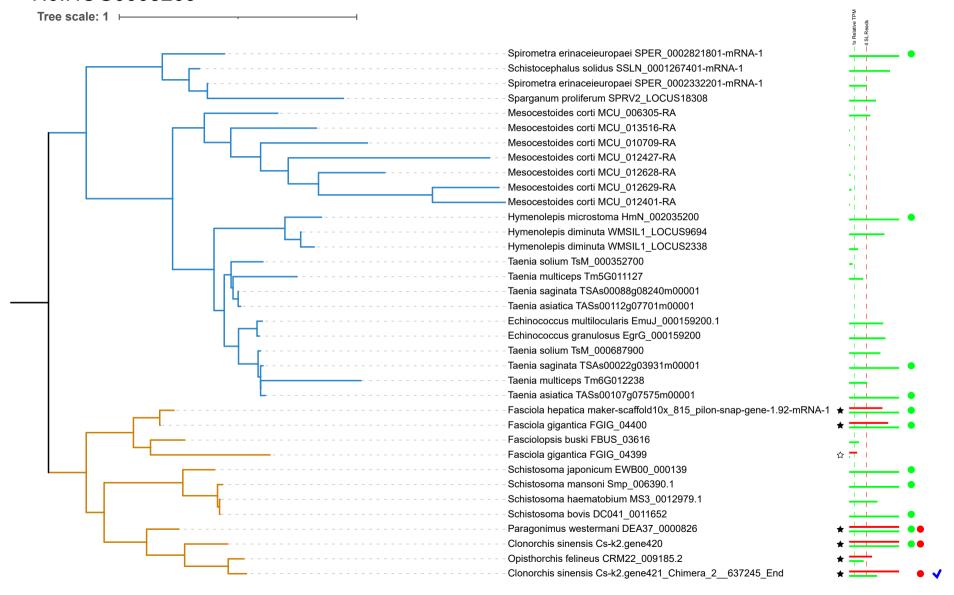
- Cestoda
- Trematoda
- ★ SL trans-spliced gene (≥4 SL bearing reads)
- ☆ SL trans-spliced gene (<4 SL bearing reads)
- Normalized TPM
- Log(N° SL Reads)
- Saturated display
- ✓ Operon Candidate



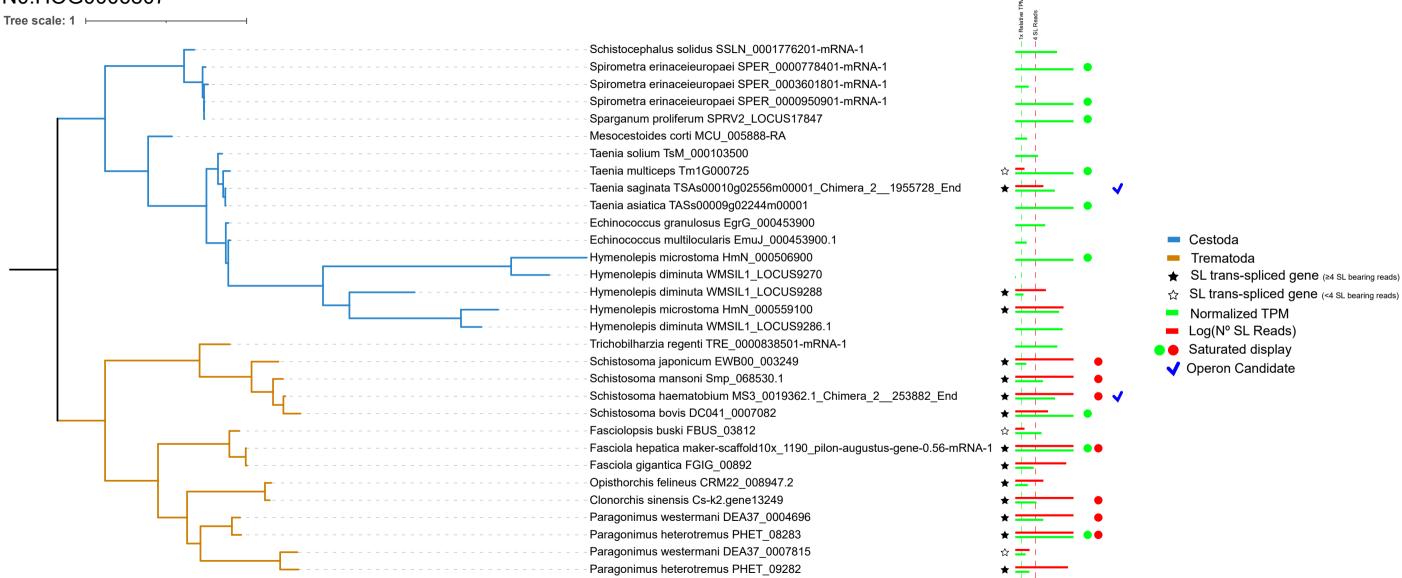


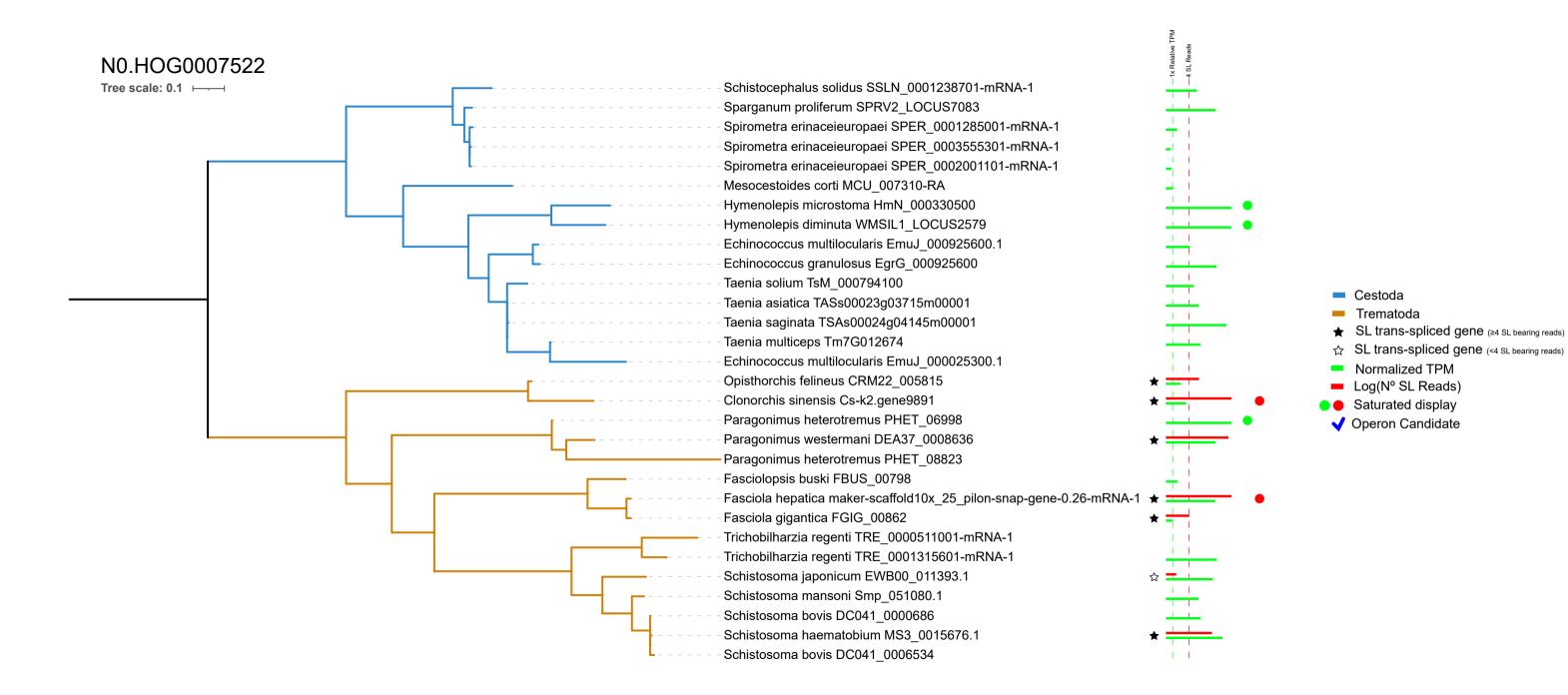




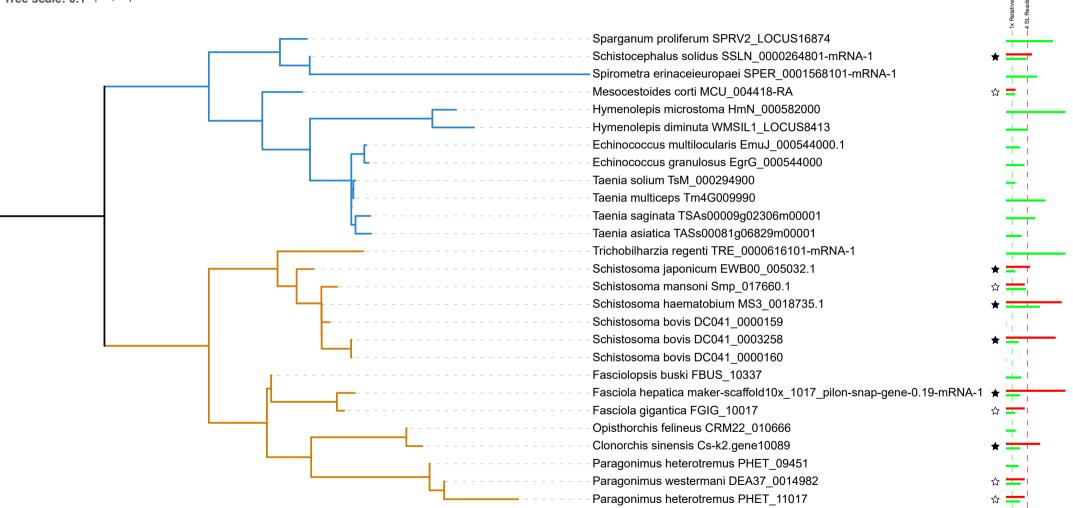


- Cestoda
- Trematoda
- ★ SL trans-spliced gene (≥4 SL bearing reads)
- ☆ SL trans-spliced gene (<4 SL bearing reads)</p>
- Normalized TPM
- Log(N° SL Reads)
- Saturated display
- ✓ Operon Candidate





Tree scale: 0.1



Cestoda

Trematoda

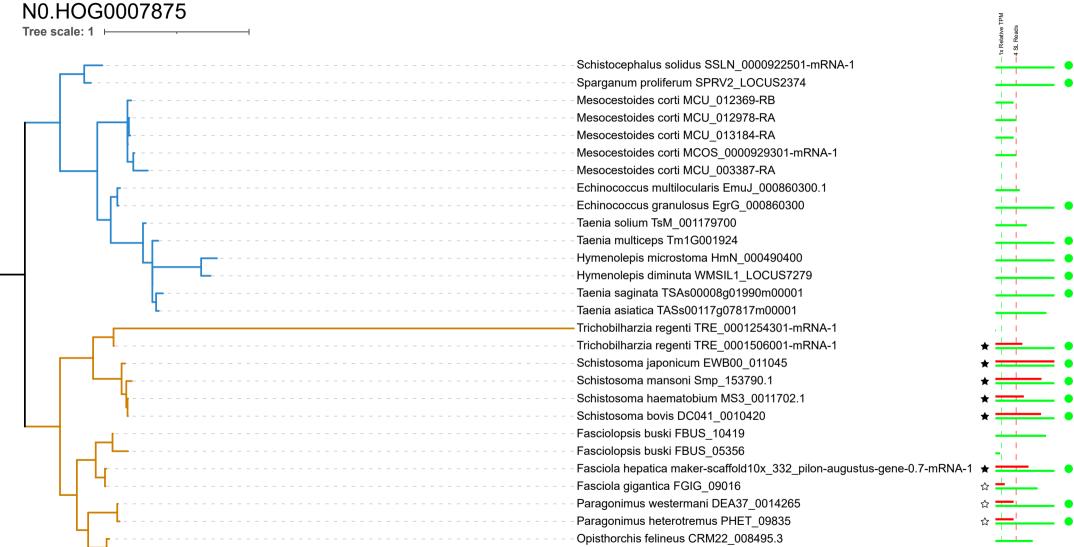
★ SL trans-spliced gene (≥4 SL bearing reads)

☆ SL trans-spliced gene (<4 SL bearing reads)

Normalized TPM

Log(N° SL Reads)

Saturated display



Clonorchis sinensis Cs-k2.gene5456

Cestoda

Trematoda

Normalized TPM

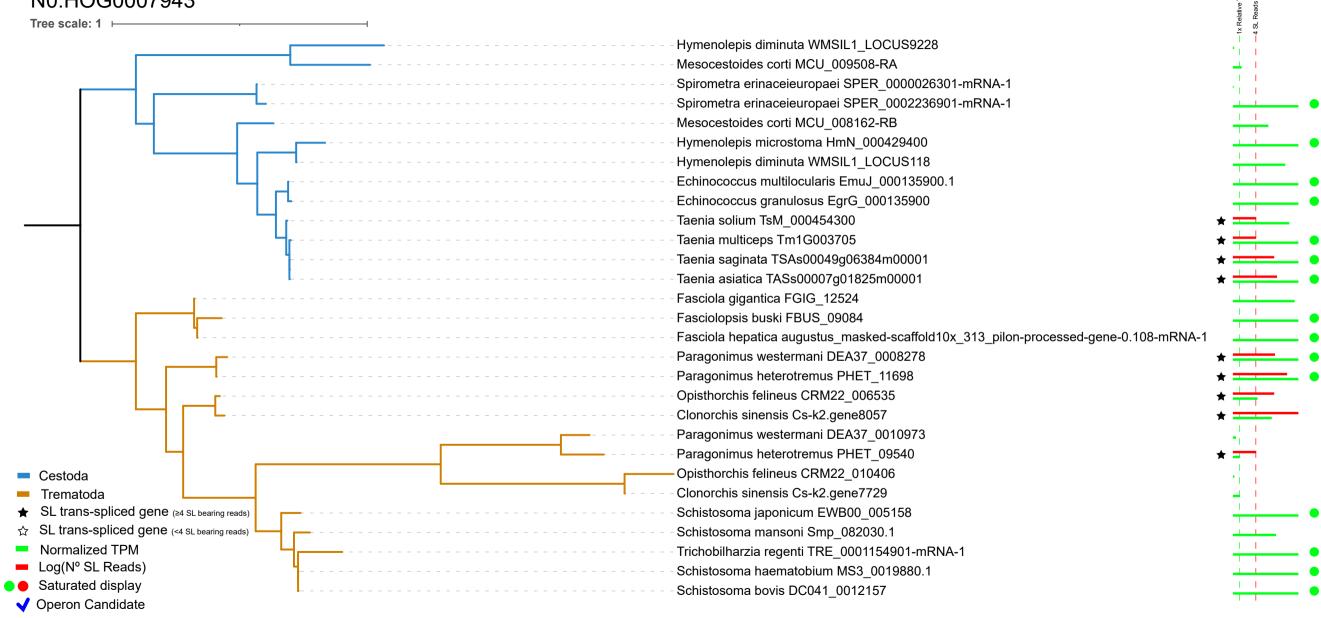
Log(N° SL Reads)

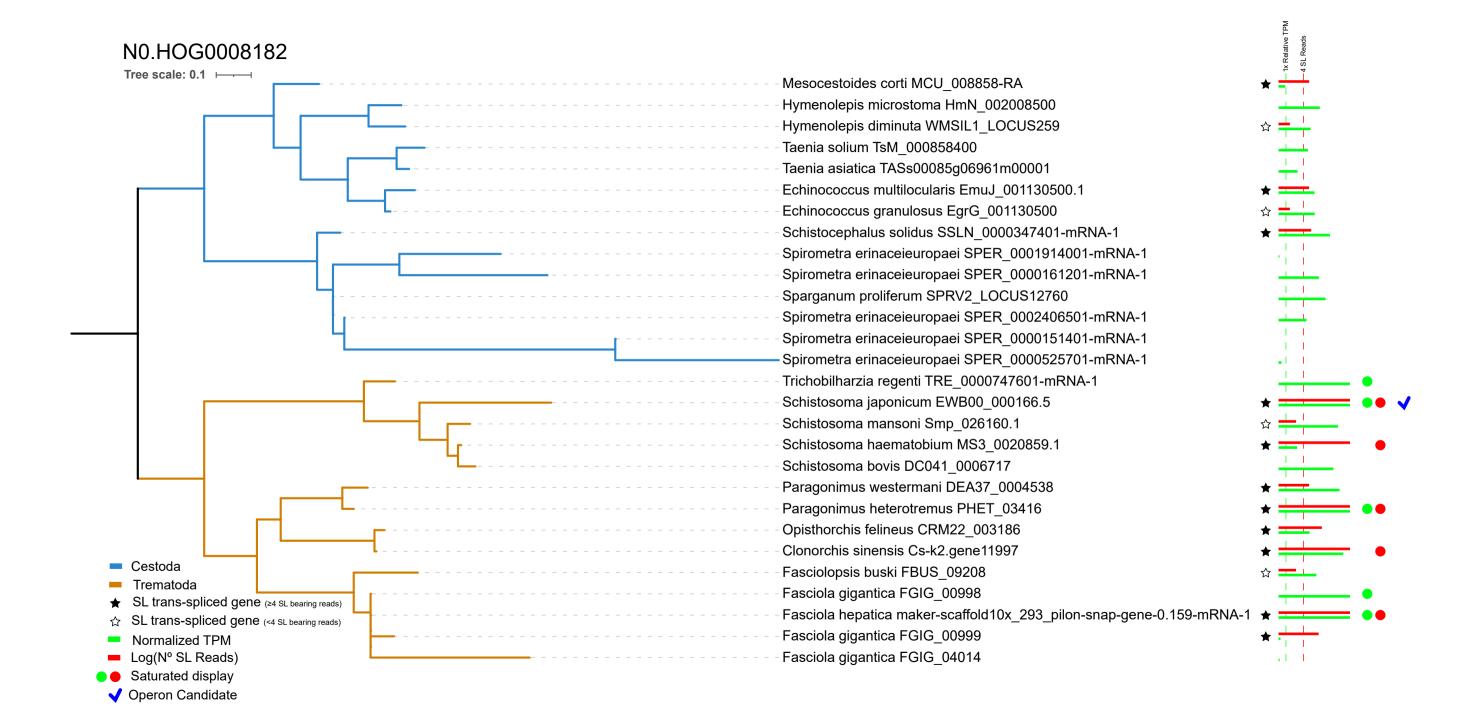
✓ Operon Candidate

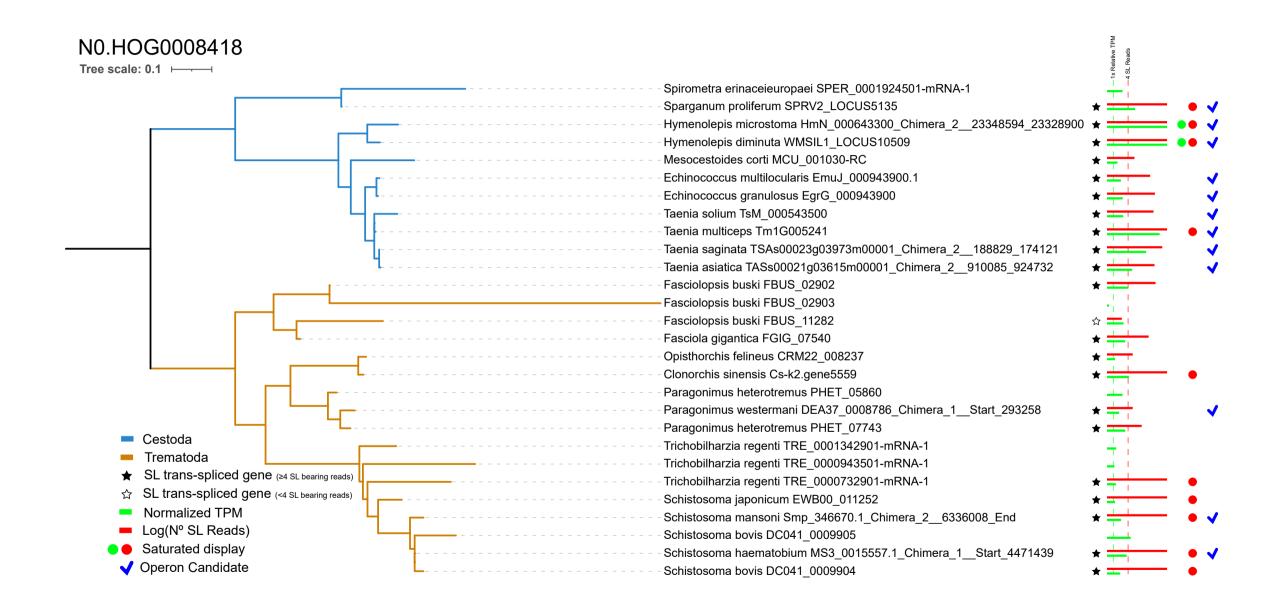
Saturated display

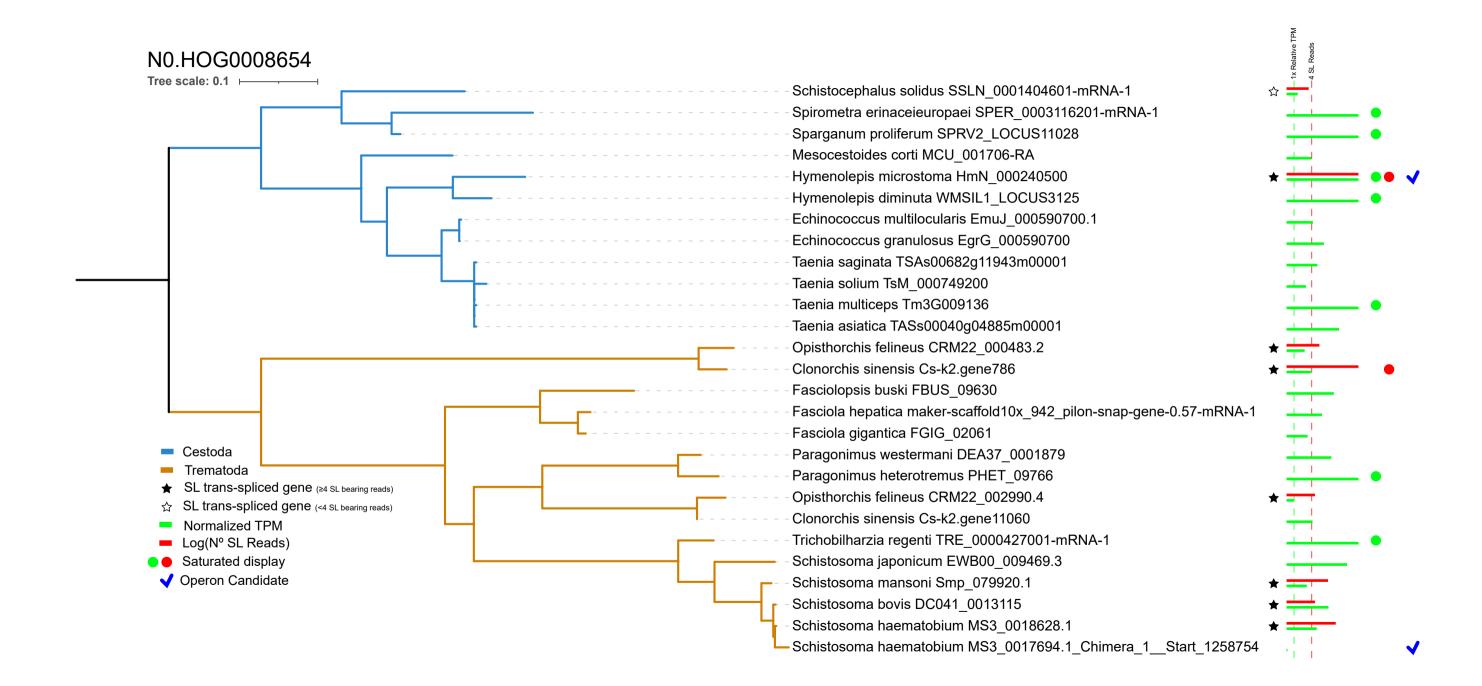
★ SL trans-spliced gene (≥4 SL bearing reads)

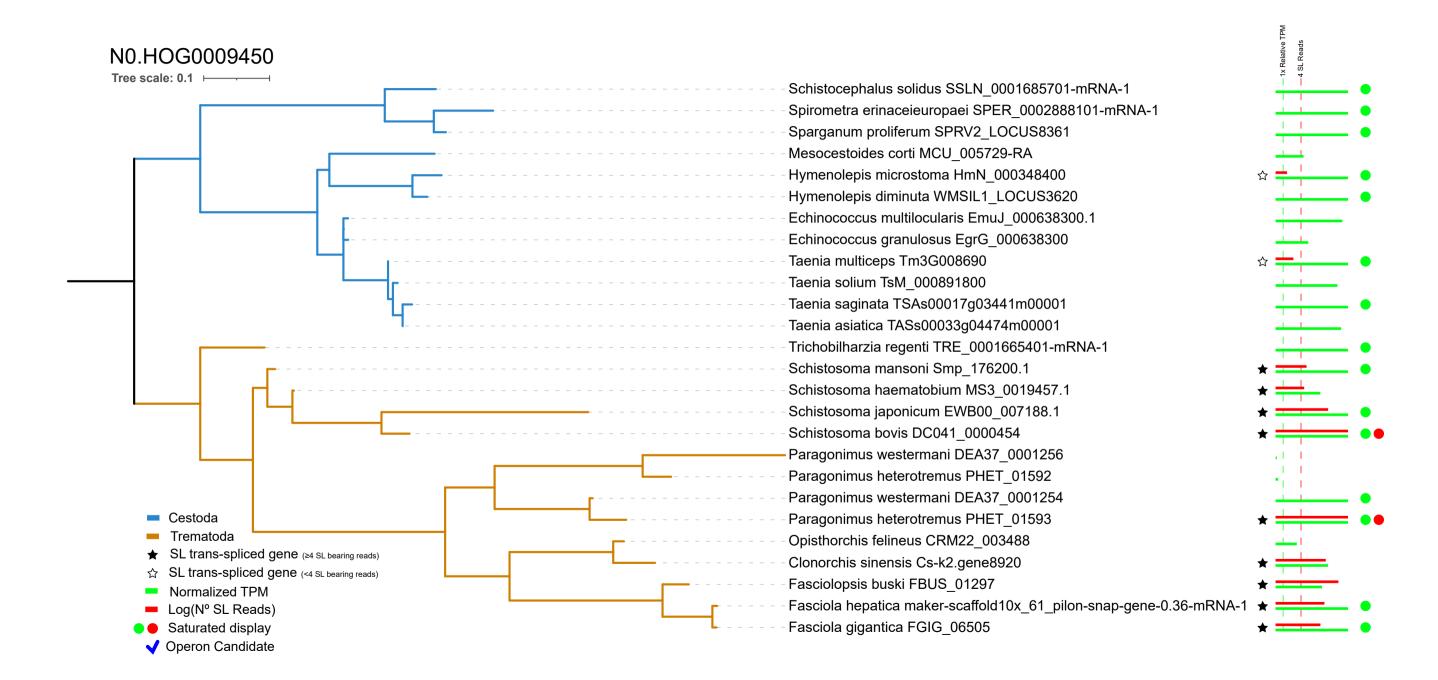
☆ SL trans-spliced gene (<4 SL bearing reads)



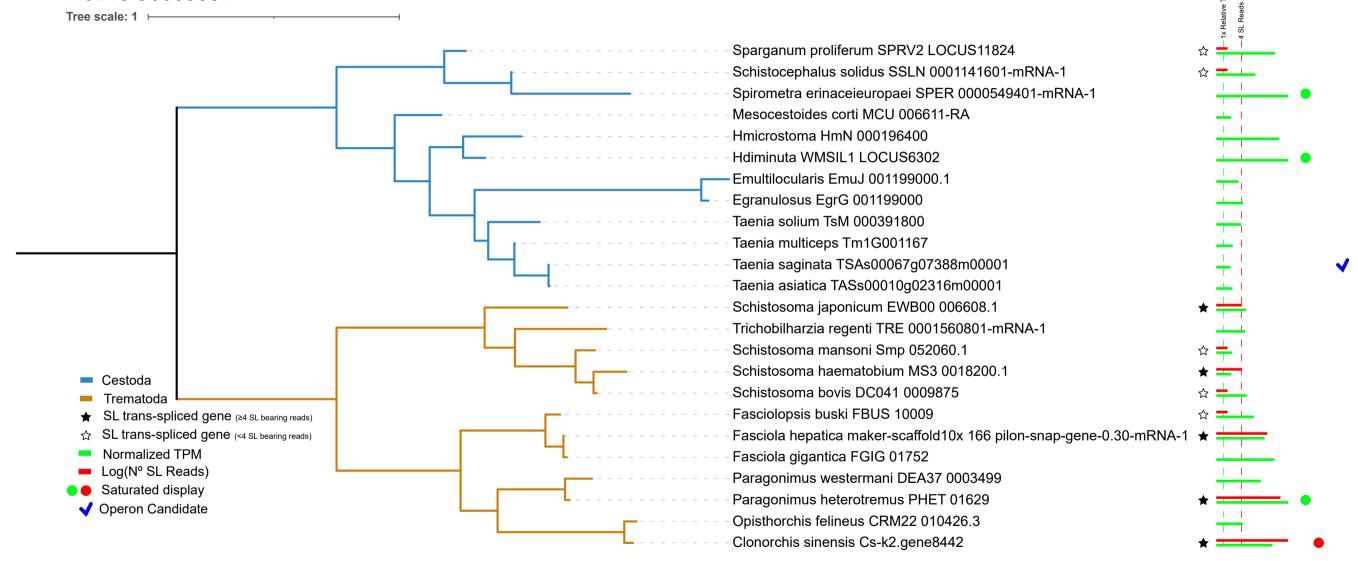


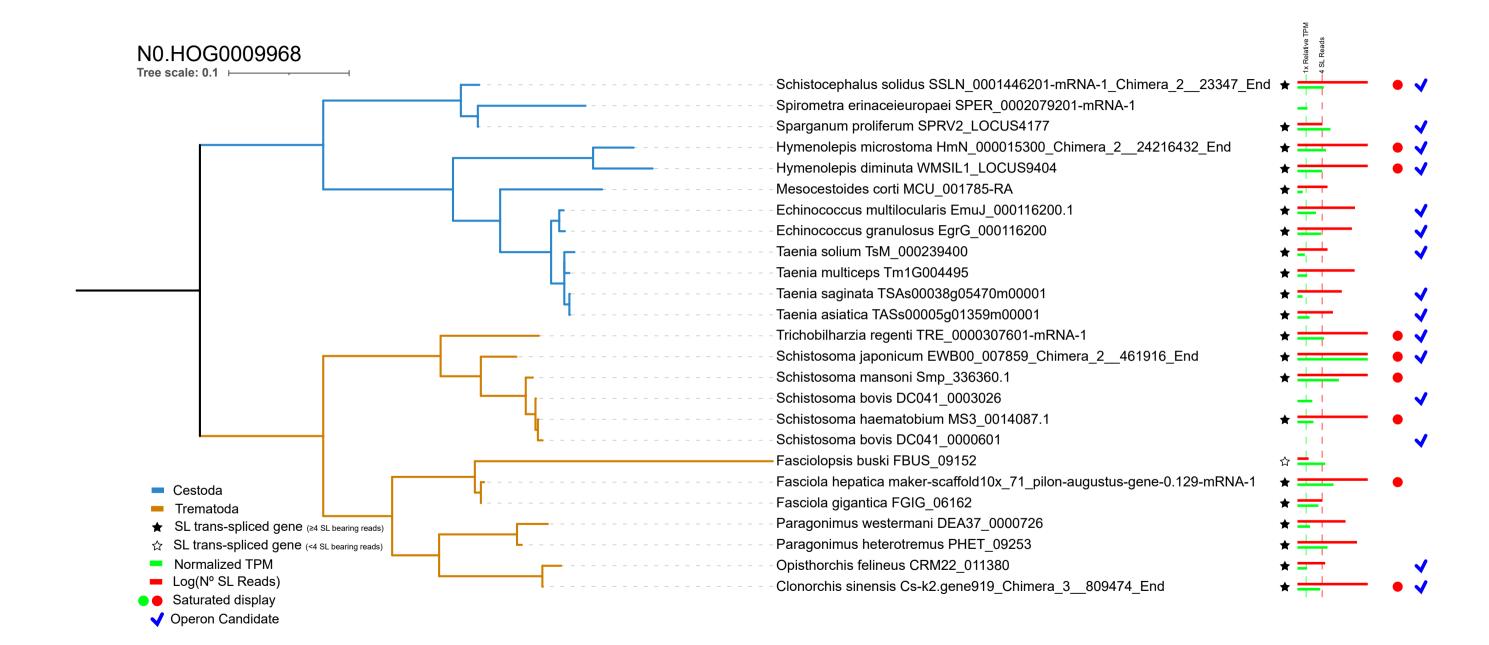


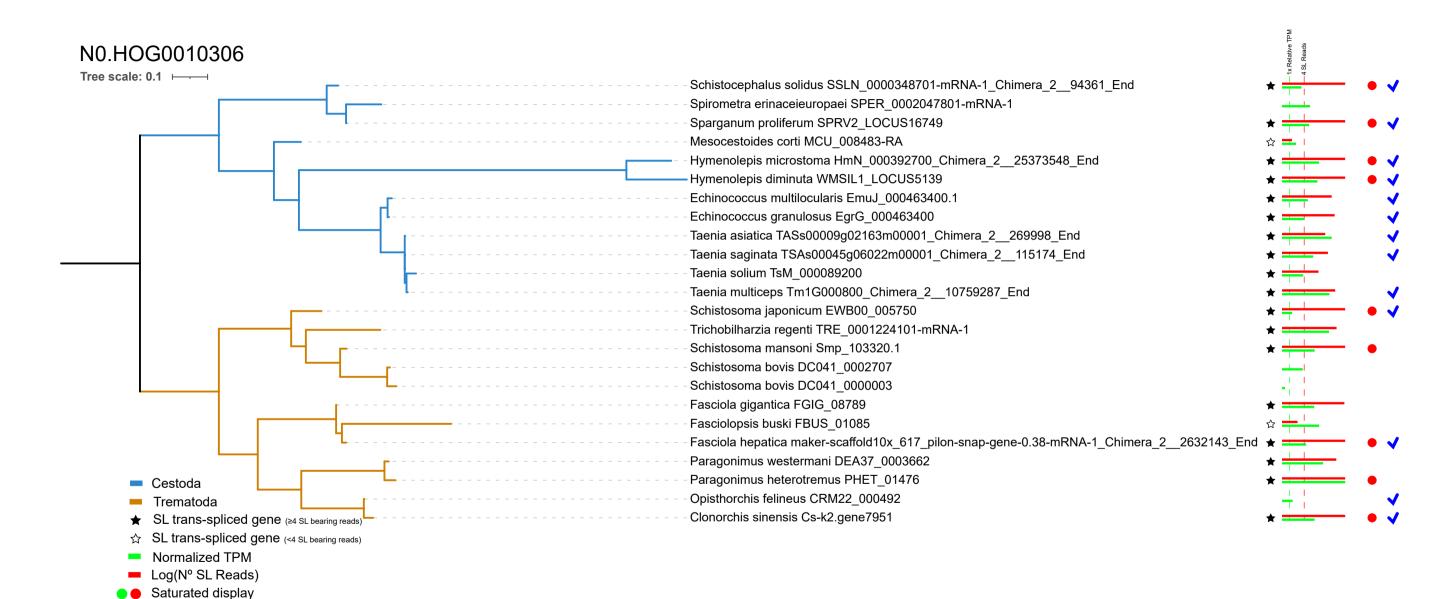


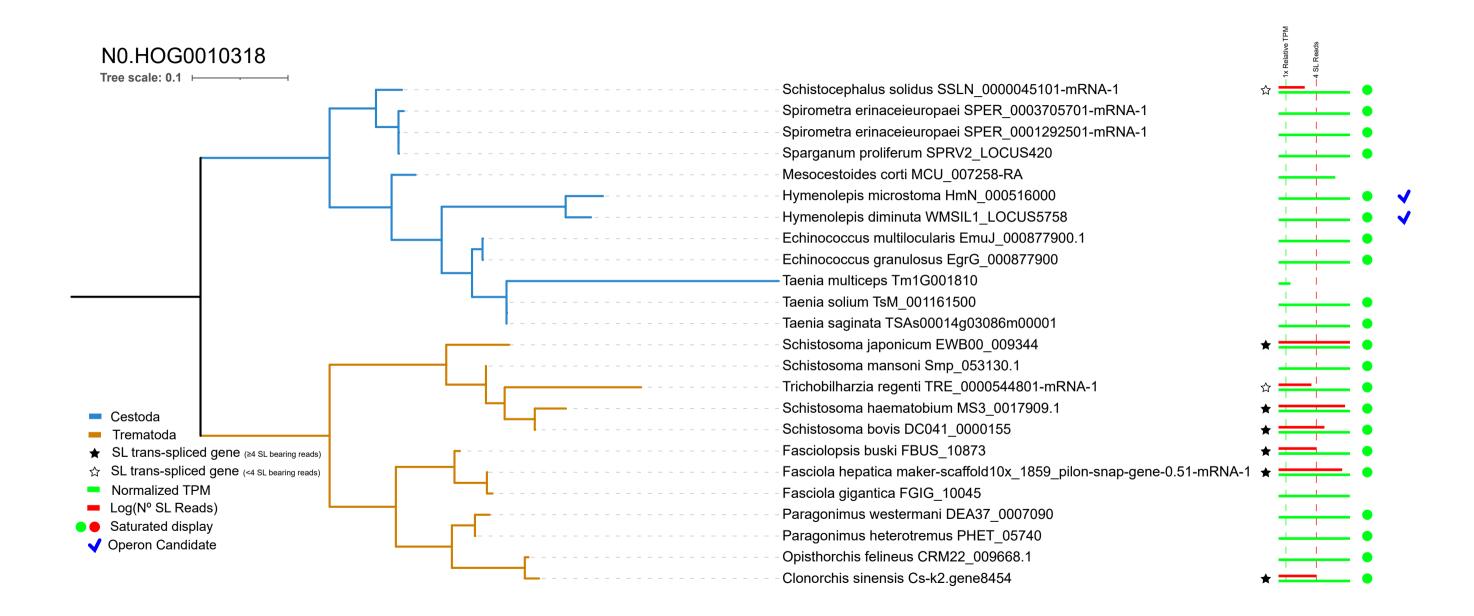


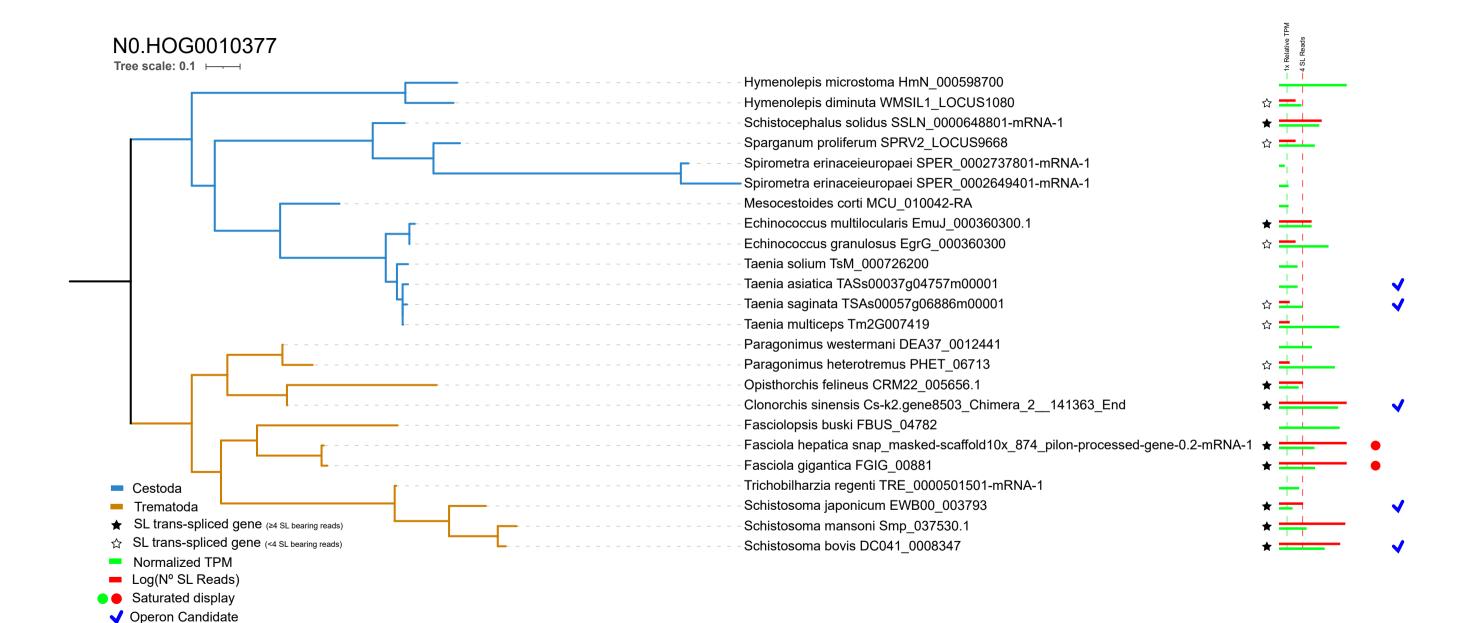


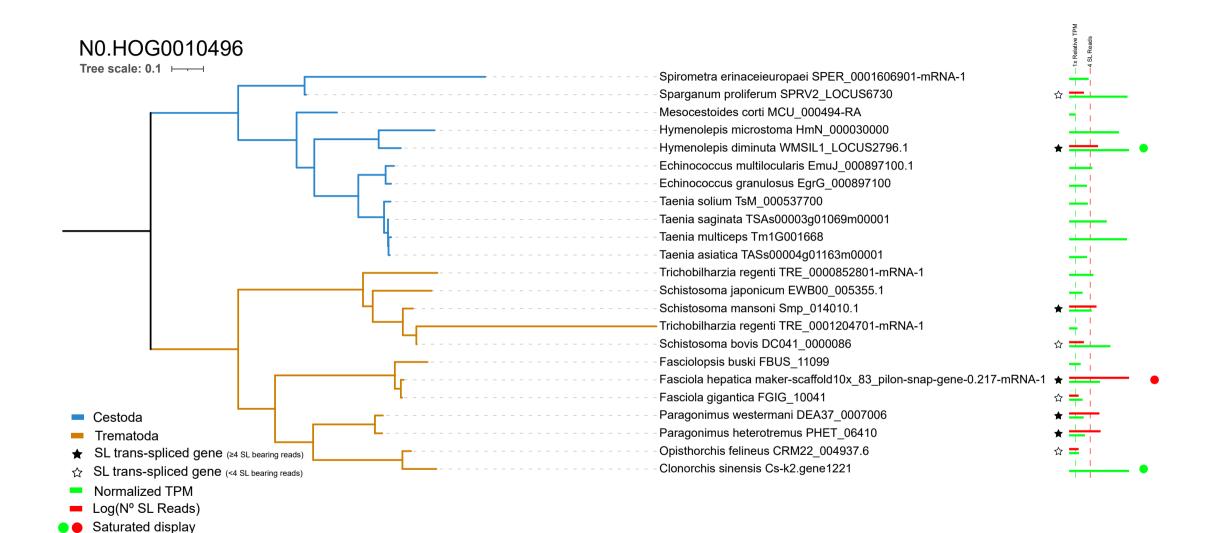




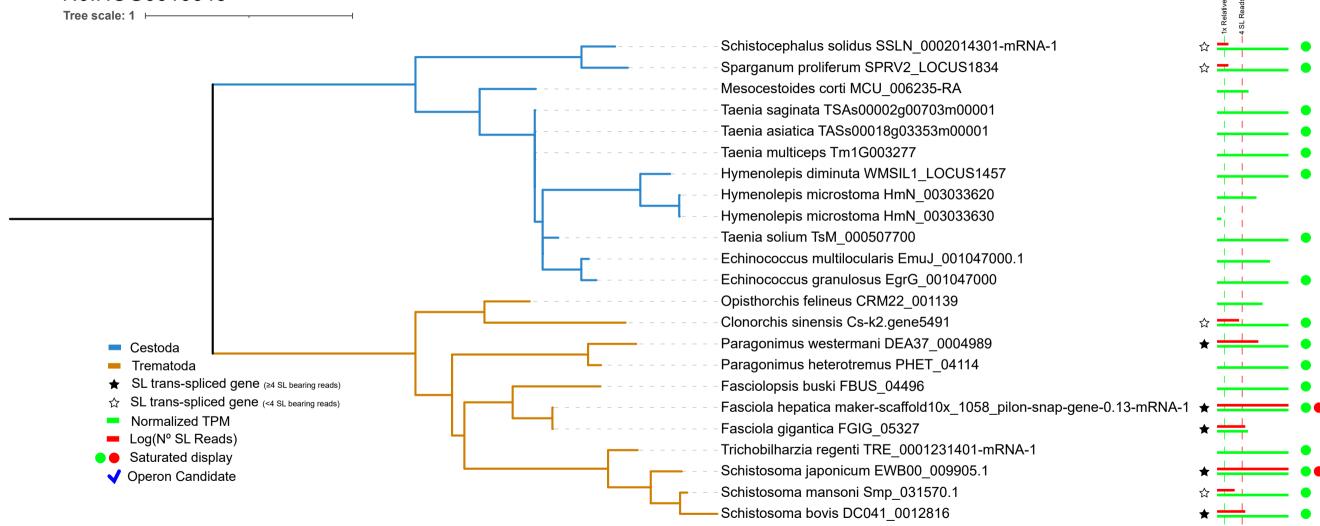


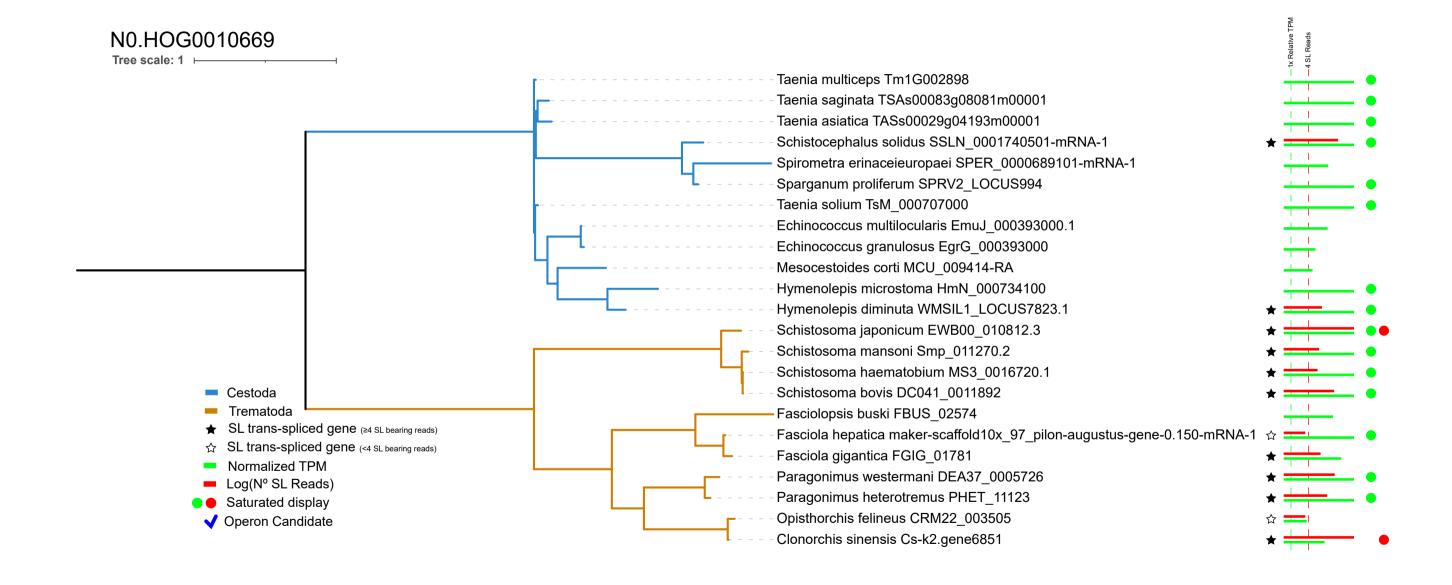


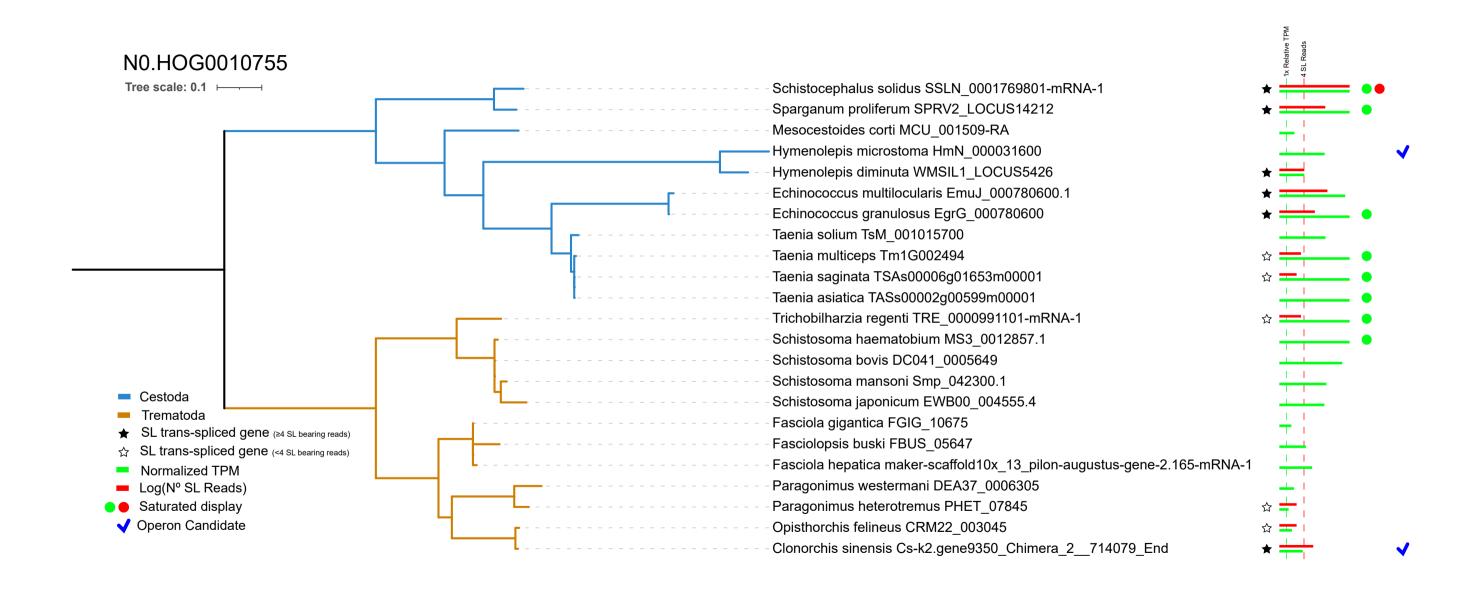




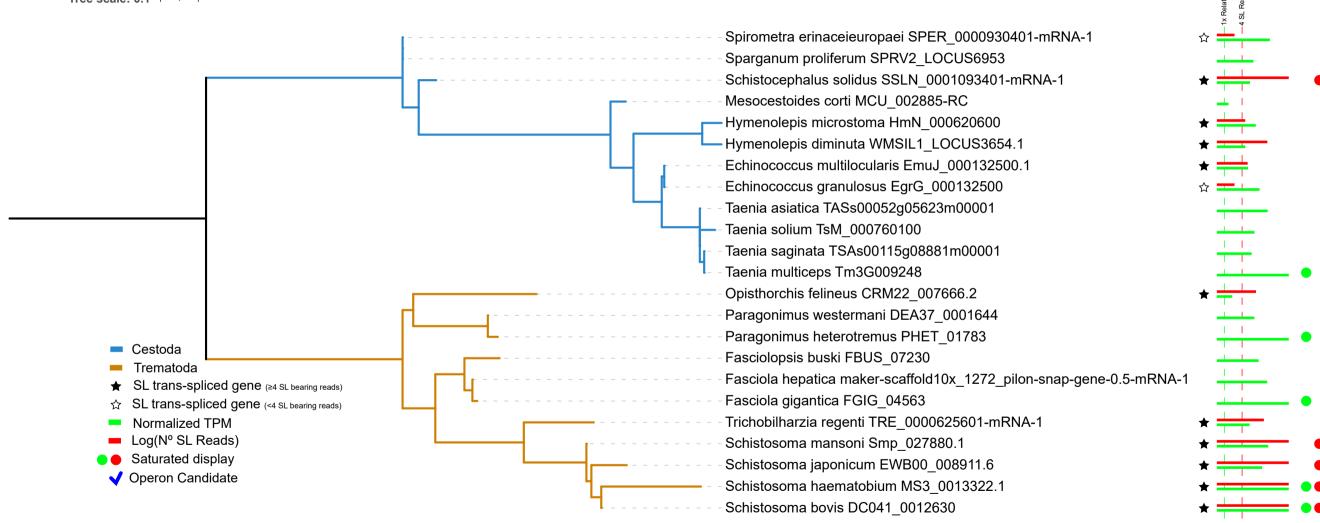




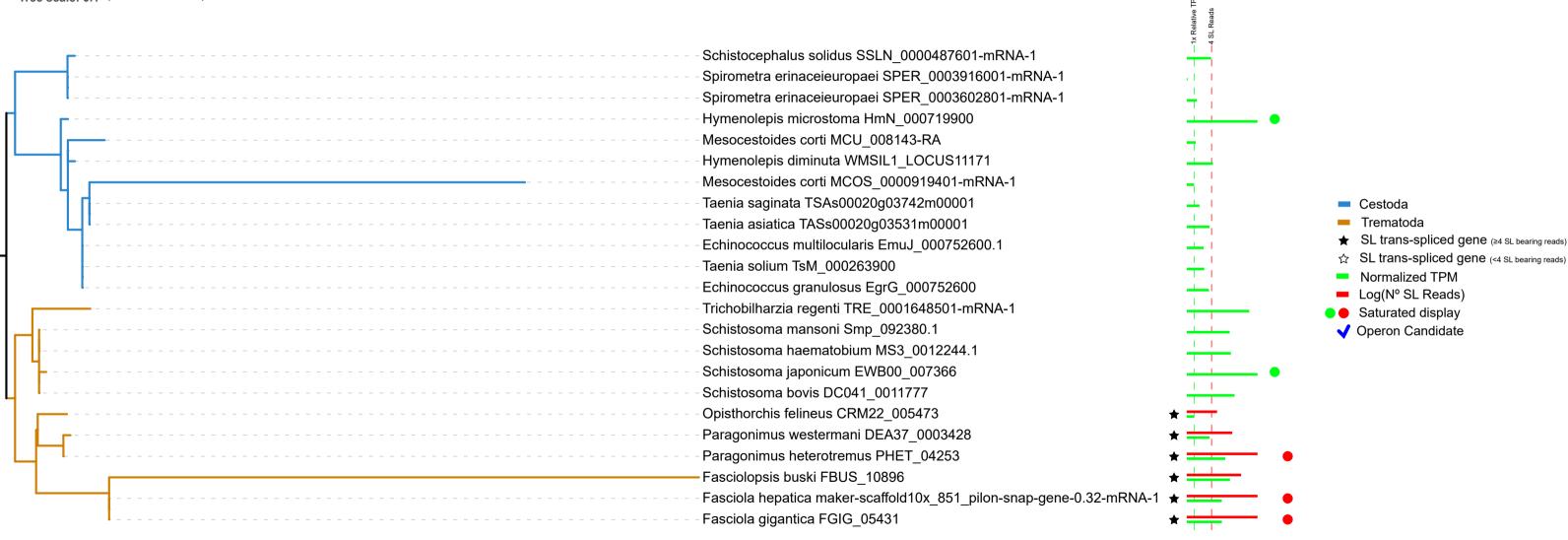


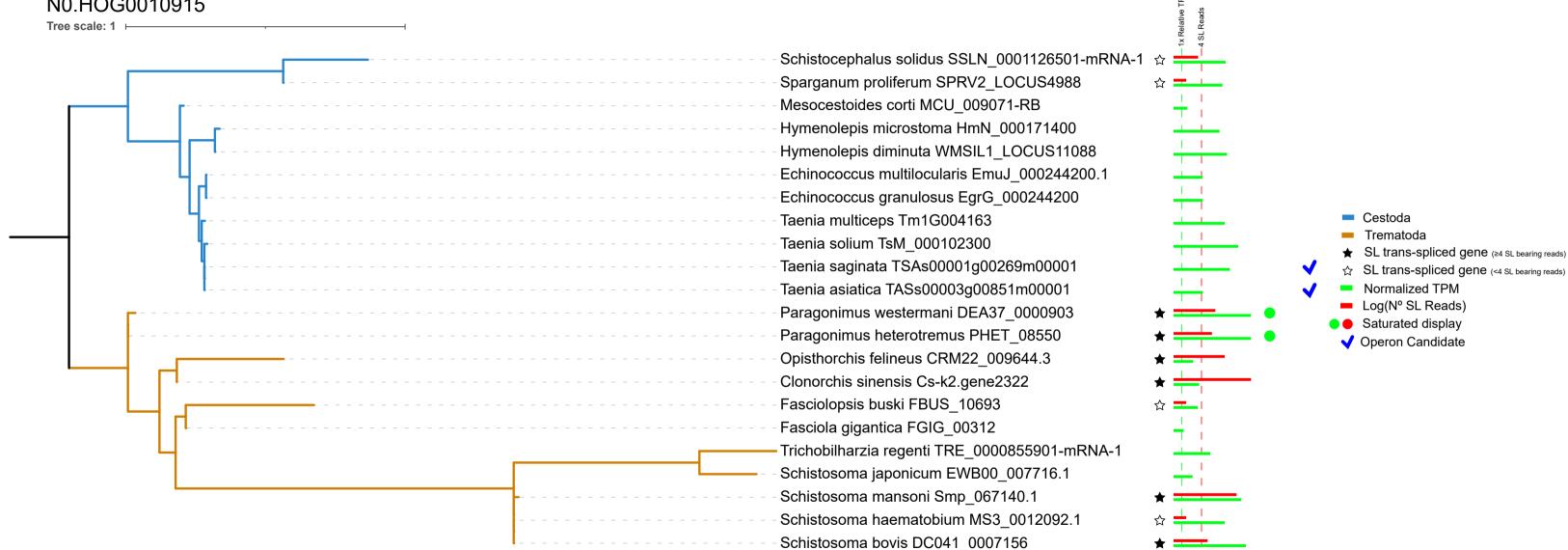


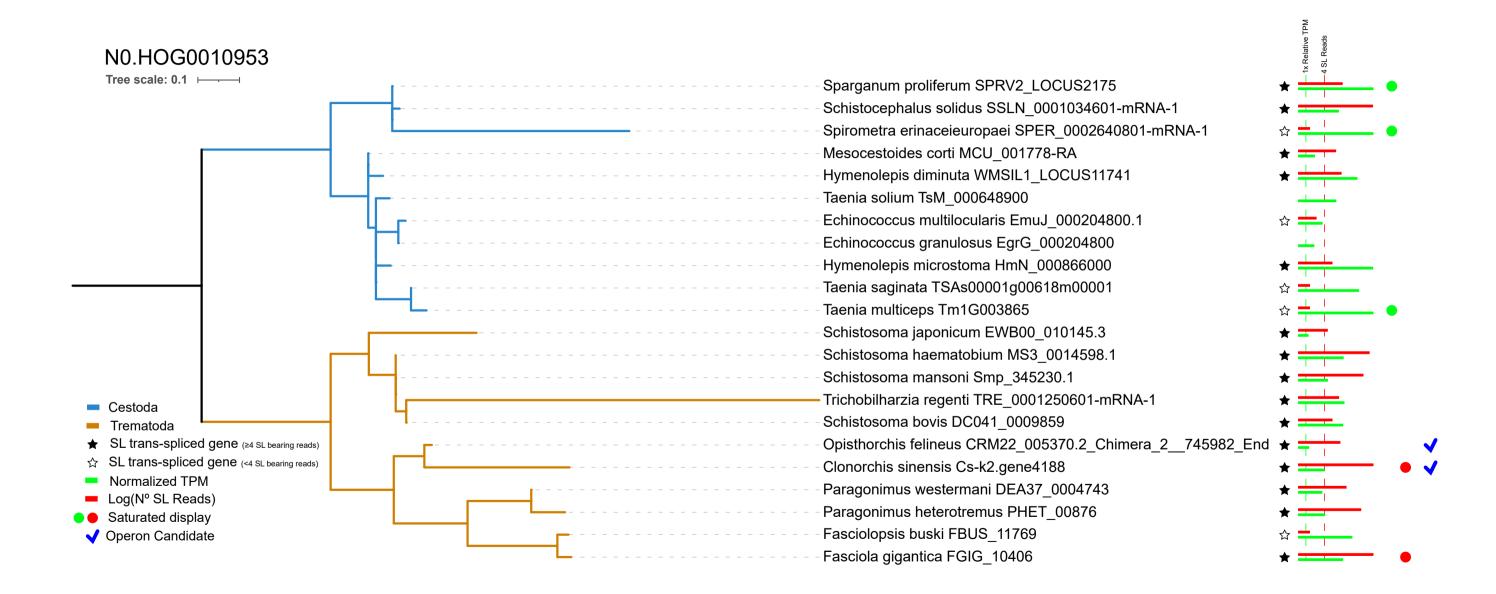
Tree scale: 0.1



Tree scale: 0.1







N0.HOG0011542 Tree scale: 0.1 Schistocephalus solidus SSLN 0001109701-mRNA-1 Spirometra erinaceieuropaei SPER 0002897301-mRNA-1 Spirometra erinaceieuropaei SPER_0002878901-mRNA-1 Mesocestoides corti MCU 011893-RA Hymenolepis microstoma HmN 000224900 Hymenolepis diminuta WMSIL1 LOCUS9149 Echinococcus multilocularis EmuJ 000074500.1 Echinococcus granulosus EgrG 000074500 Taenia saginata TSAs00081g08026m00001 Taenia solium TsM 000564100 Taenia asiatica TASs00069g06337m00001 Opisthorchis felineus CRM22 007120.6 Fasciolopsis buski FBUS_03870_Chimera_1__Start_194699 - Fasciola hepatica maker-scaffold10x_935_pilon-snap-gene-0.41-mRNA-1_Chimera_1__Start_1006378 🖈 ==== Fasciola gigantica FGIG 05463 Trichobilharzia regenti TRE 0000050901-mRNA-1 Chimera 1 Start 1207 Cestoda Schistosoma japonicum EWB00_002371 Trematoda

★ SL trans-spliced gene (≥4 SL bearing reads)

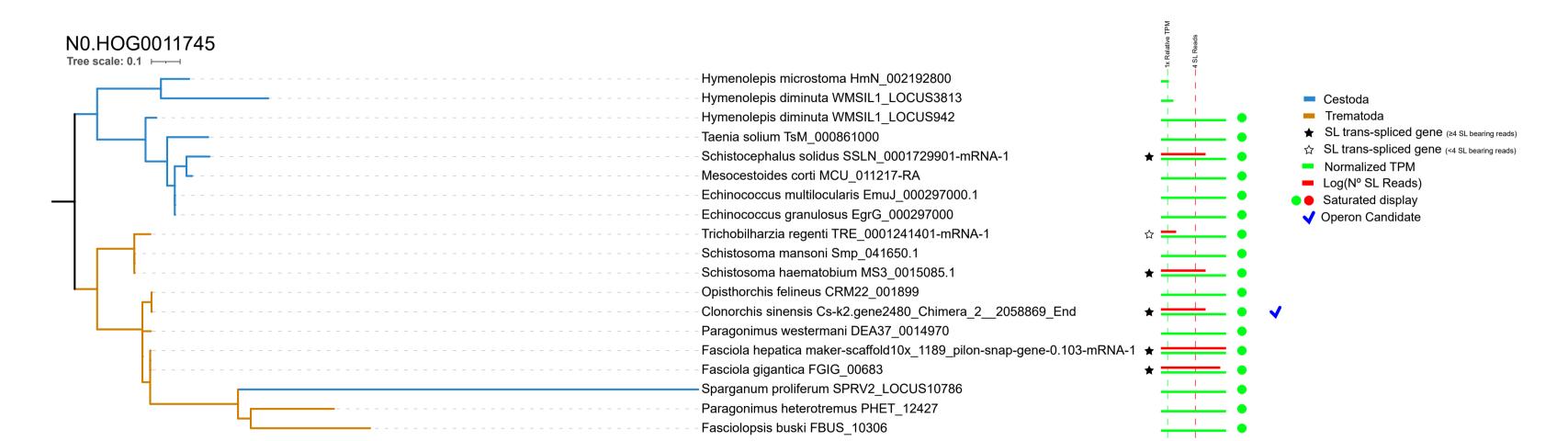
☆ SL trans-spliced gene (<4 SL bearing reads)

Normalized TPM
Log(N° SL Reads)
Saturated display
Operon Candidate

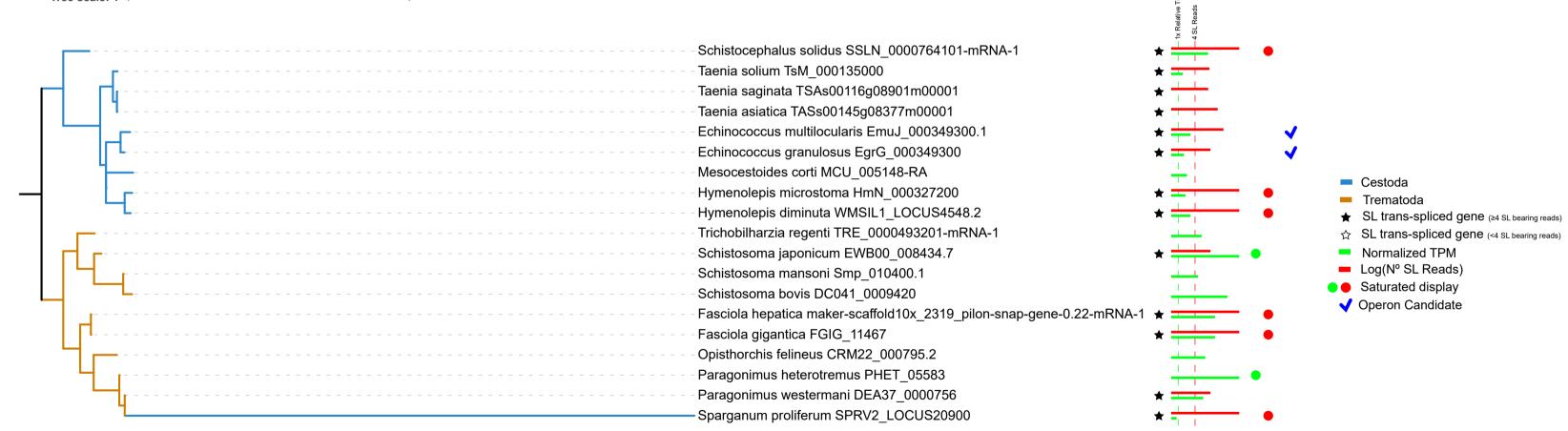
Schistosoma mansoni Smp_024120.1

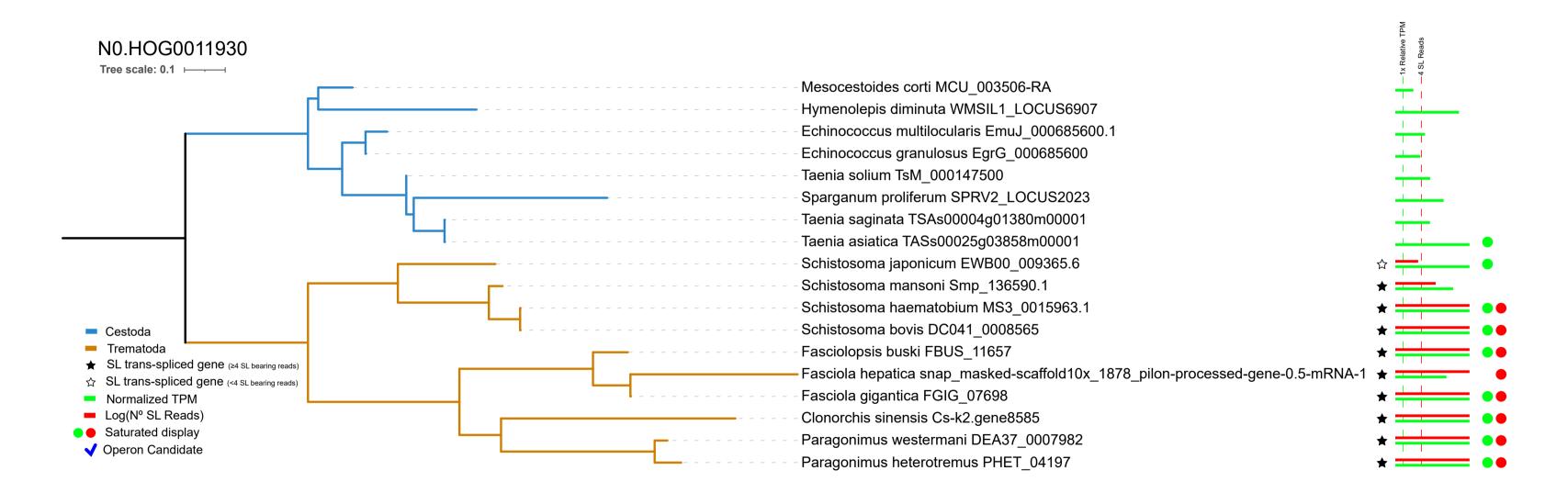
Schistosoma bovis DC041 0003731

Schistosoma haematobium MS3 0016028.1 Chimera 1 Start 4239267



Tree scale: 1





N0.HOG0012348 Tree scale: 0.1 Sparganum proliferum SPRV2 LOCUS7986 Mesocestoides corti MCU 011410-RA Hymenolepis microstoma HmN_000107400 Hymenolepis diminuta WMSIL1 LOCUS14892 Taenia solium TsM 000865000 Taenia multiceps Tm1G002324 ☆ === Echinococcus multilocularis EmuJ 000803300.1 Echinococcus granulosus EgrG 000803300 . accionopsis puski FBUS_06958 Fasciola hepatica snap_masked-scaffold10x_273_pilon-processed-gene-0.19-mRNA-1 Trichobilharzia regenti TRE_0001137601-mRNA-1 Schistosoma japonicum EWB00_008183 chistosoma mansoni Smp_136990.1 histosoma bovis DC041_0012200 Cestoda Trematoda ★ SL trans-spliced gene (≥4 SL bearing reads) ☆ SL trans-spliced gene (<4 SL bearing reads) Schistosoma bovis DC041_0002644 Normalized TPM

Log(N° SL Reads)Saturated displayOperon Candidate

