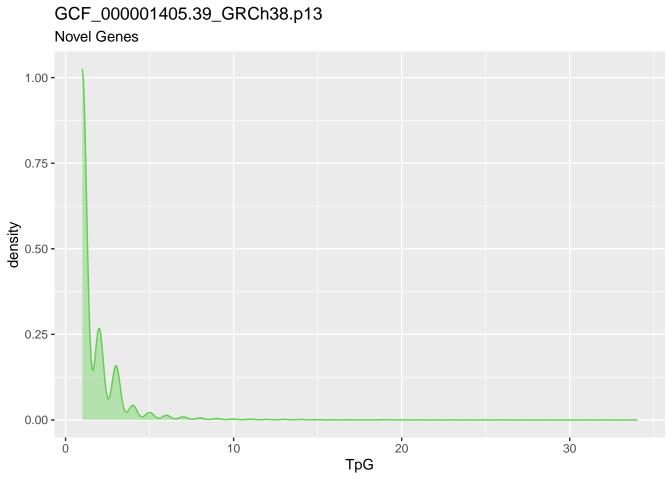
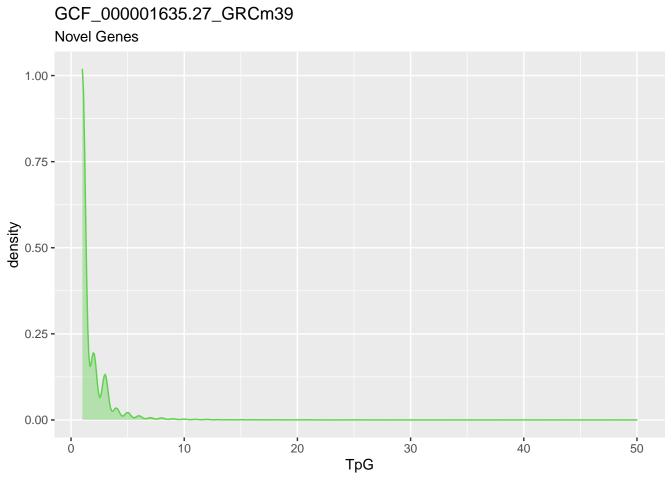
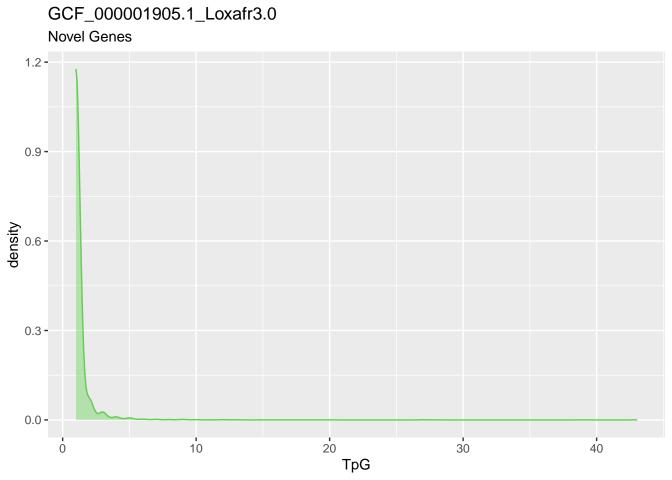
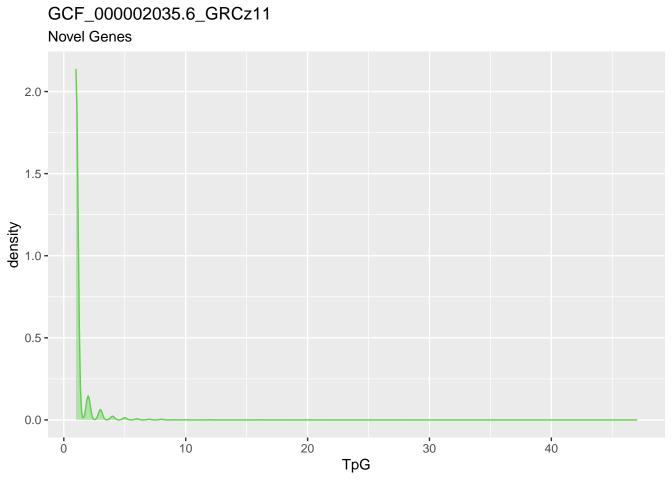
NOVEL GENE DENSITY PLOTS

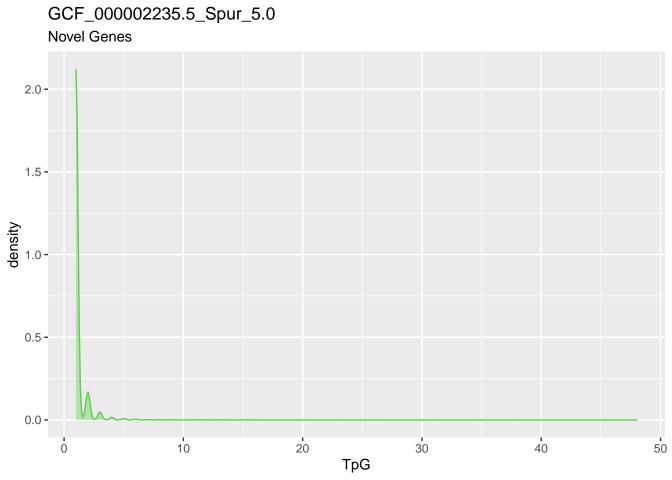
Figures below are density plots for all novel gene distributions (light green) for complexity metrics (TpG, EpT, and EpG) with densities on the y-axis and genetic element counts on the x-axis. Plots include every organism used in this study.

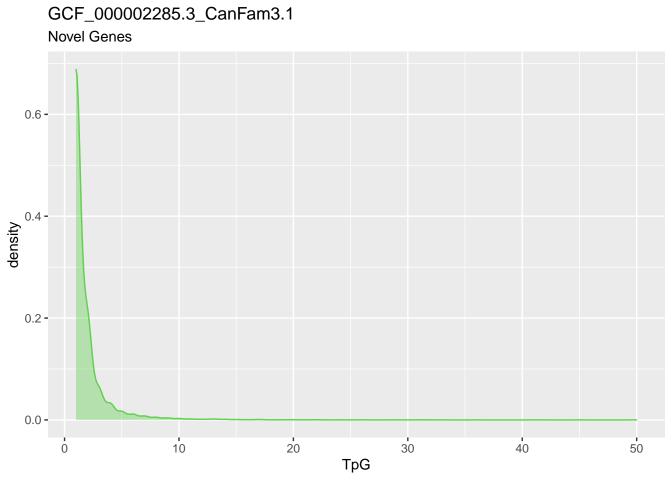






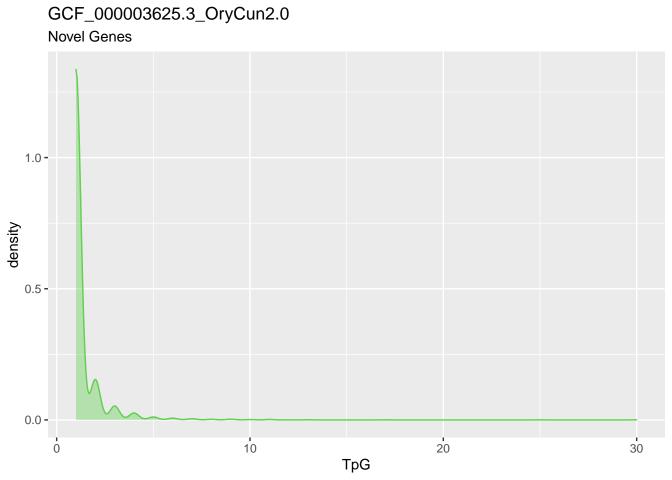


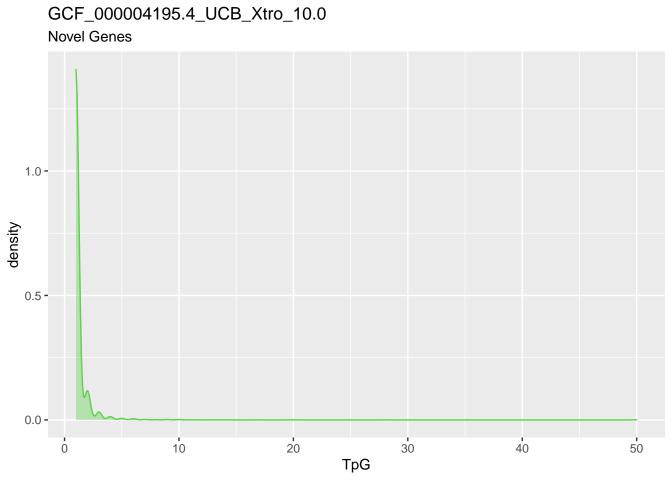


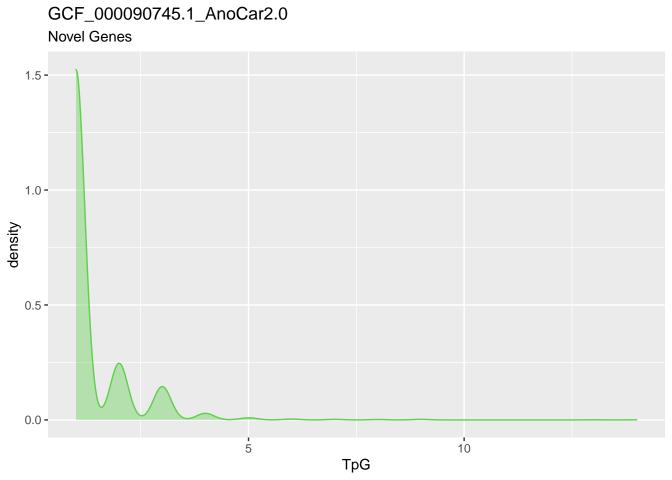


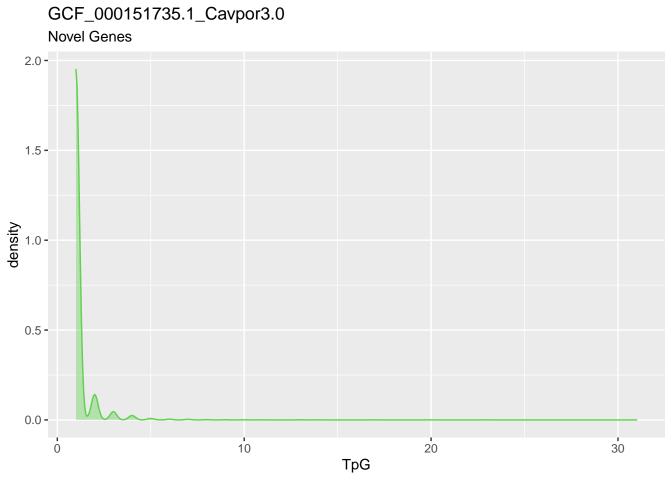
GCF_000002295.2_MonDom5 **Novel Genes** 1.0 density 0.5 -0.0 -10 20 30 40 50 TpG

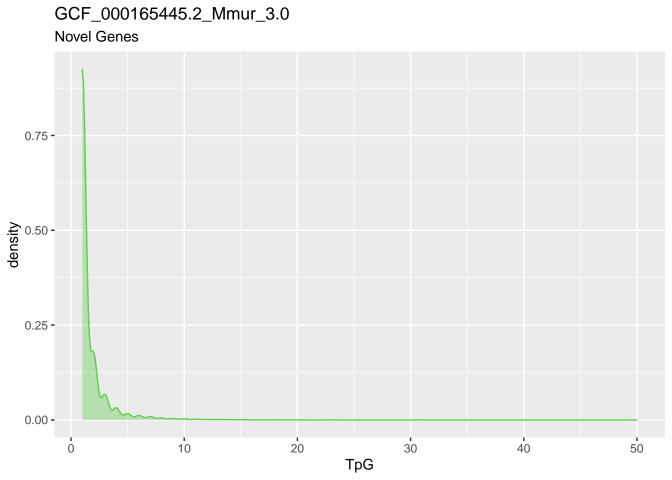
GCF_000003025.6_Sscrofa11.1 **Novel Genes** 0.6 density 0.2 -0.0 -10 20 30 40 50 TpG

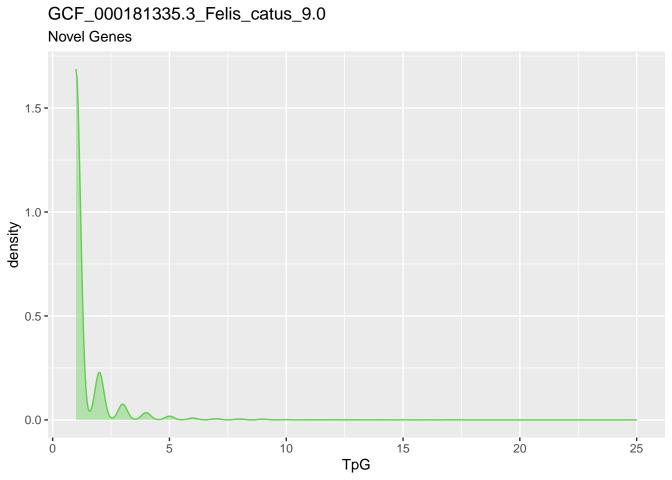


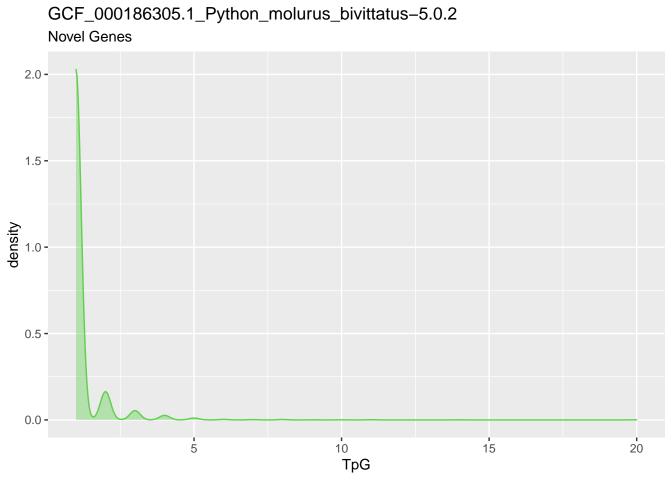


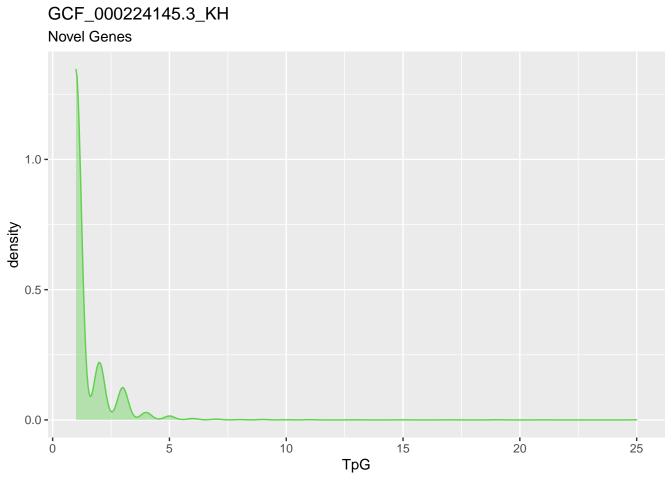


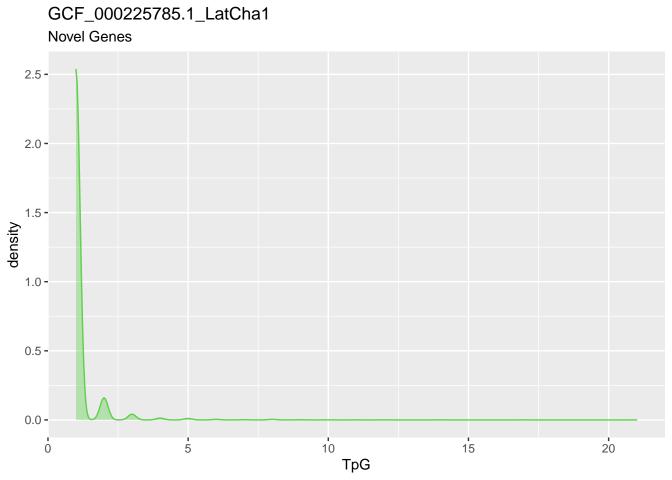


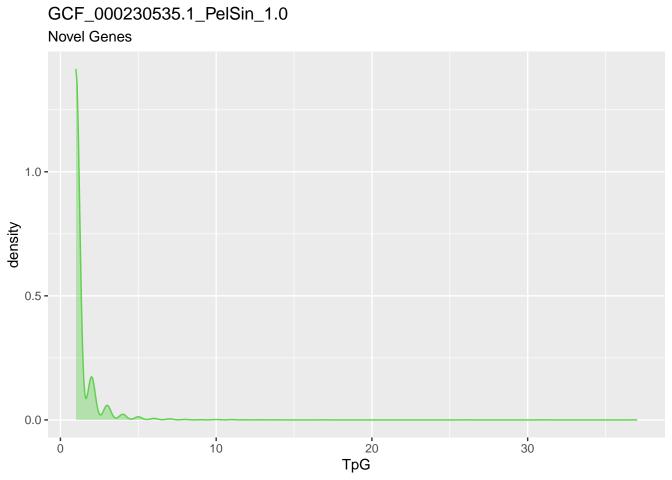


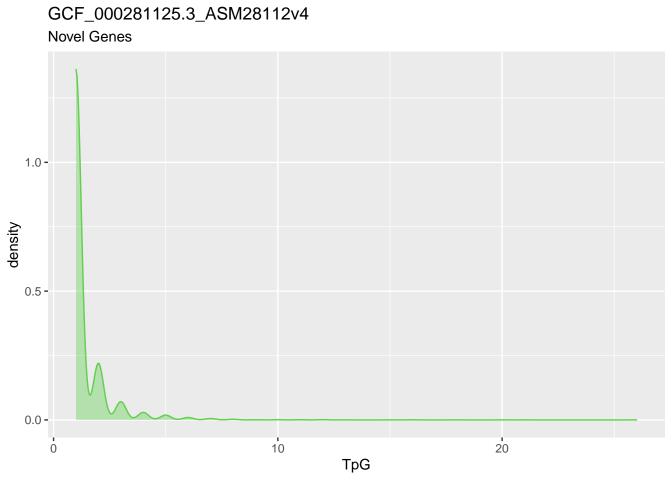


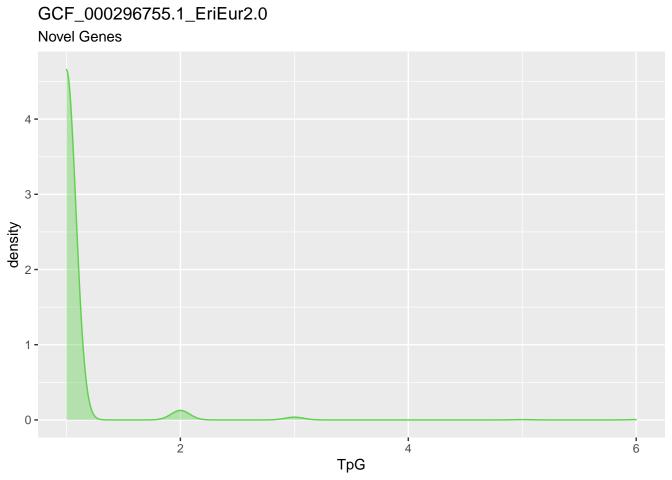


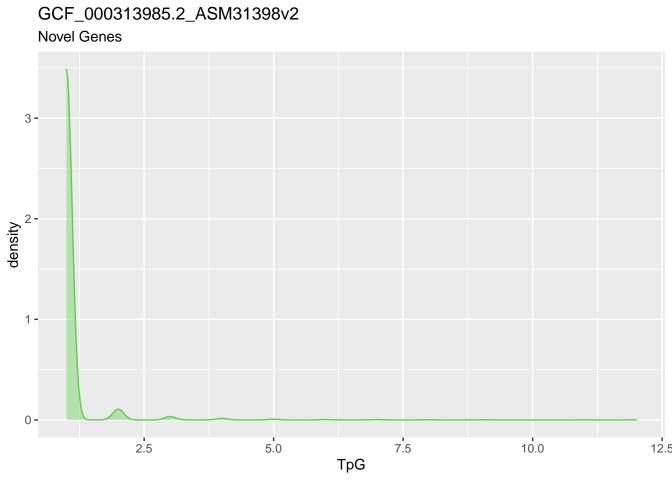


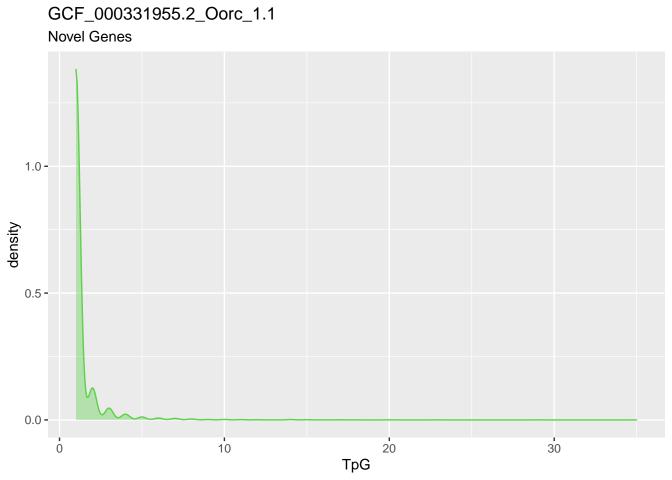


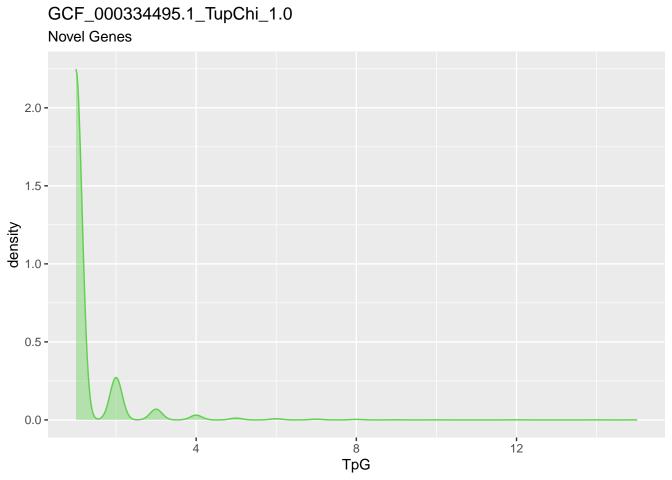


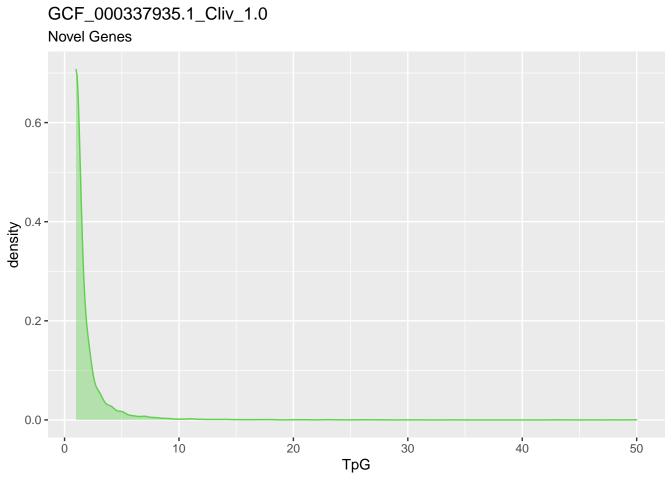


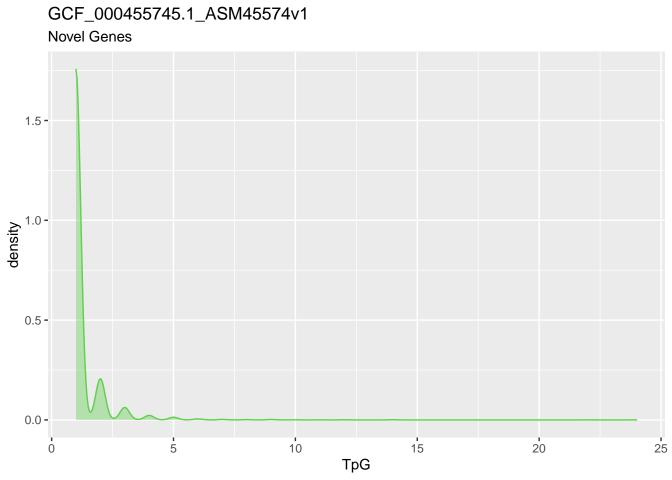


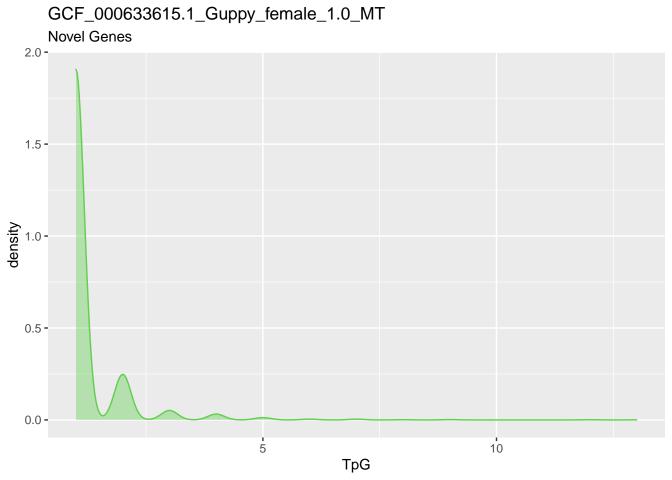


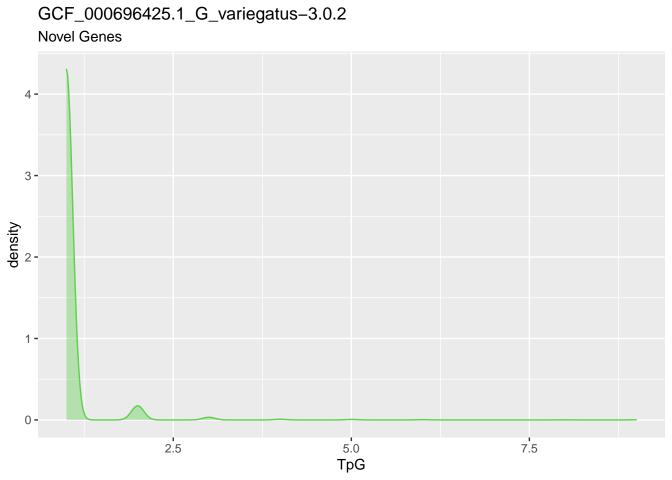


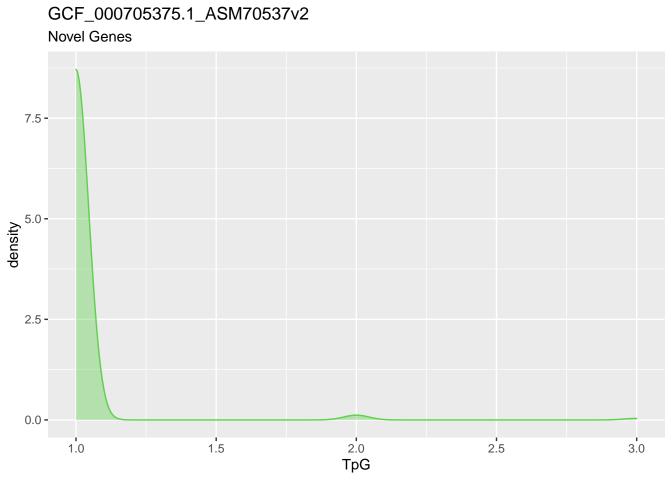


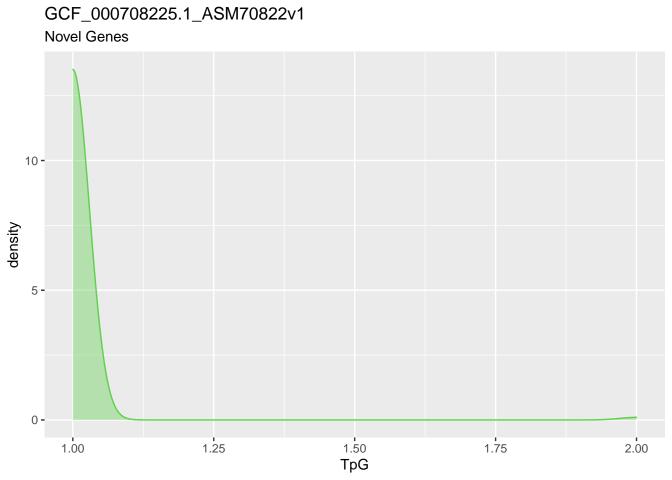


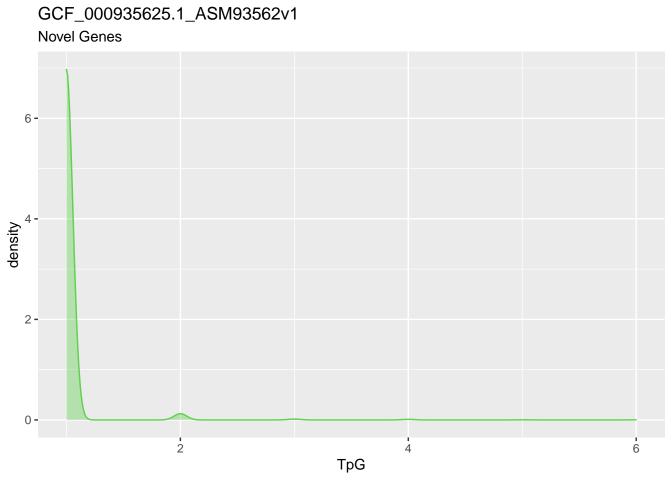


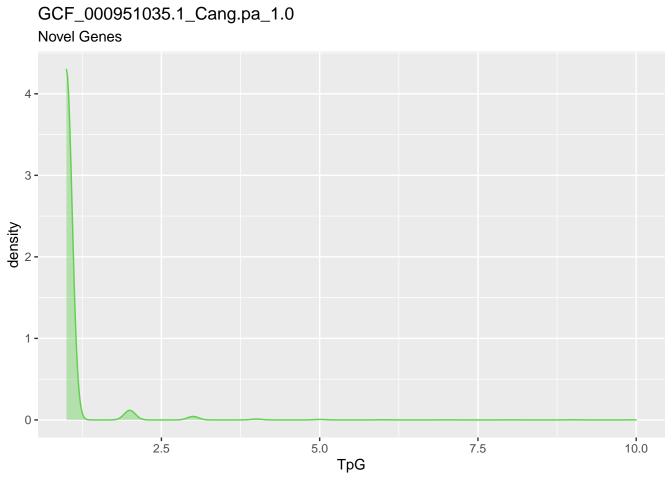


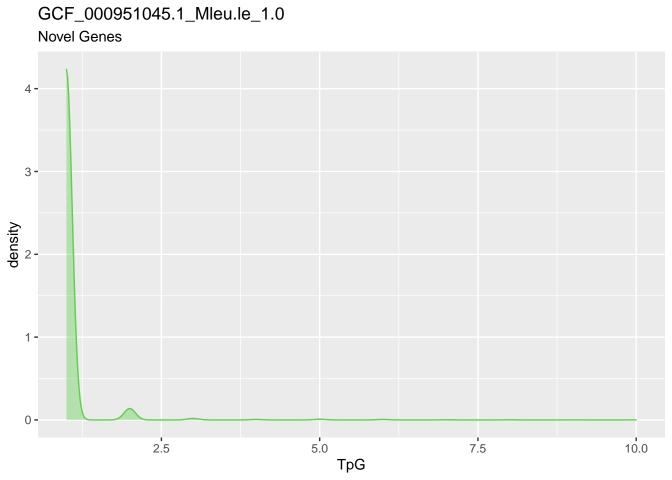


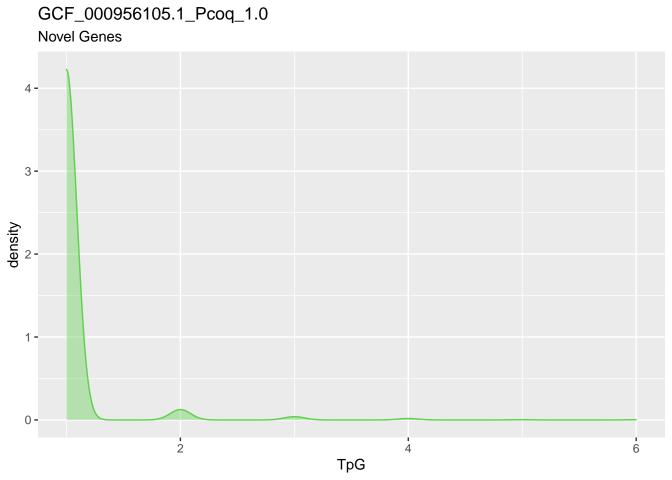


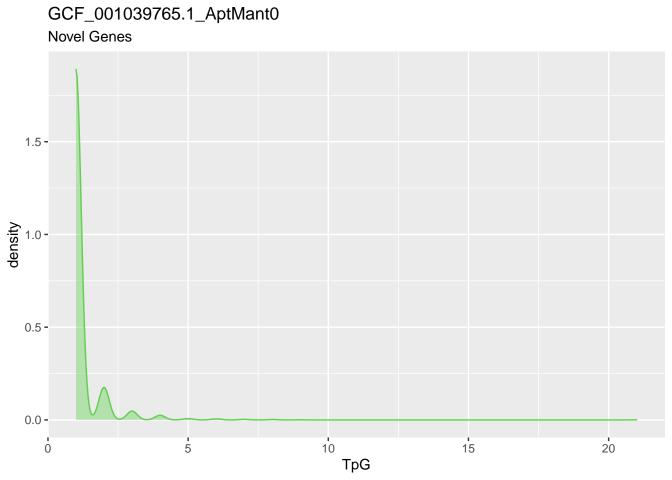


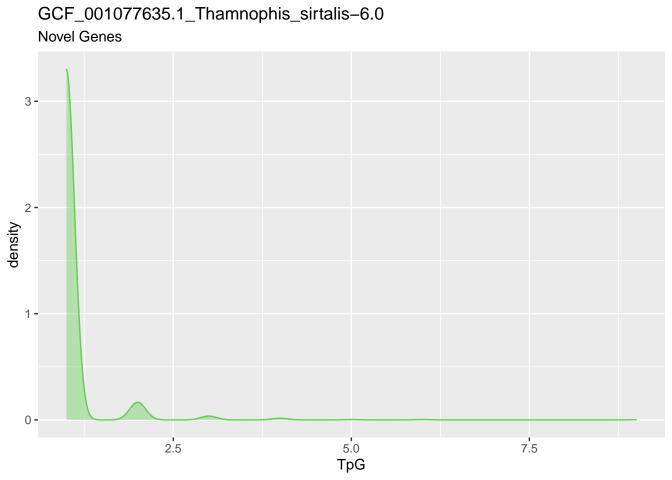


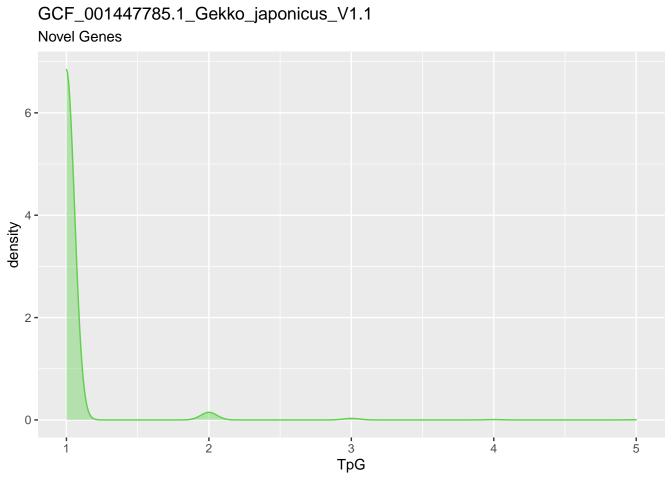




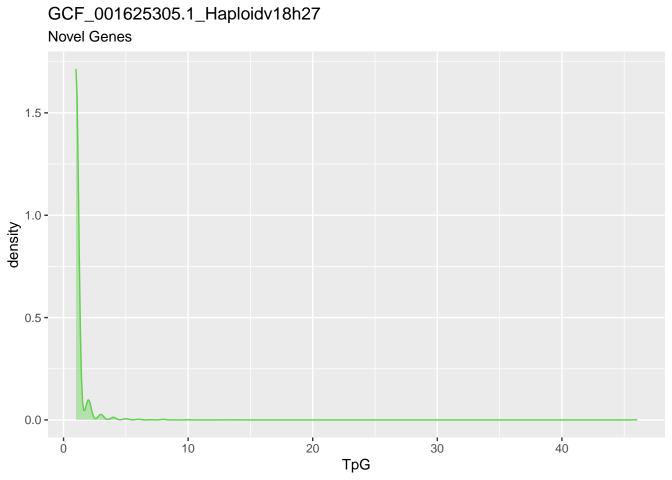


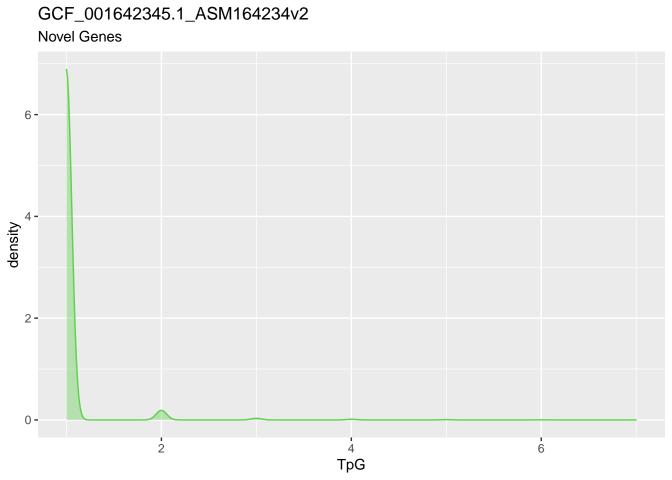


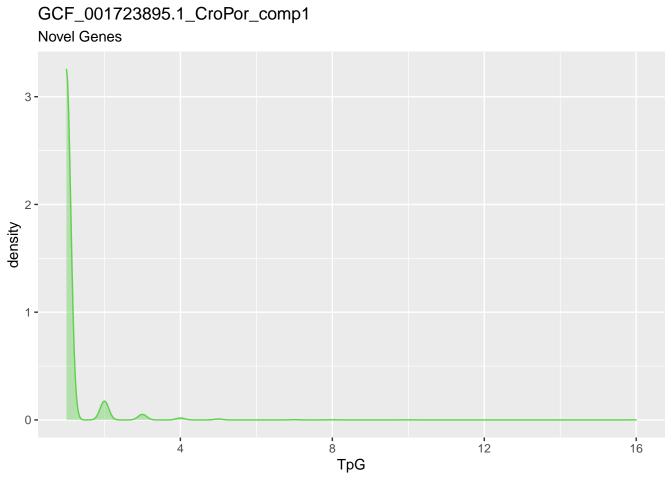


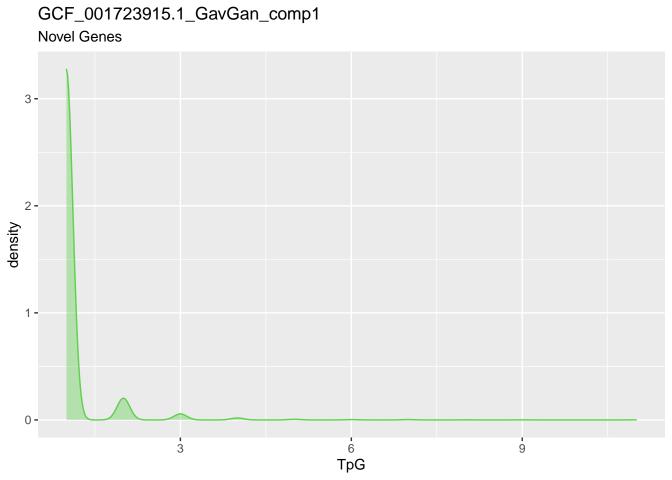


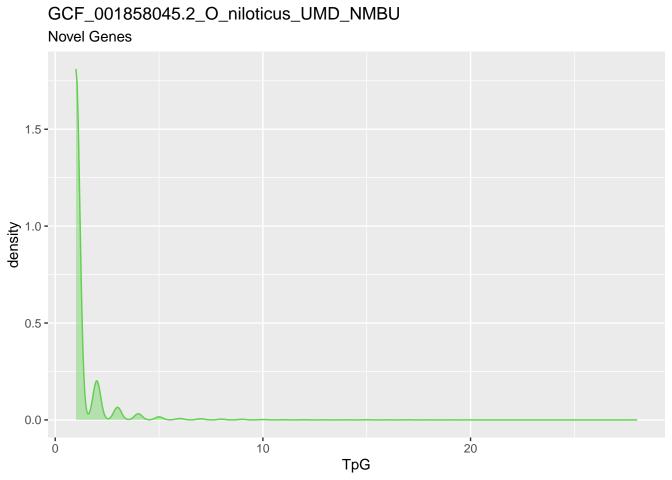
GCF_001522545.3_Parus_major1.1 **Novel Genes** 1.00 -0.75 density 0.50 -0.25 -0.00 -10 15 20 TpG

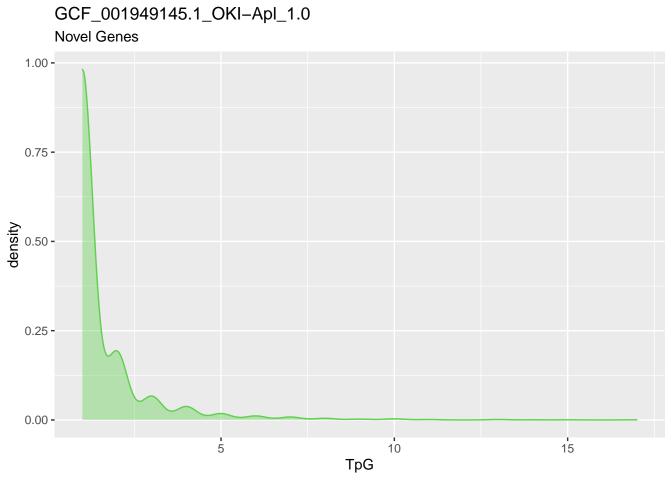


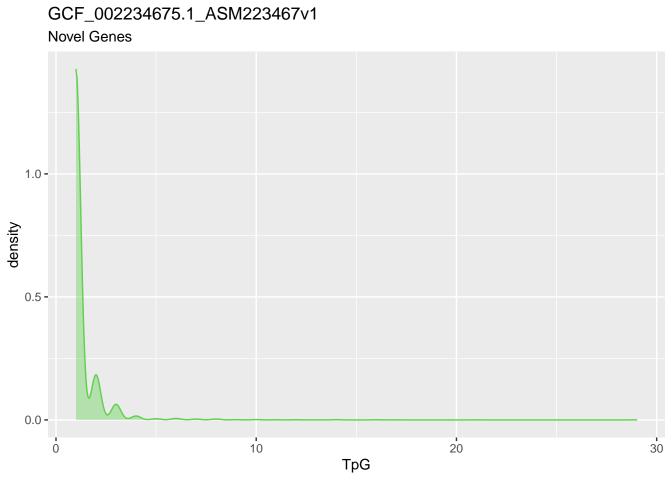


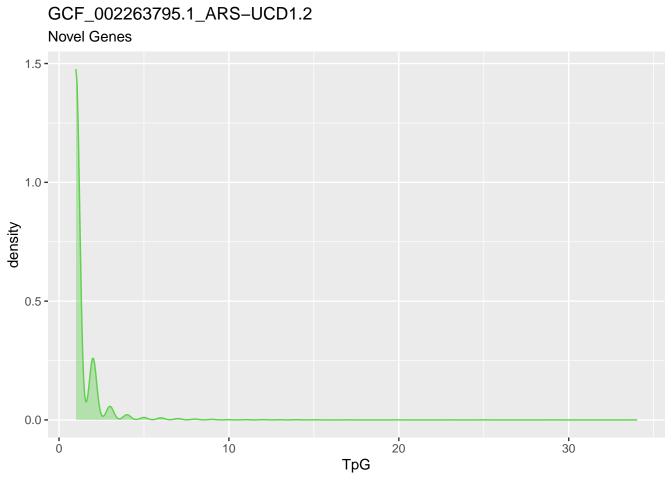


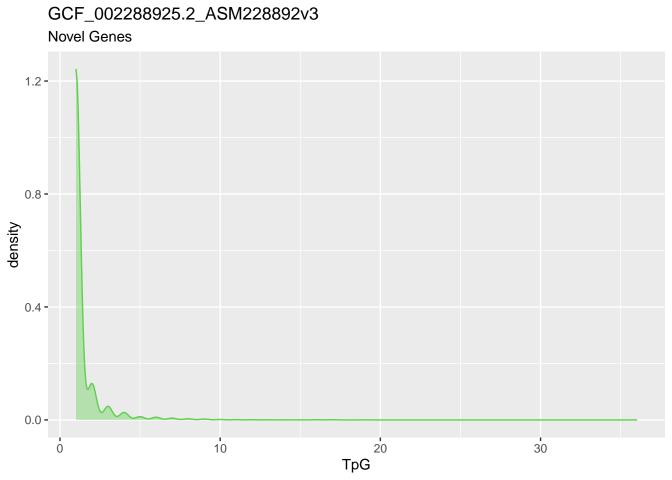




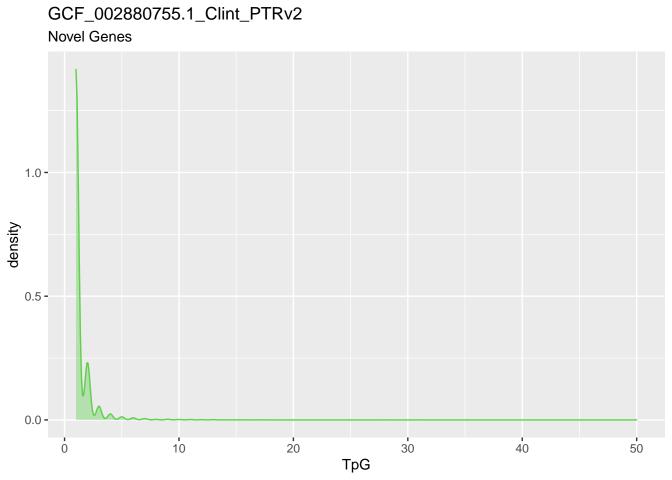


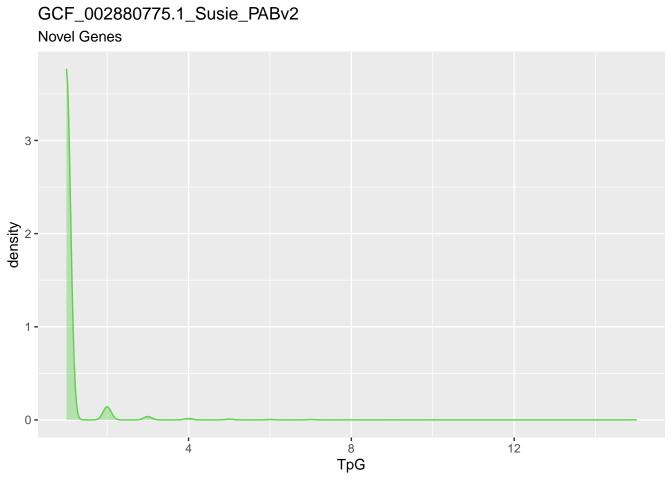


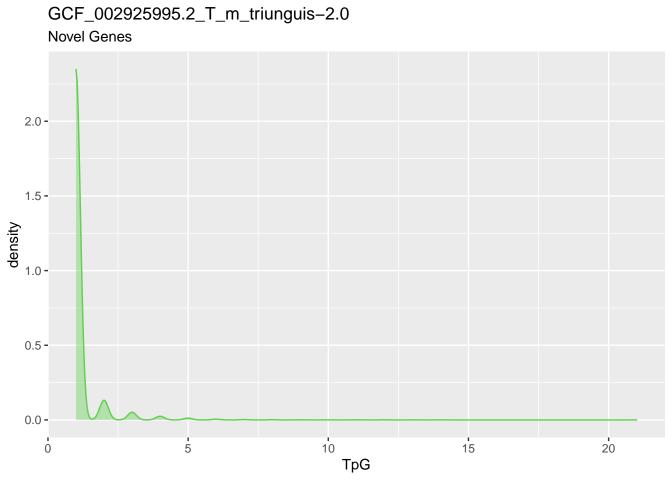


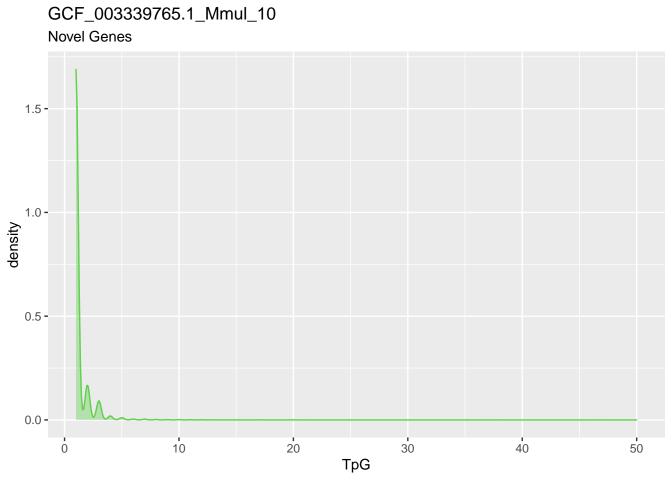


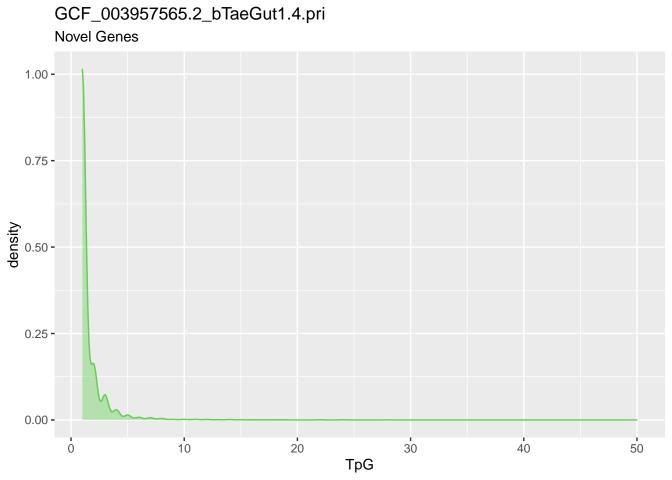
GCF_002863925.1_EquCab3.0 **Novel Genes** 1.00 -0.75 density 0.25 -0.00 -10 20 30 50 40 TpG

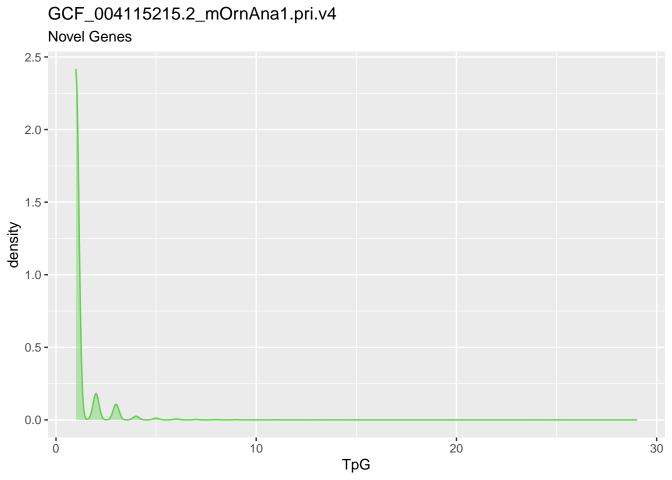


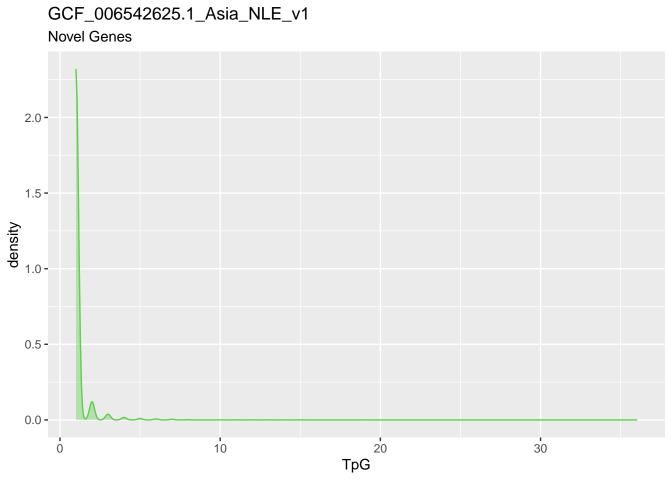


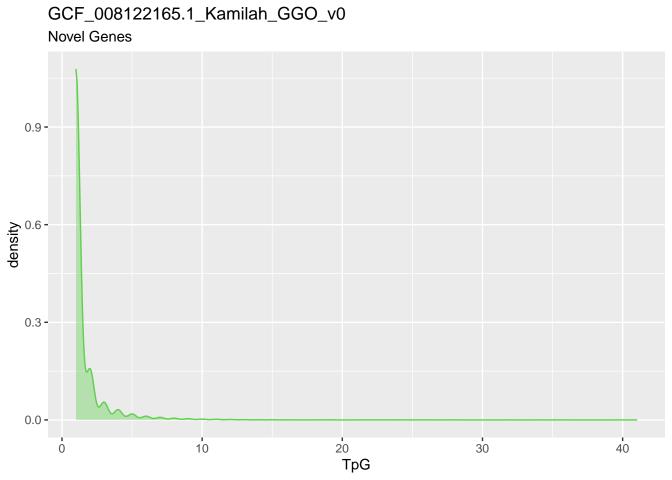




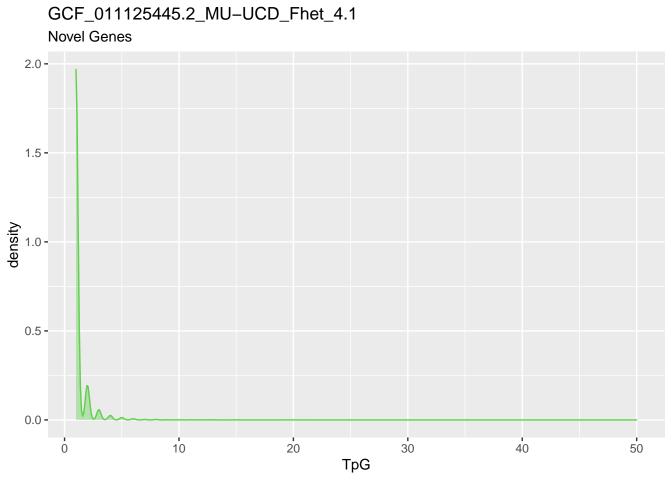


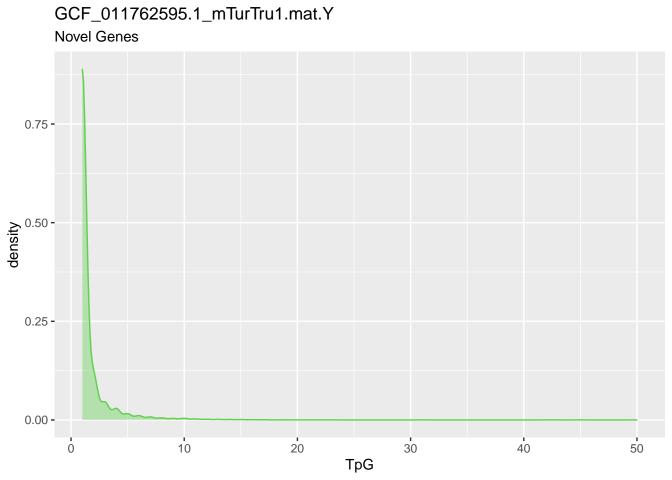


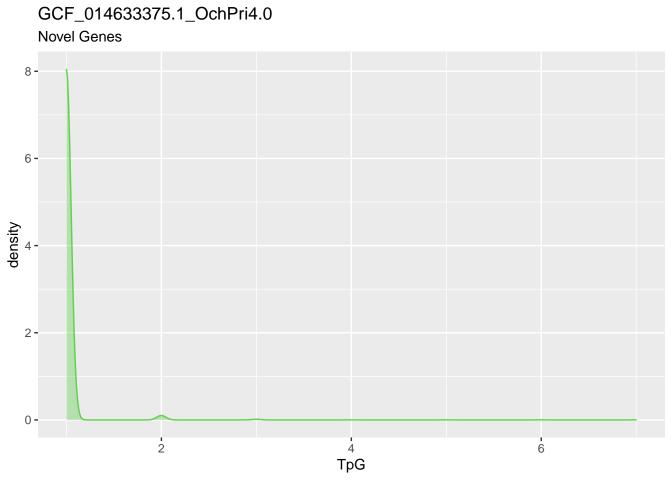




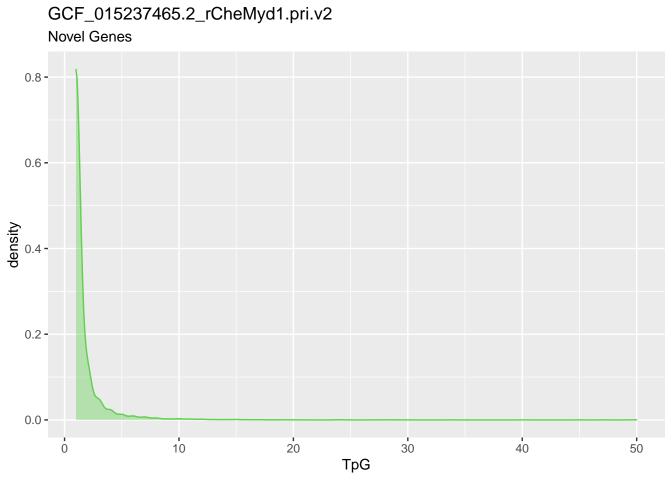
GCF_009663435.1_Callithrix_jacchus_cj1700_1.1 **Novel Genes** 0.75 density 0.25 -0.00 -10 20 30 40 50 TpG

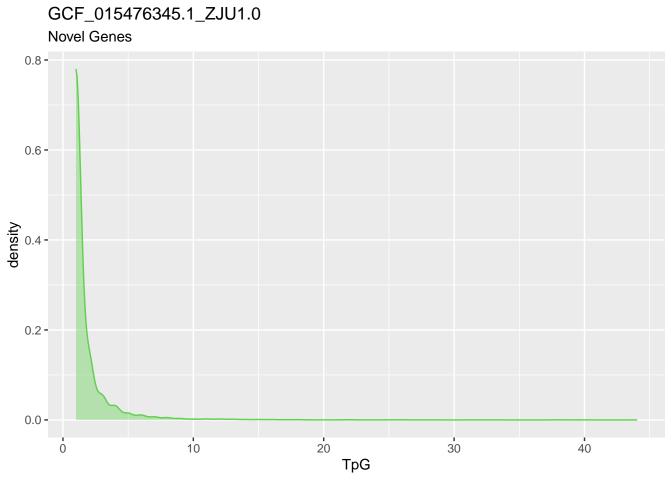






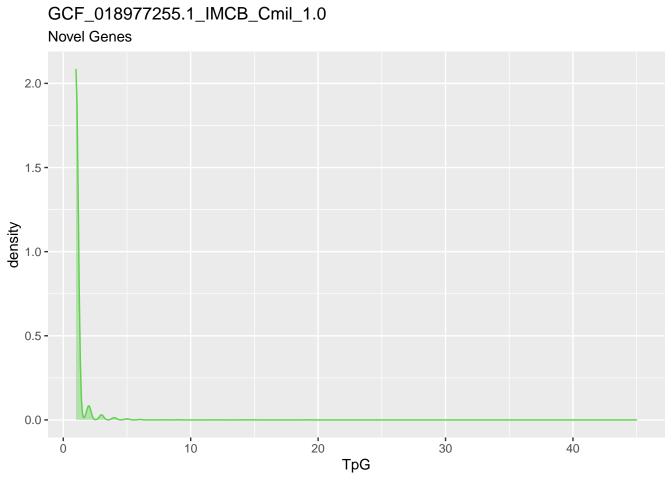
GCF_015227675.2_mRatBN7.2 **Novel Genes** 0.8 -0.6 density 0.2 -0.0 -10 20 30 40 50 TpG

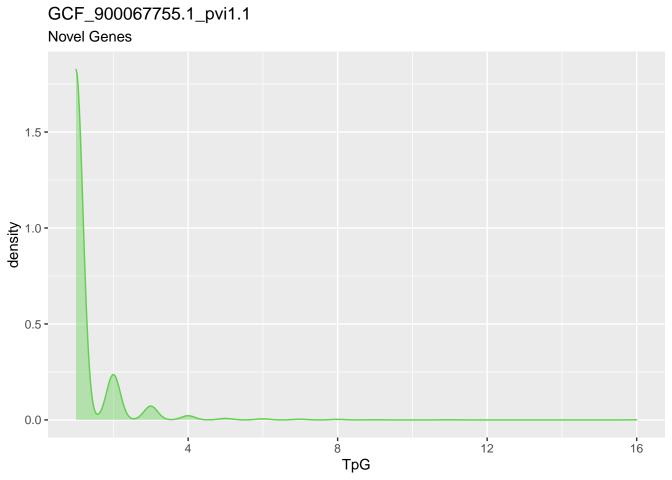




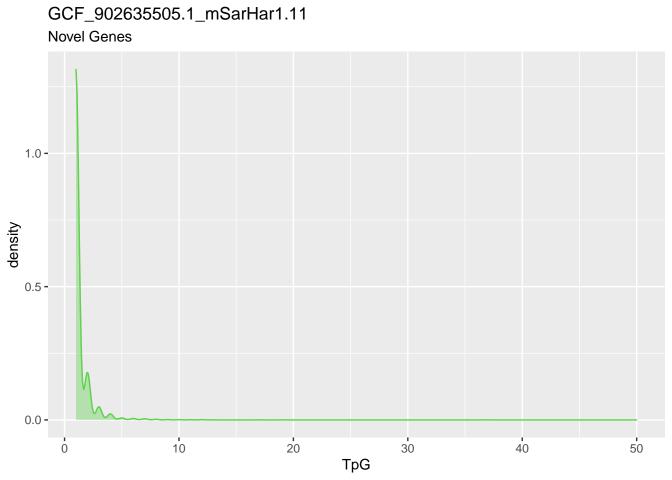
GCF_016699485.2_bGalGal1.mat.broiler.GRCg7b **Novel Genes** 0.4 density 0.2 -0.0 -10 20 30 40 50

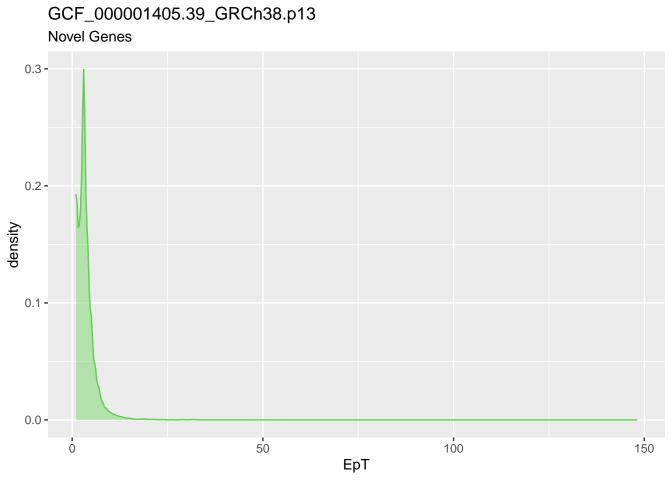
TpG

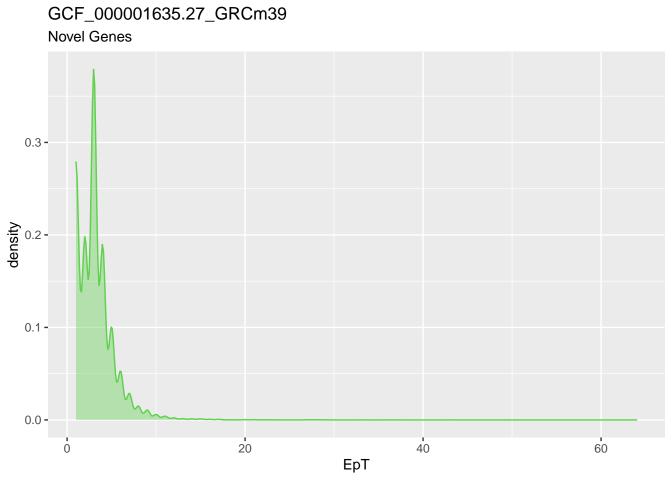


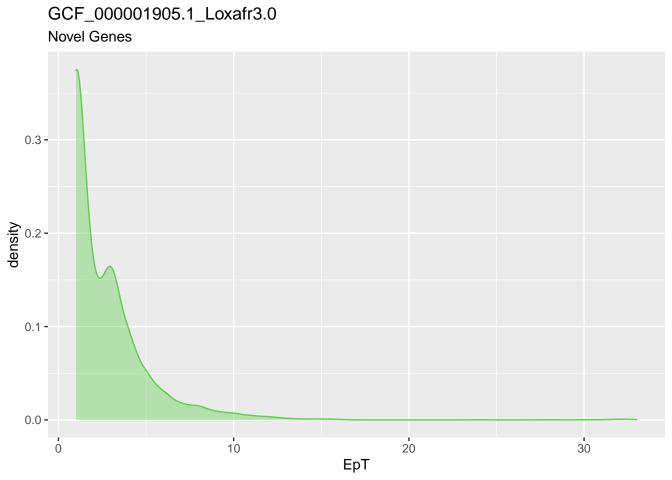


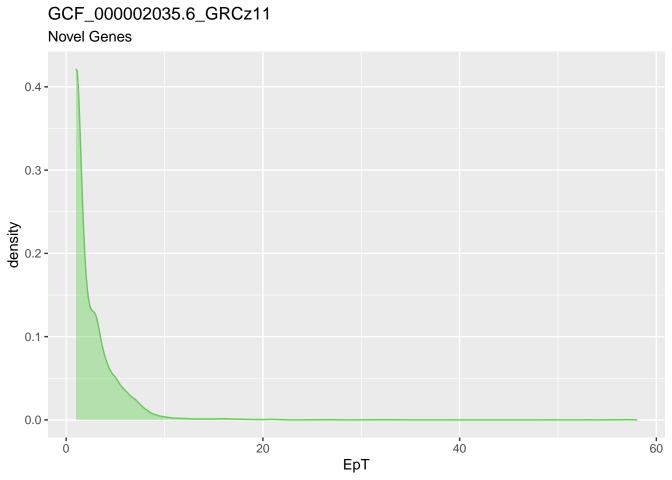
GCF_901000725.2_fTakRub1.2 **Novel Genes** 1.25 -1.00 -0.75 density 0.50 -0.25 -0.00 -10 20 30 40 TpG

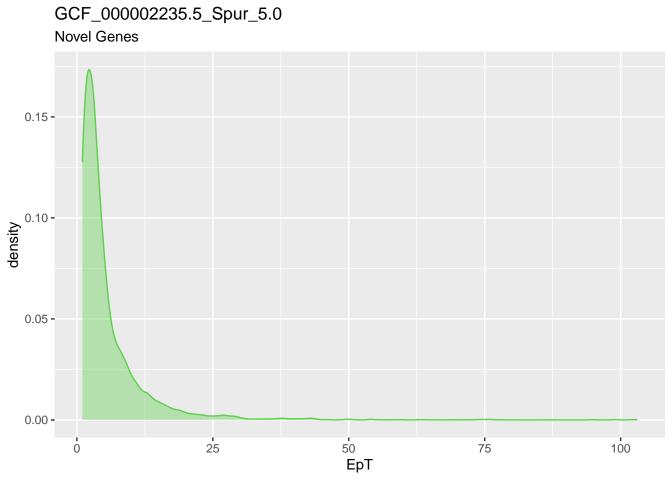


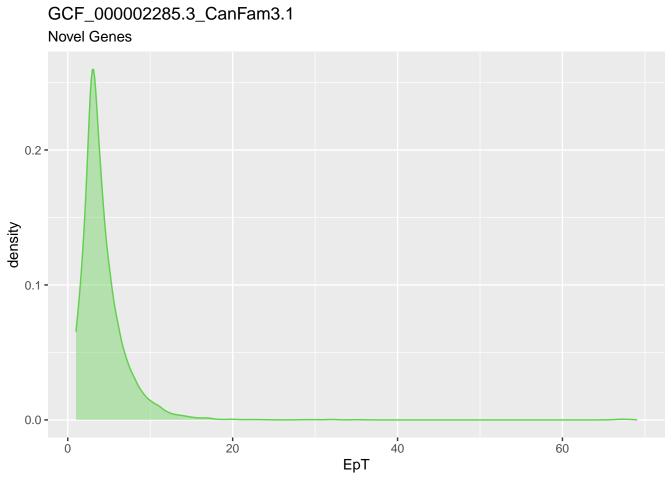


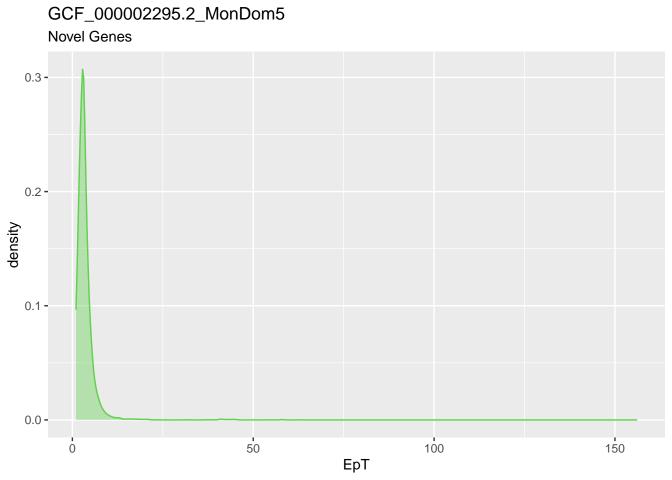


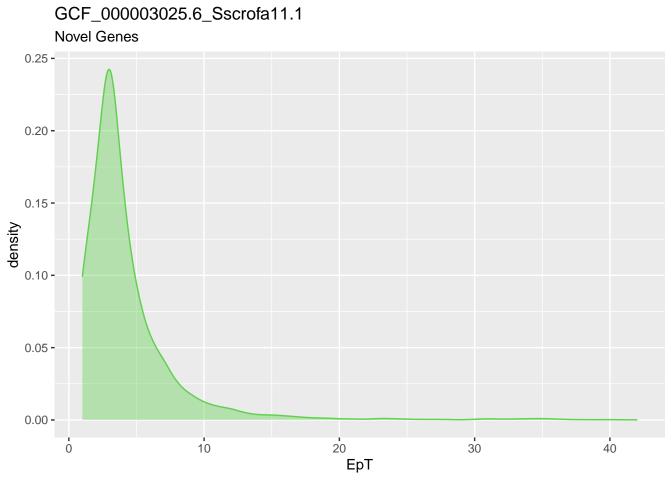


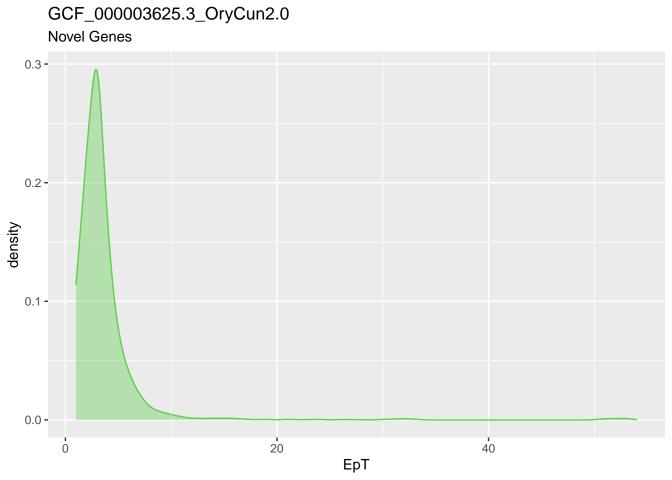


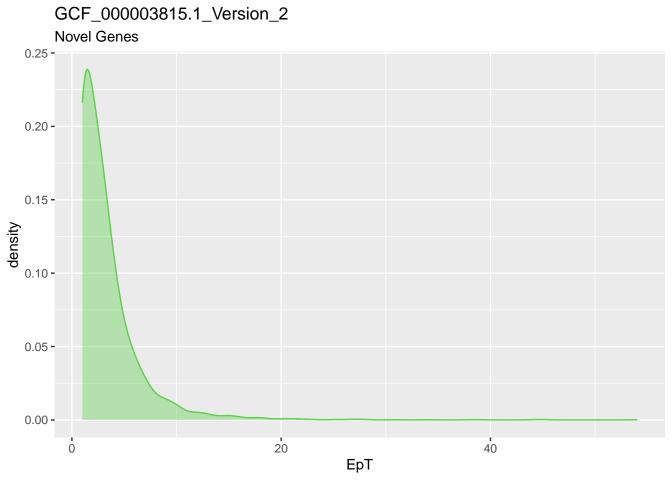


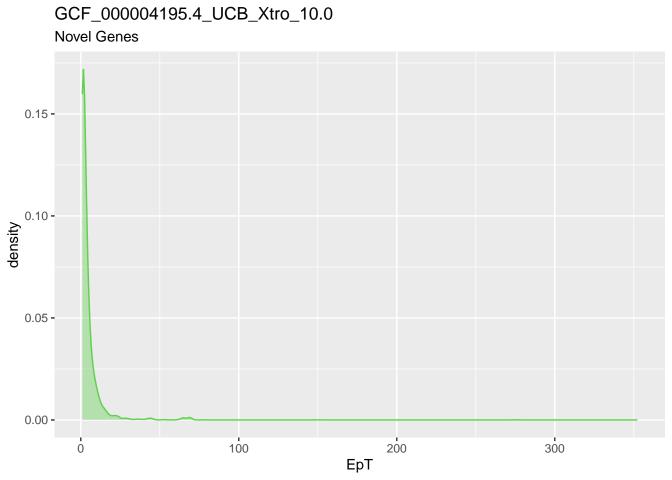


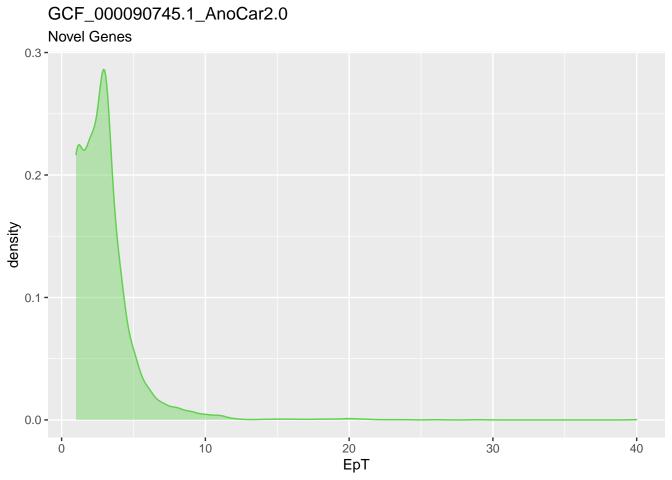




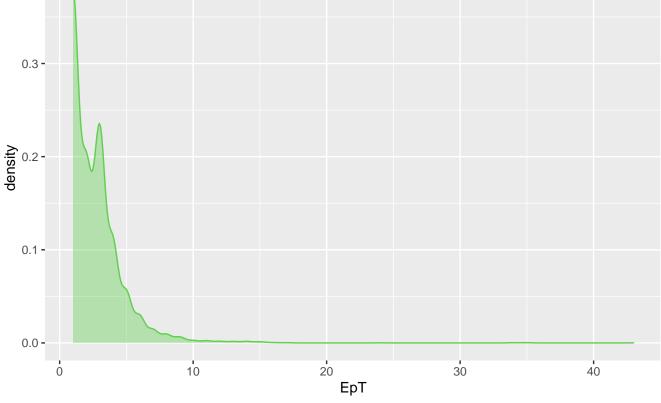


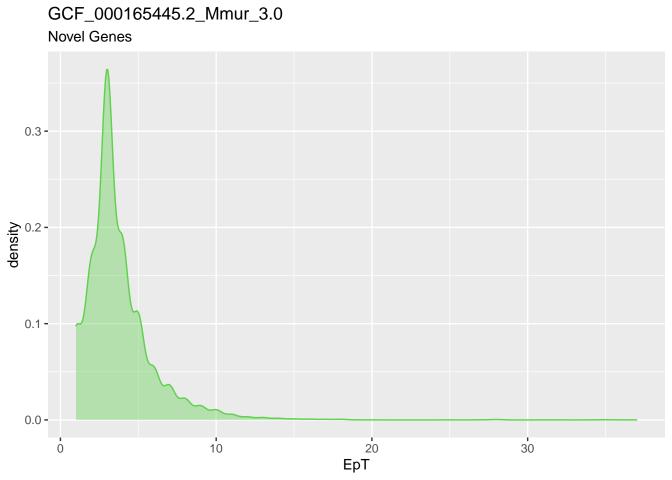


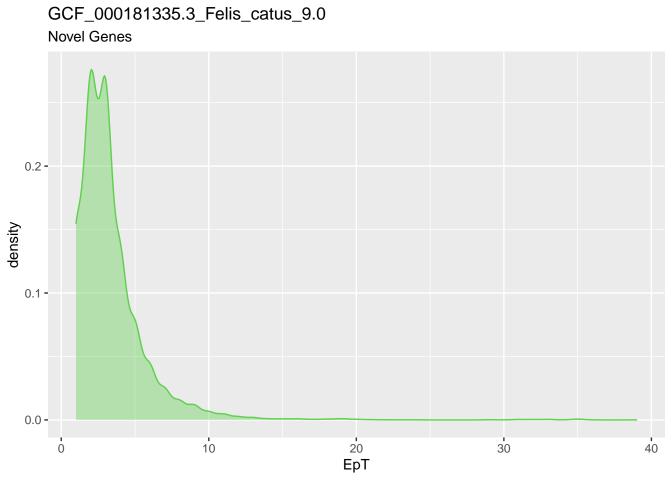


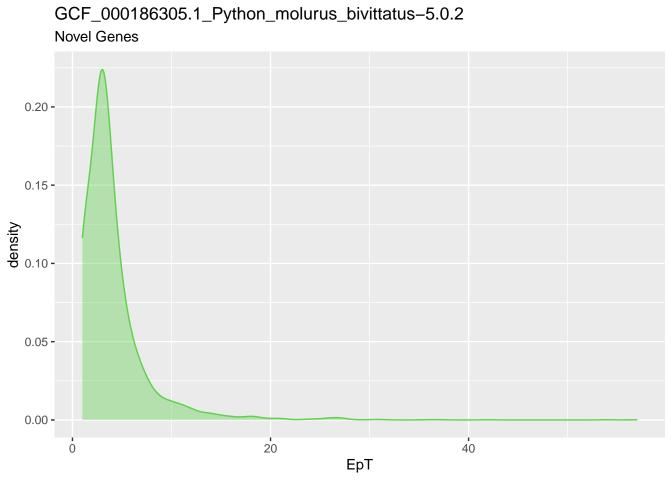


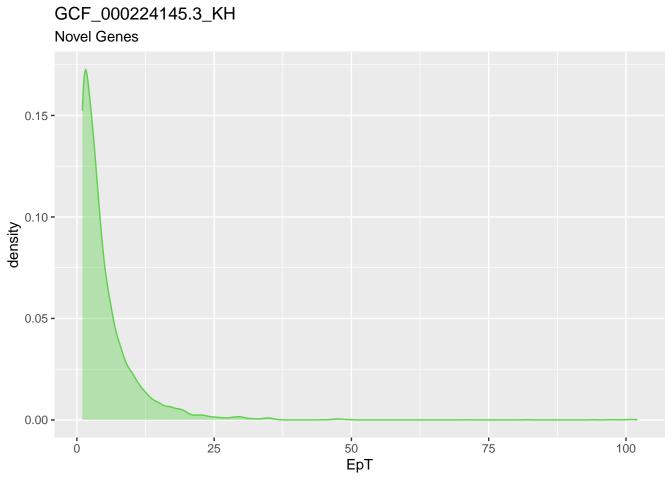
GCF_000151735.1_Cavpor3.0 **Novel Genes** 0.3 density 0.1 -

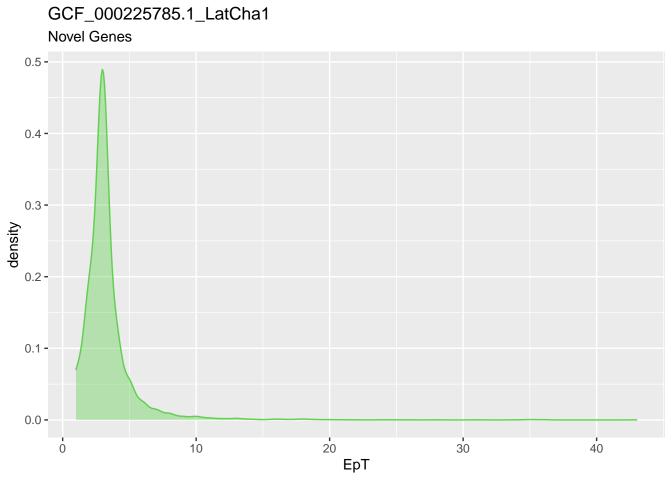


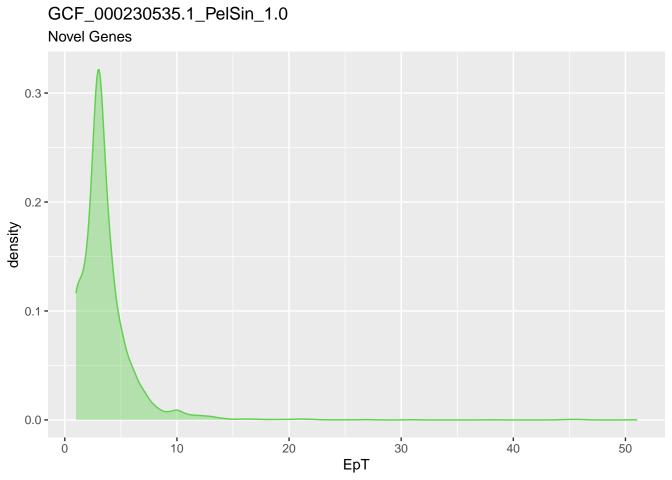


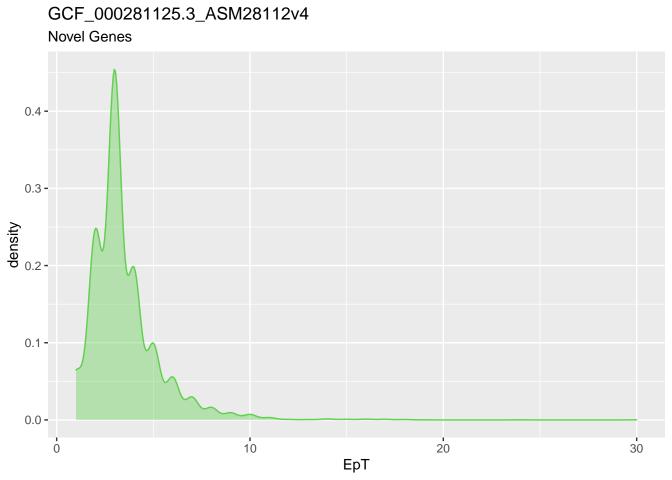


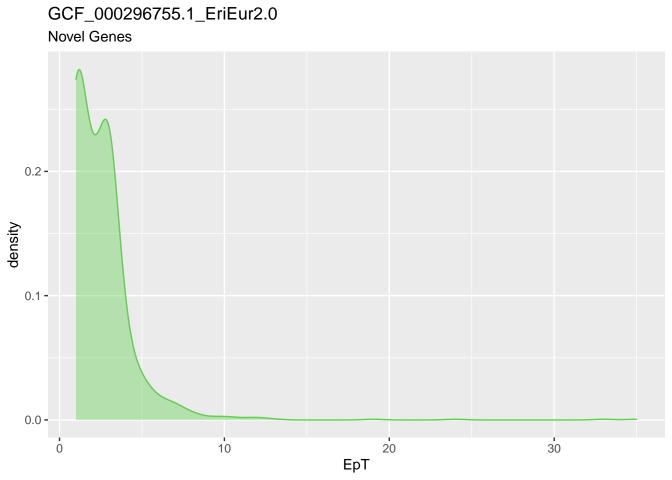


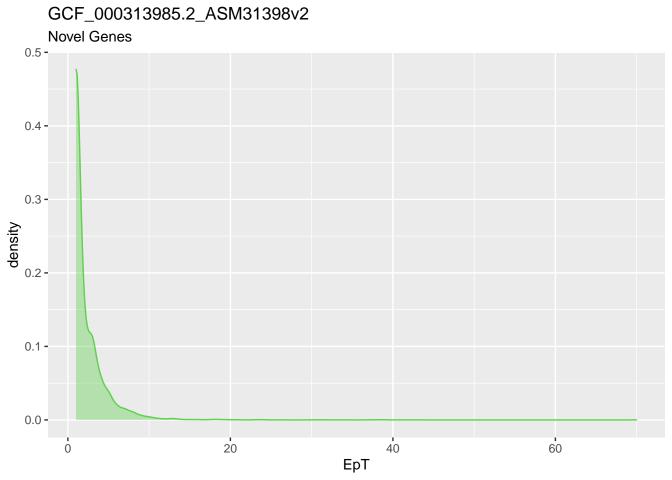


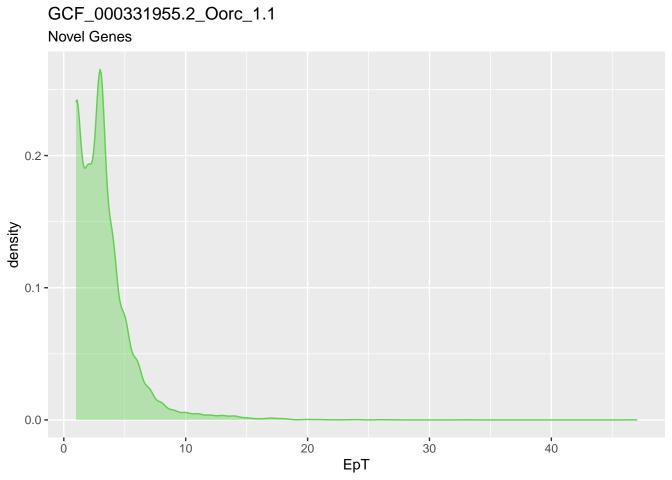


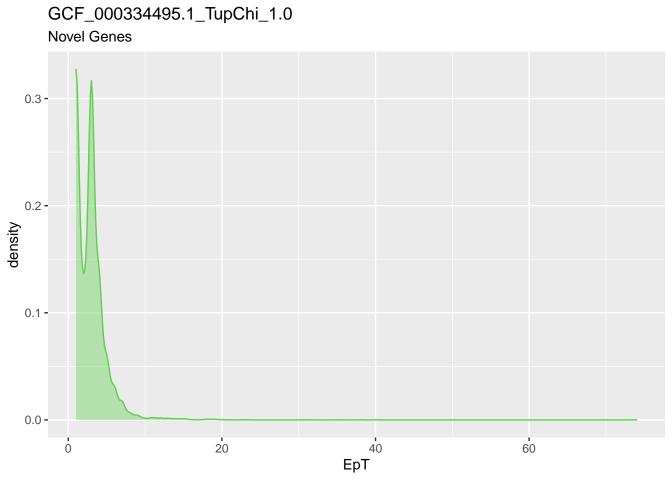


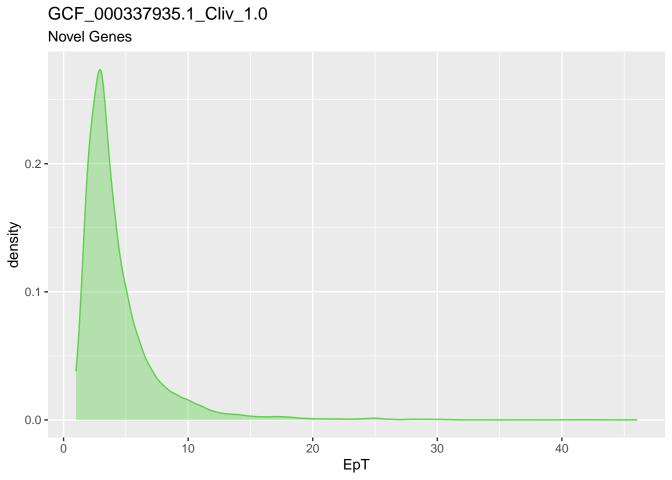


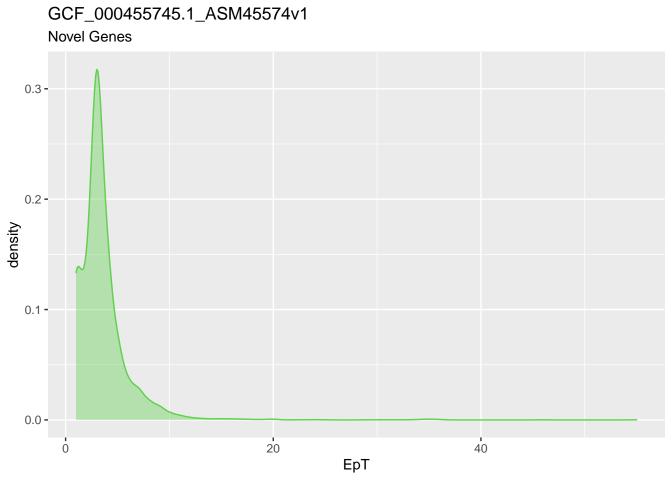


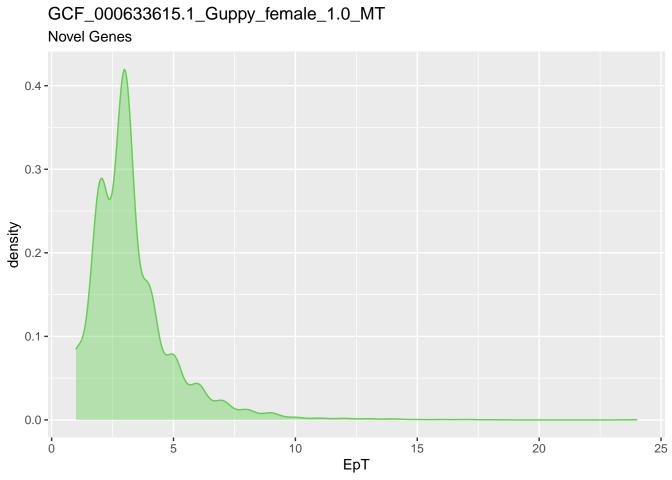


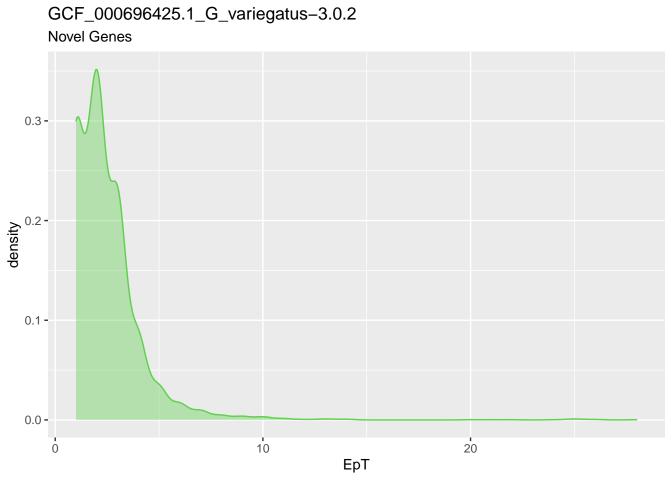


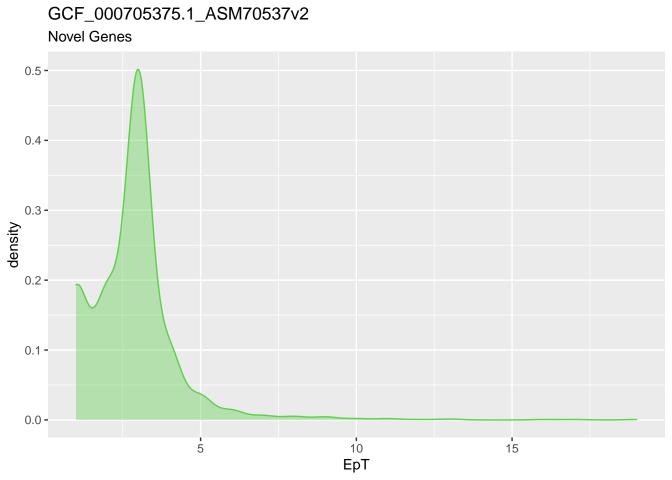


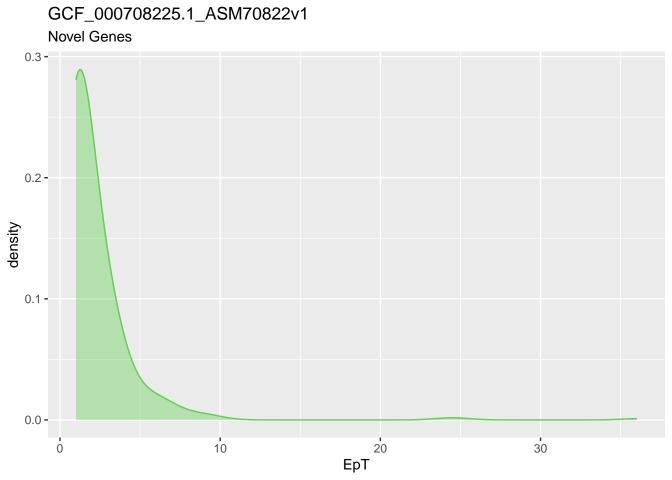


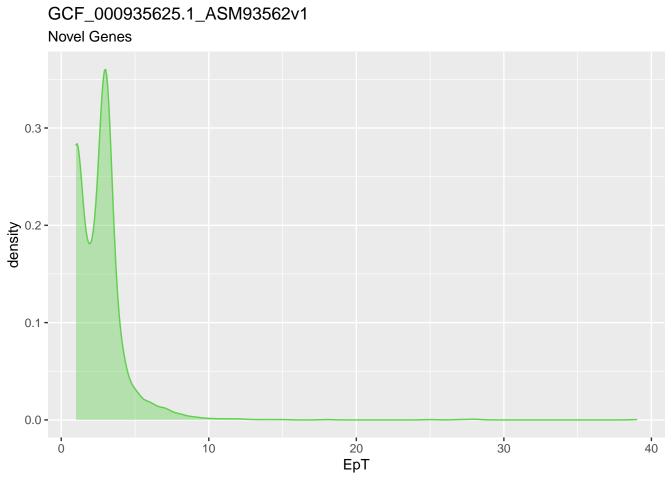


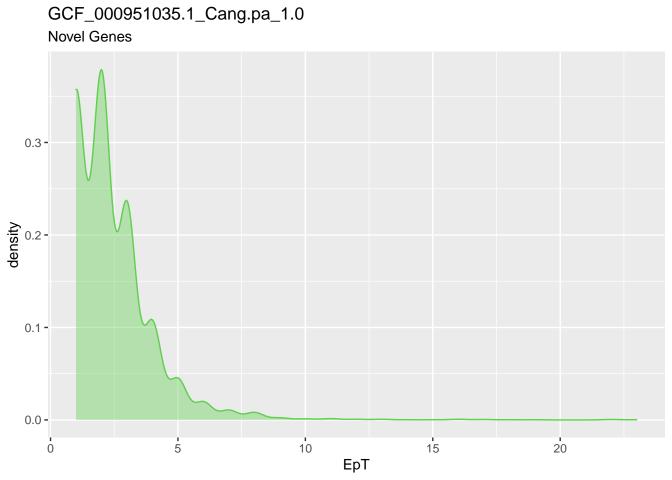


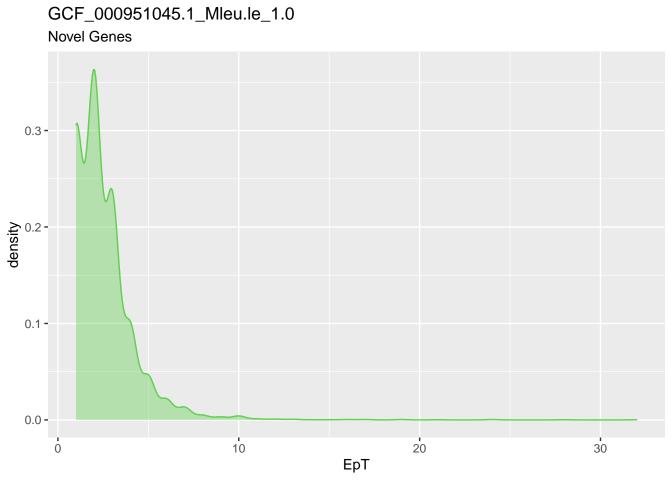


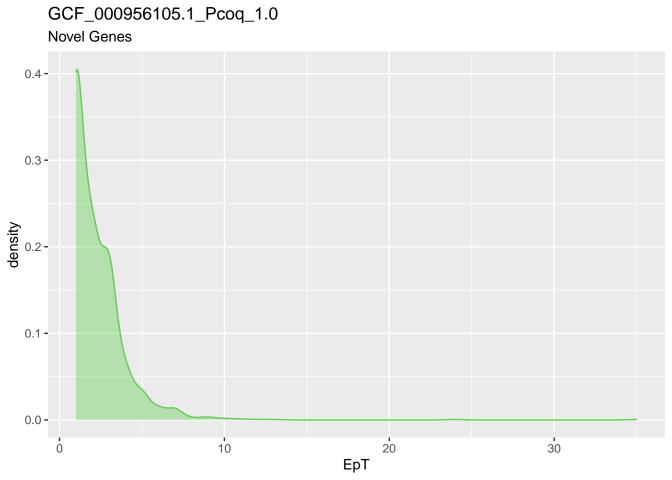


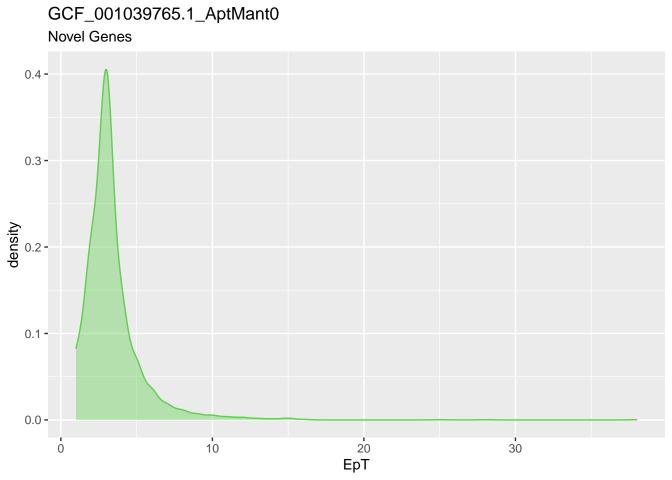


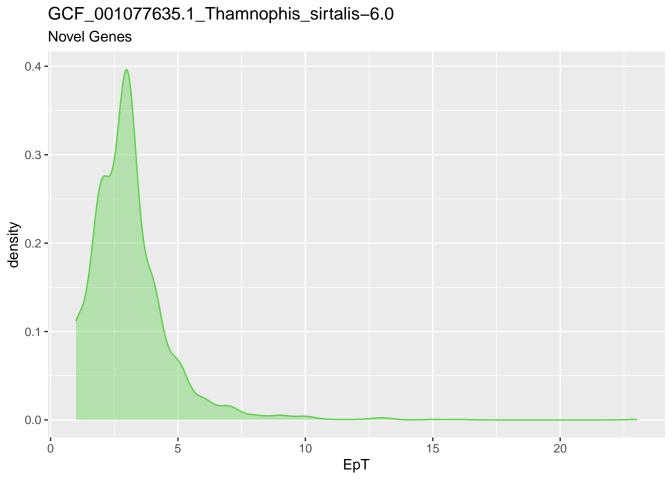


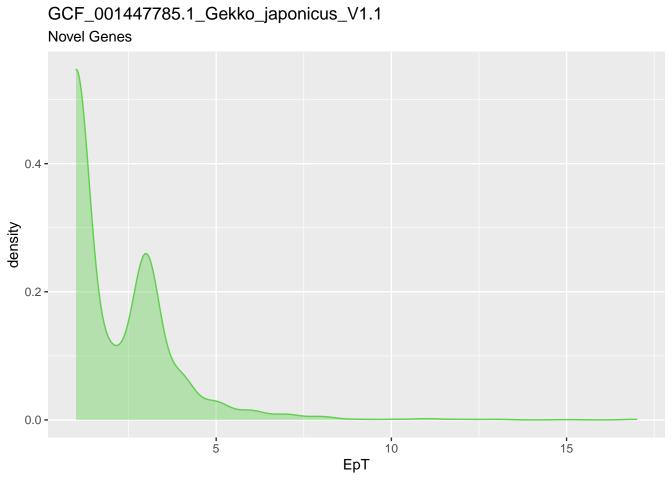


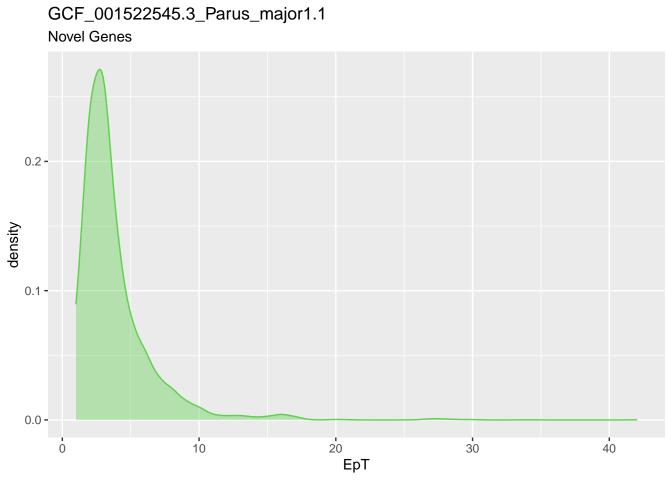


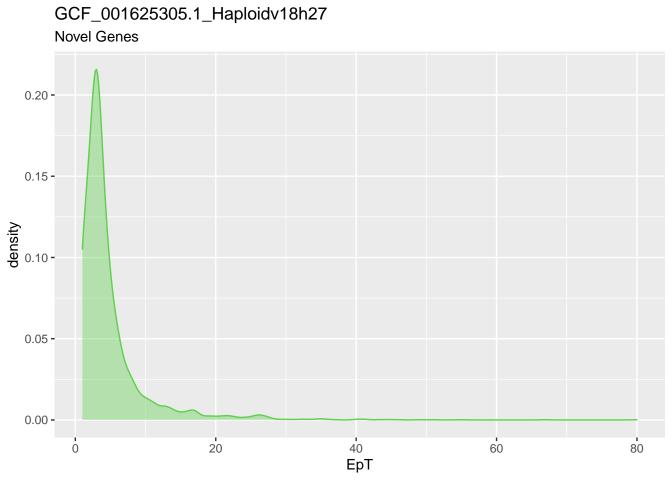


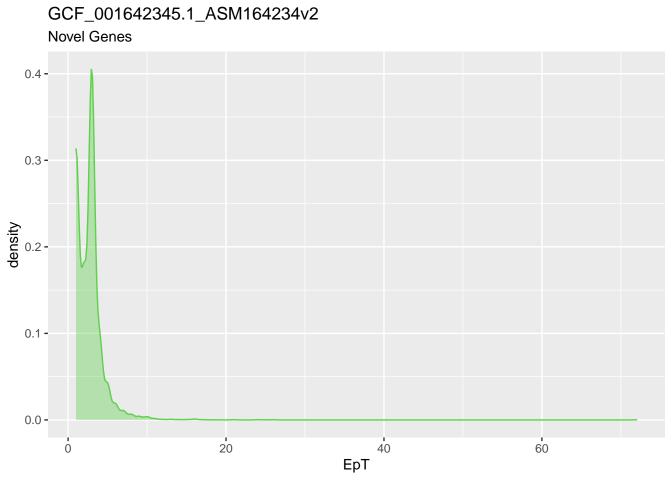


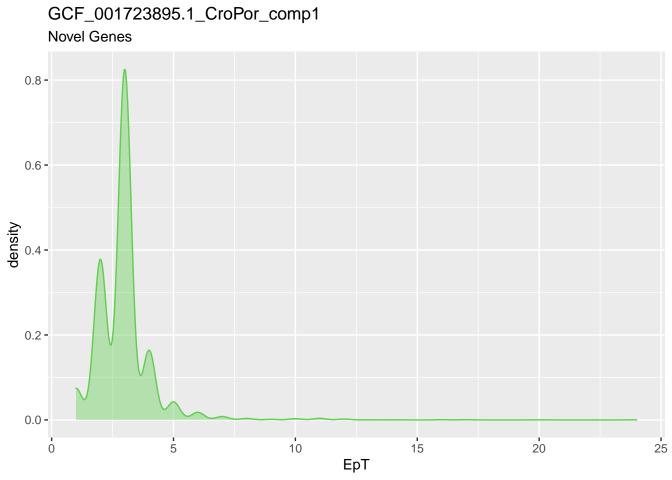


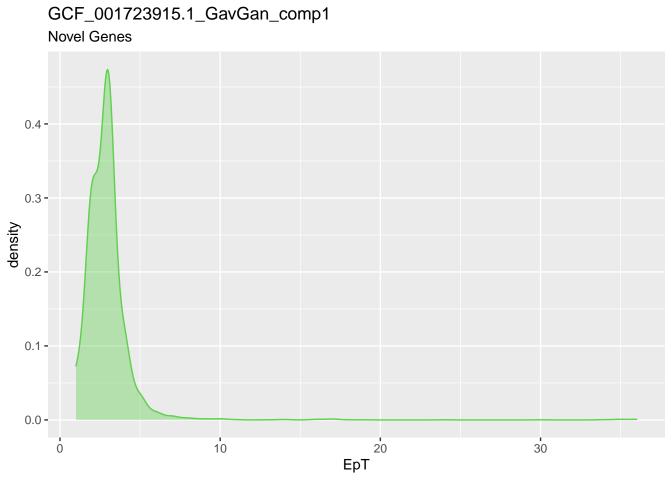


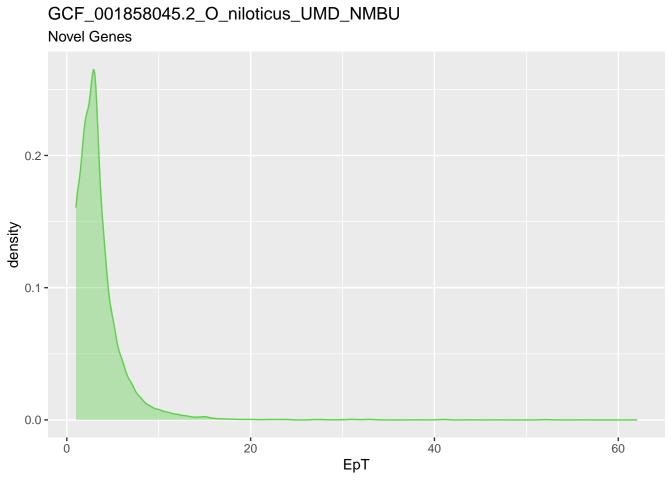


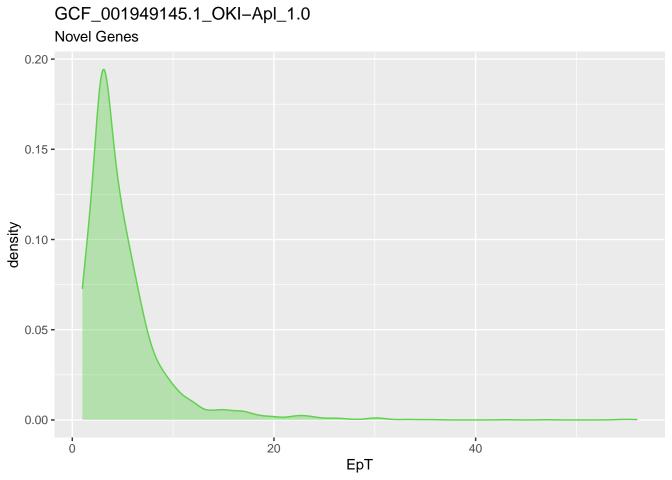




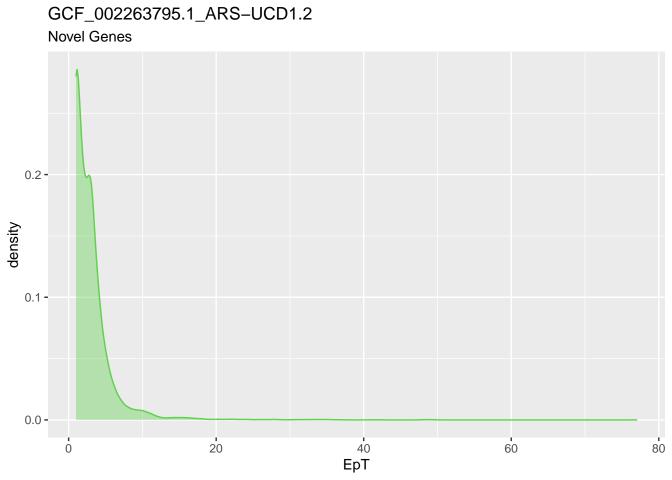


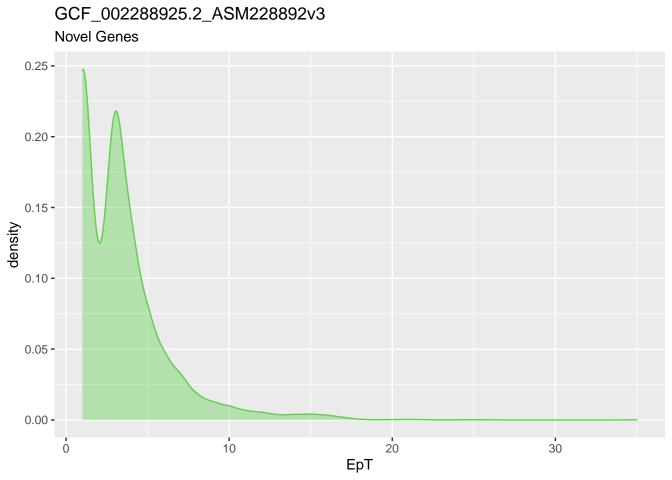


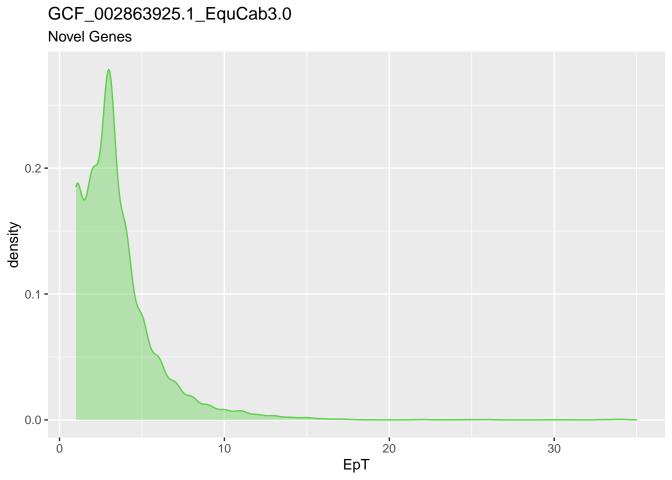


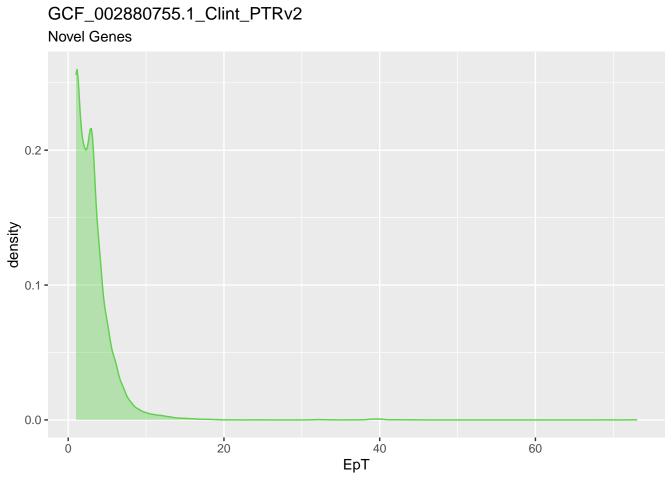


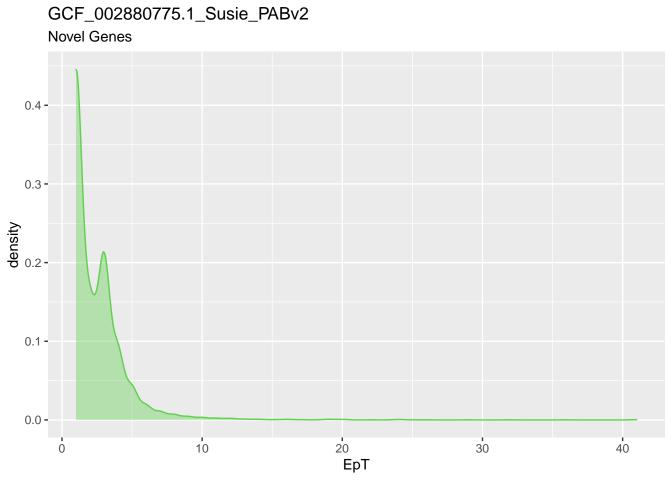
GCF_002234675.1_ASM223467v1 **Novel Genes** 0.25 -0.20 -0.15 density 0.10 -0.05 -0.00 -10 20 30 40 50 EpT

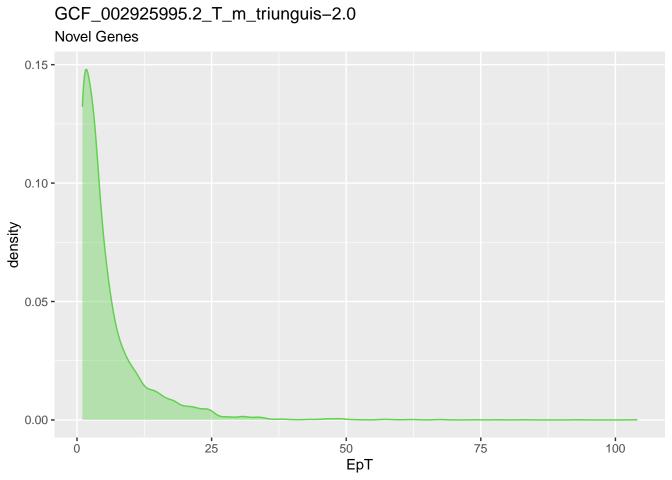


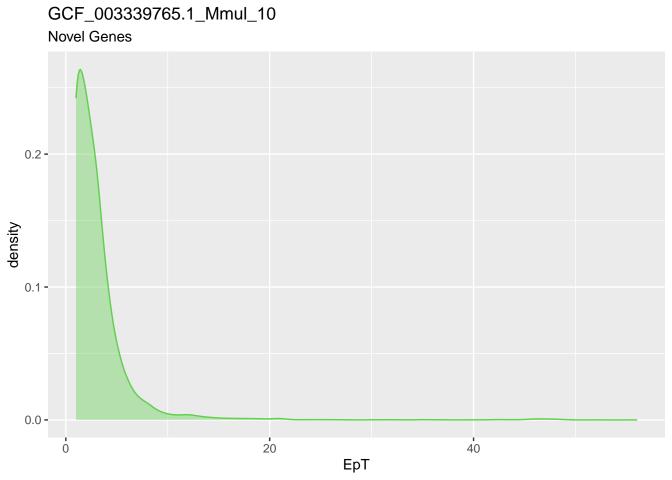


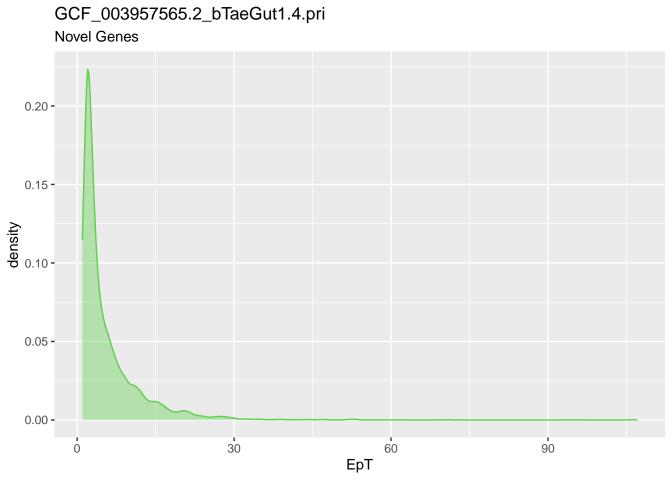


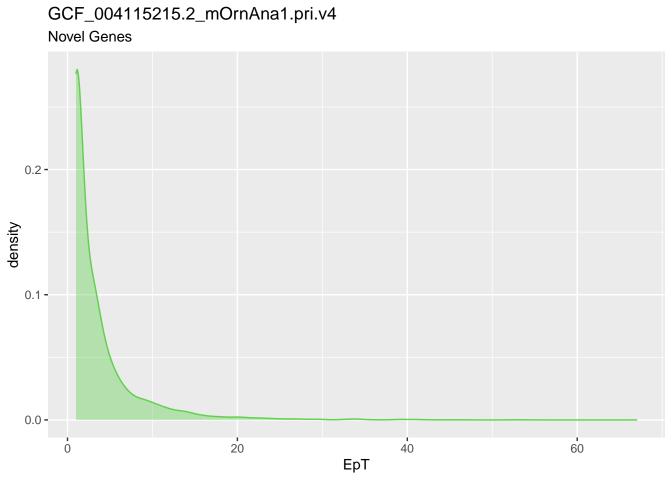


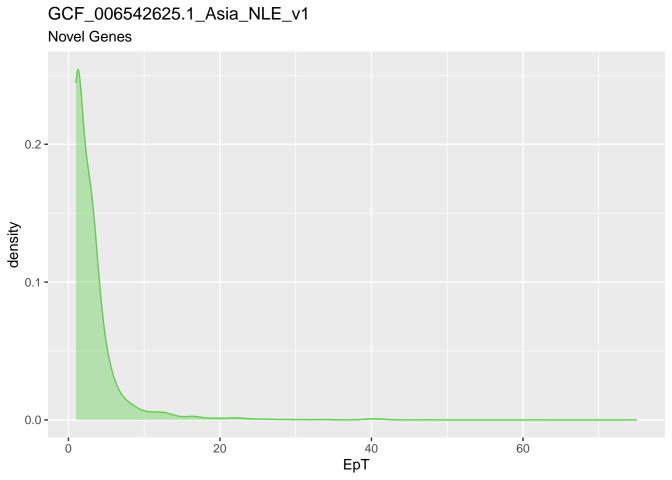


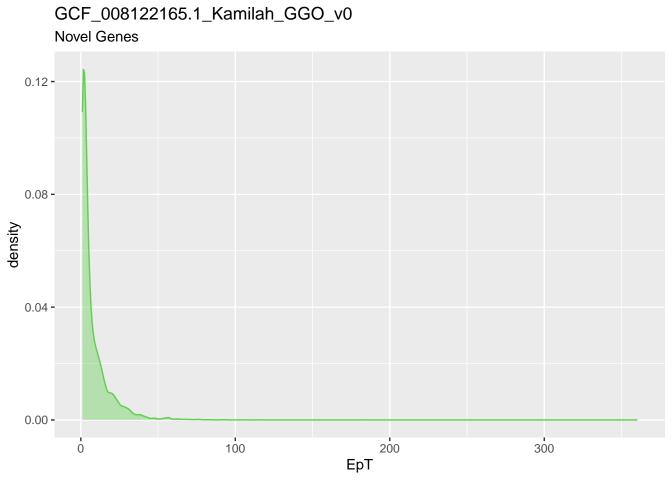




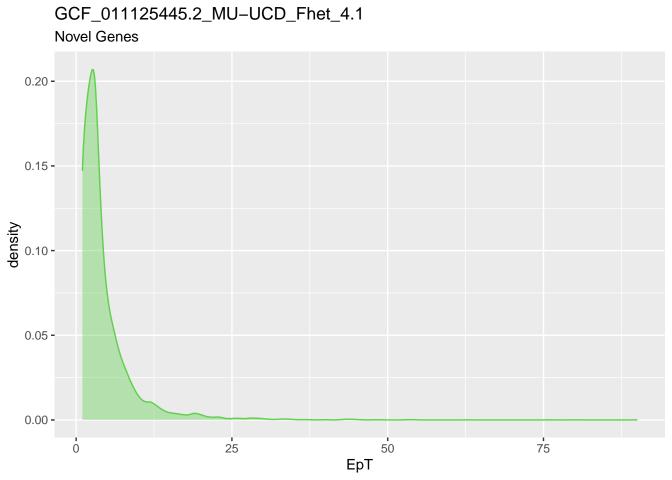




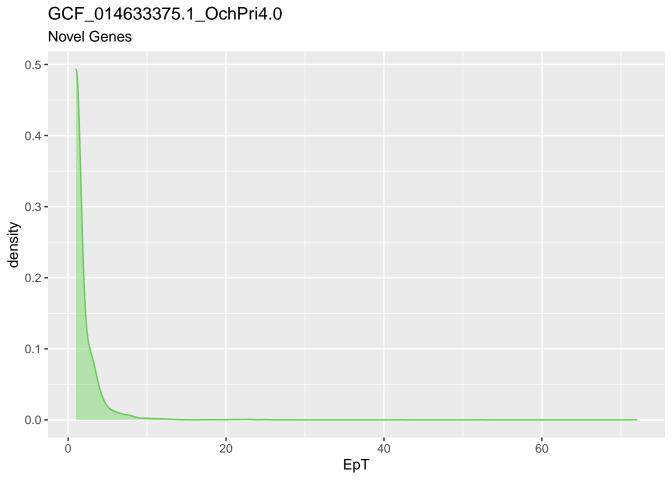


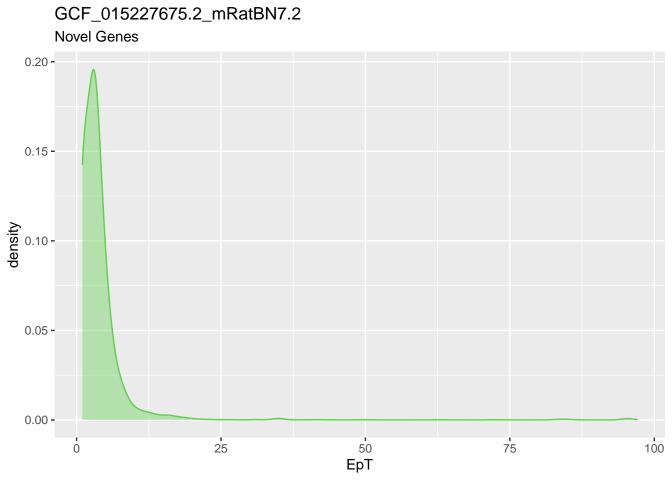


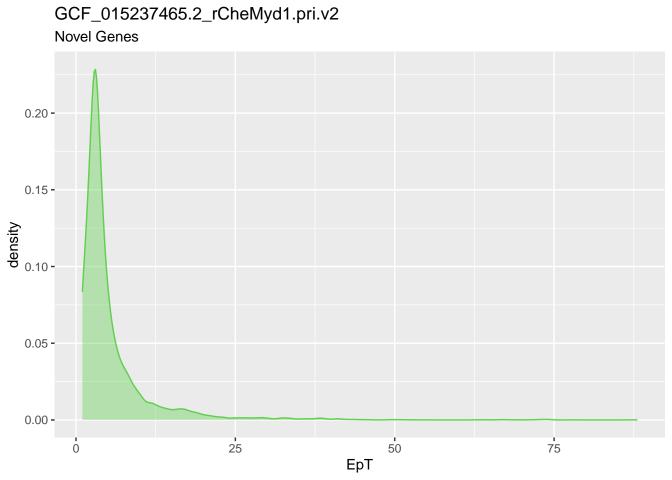
GCF_009663435.1_Callithrix_jacchus_cj1700_1.1 **Novel Genes** 0.15 density 0.05 -0.00 -20 40 60 80 EpT

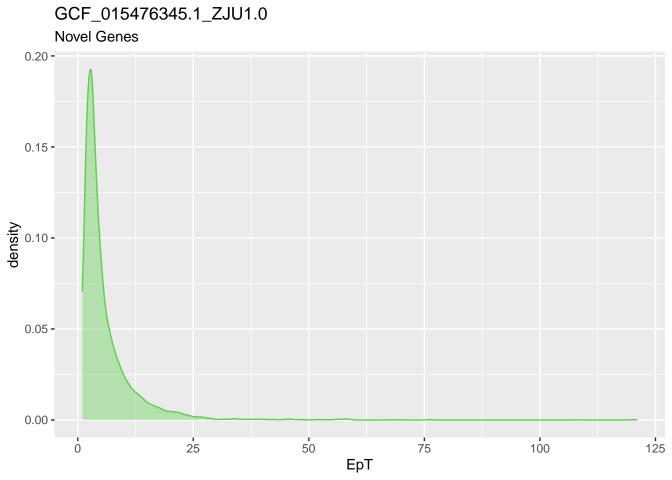


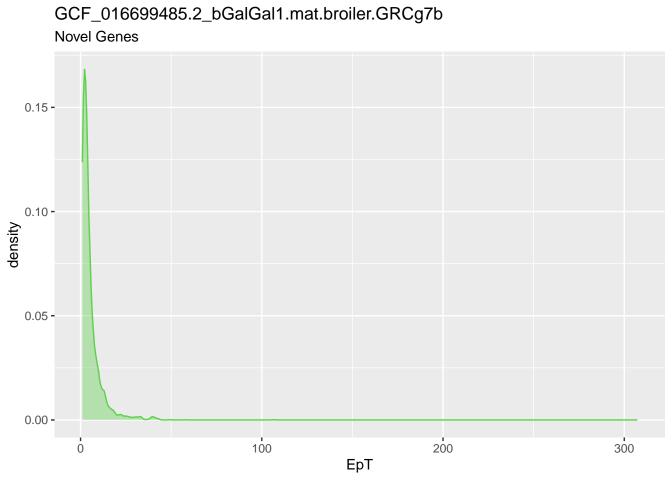
GCF_011762595.1_mTurTru1.mat.Y **Novel Genes** 0.15 -0.10 density 0.05 -0.00 -25 50 75 100 EpT

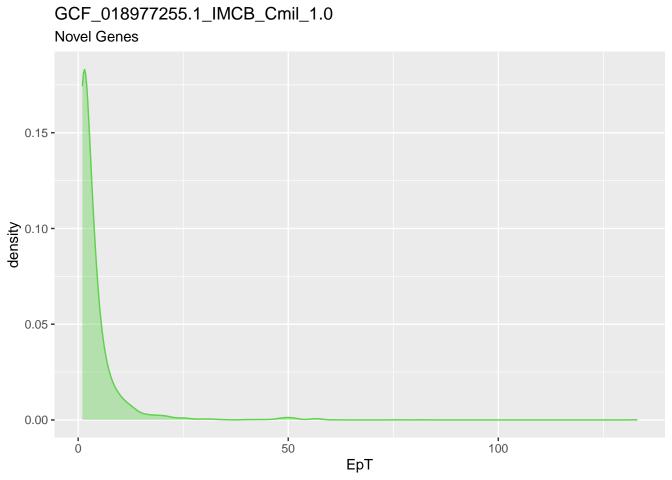


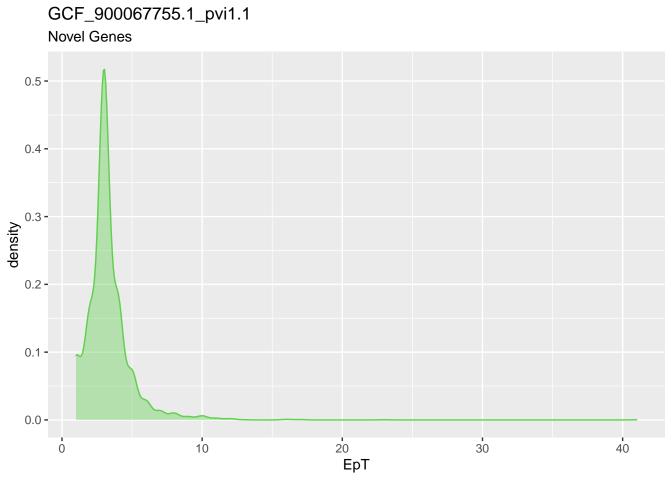


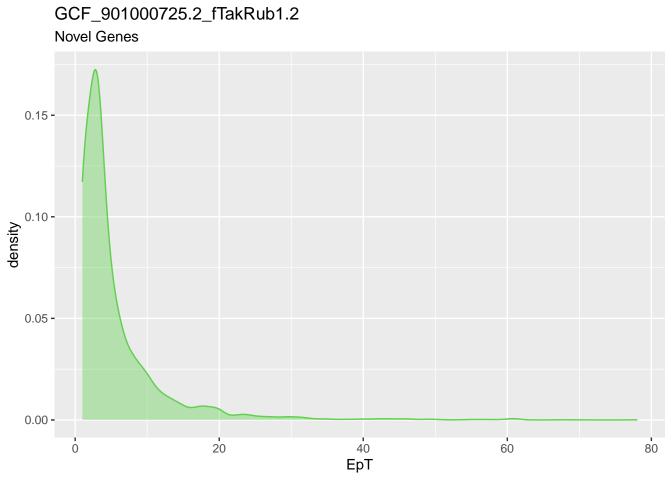


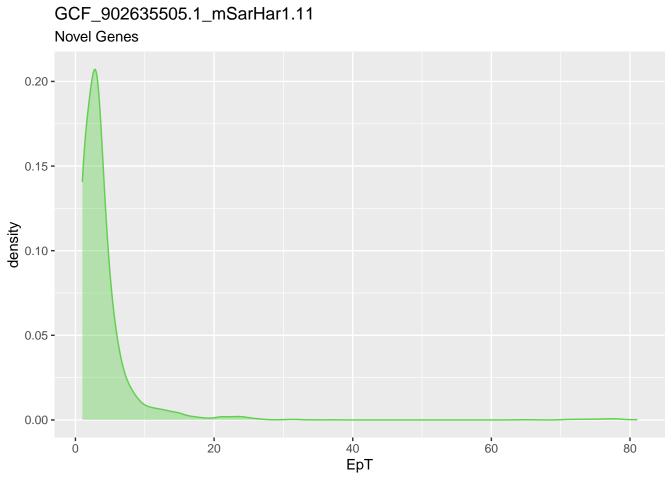


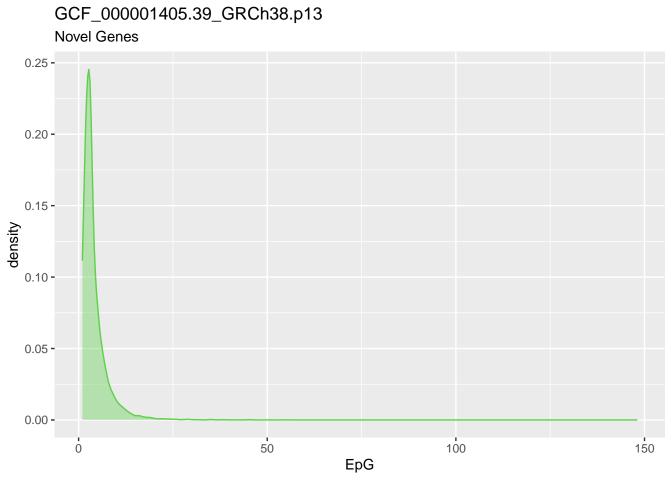


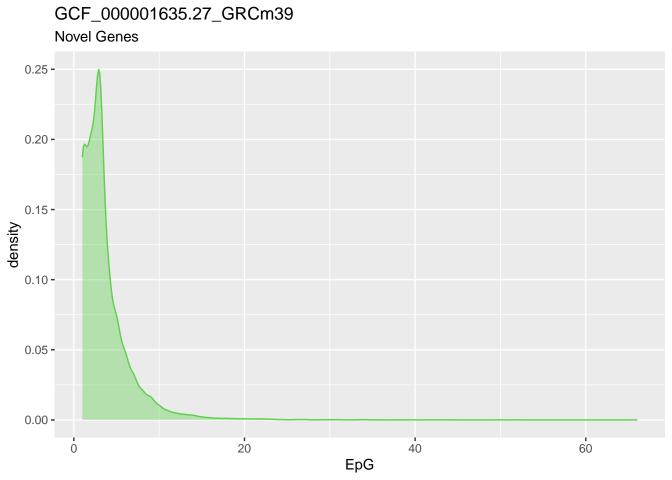


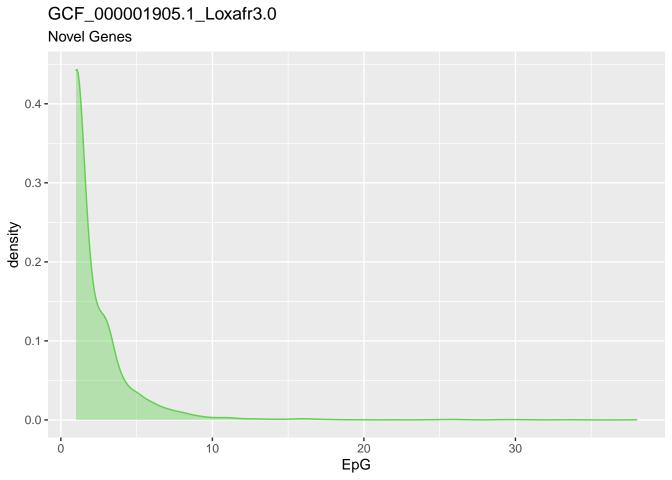


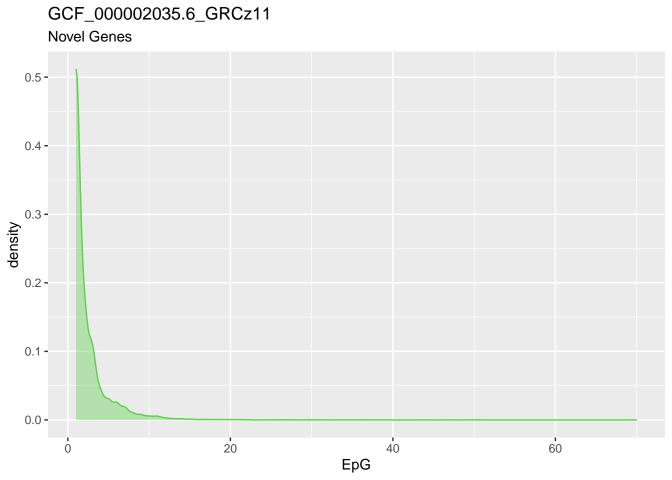


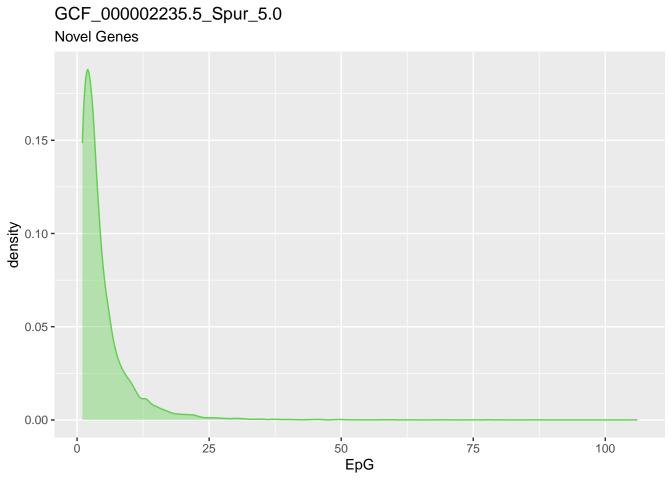




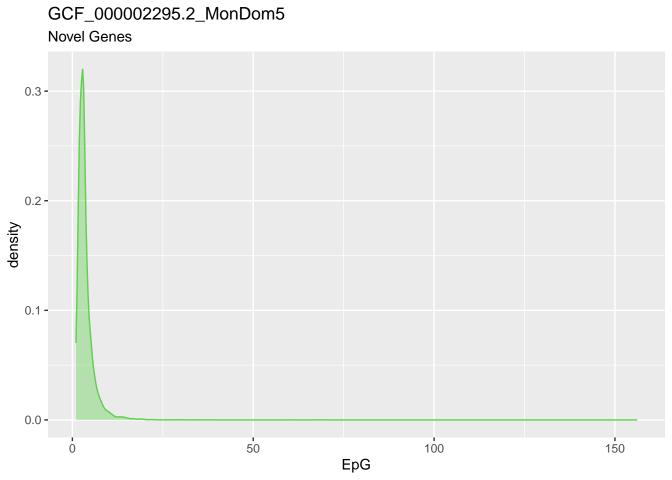


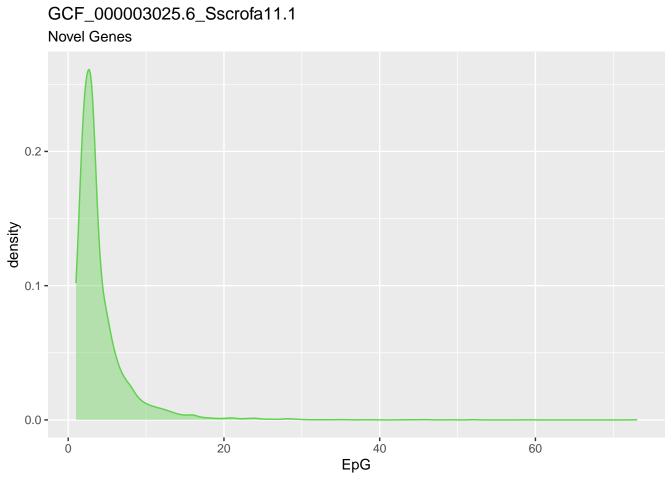


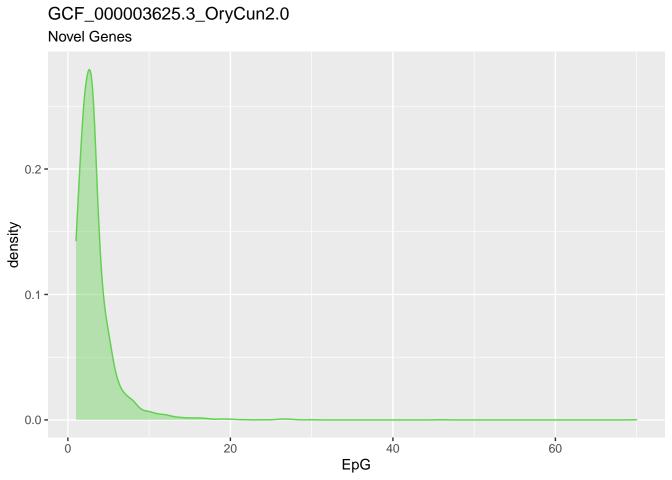


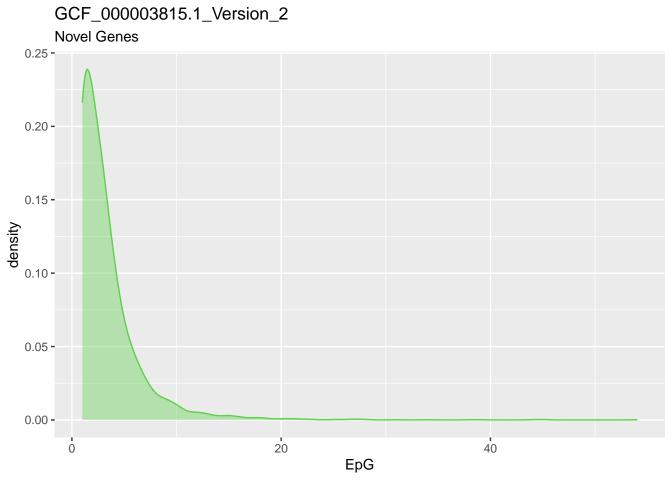


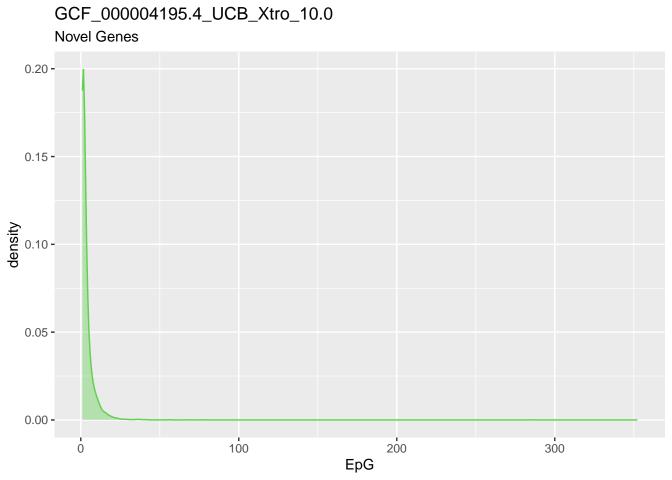
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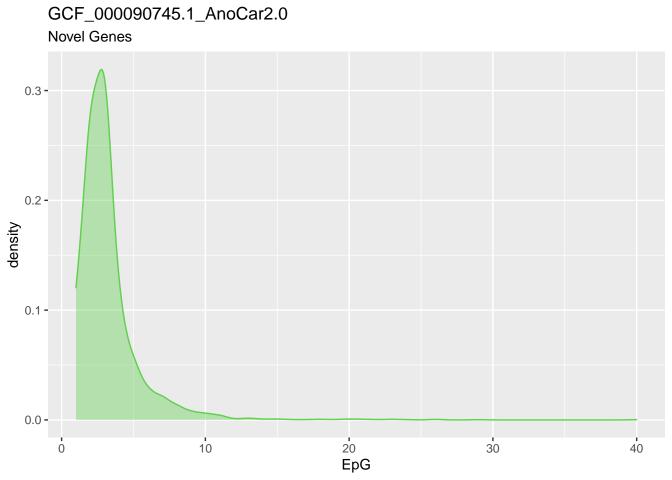




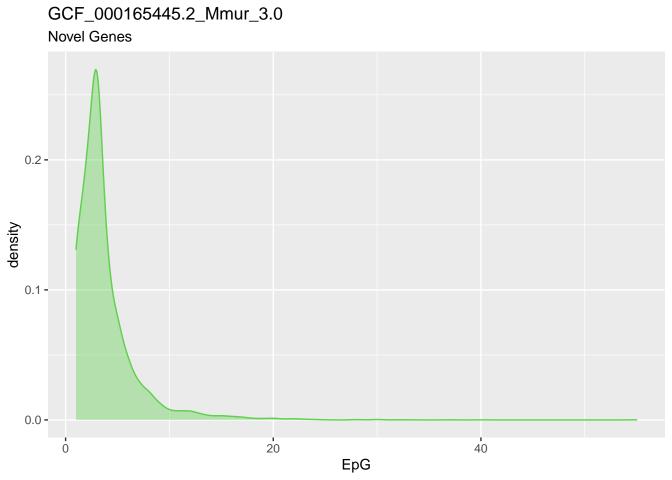


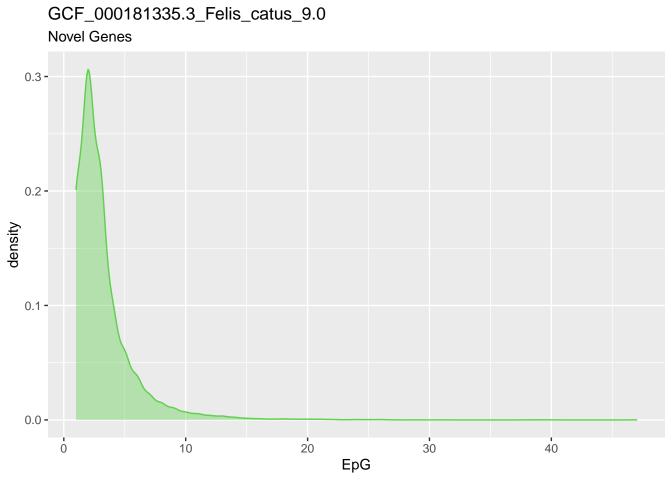


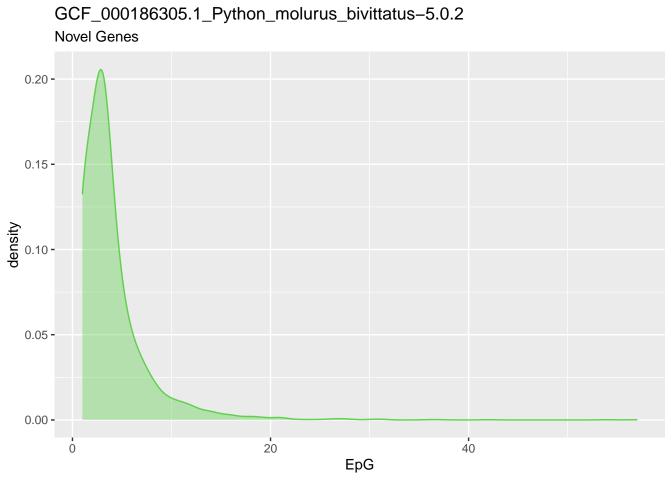


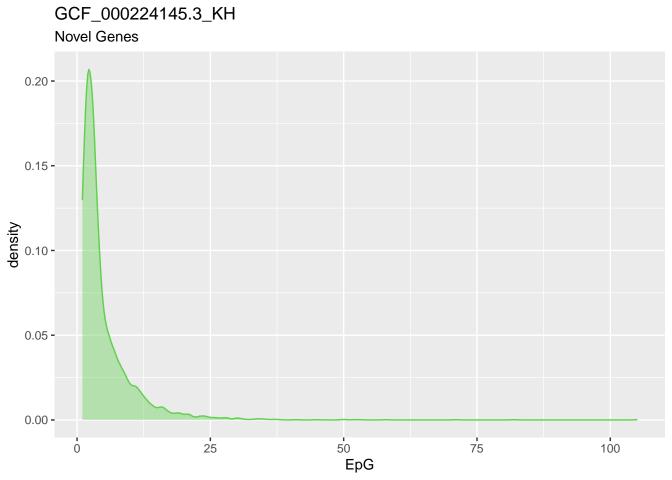


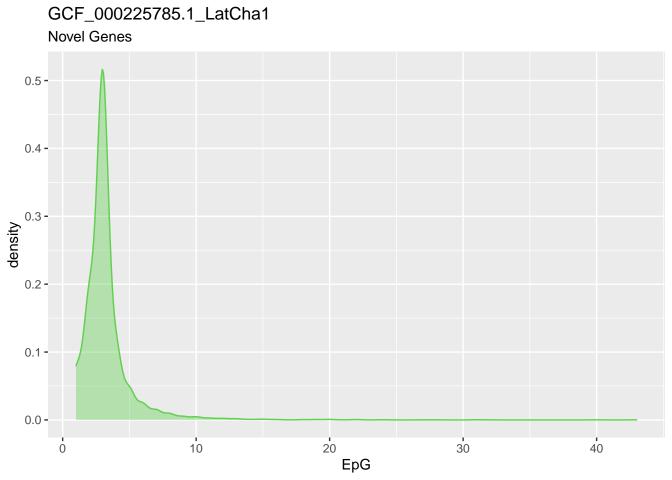
GCF_000151735.1_Cavpor3.0 **Novel Genes** 0.4 -0.3 density 0.1 -0.0 -10 20 30 40 EpG

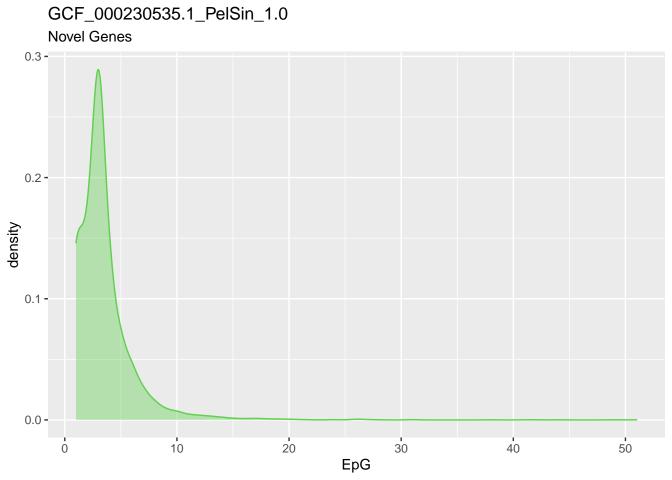


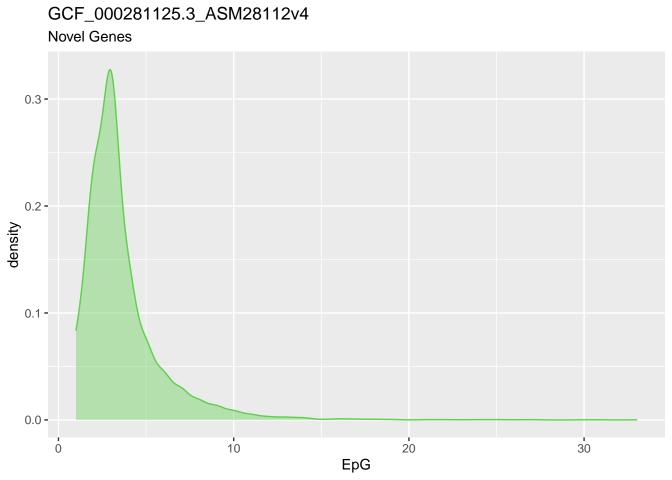


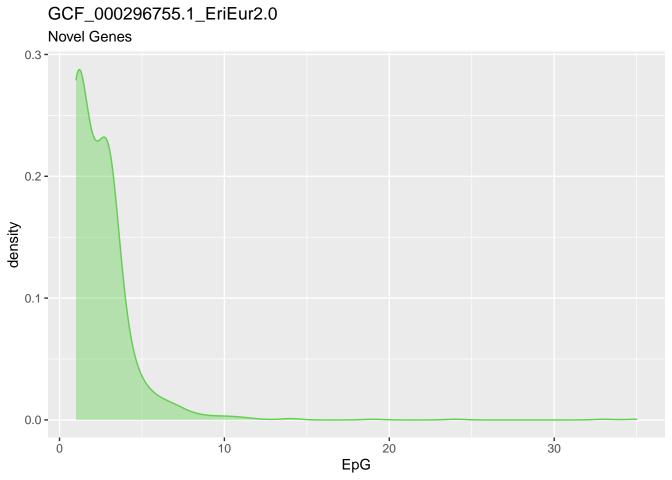


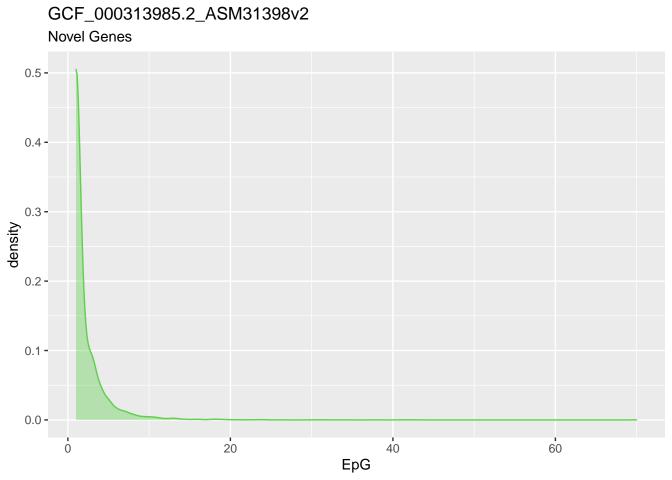


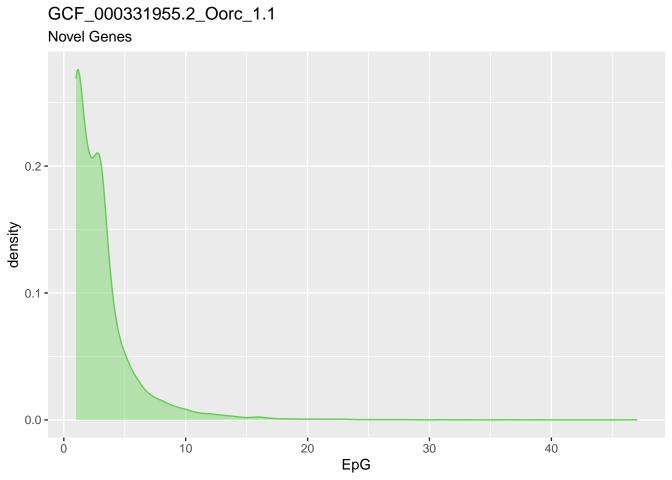


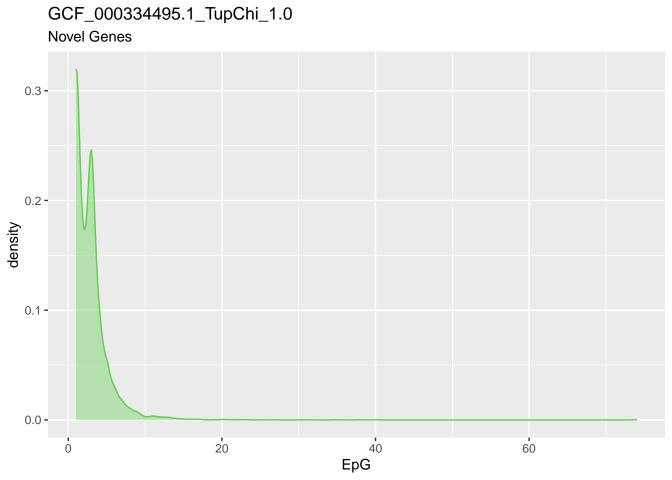


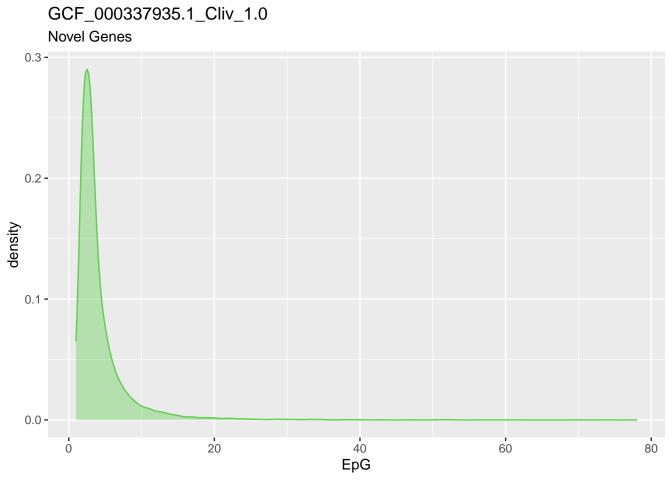


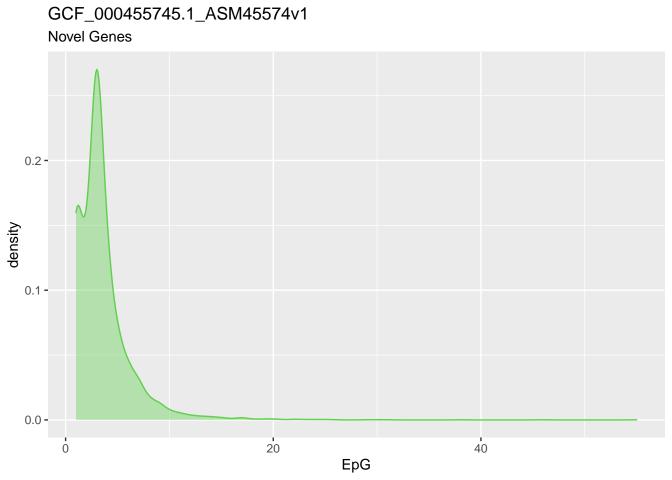


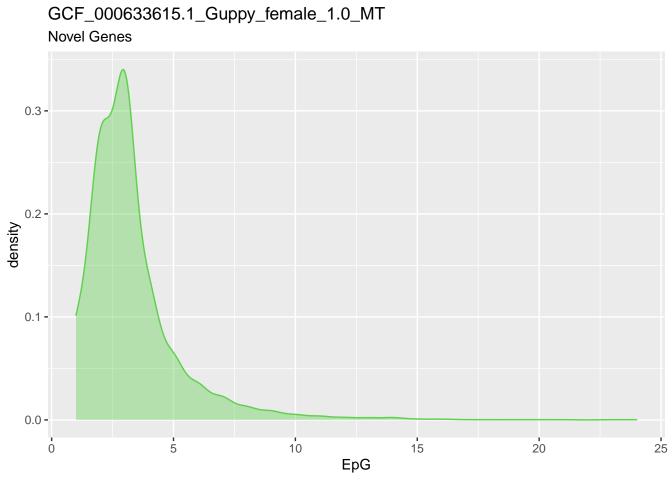




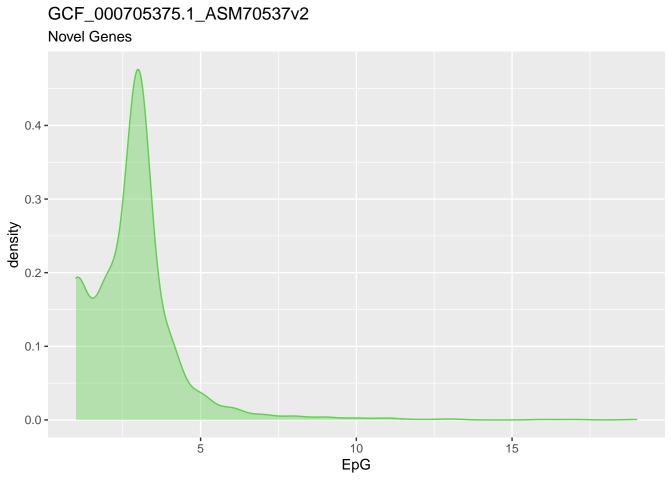




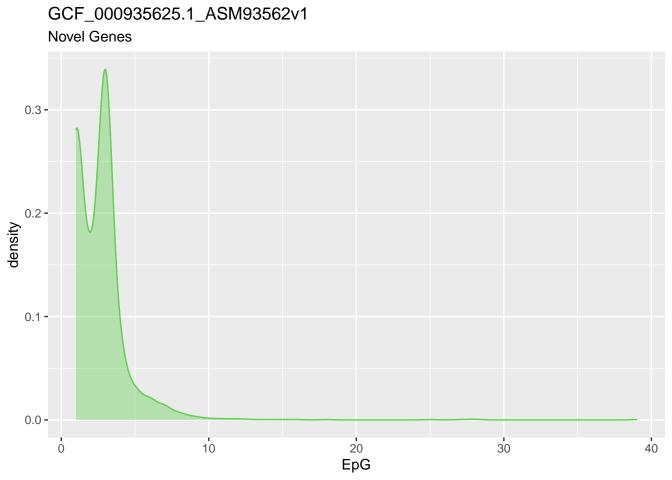


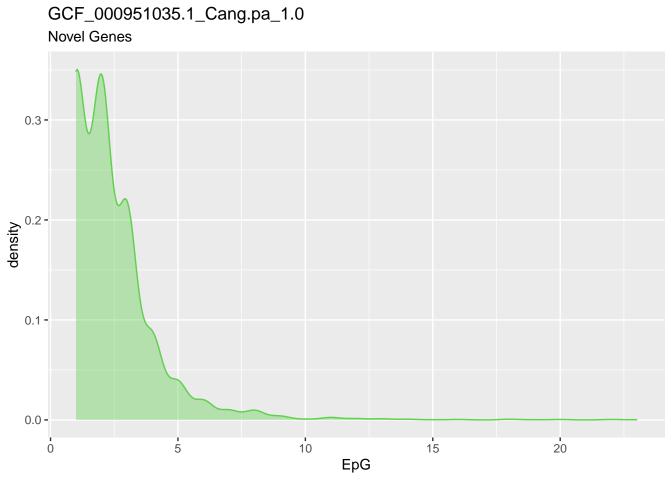


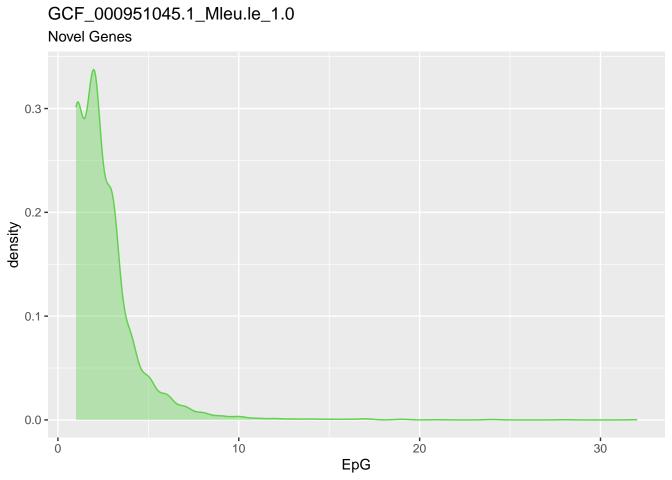
GCF_000696425.1_G_variegatus-3.0.2 **Novel Genes** 0.3 density 0.1 -0.0 -10 20 30 EpG

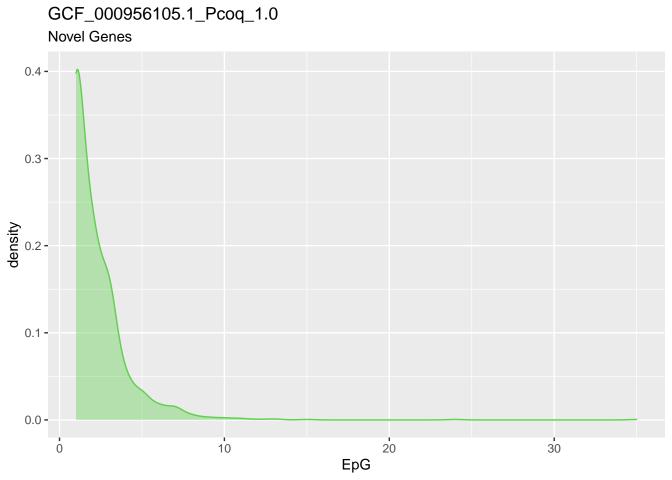


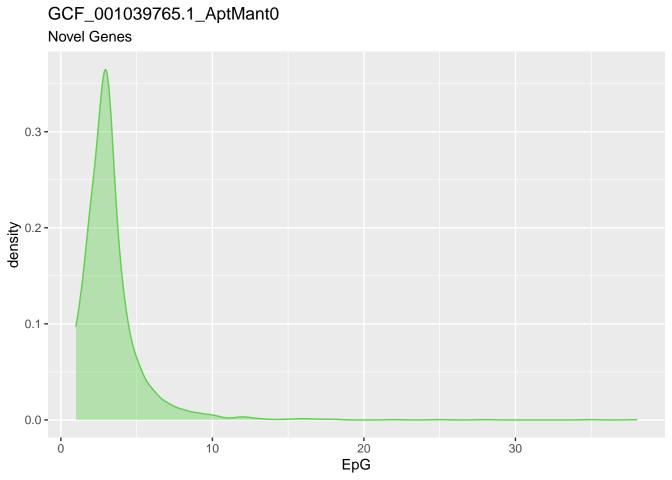
GCF_000708225.1_ASM70822v1 **Novel Genes** 0.3 -0.2 density 0.1 -0.0 -10 20 30 EpG

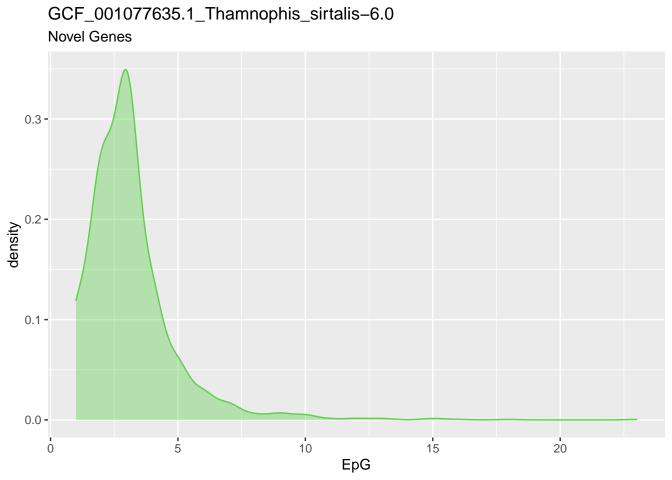


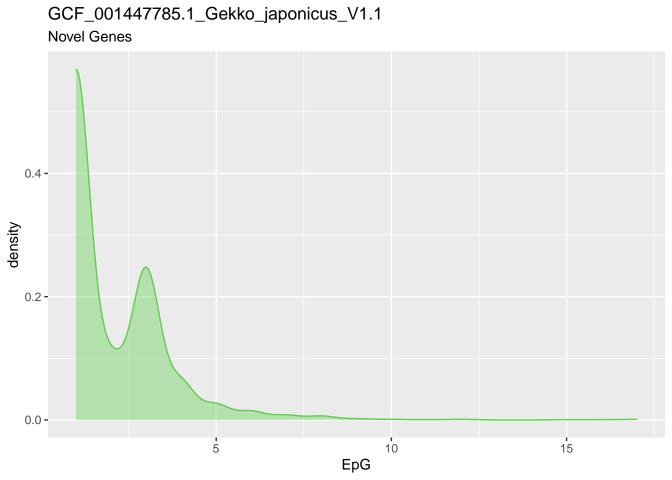


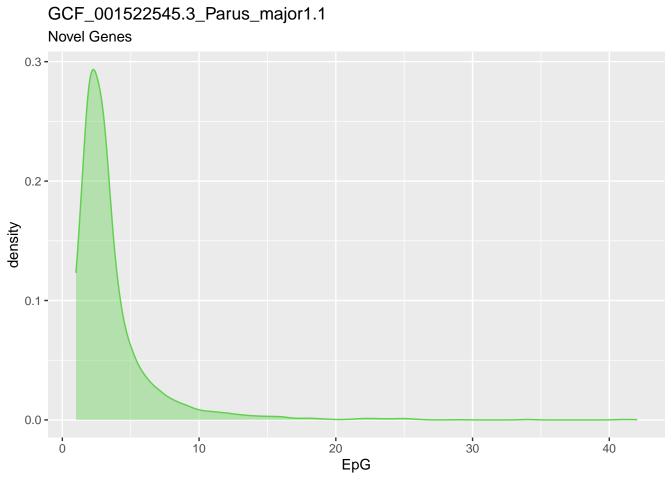






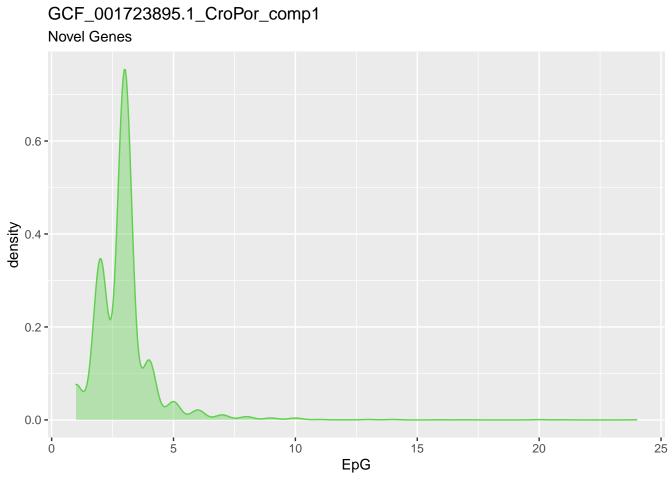


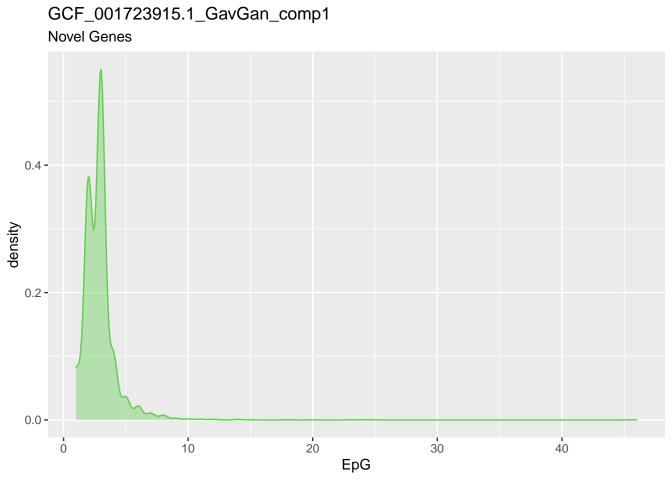


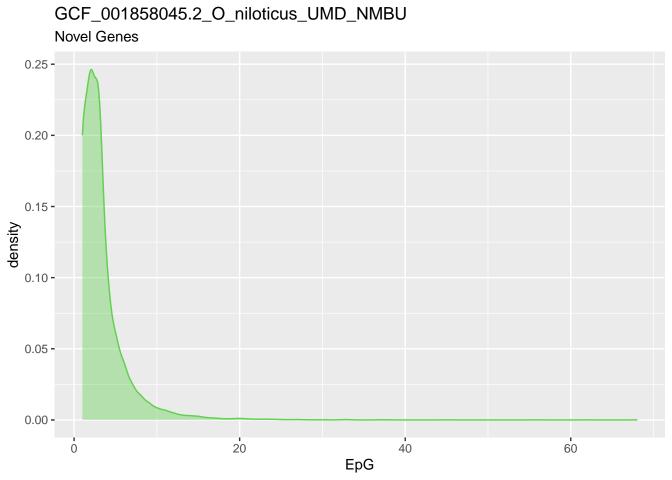


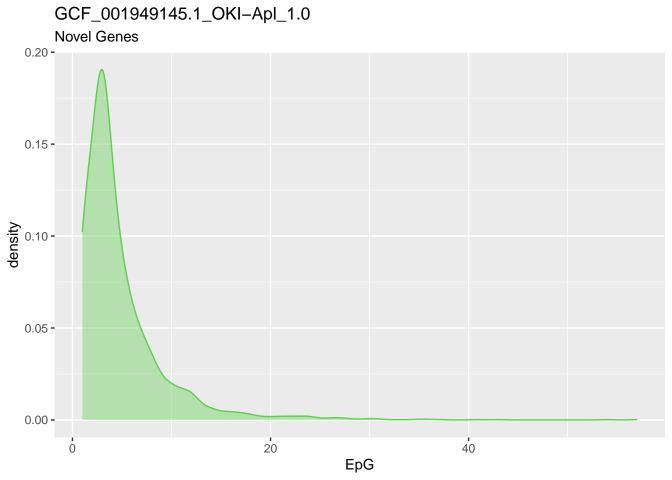
GCF_001625305.1_Haploidv18h27 **Novel Genes** 0.20 -0.15 density 0.10 -0.05 -0.00 -40 20 60 80 EpG

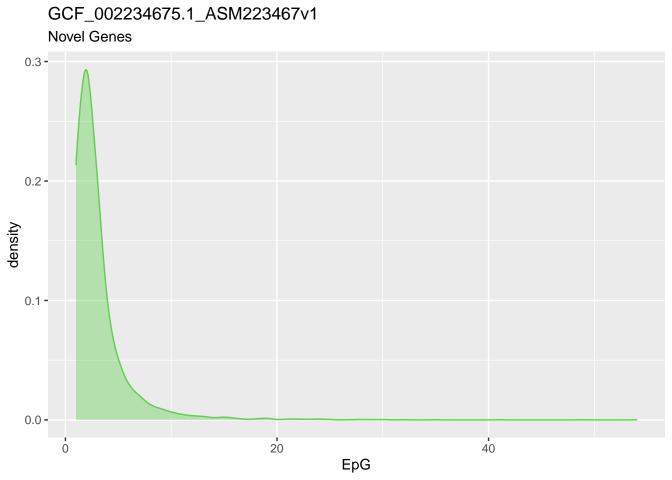
GCF_001642345.1_ASM164234v2 **Novel Genes** 0.4 -0.3 density 0.1 -0.0 -20 60 40 EpG

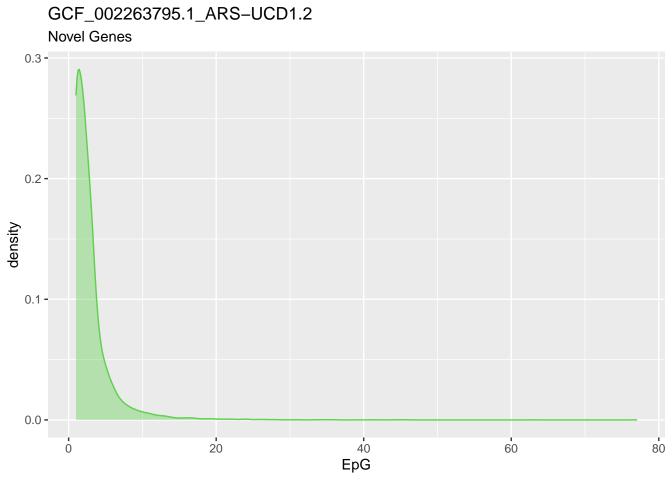












GCF_002288925.2_ASM228892v3 **Novel Genes** 0.3 -0.2 density 0.1 -

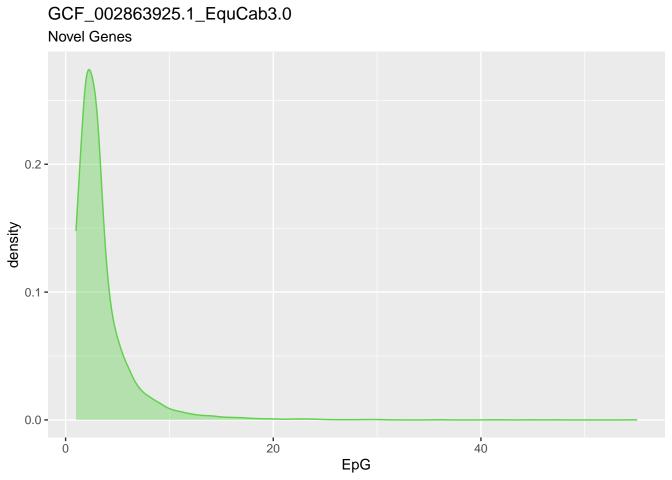
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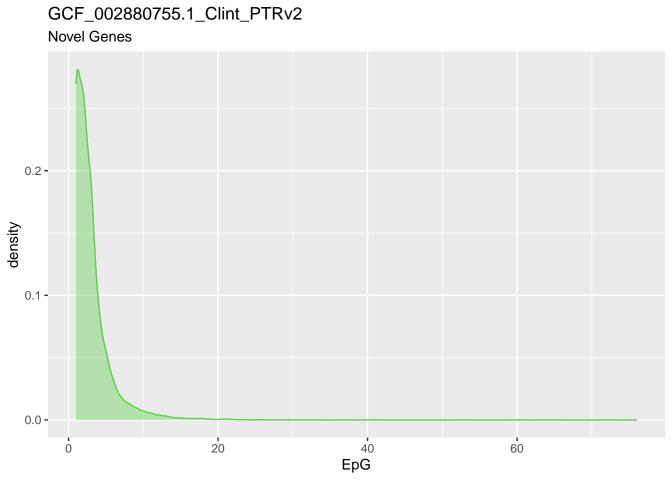
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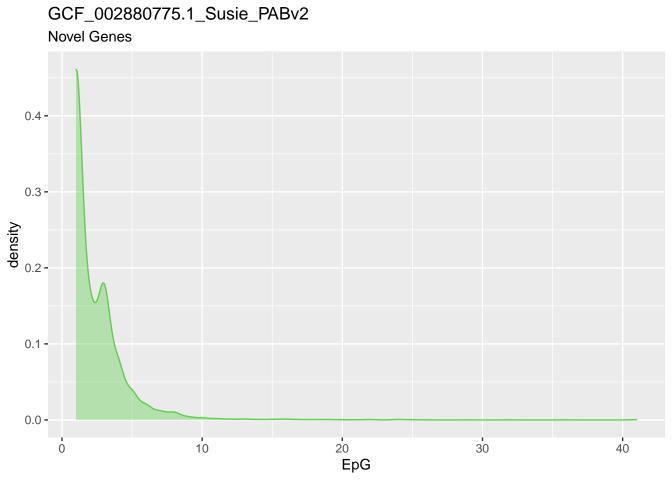
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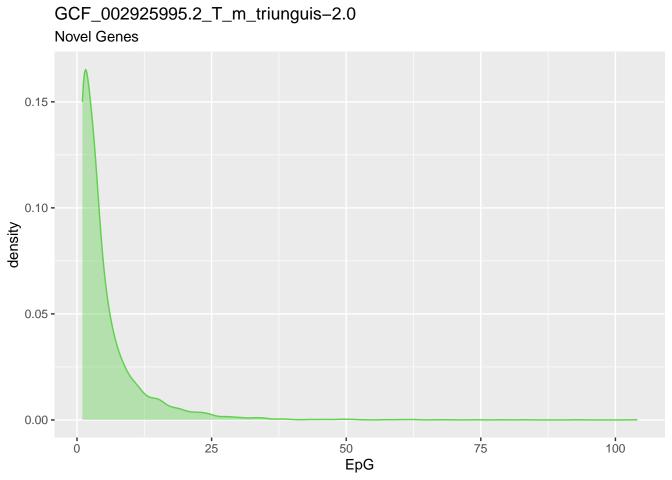
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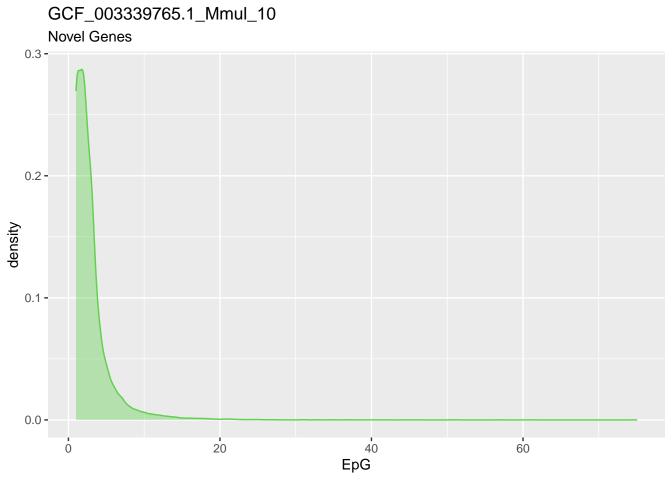
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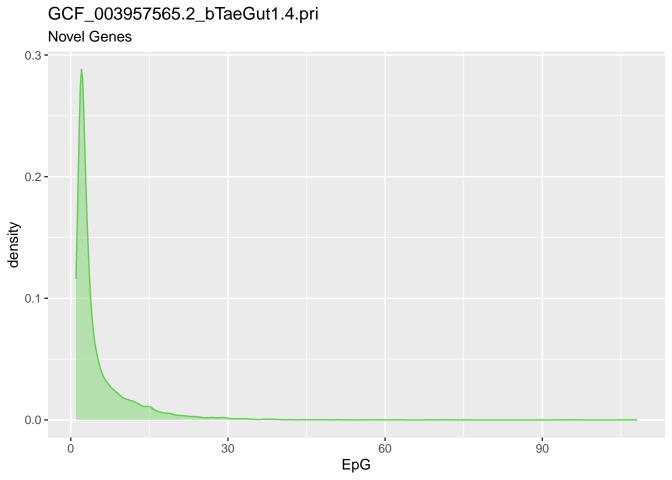


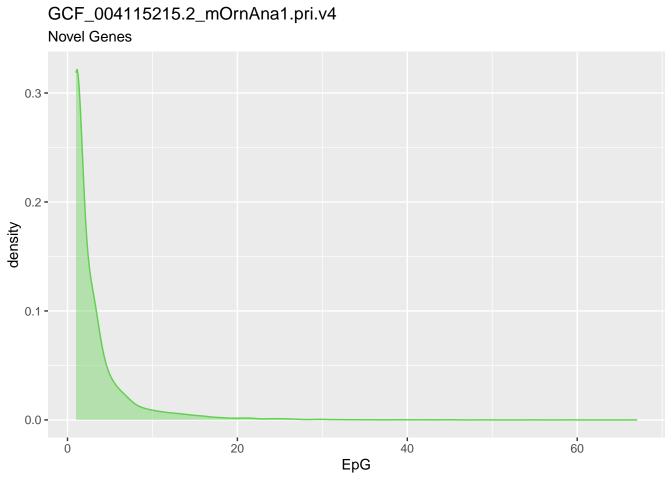


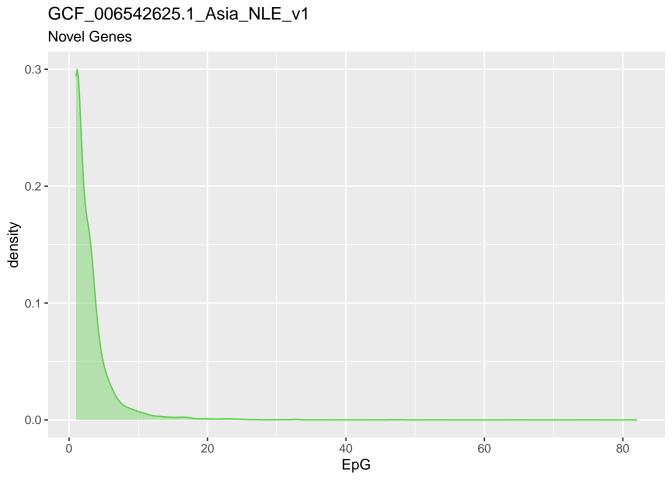


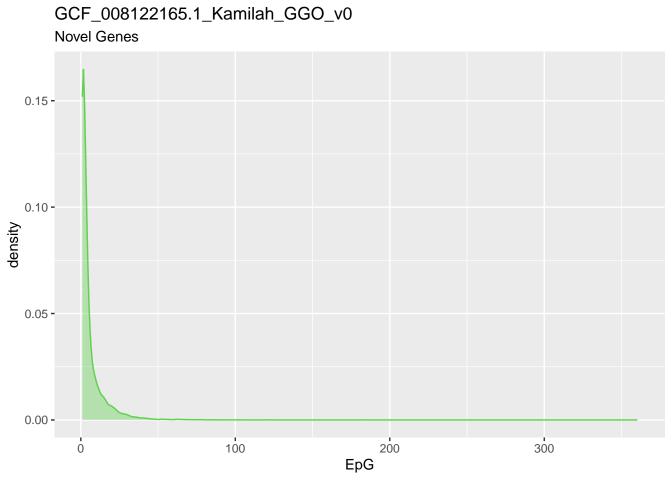


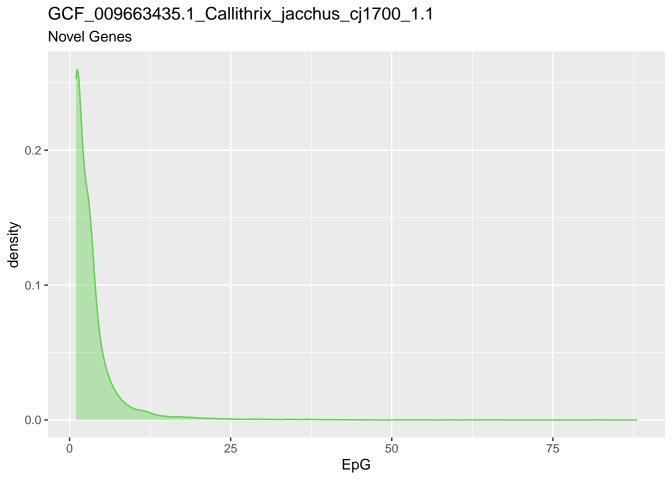






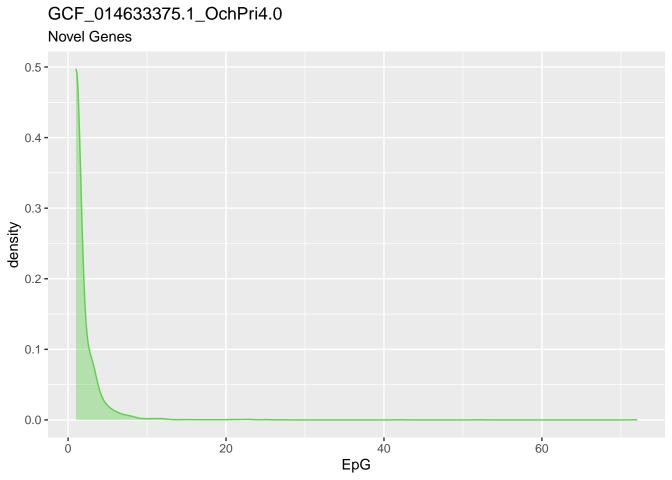


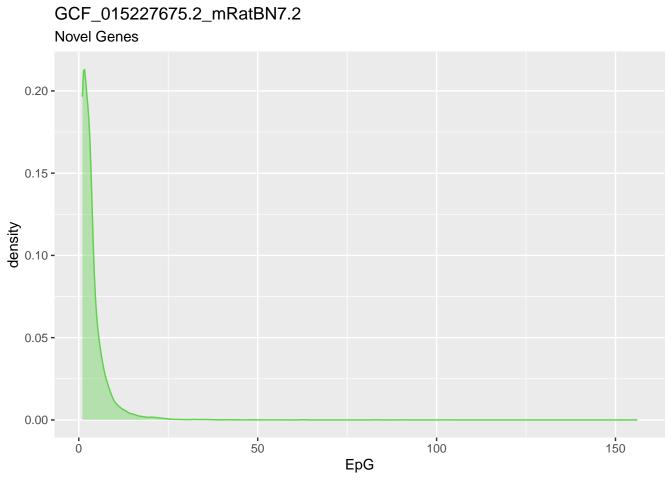


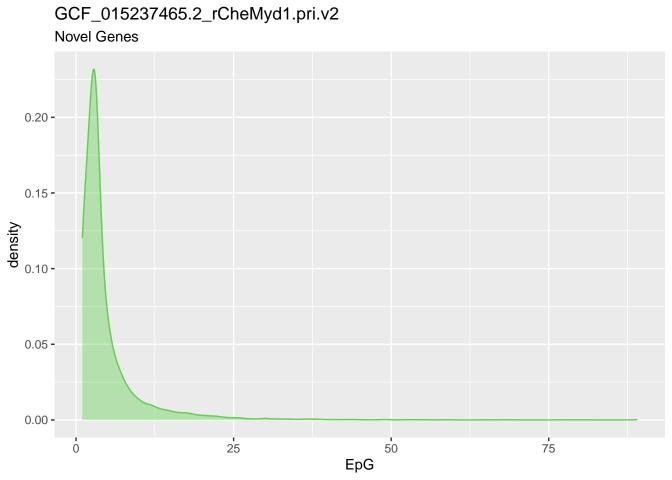


GCF_011125445.2_MU-UCD_Fhet_4.1 **Novel Genes** 0.20 -0.15 density 0.05 -0.00 -25 50 75 EpG

GCF_011762595.1_mTurTru1.mat.Y **Novel Genes** 0.15 density 0.05 -0.00 -25 50 75 100 125 EpG

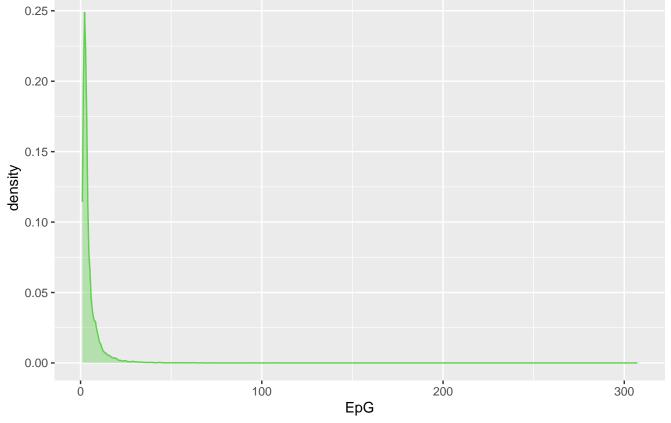


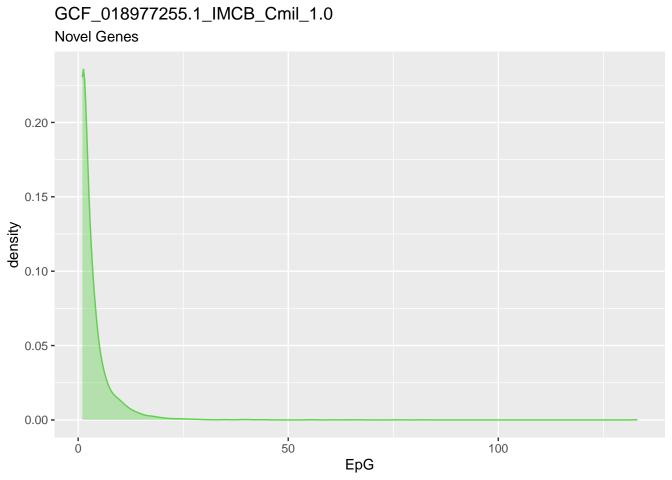


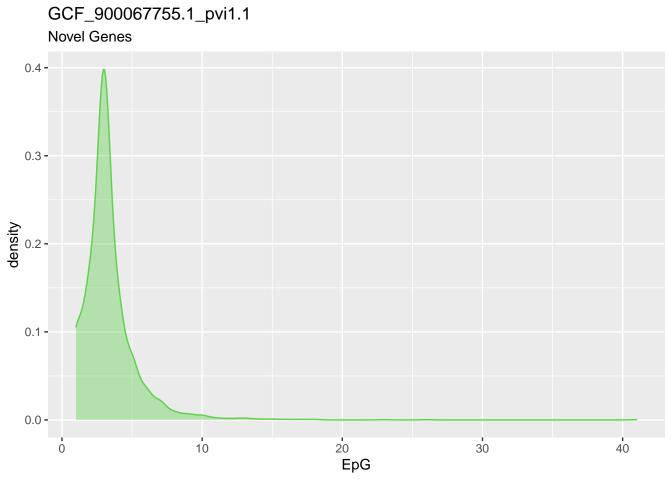


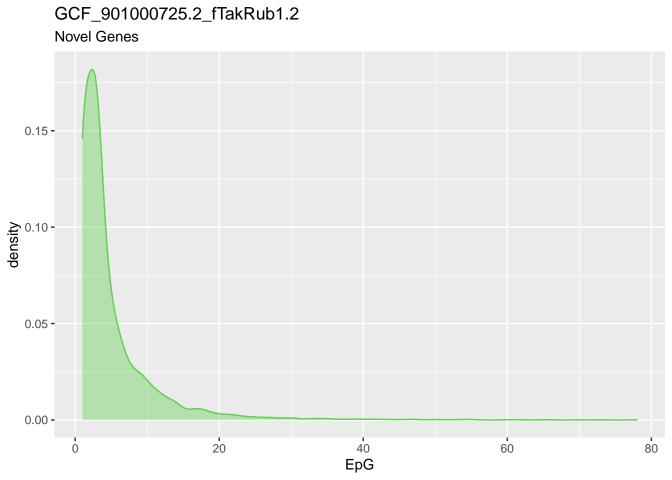
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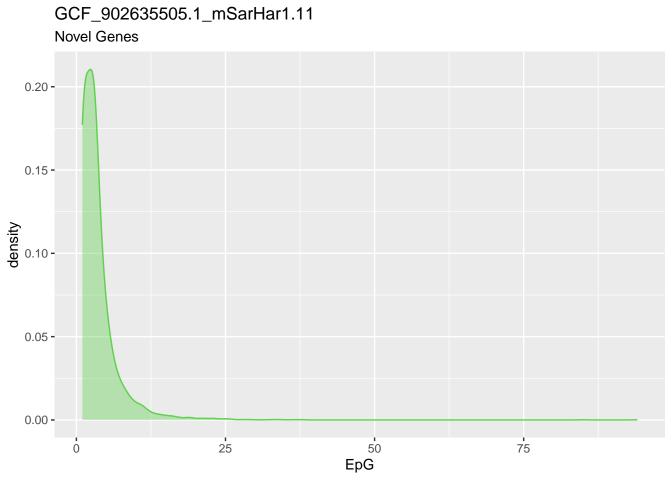
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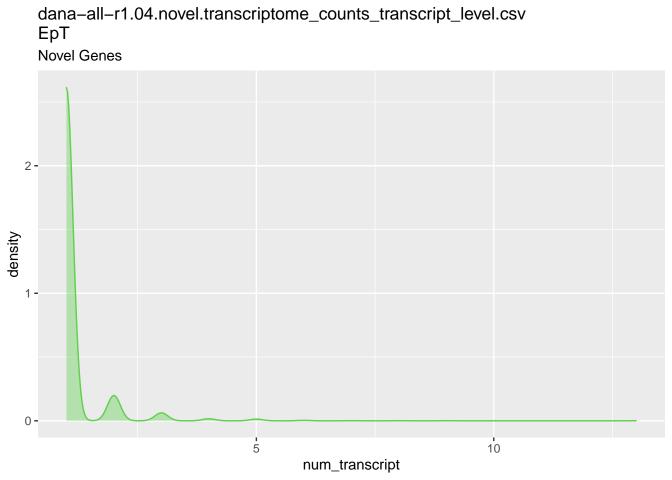


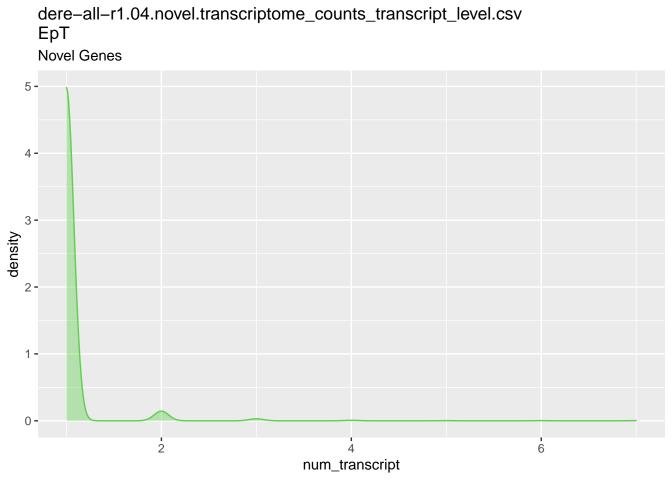




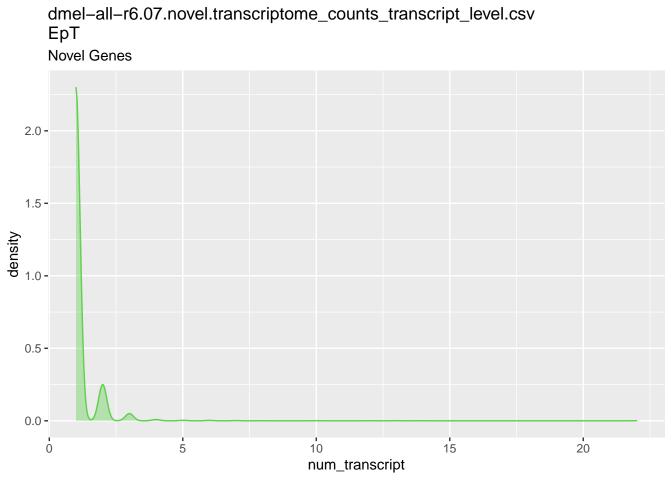


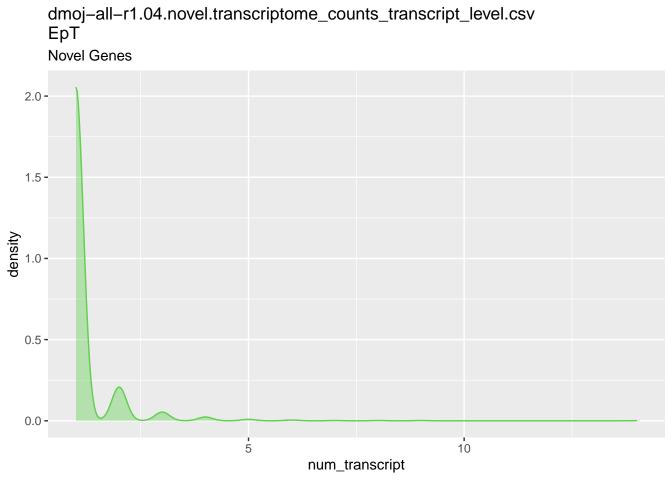






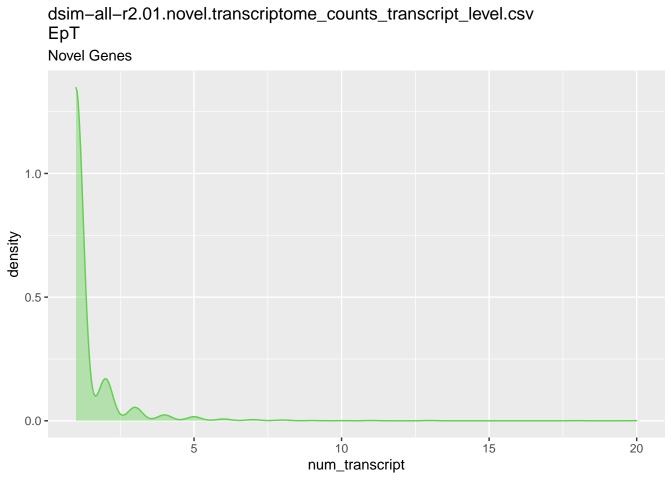
	dgri-all-r1.3.novel.transcriptome_counts_transcript_level.csv EpT
	Novel Genes
Ę	
density	
0	
	num_transcript

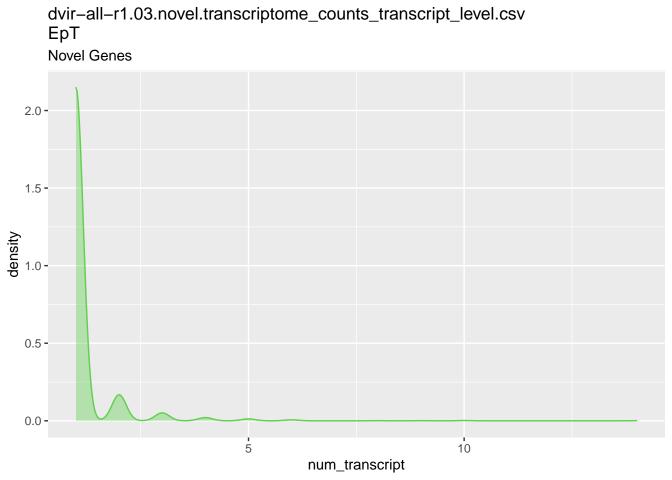


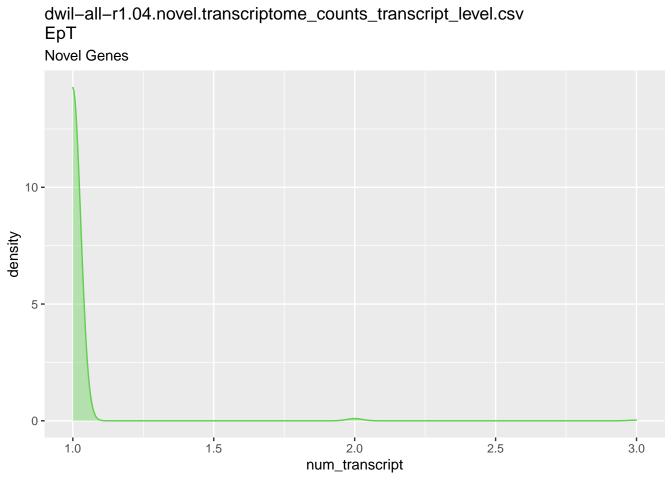


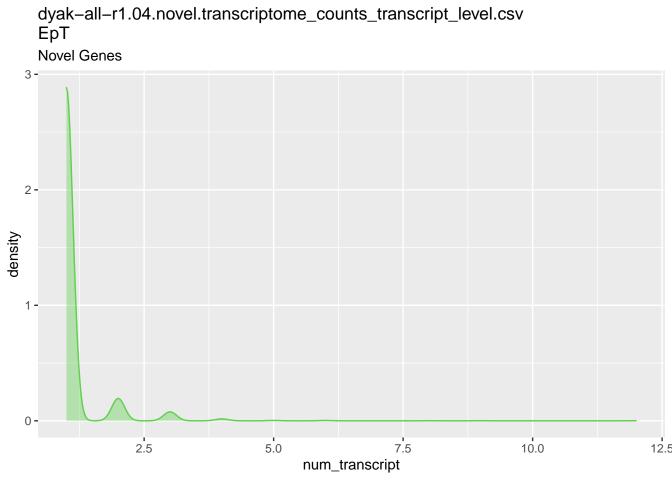
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density	num_transcript
	num_transcript

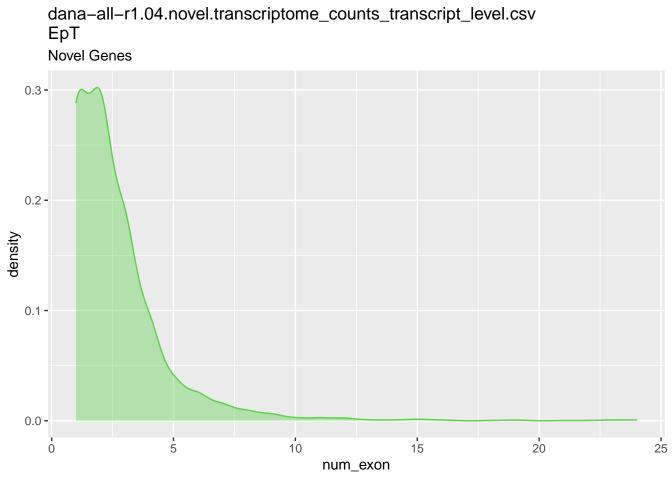
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density	Novel Genes
	num_transcript











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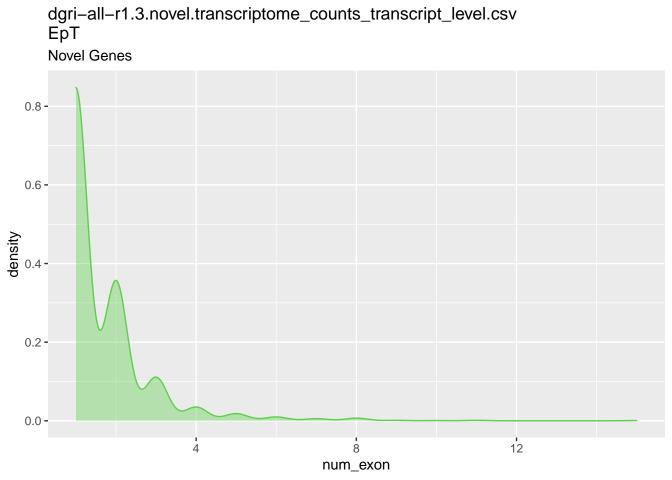
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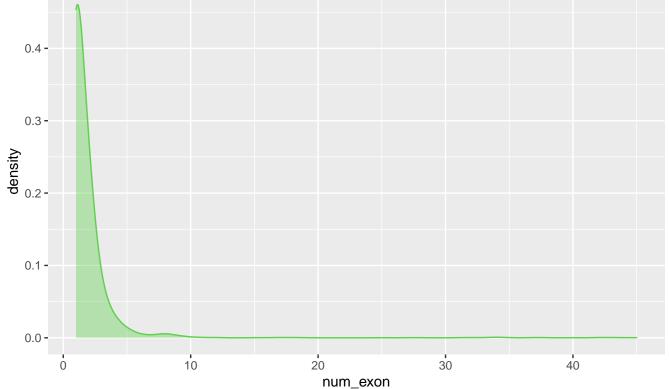
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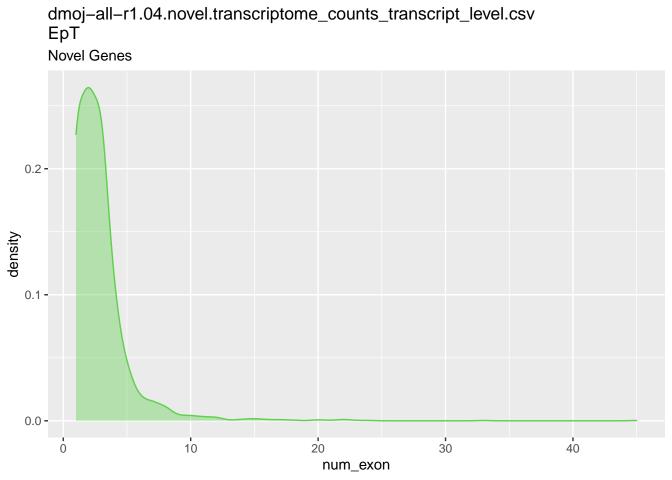
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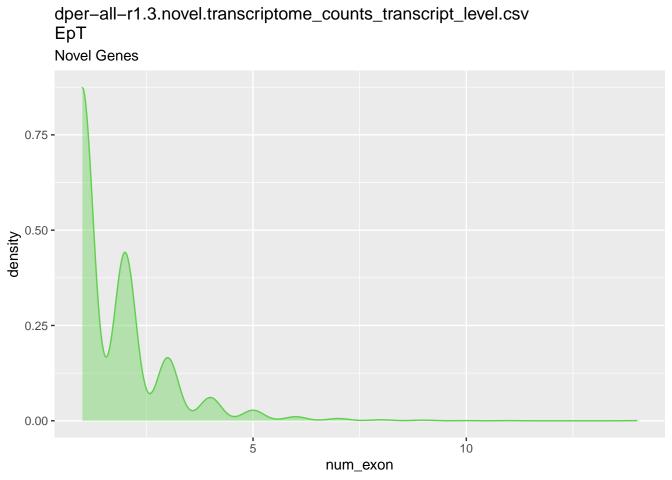
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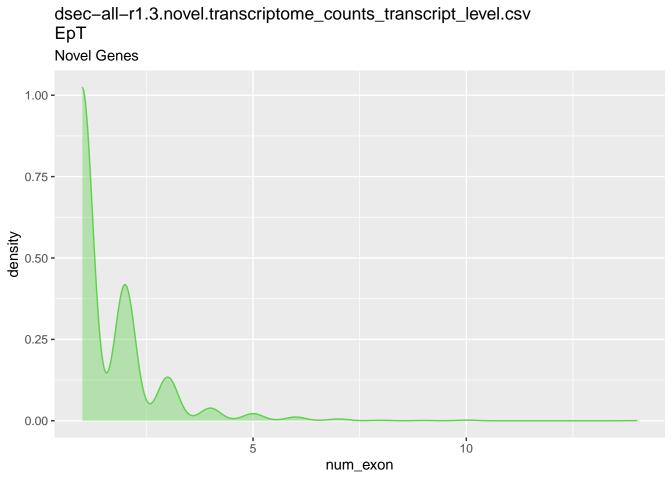


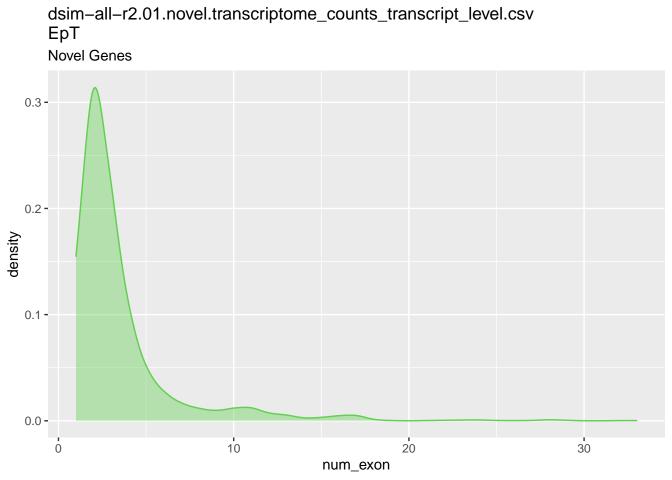
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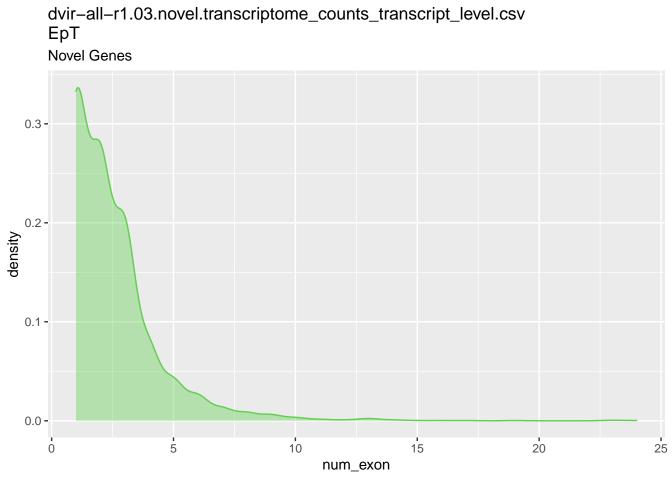


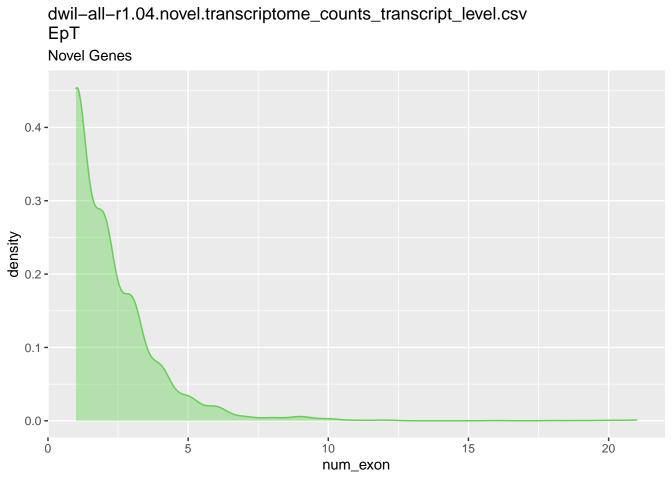












dyak-all-r1.04.novel.transcriptome_counts_transcript_level.csv EpT **Novel Genes** 0.3 -0.2 density 0.1 -0.0 -20 40 60 num_exon

dana-all-r1.04.novel.transcriptome_counts_transcript_level.csv EpT **Novel Genes** 0.3 -0.2 density 0.1 -0.0 -10 20 30 num_uniq_exon

dere-all-r1.04.novel.transcriptome_counts_transcript_level.csv EpT **Novel Genes** 0.4 -0.3 density 0.2 0.1 -

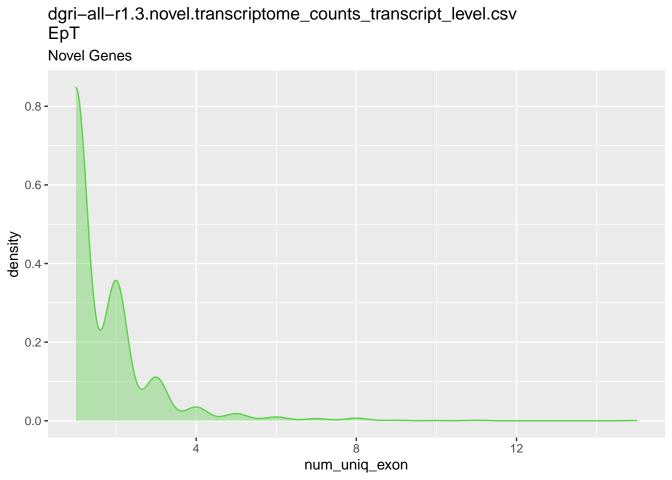
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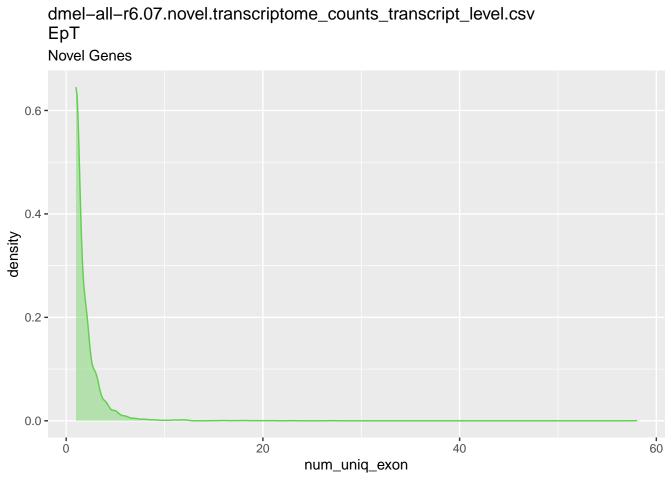
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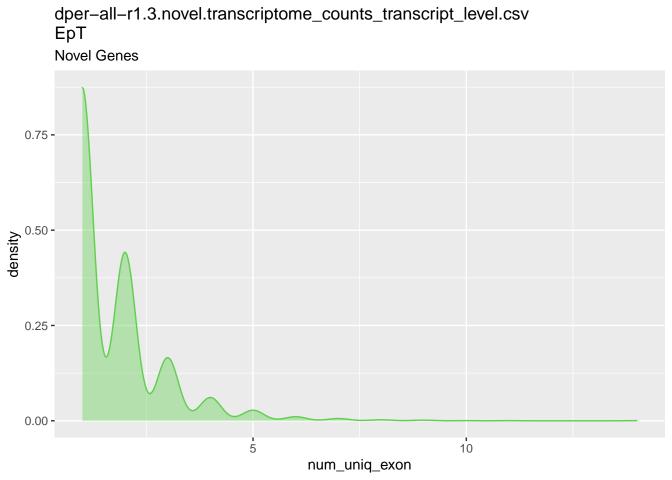
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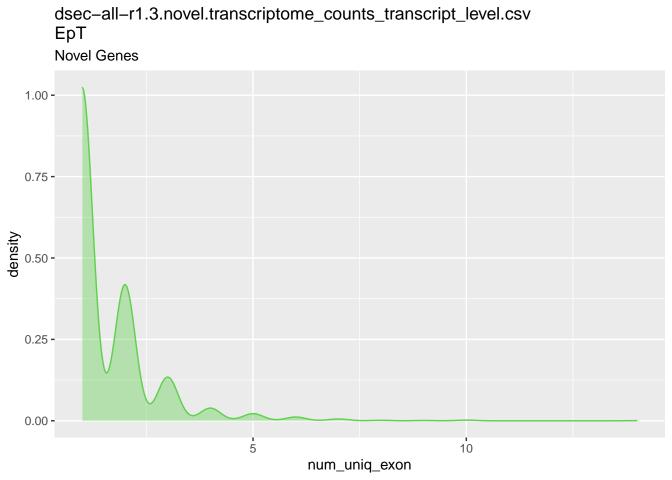
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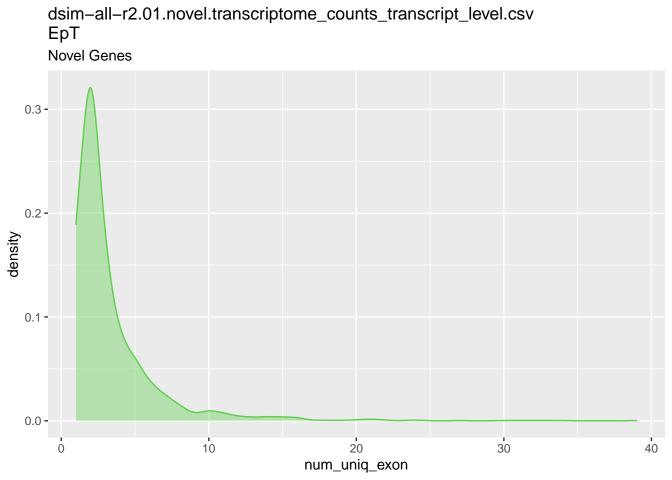




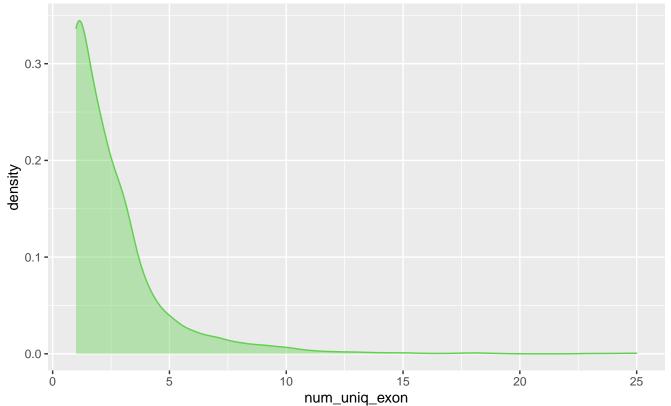
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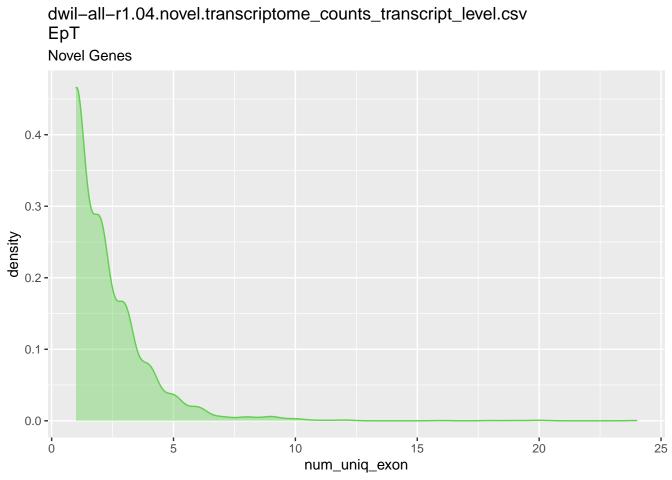


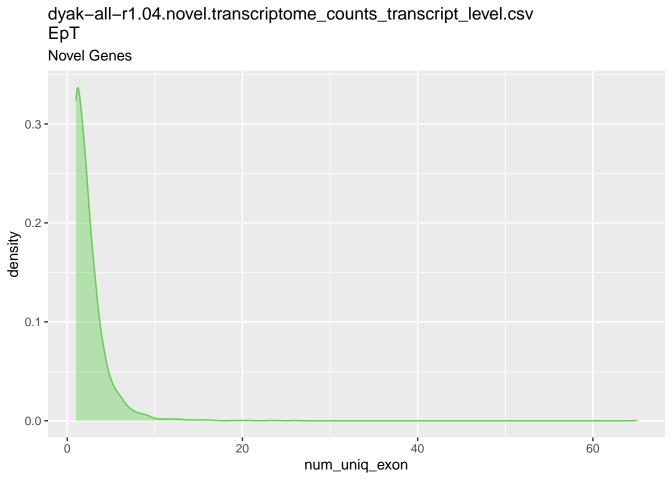


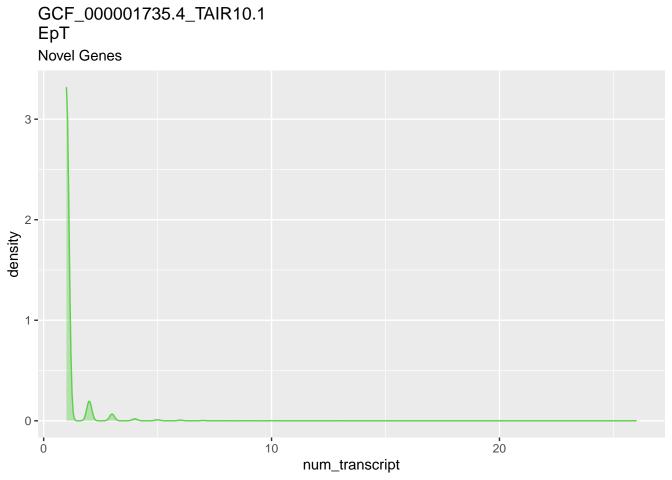


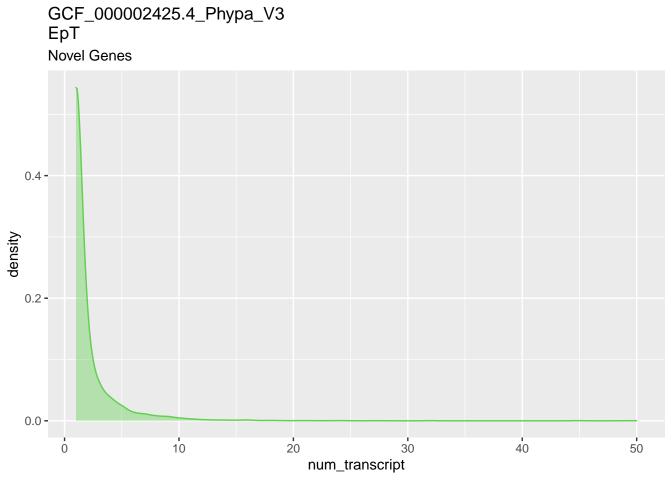
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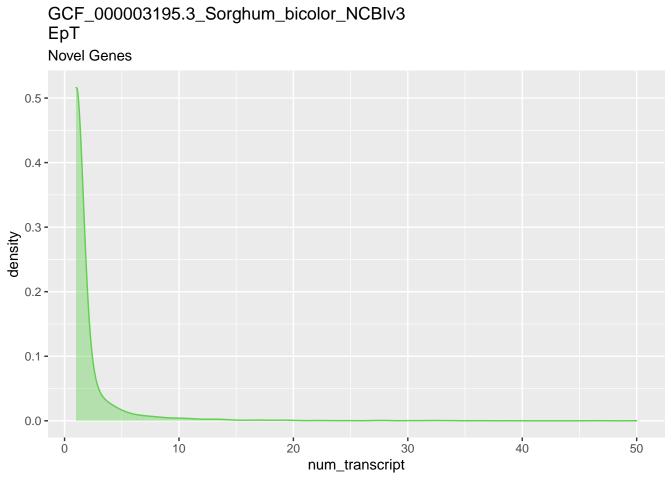


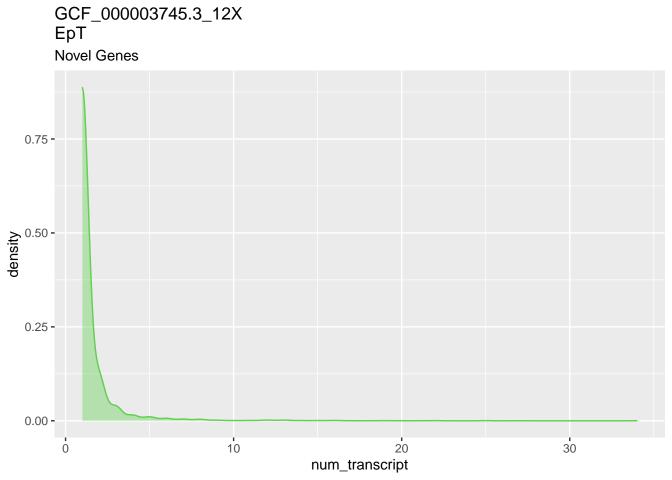


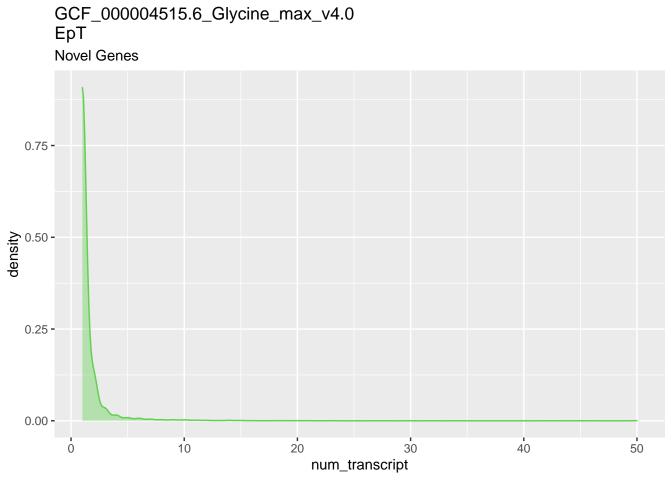


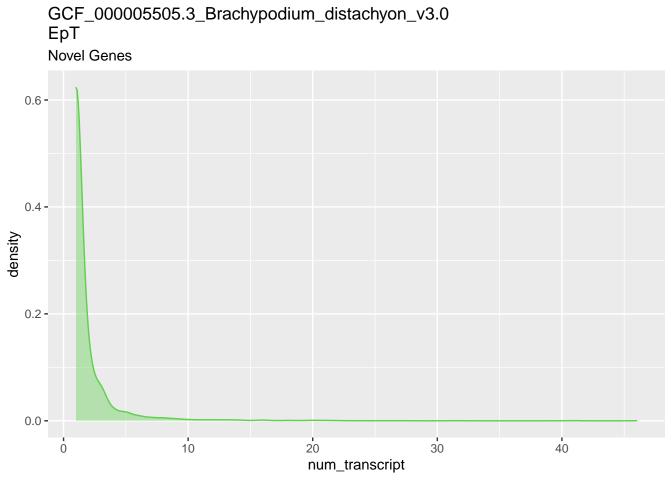


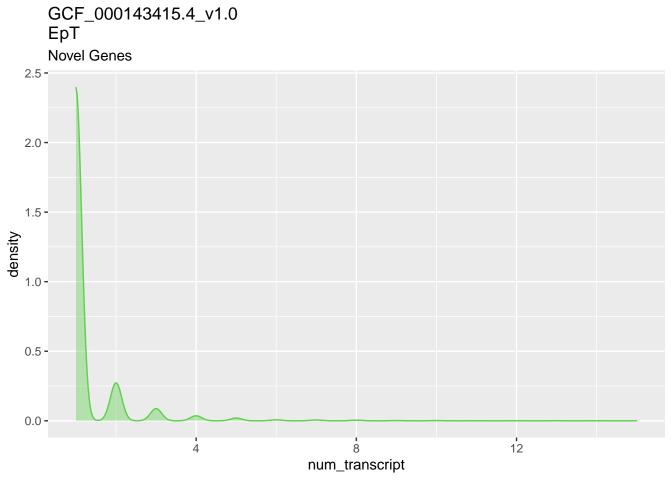


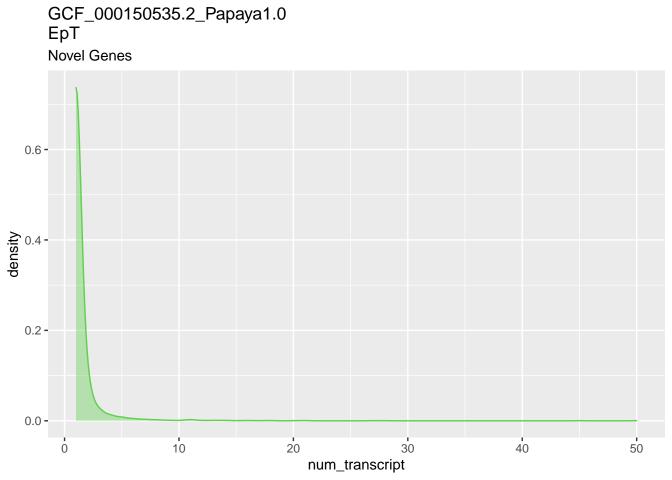


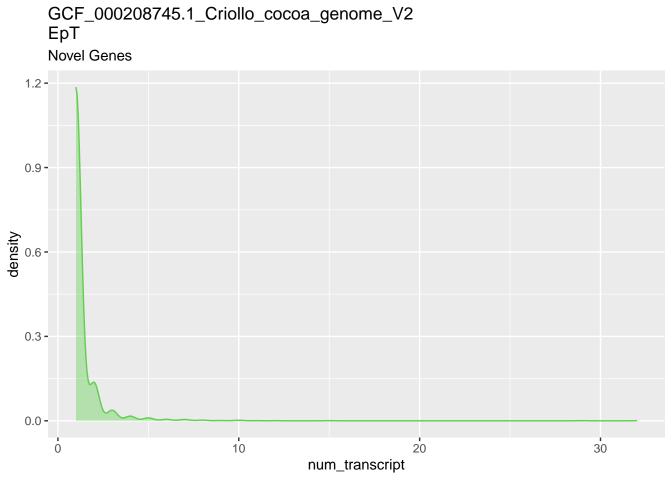


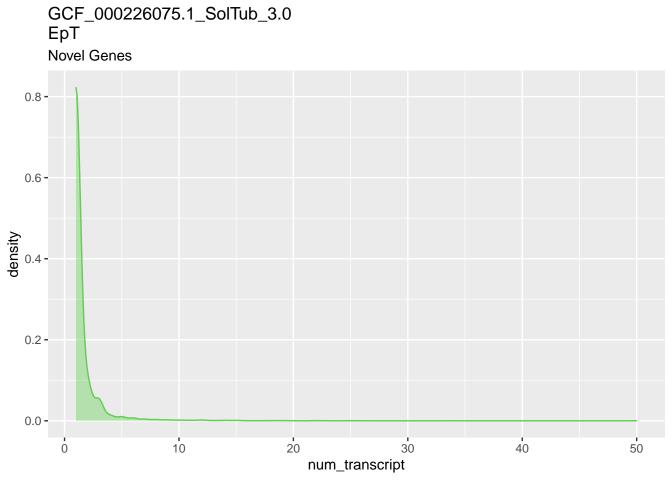


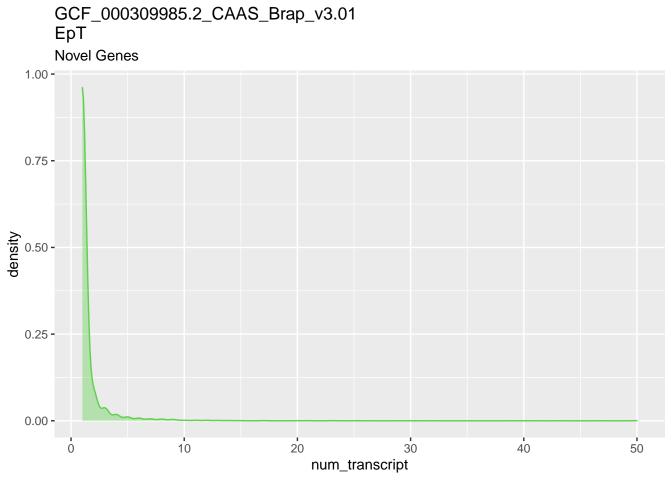


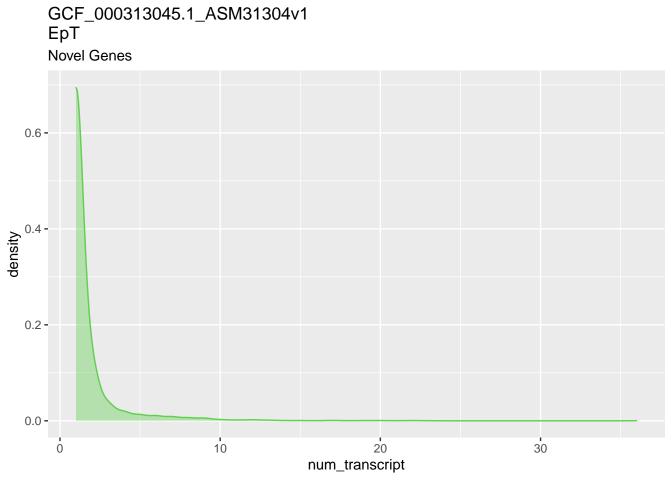


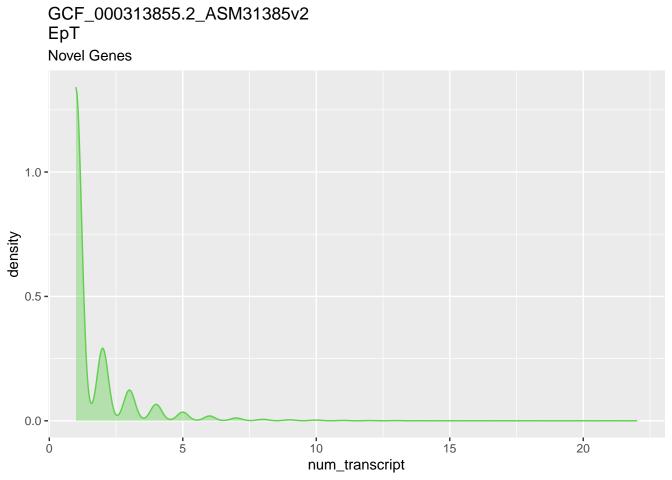


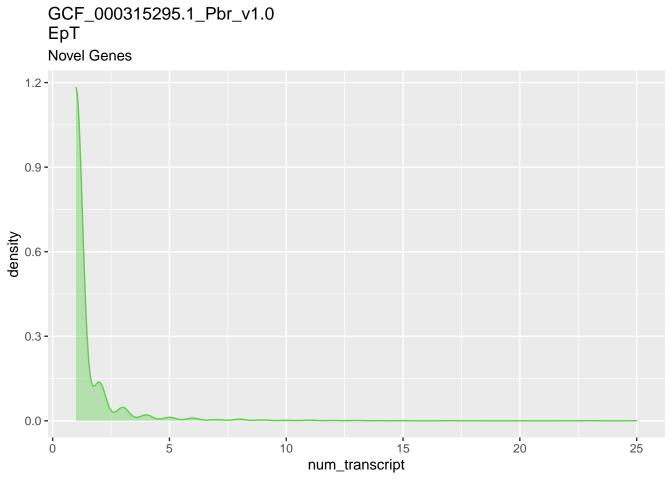


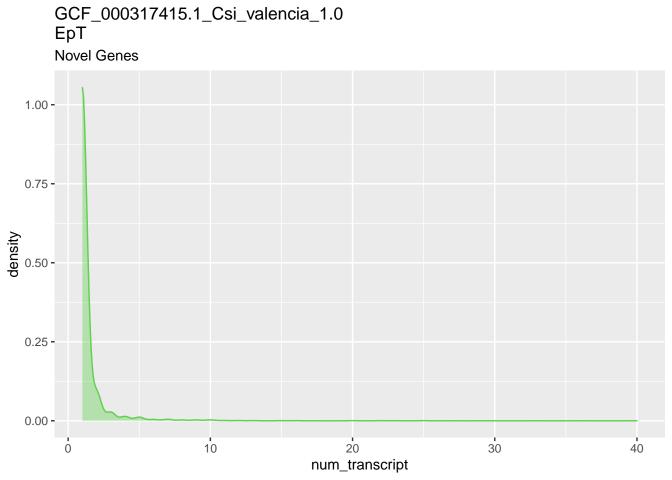


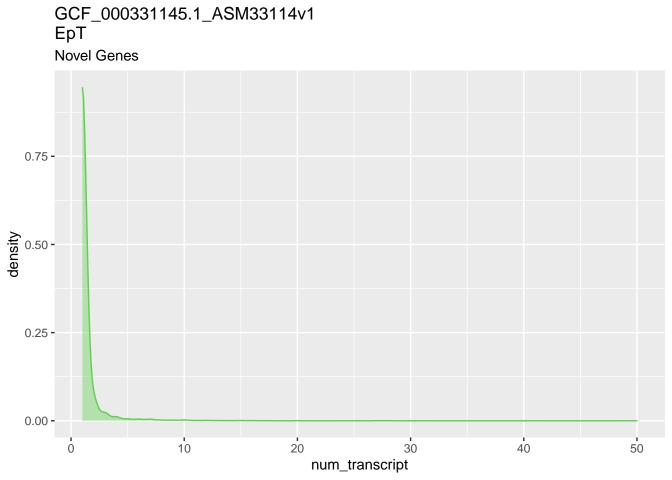


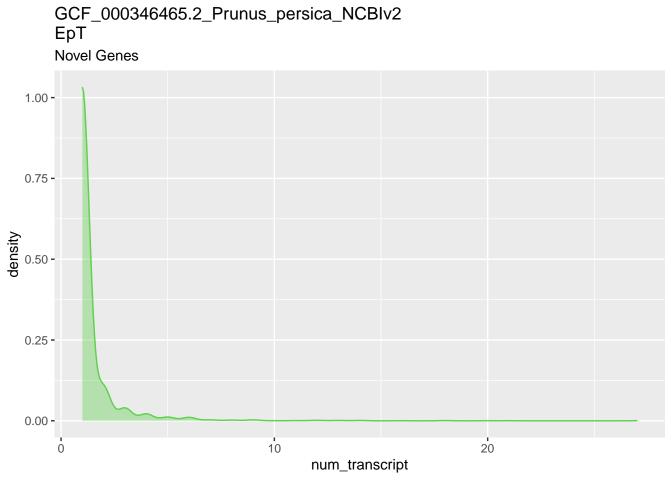


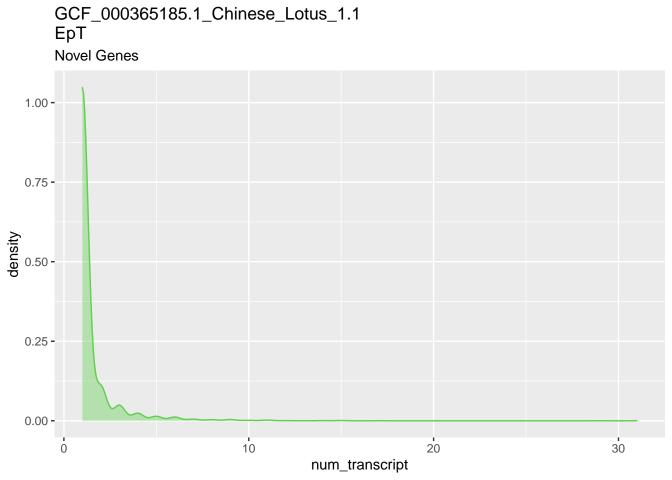


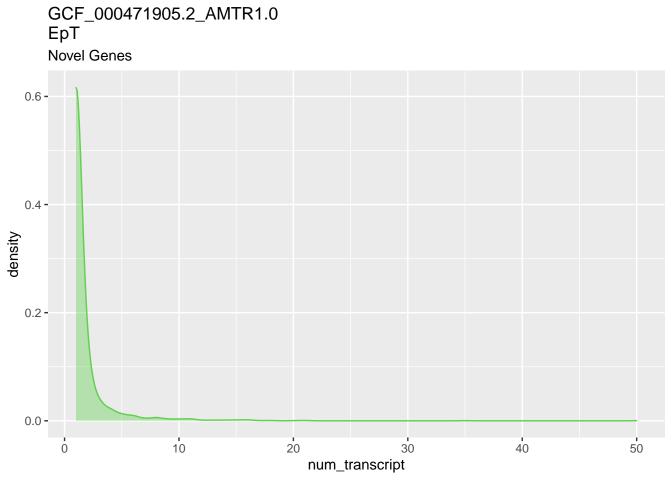


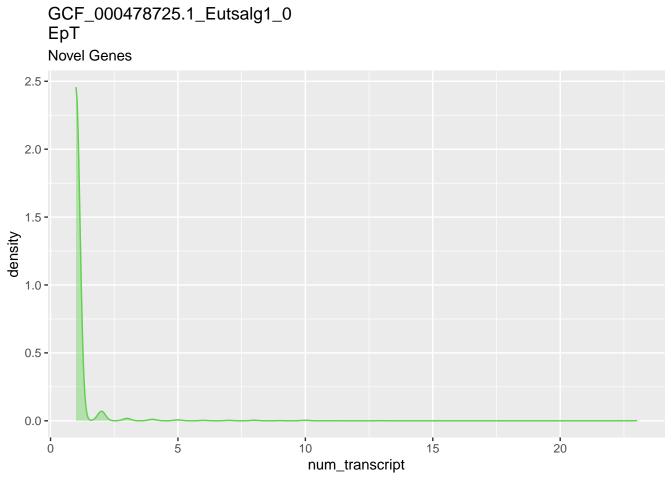


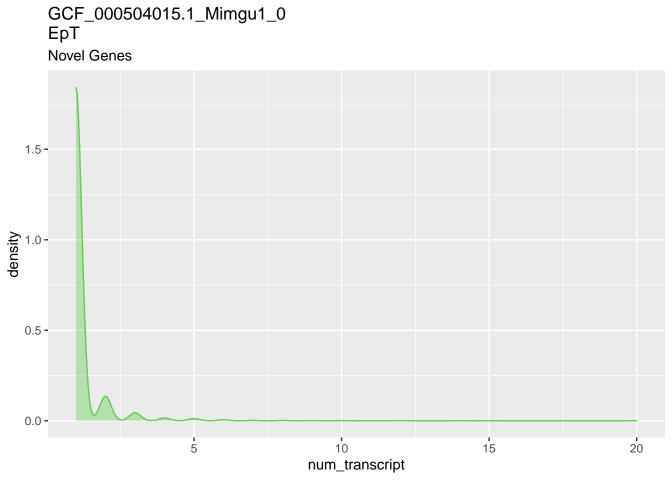


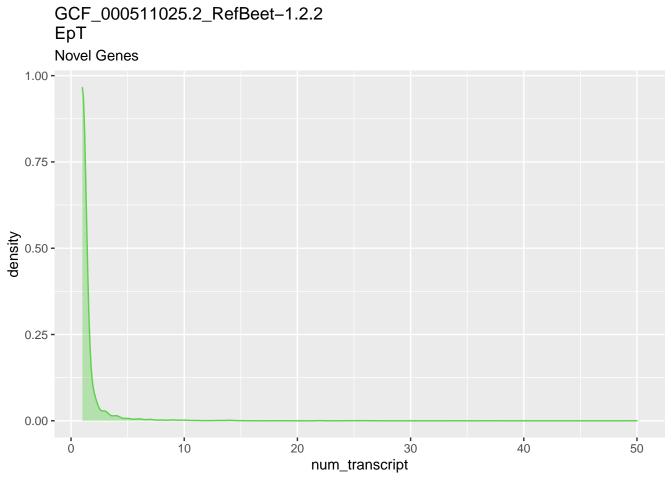


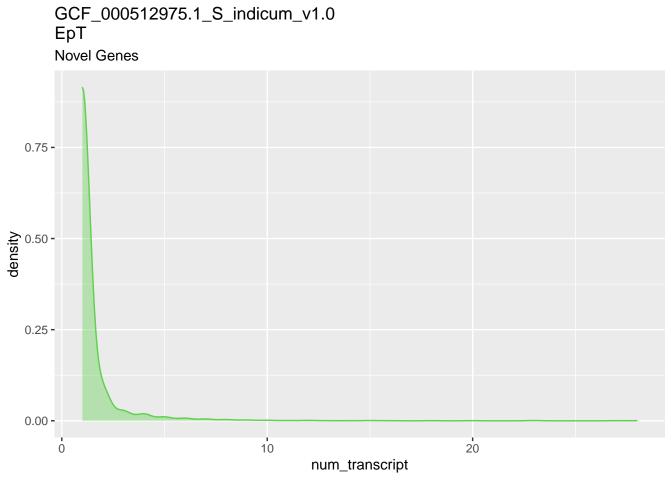


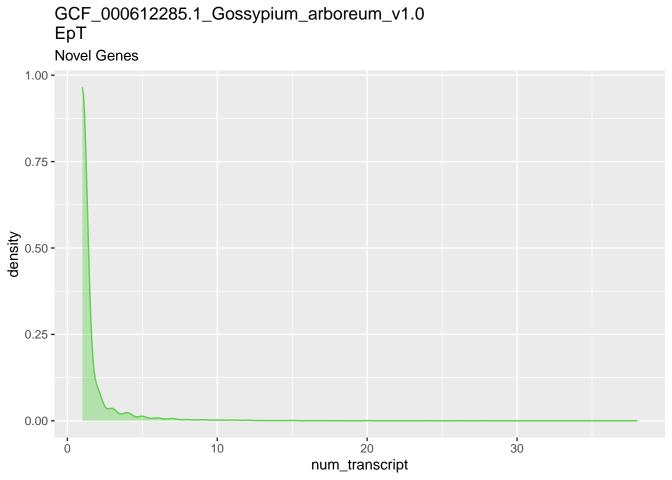


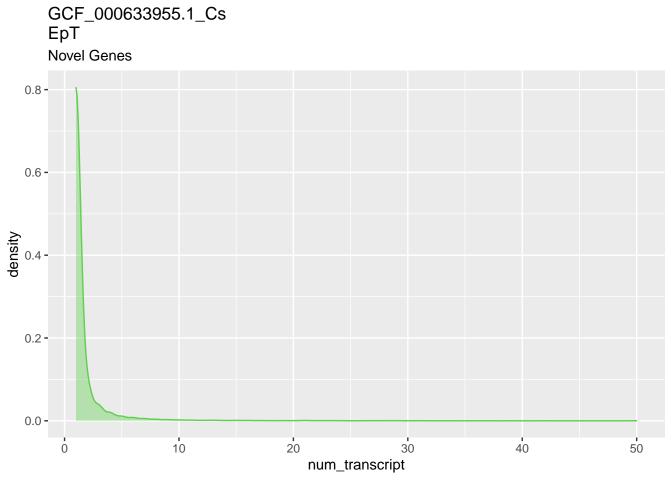


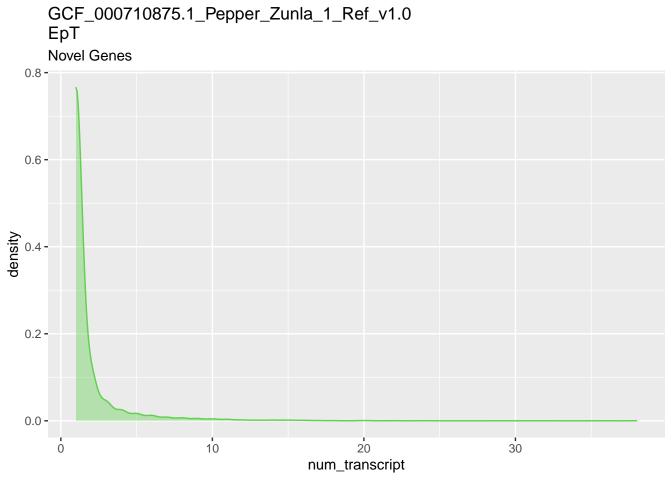


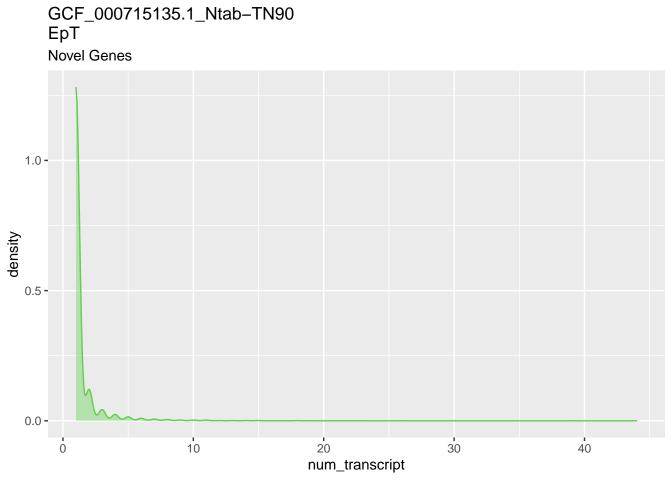


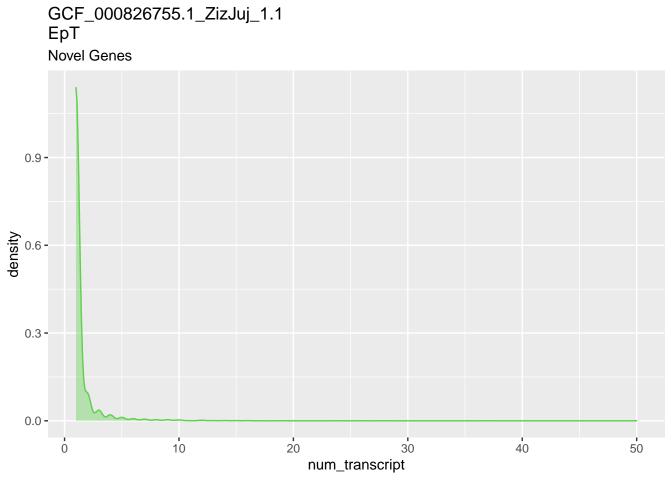


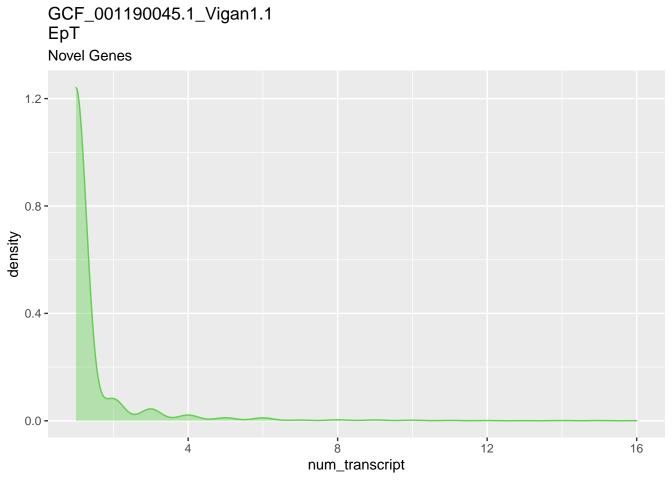


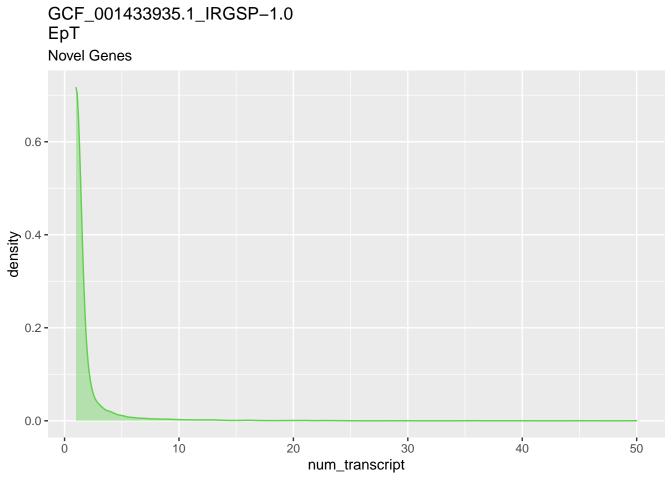


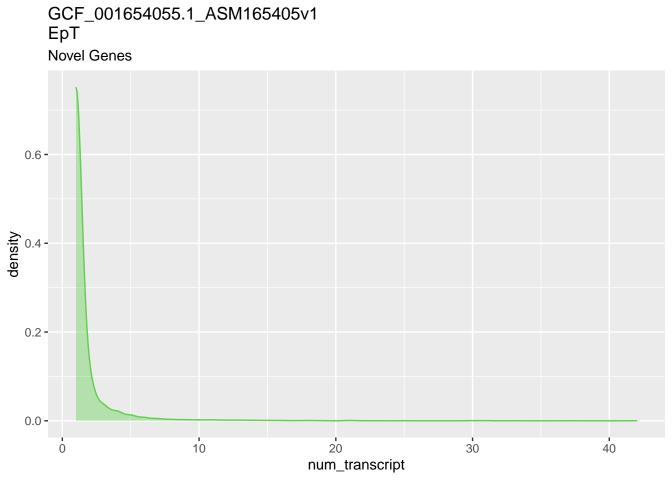


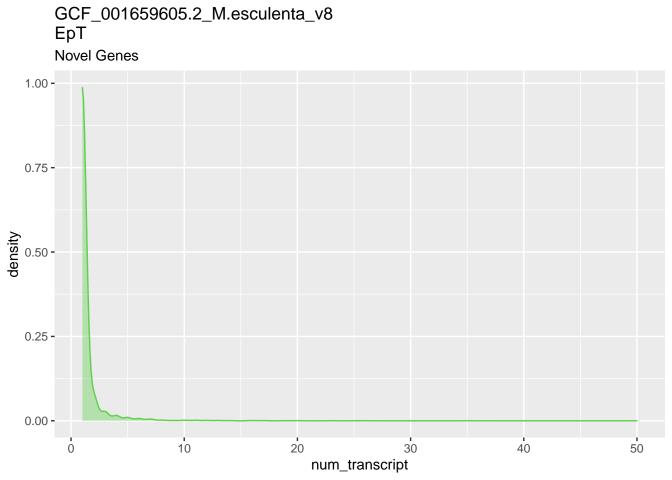


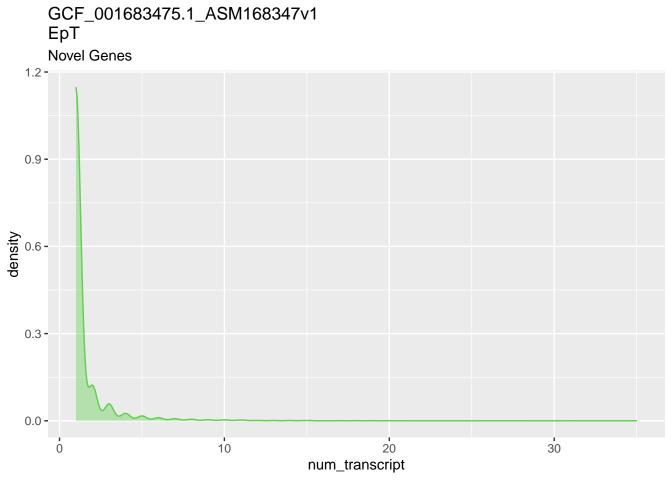


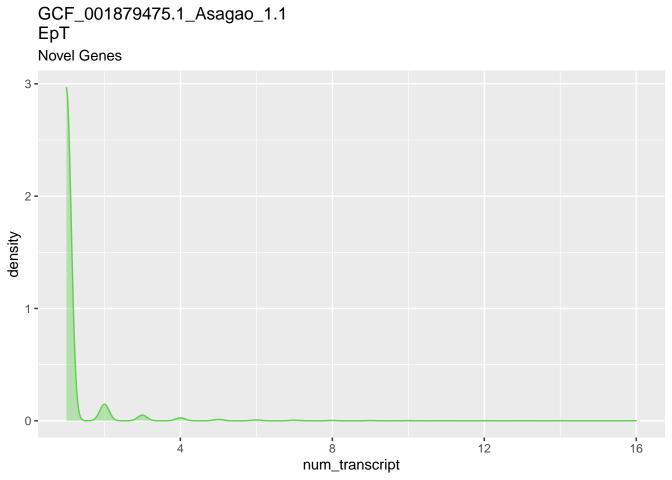


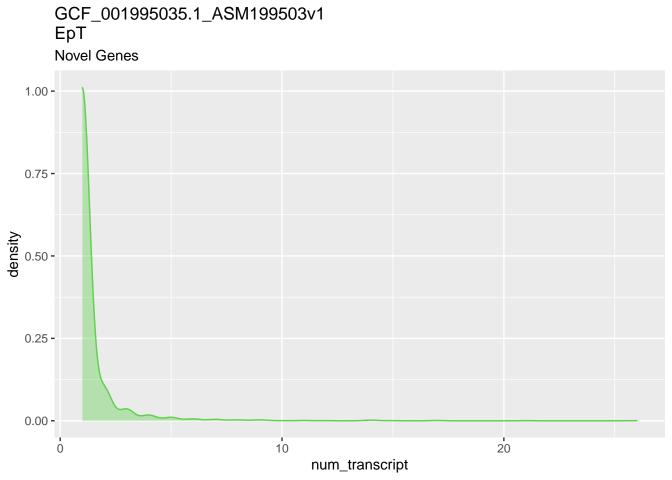


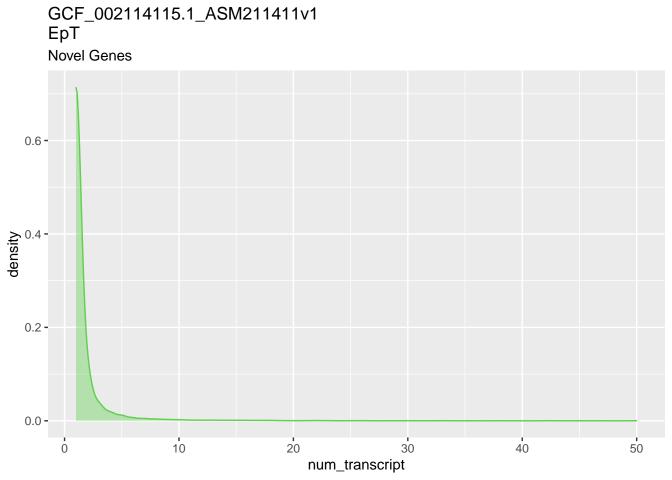


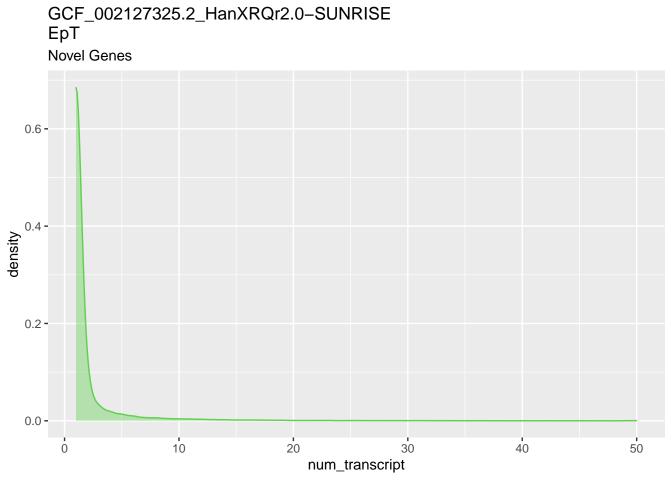


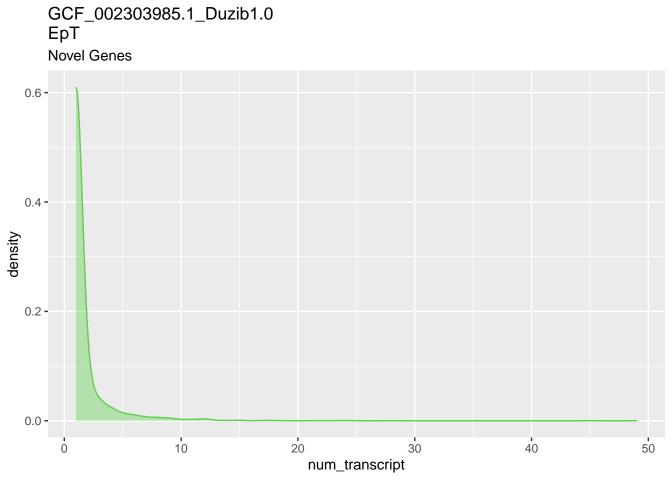


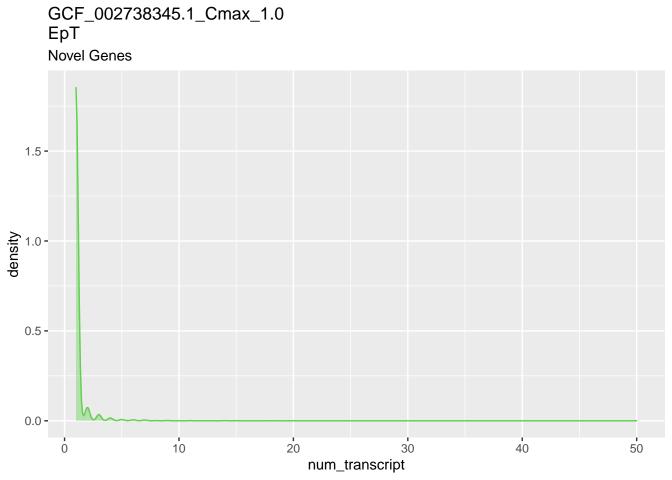


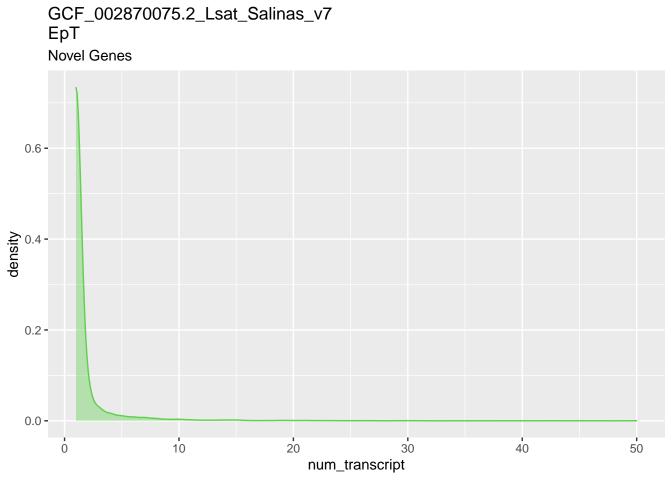


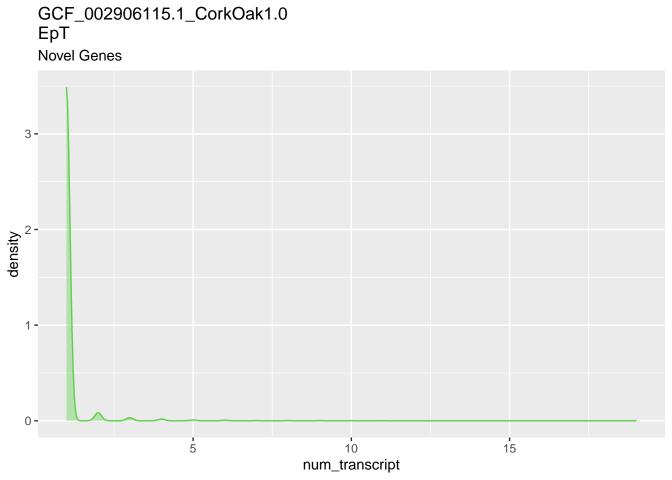


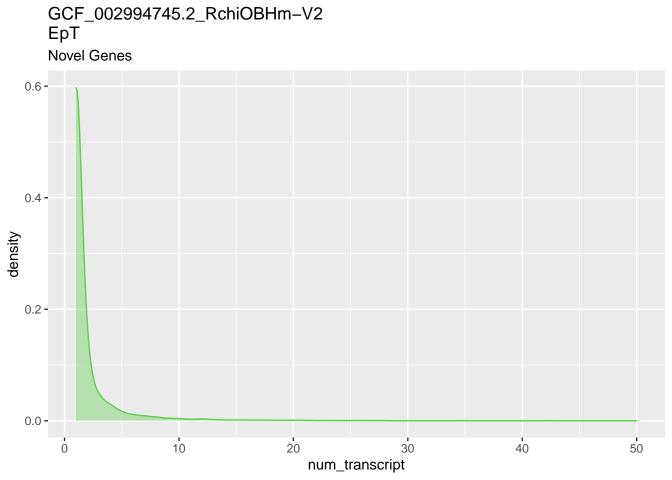




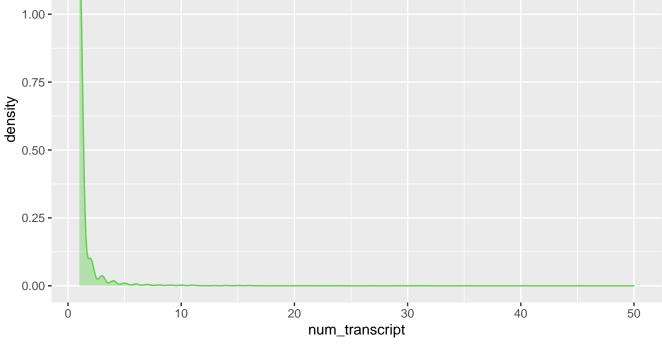


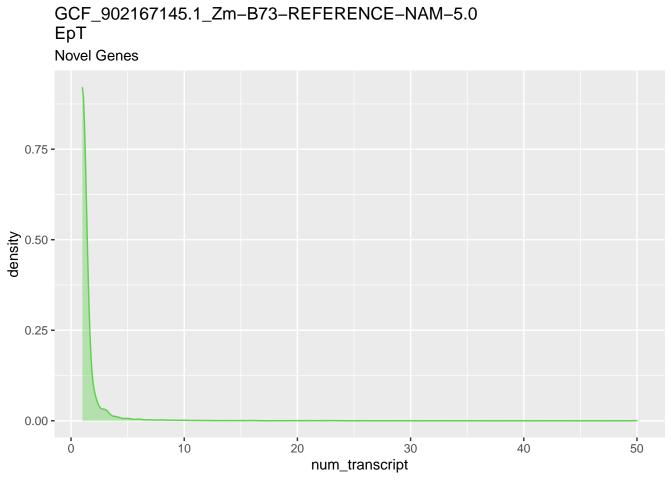


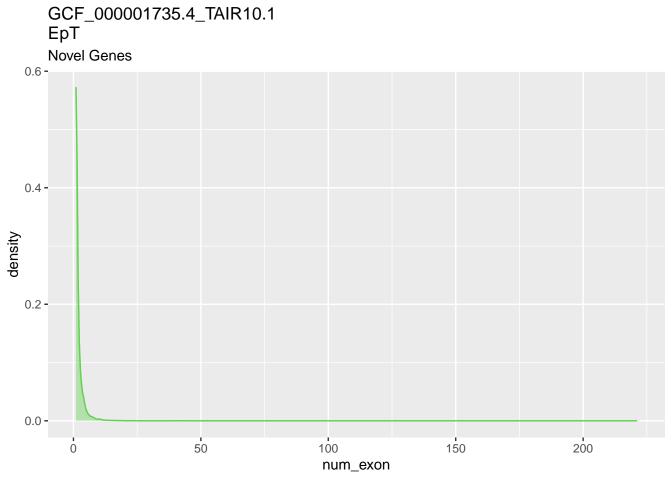


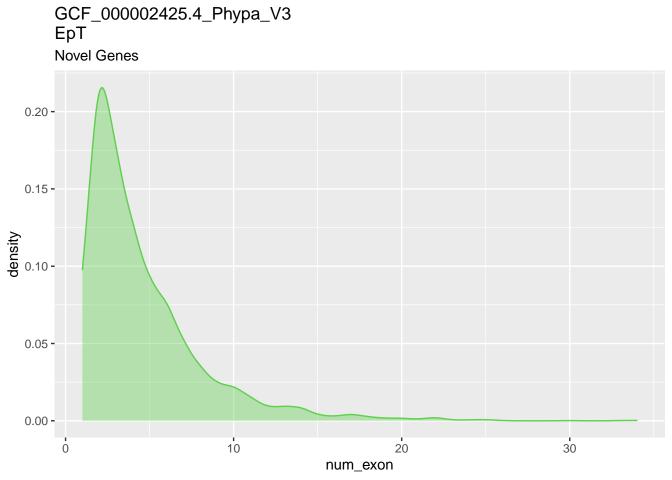


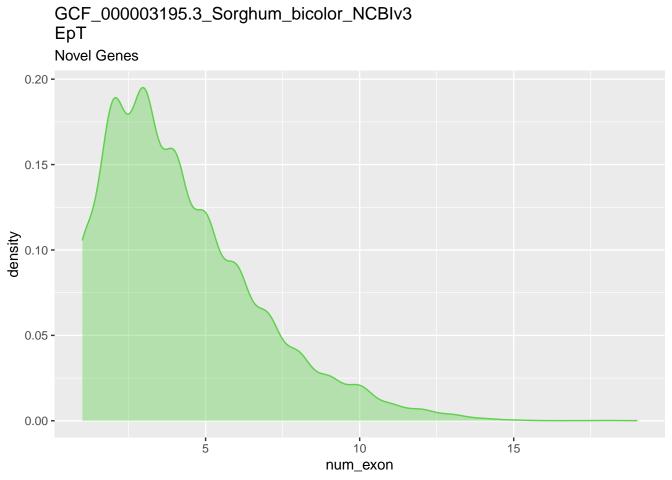
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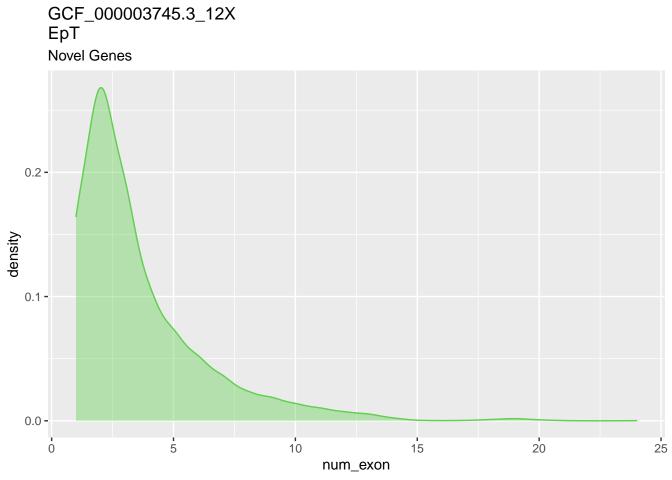


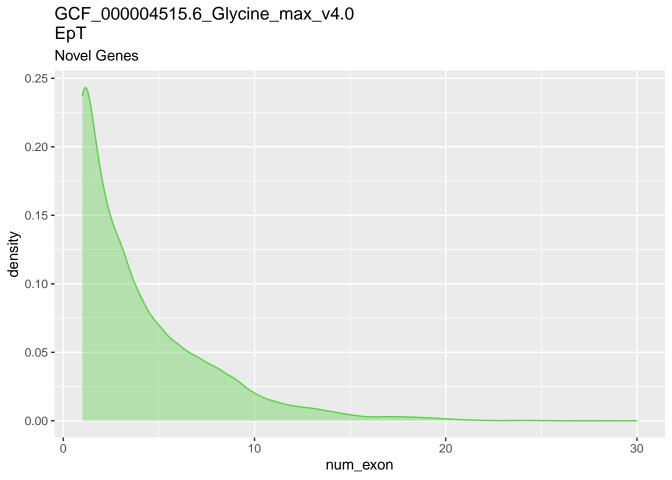




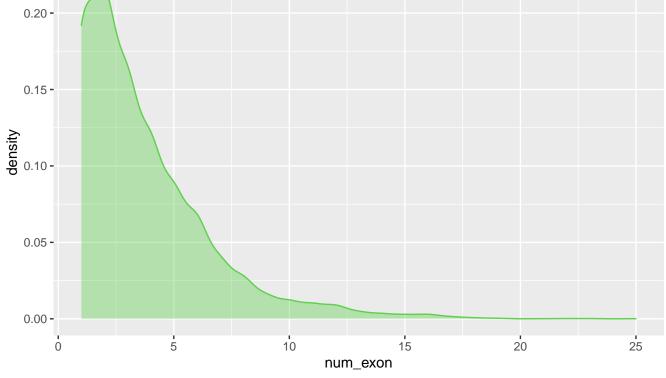


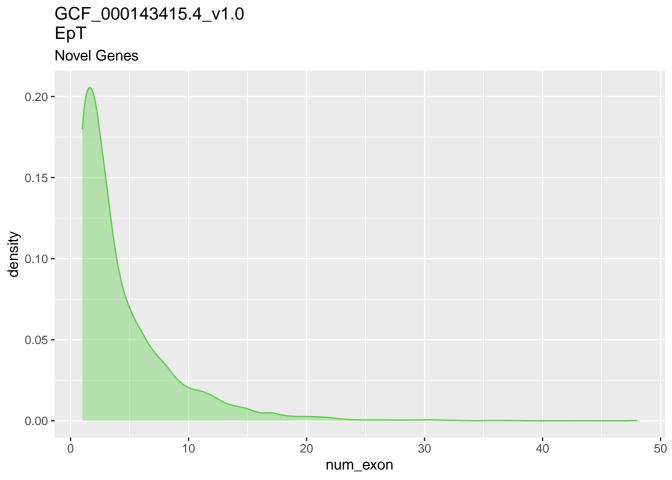


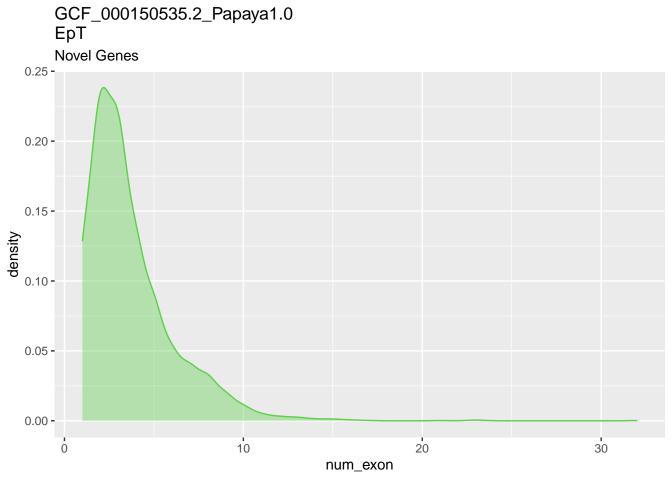


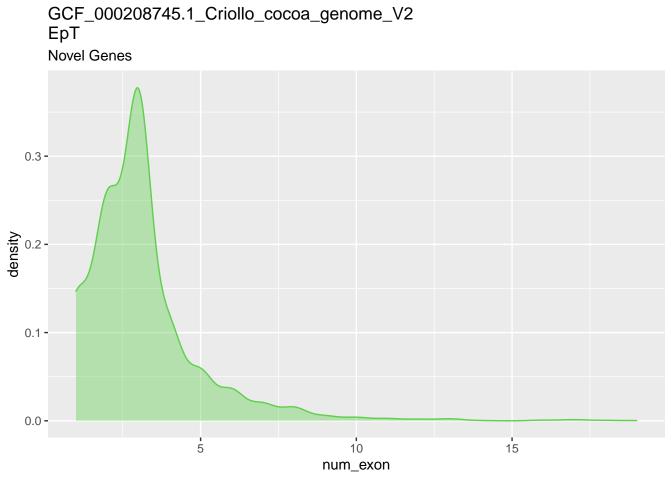


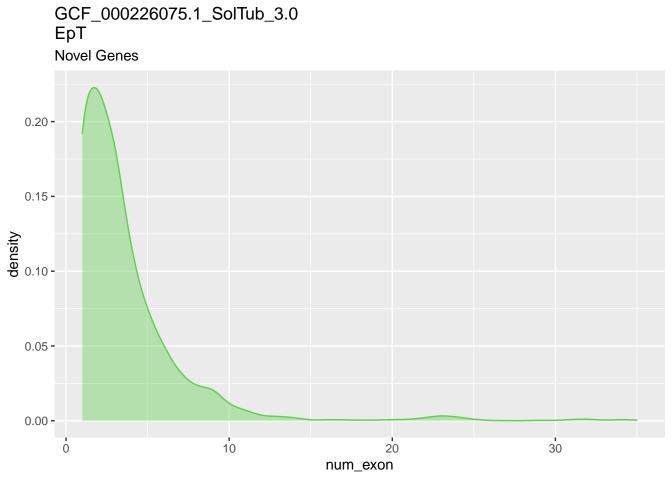
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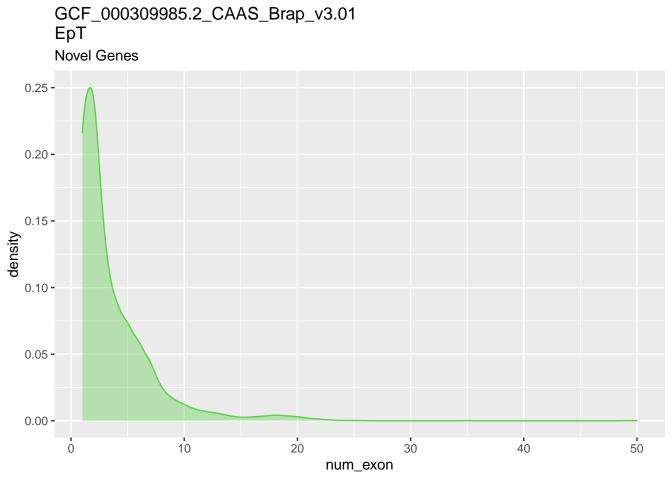


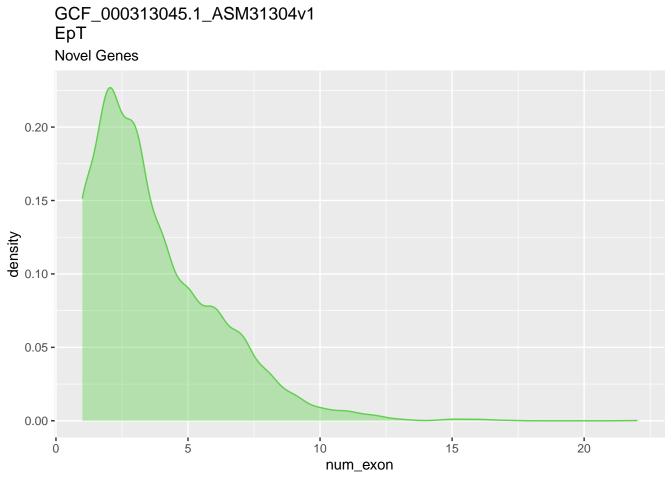


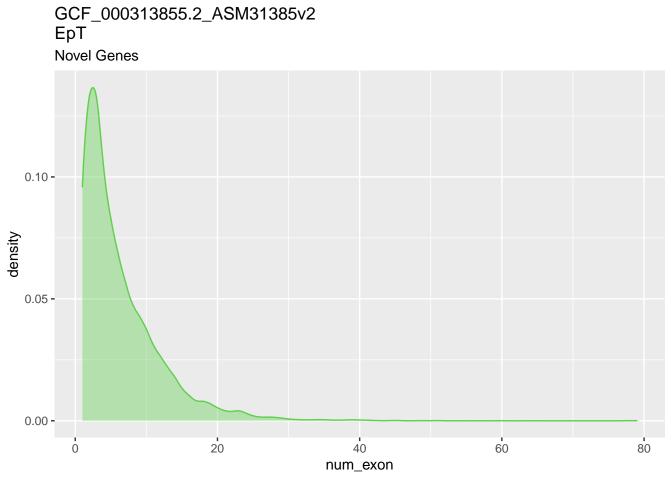


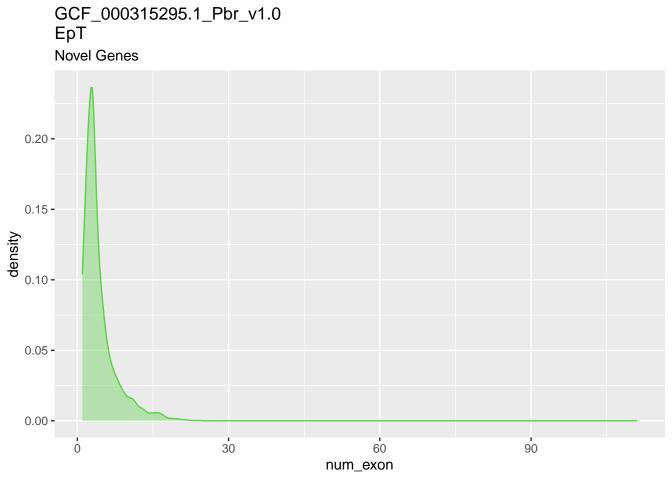


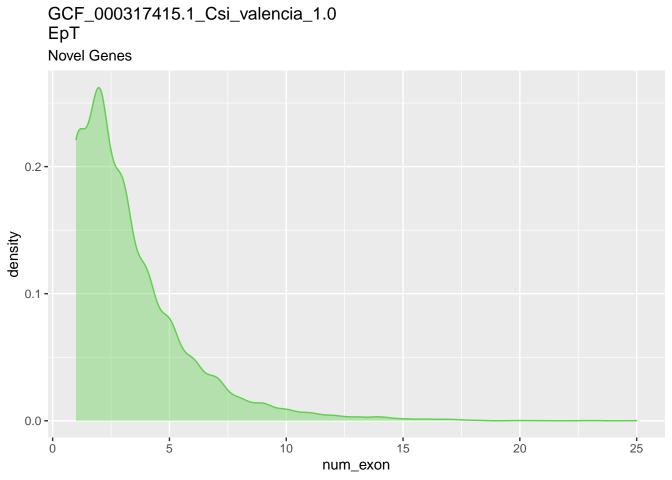


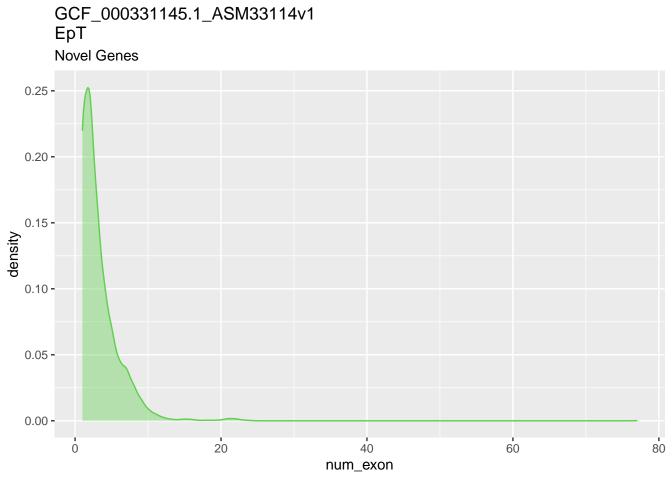


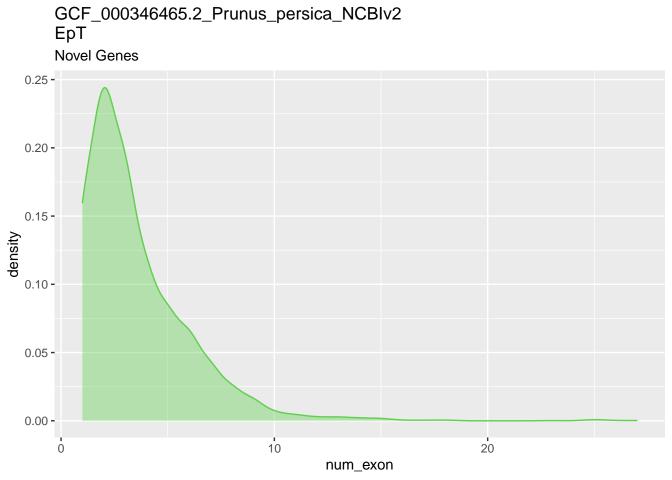


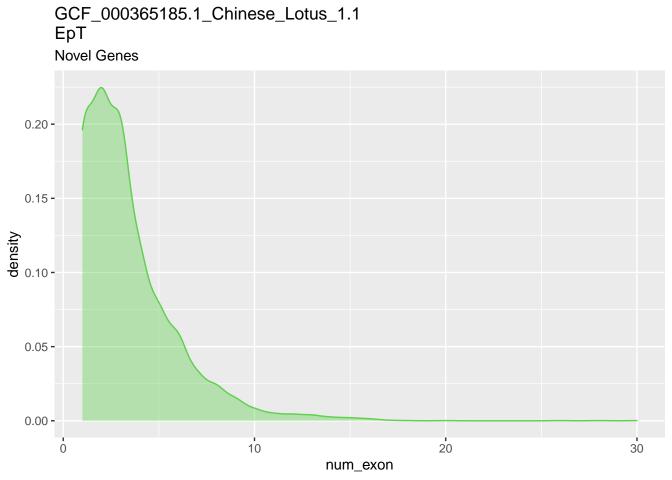


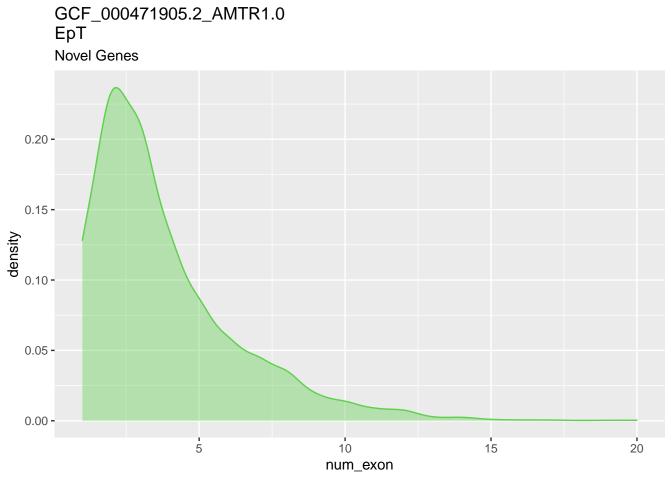


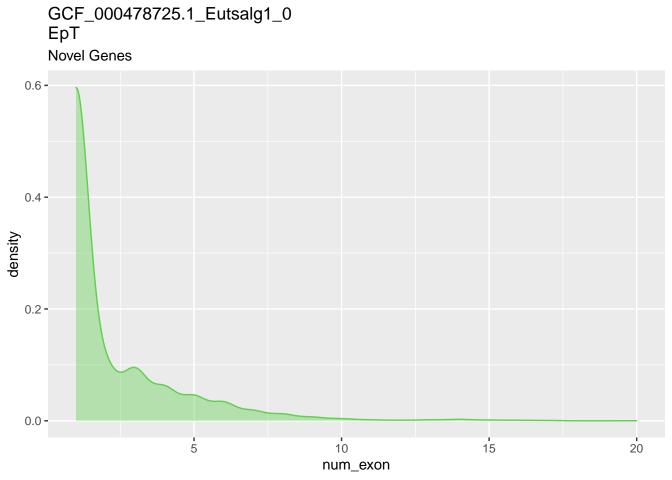


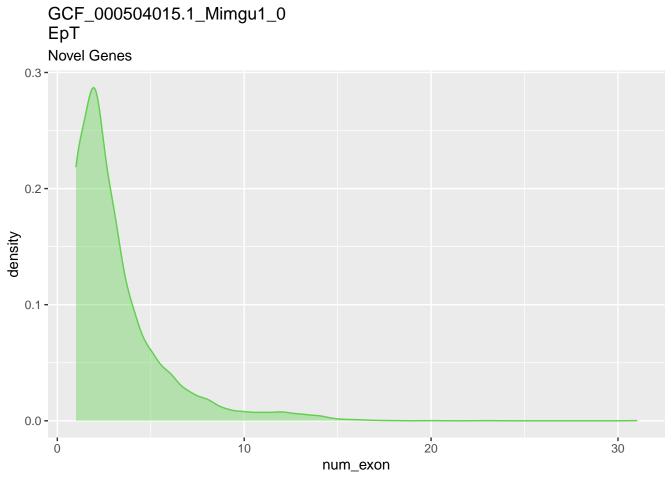


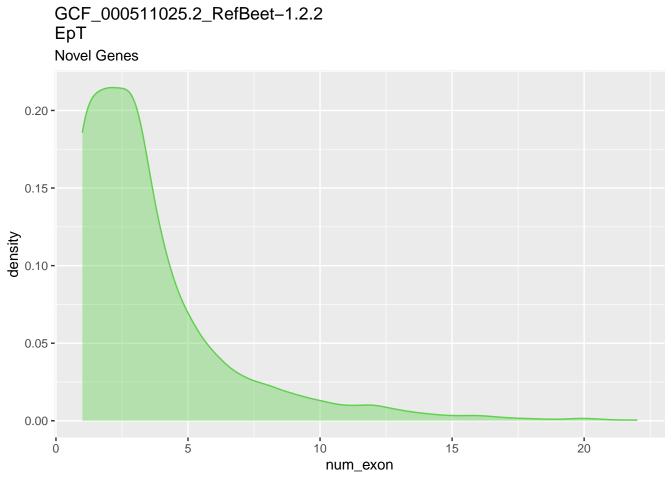


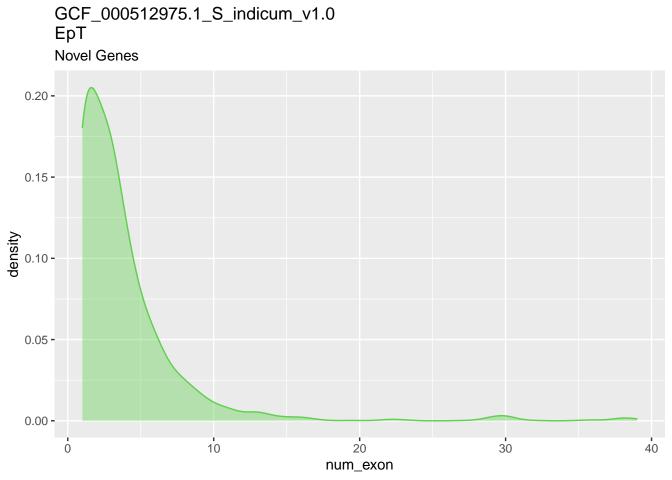


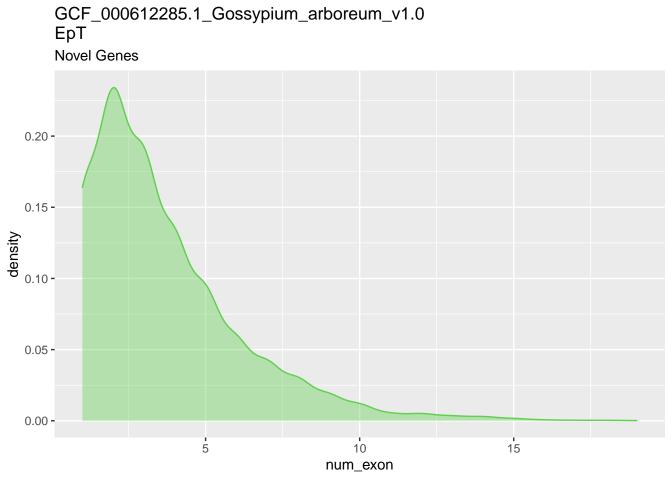


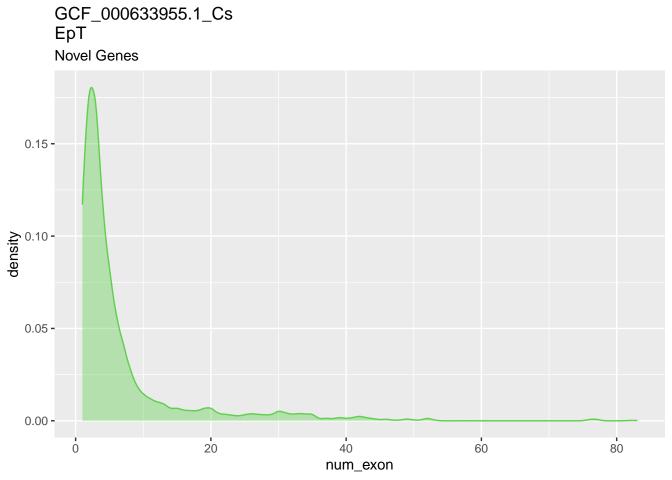


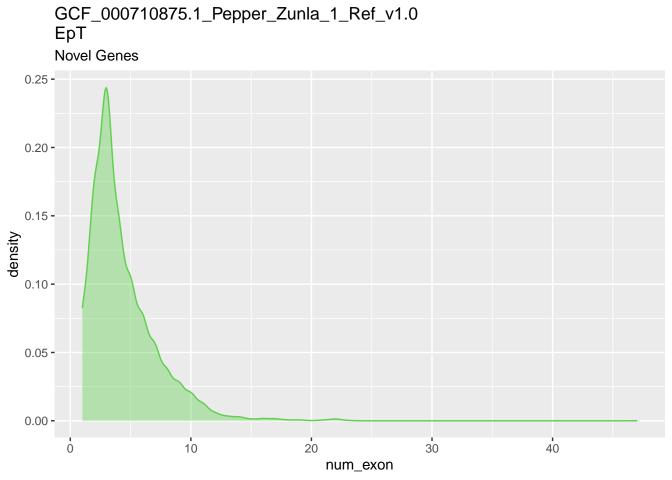


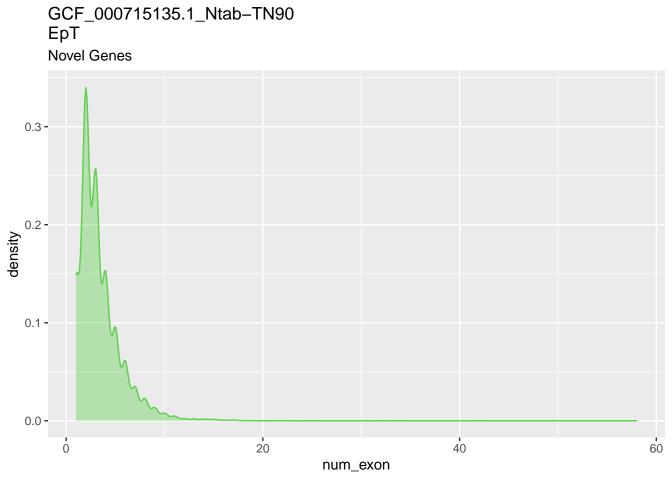


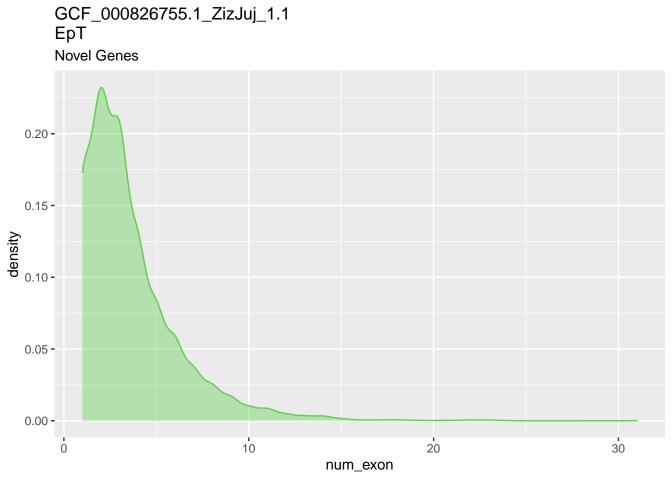


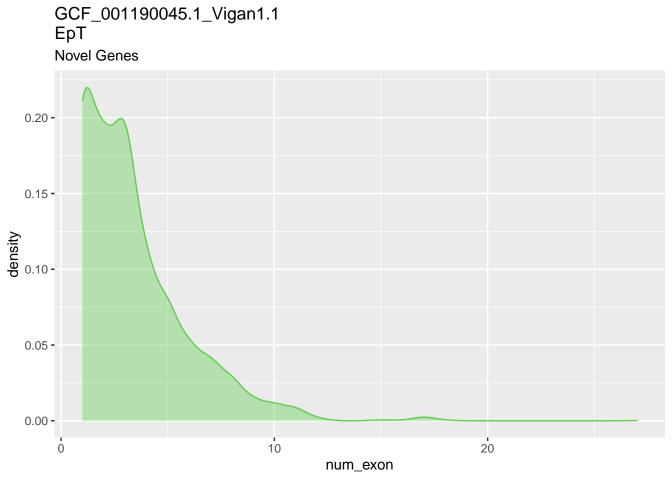


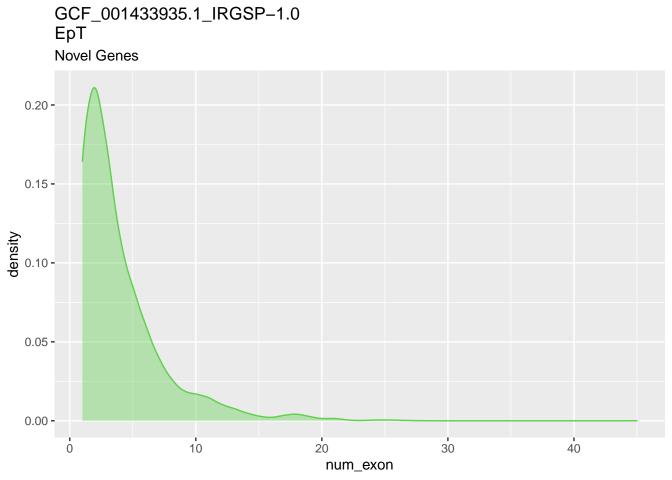


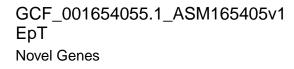


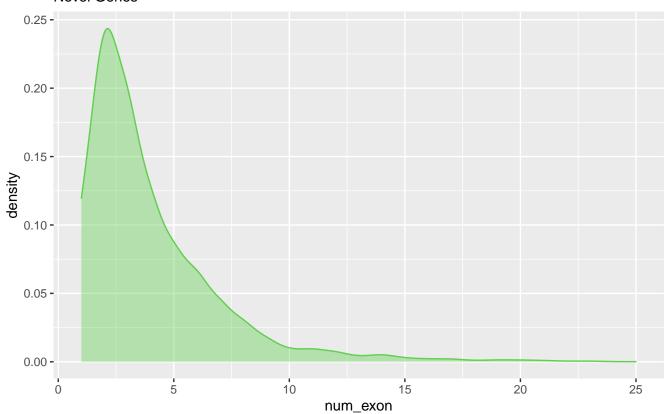


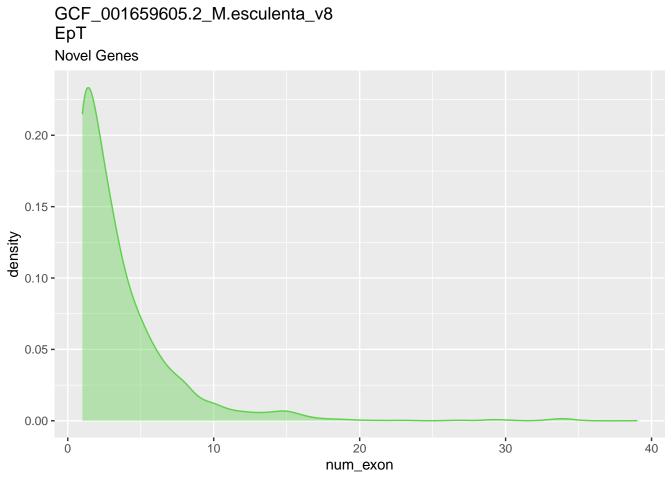


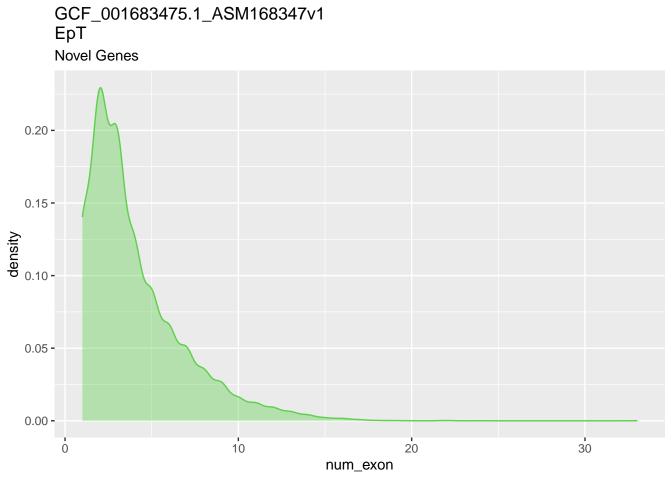


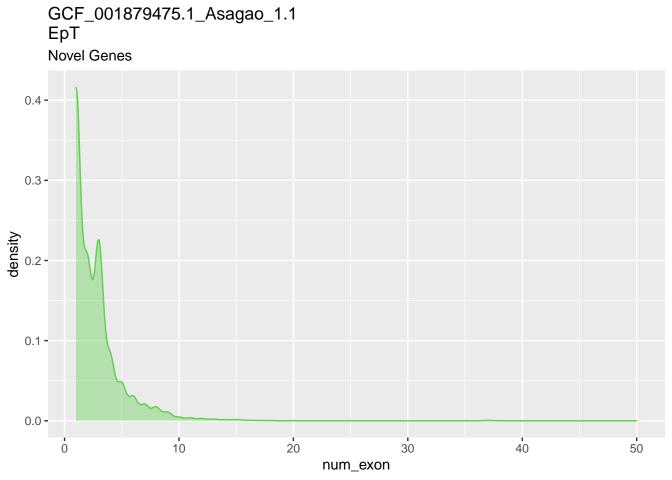


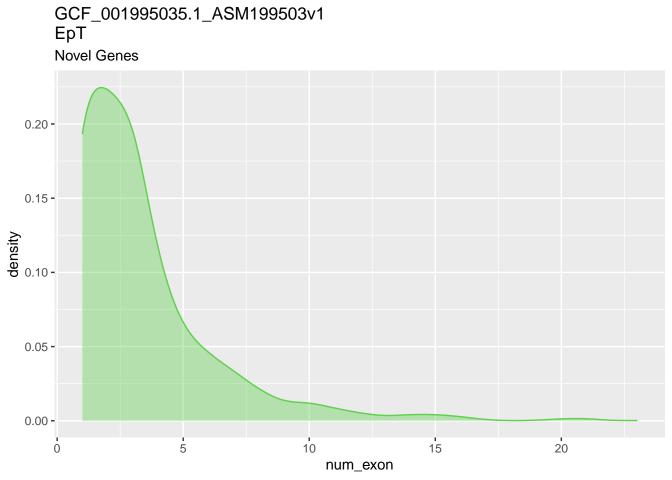


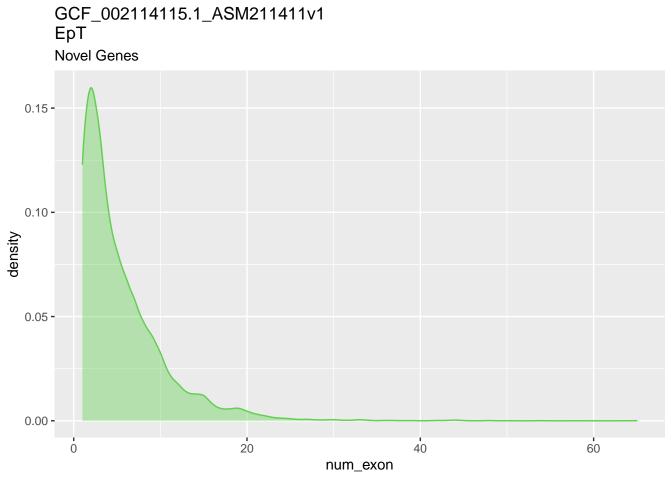


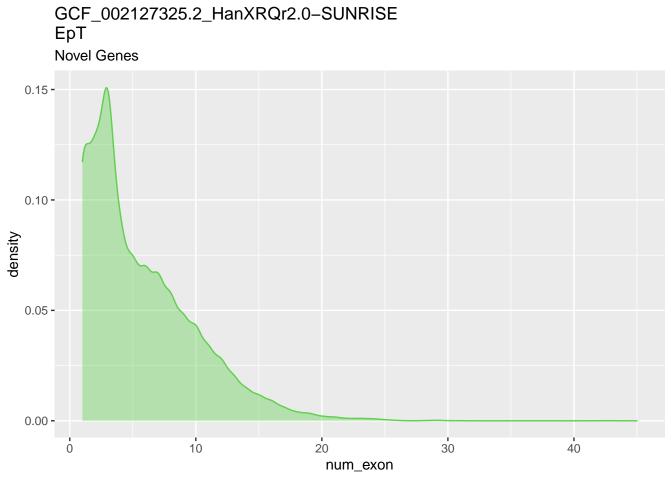


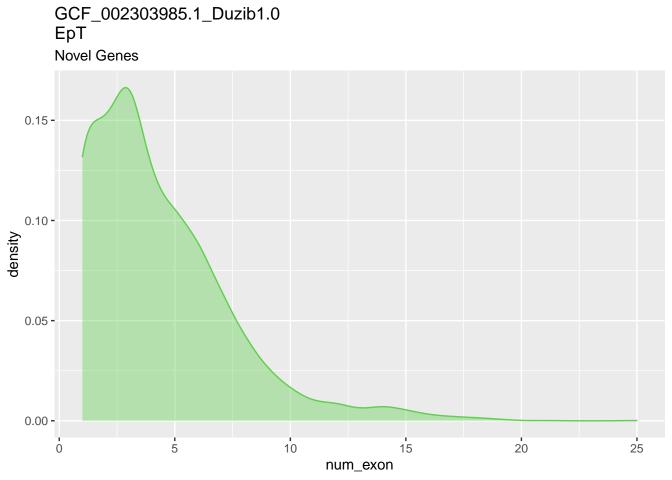


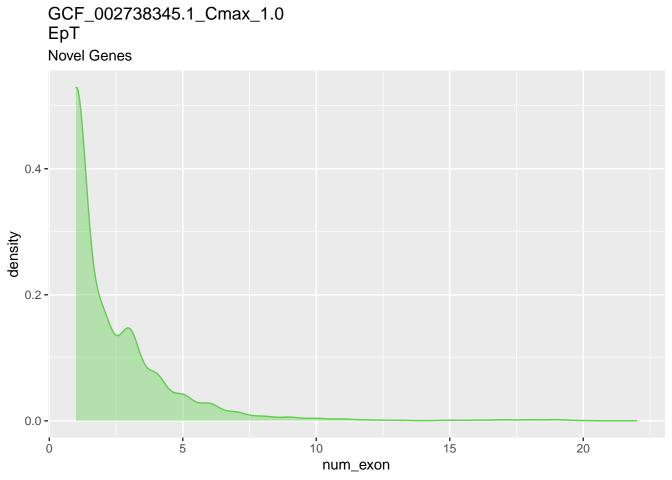


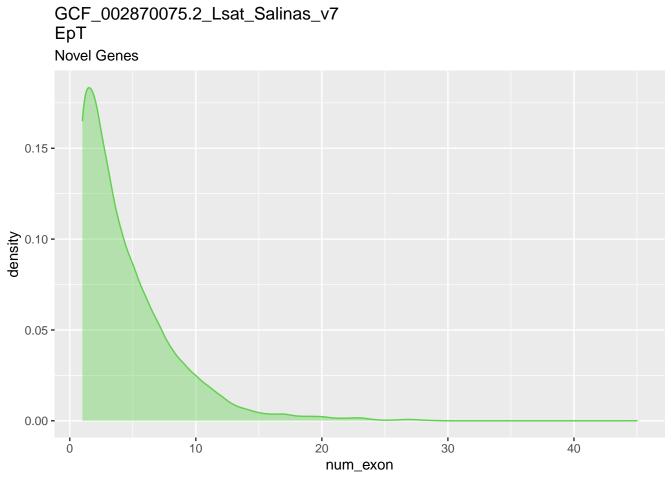


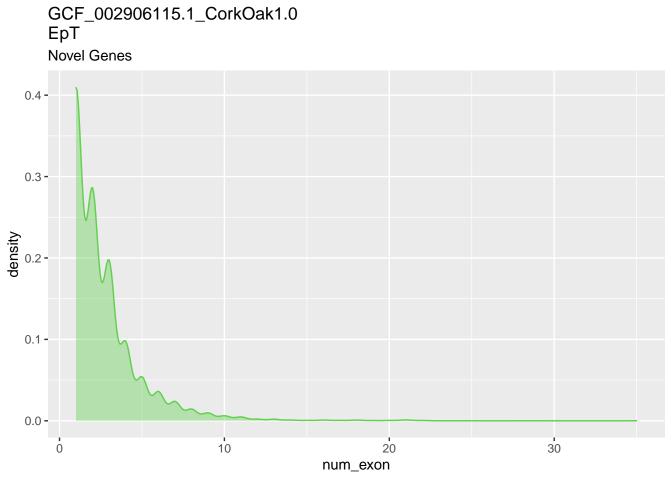


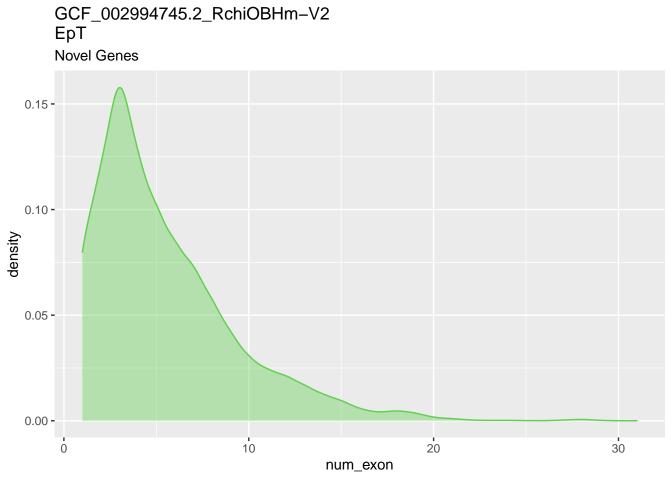


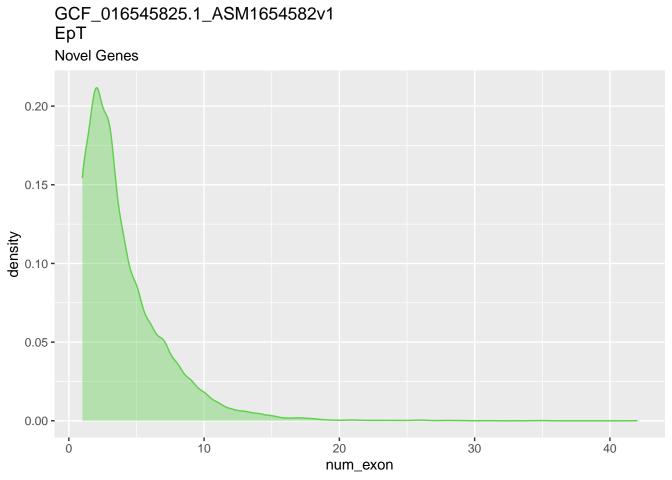


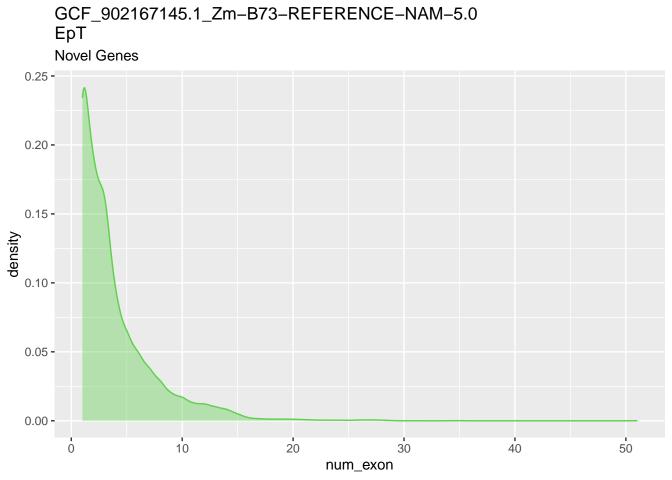


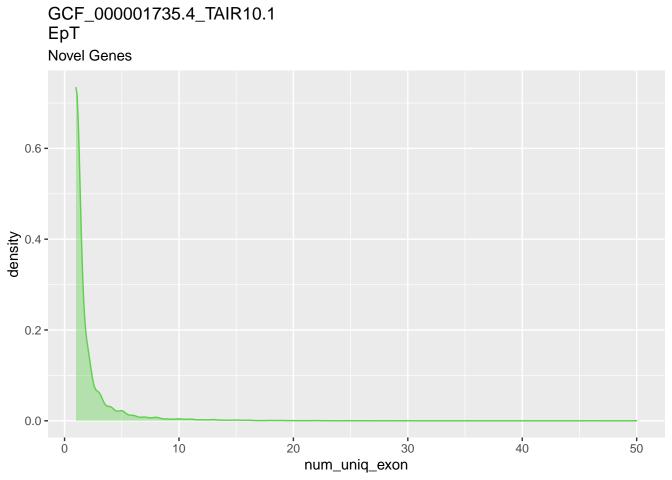


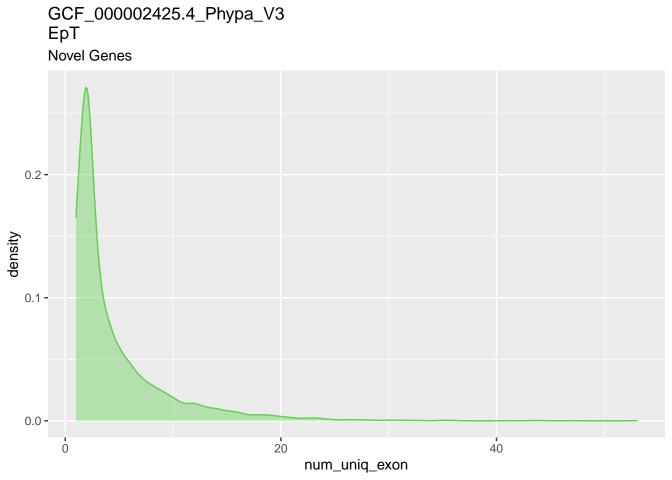


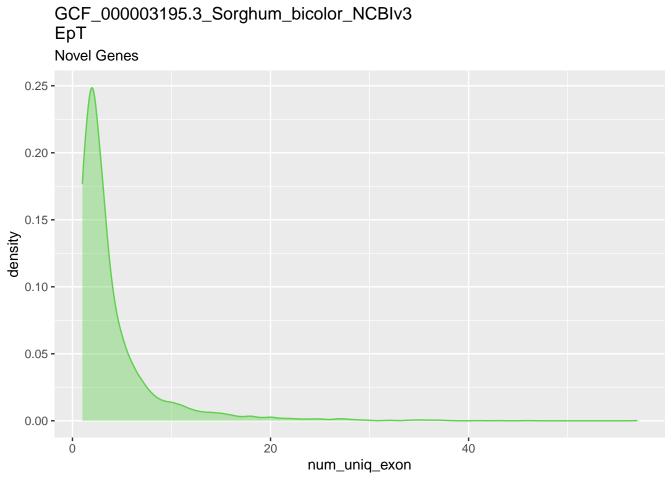


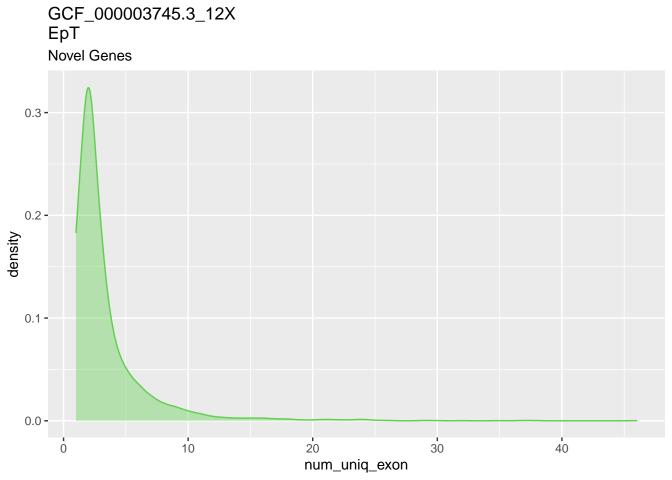


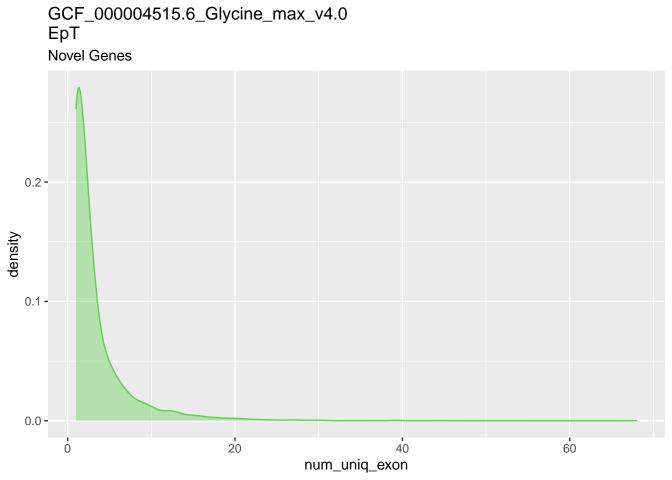


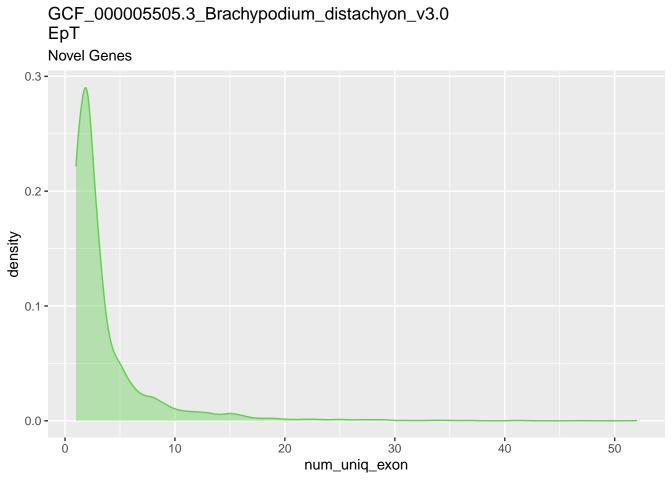


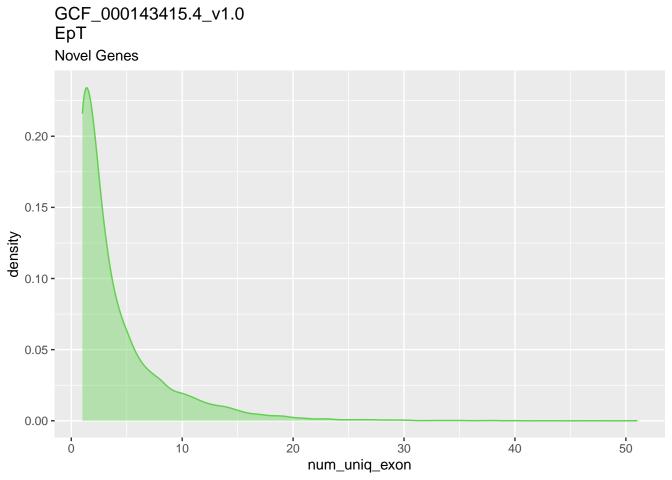


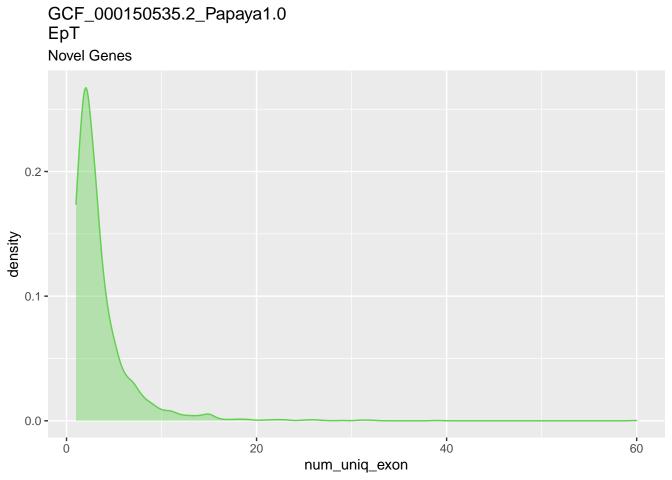


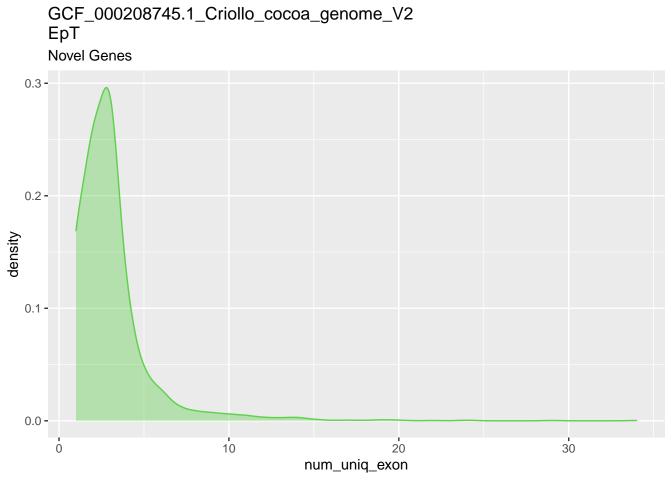


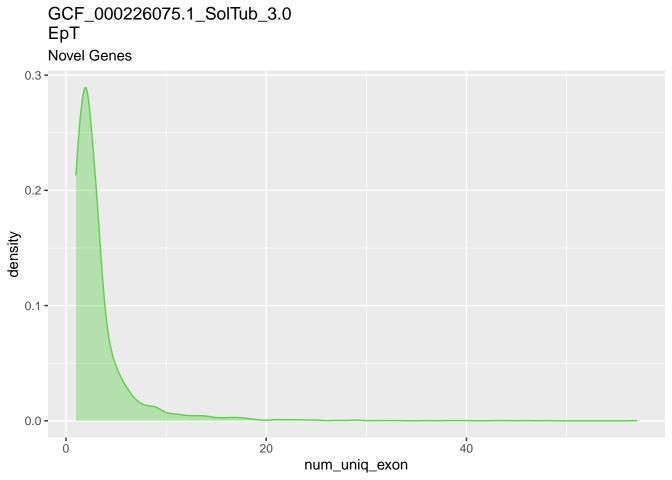


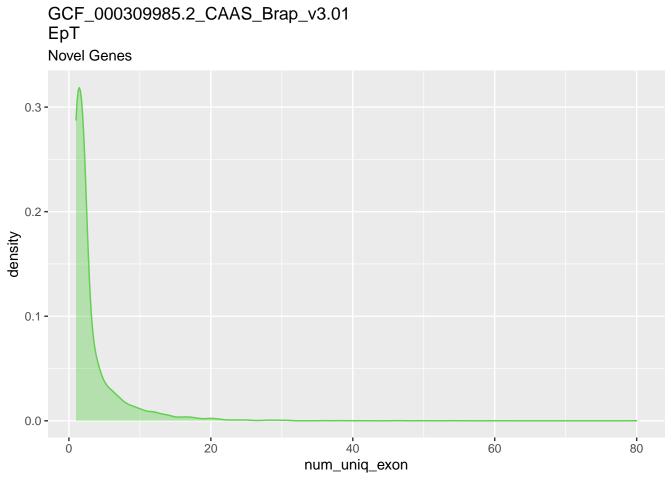


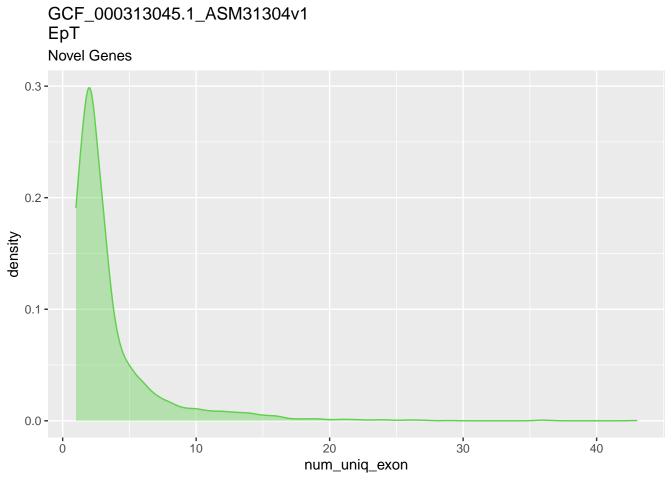


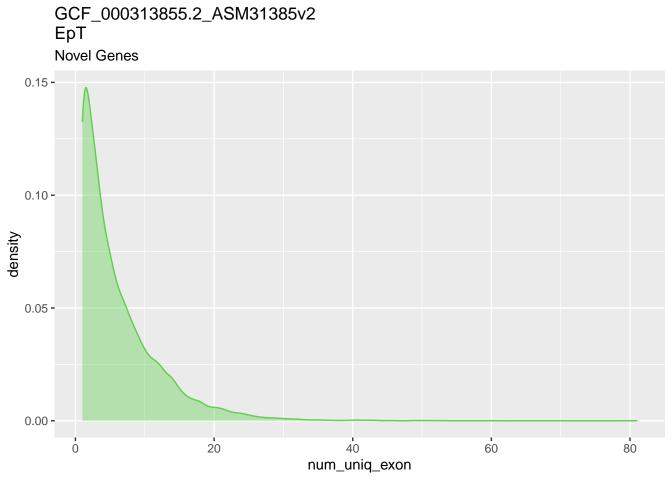


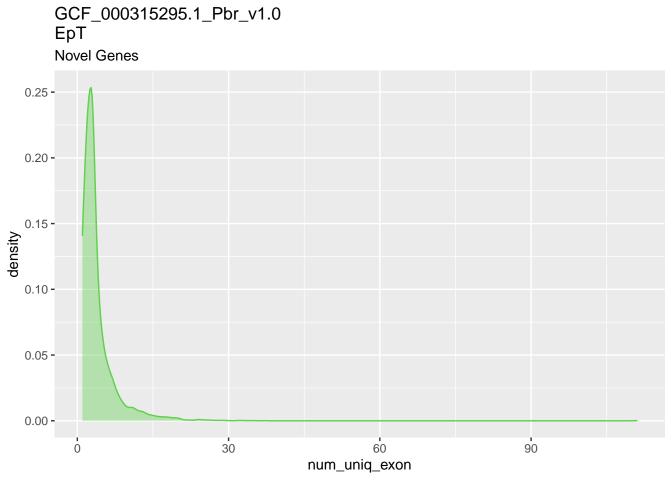


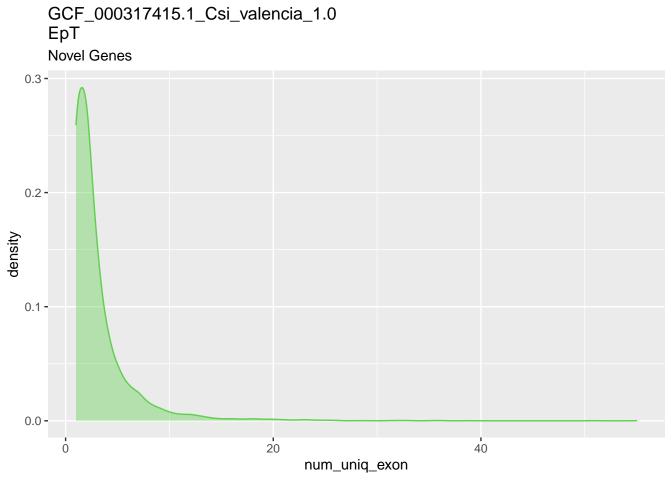


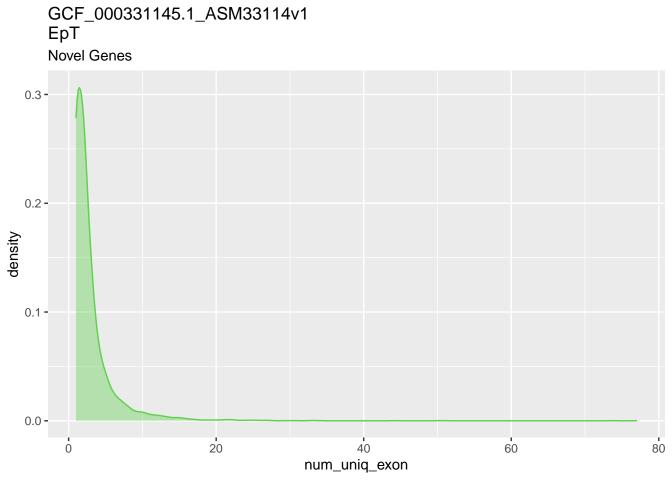


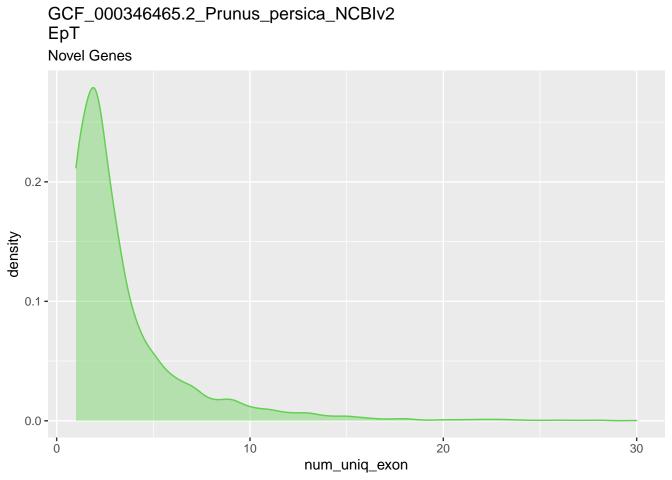


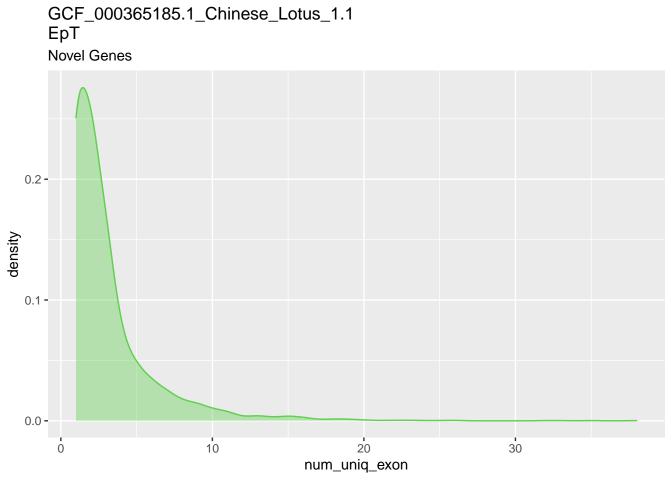


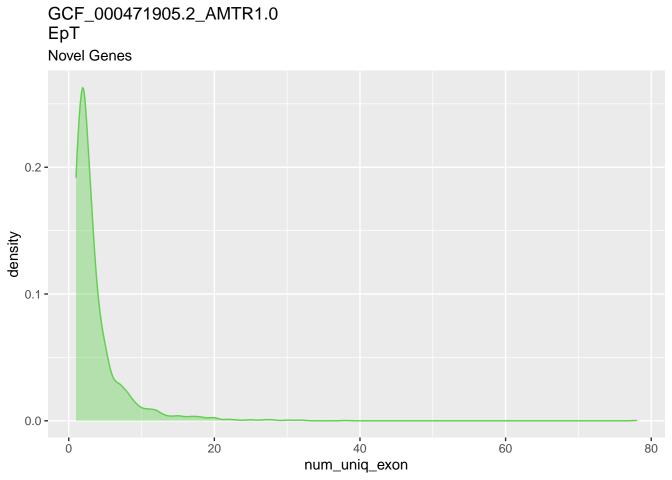


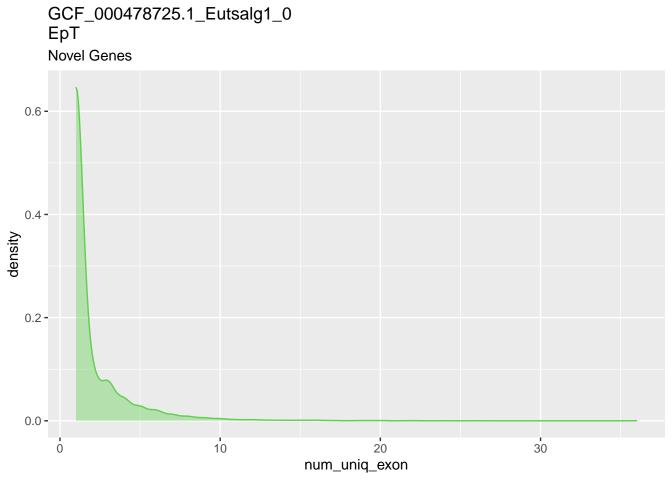


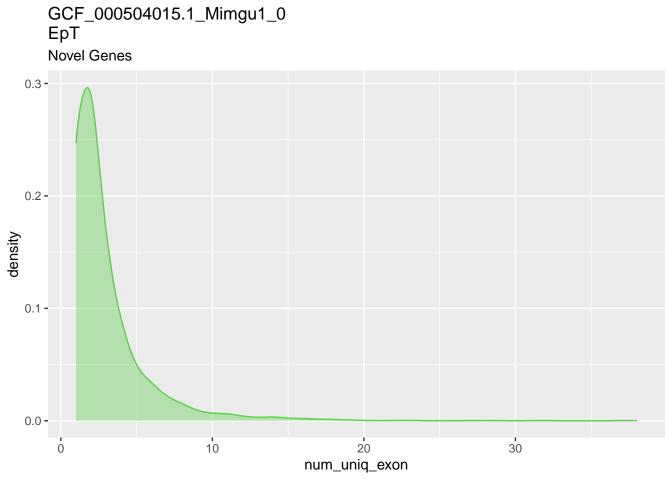


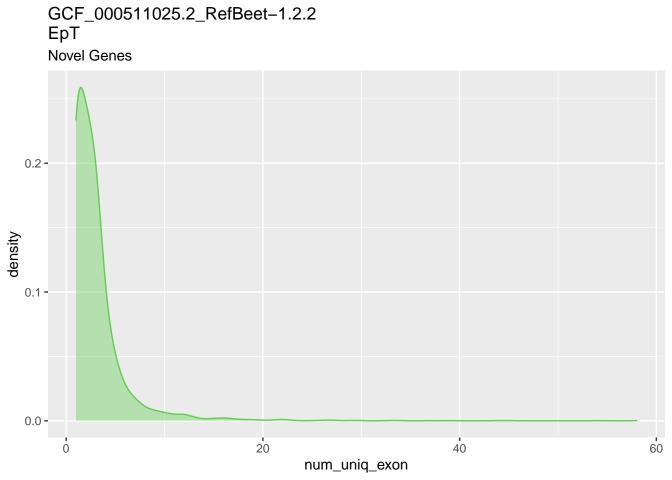


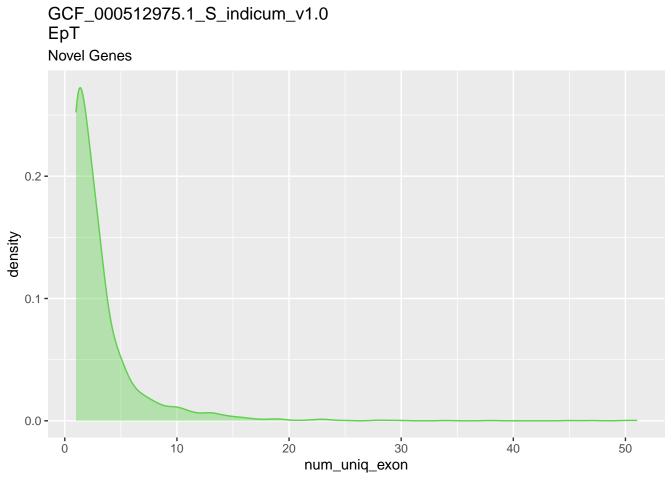


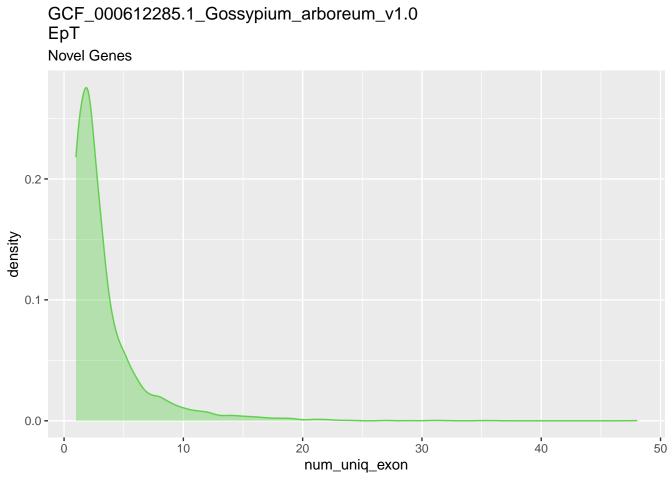


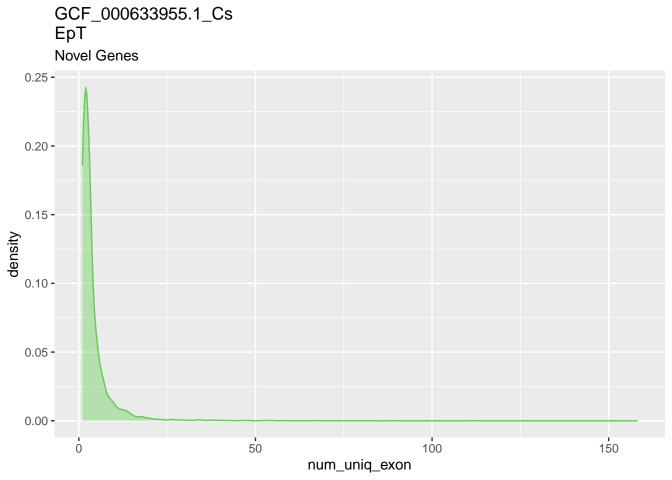


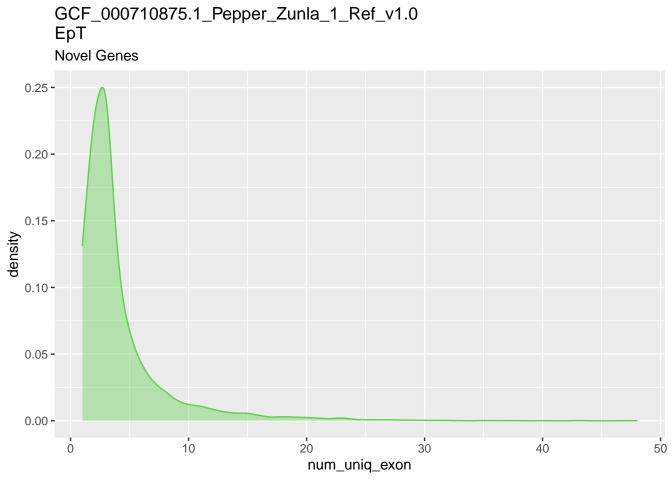


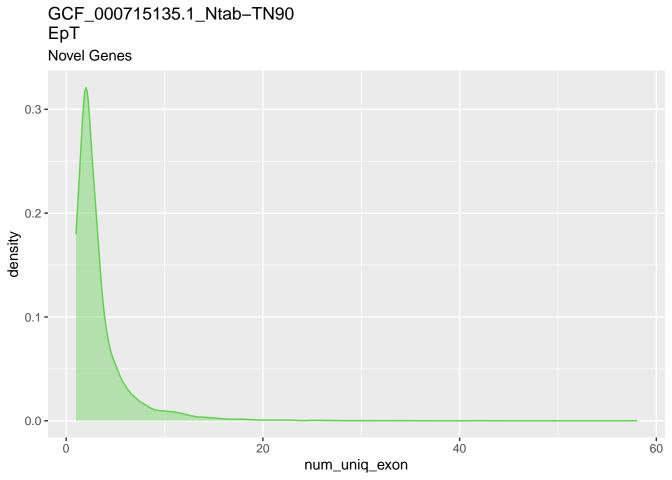


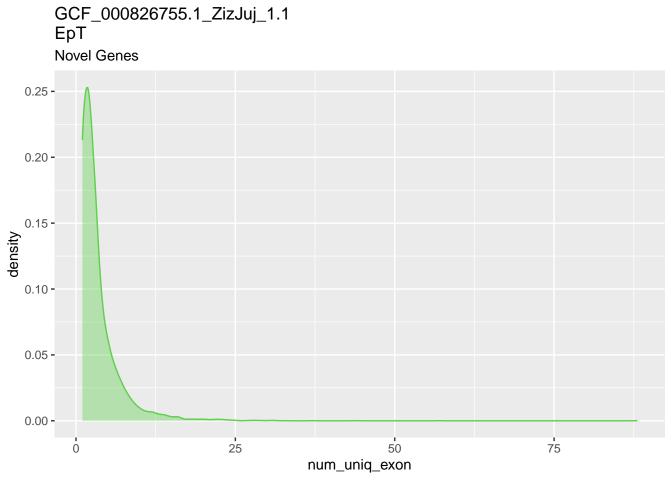


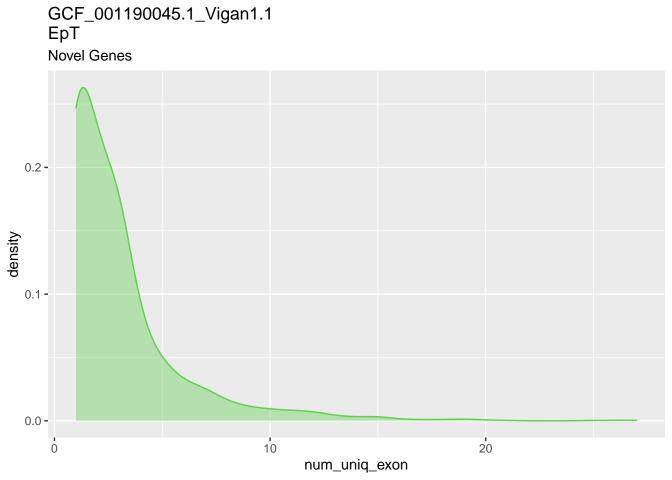


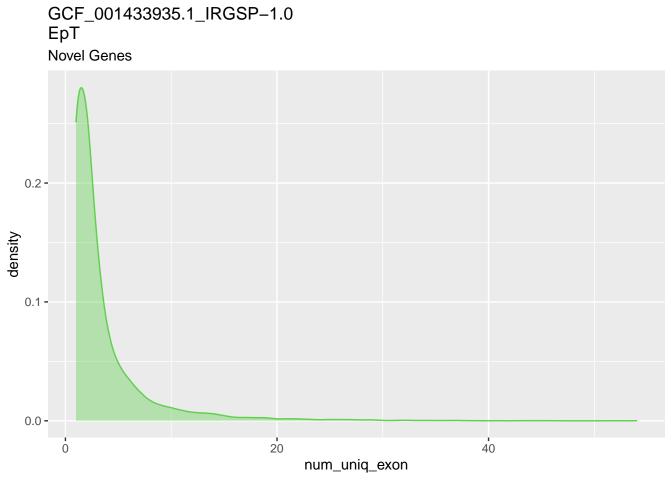


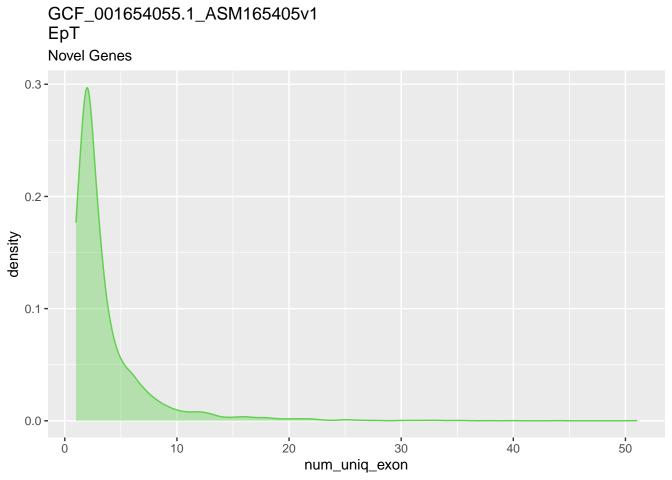


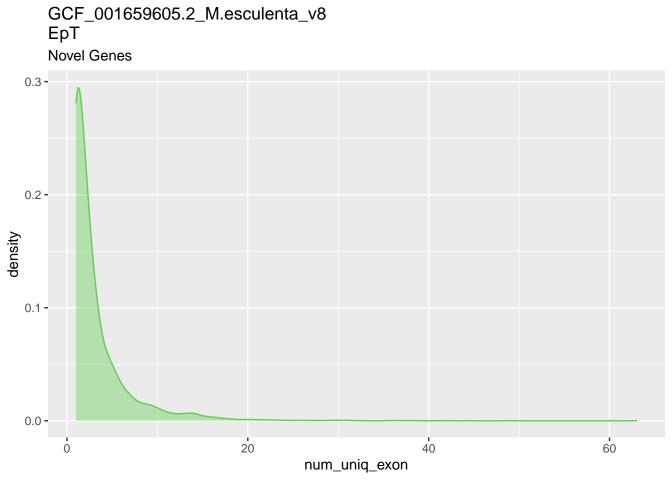


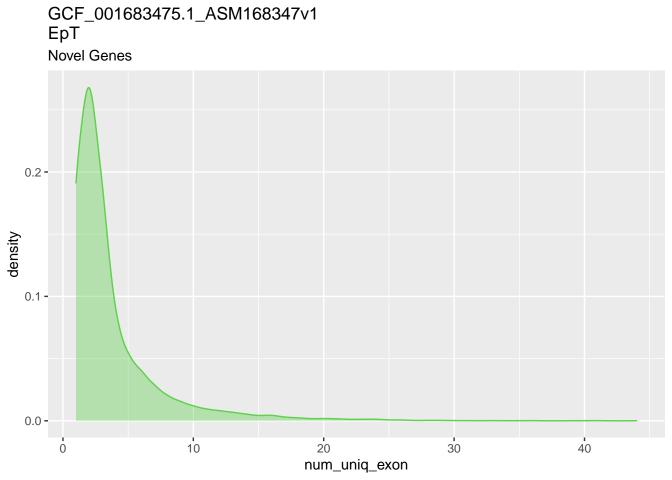


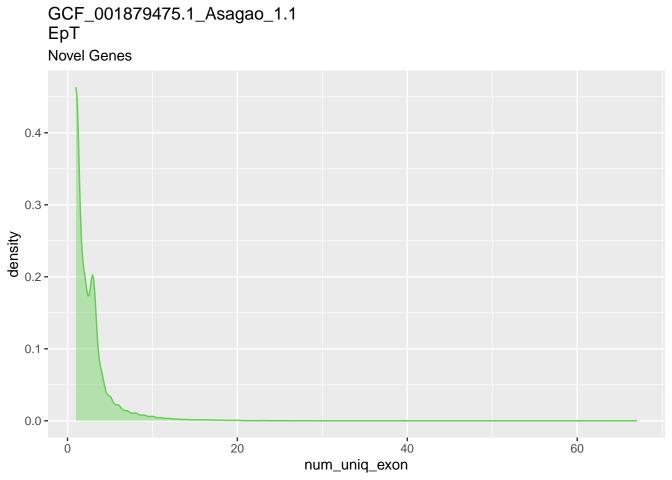


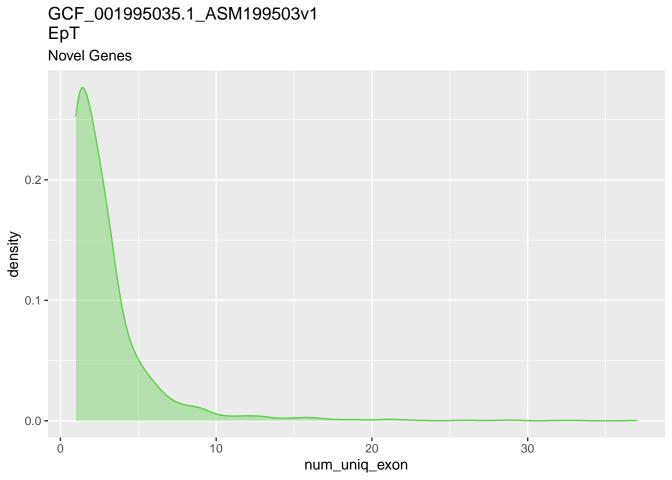


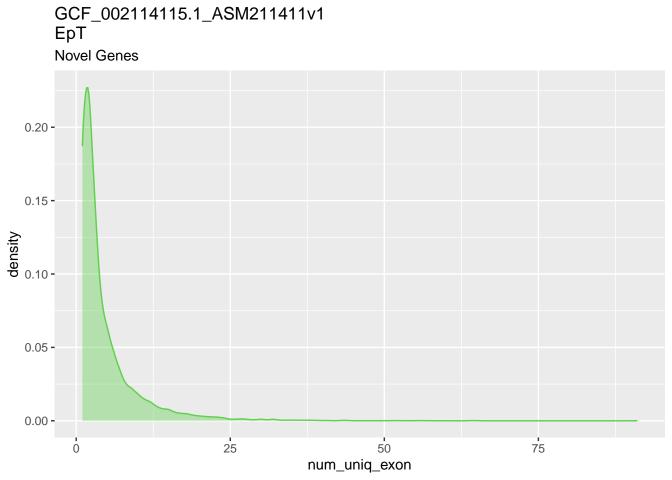


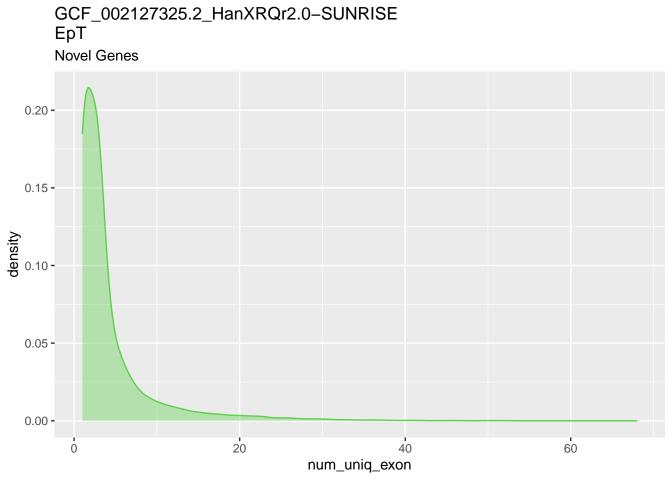


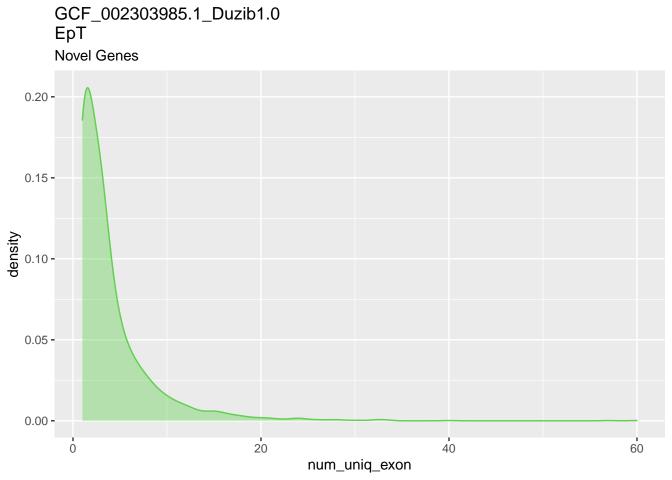


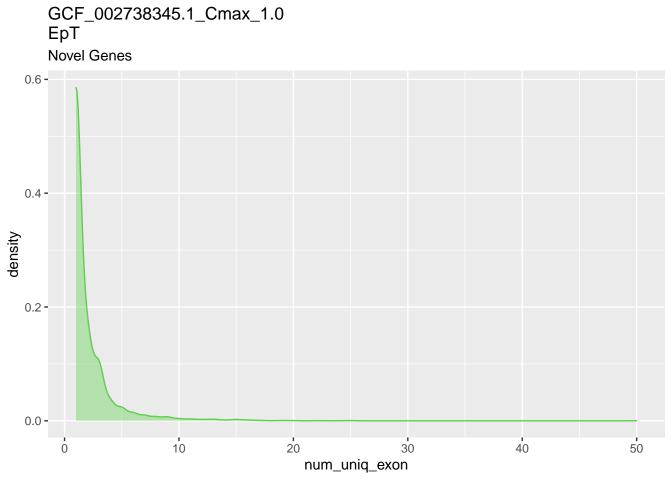


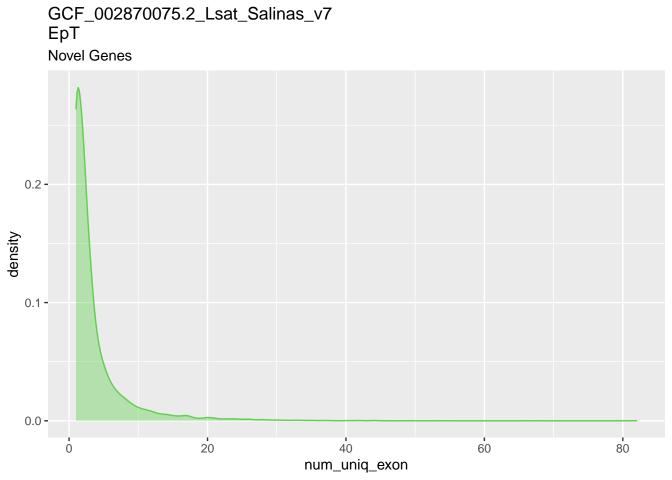


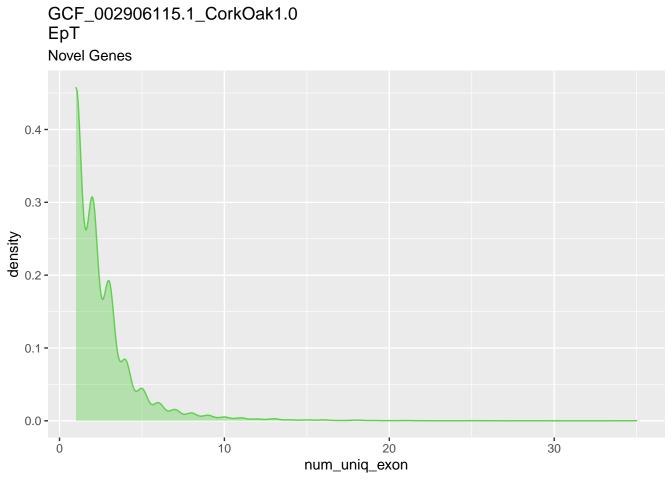


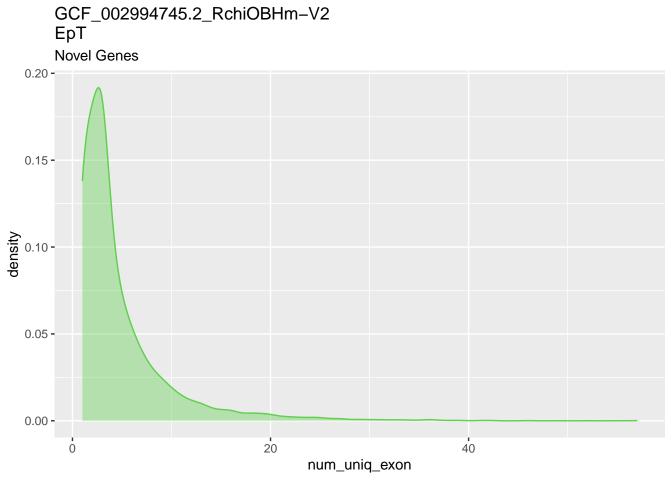


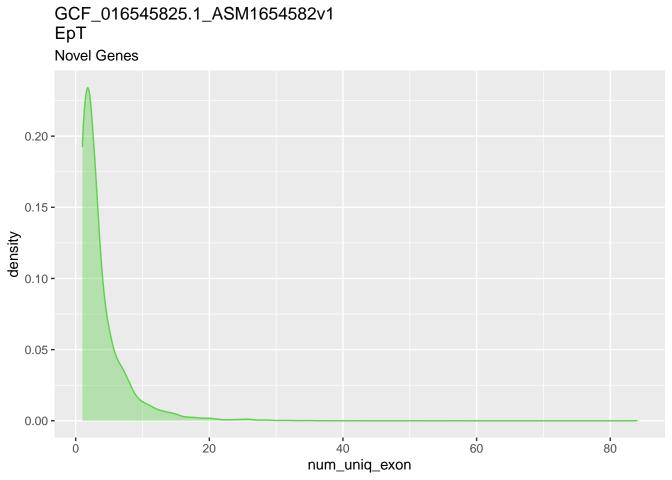




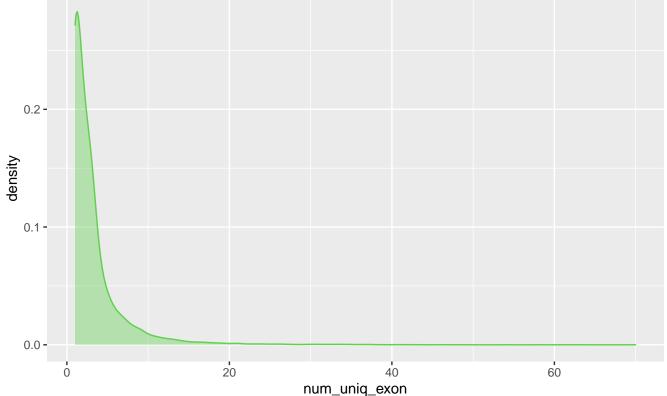






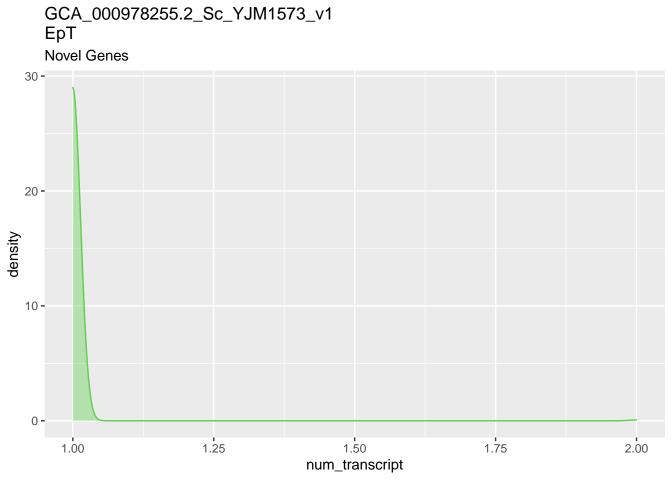


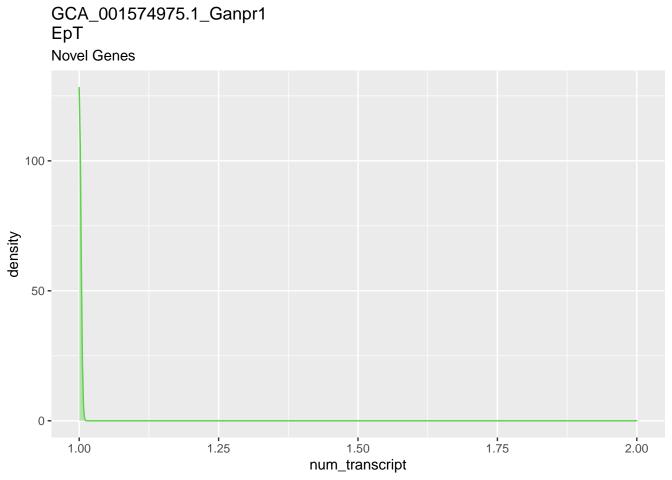
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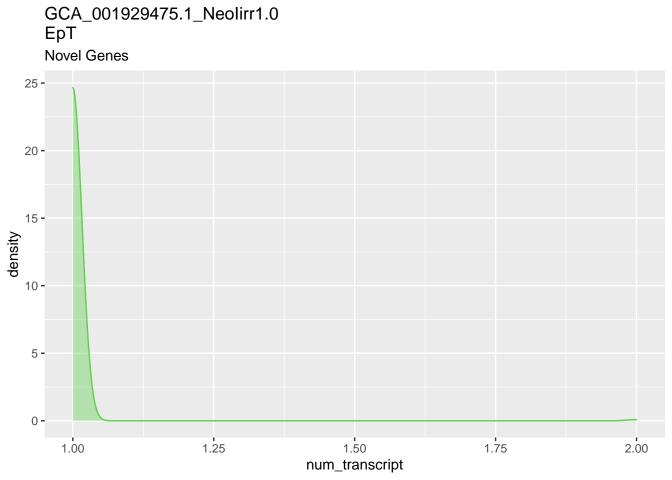


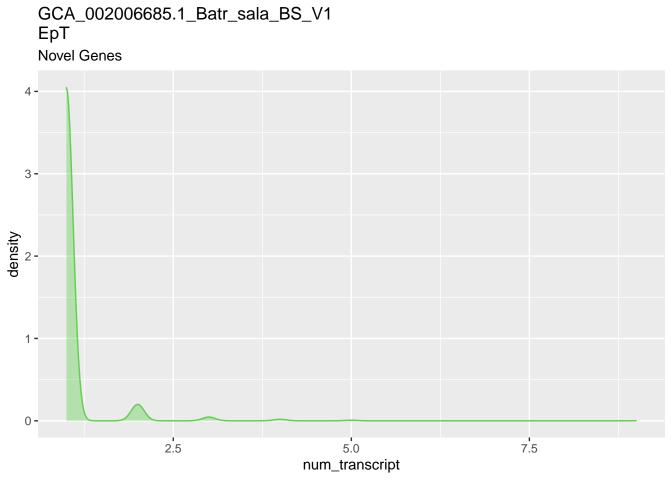
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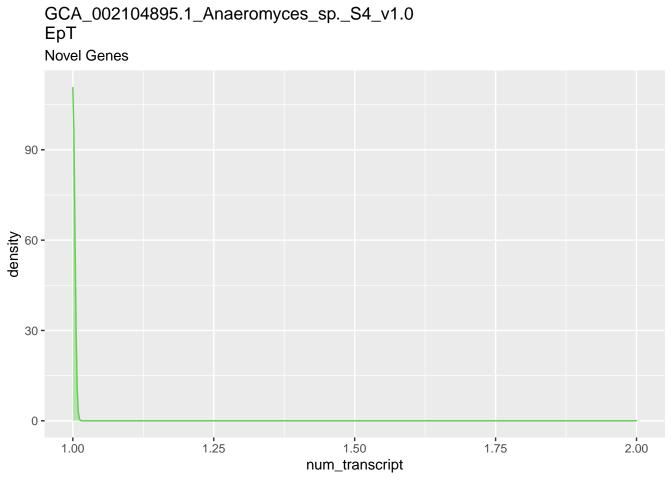
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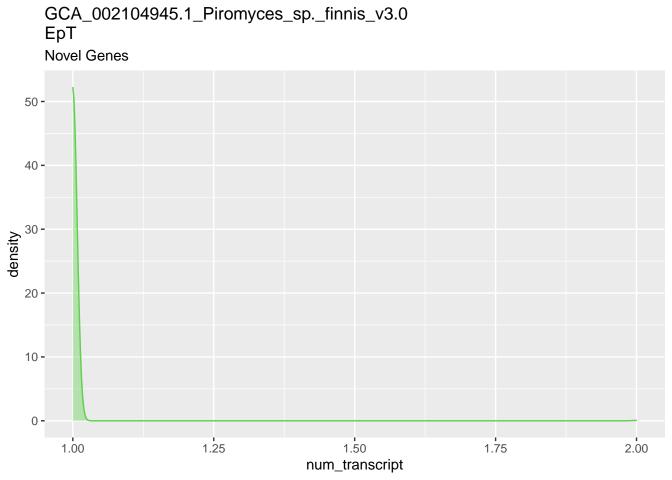
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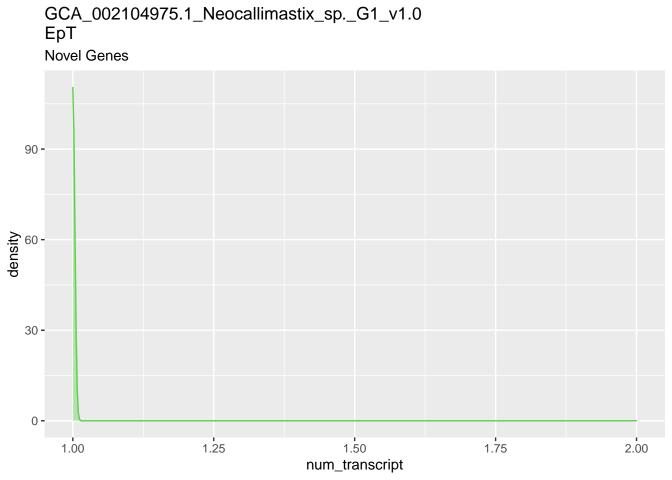
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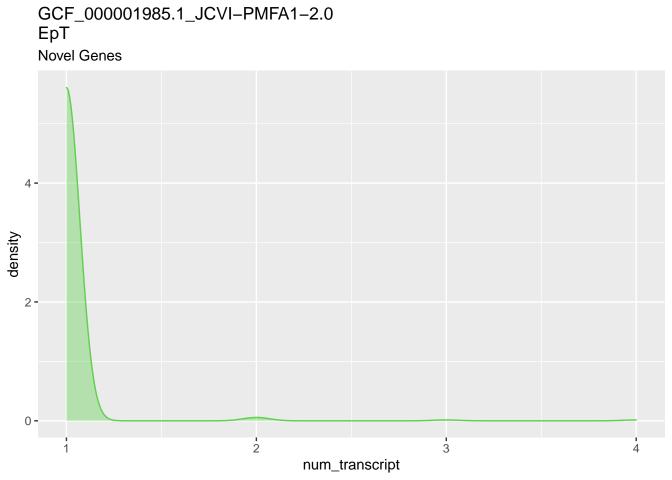


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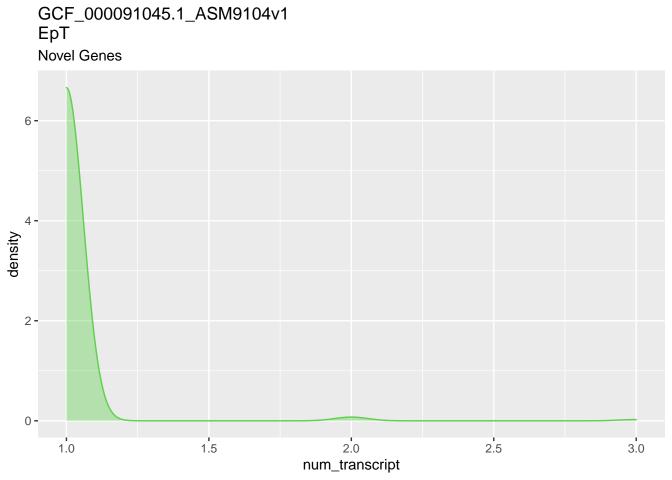
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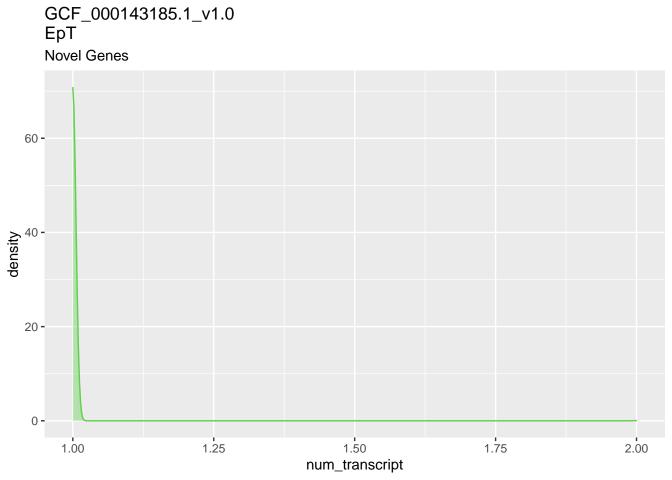
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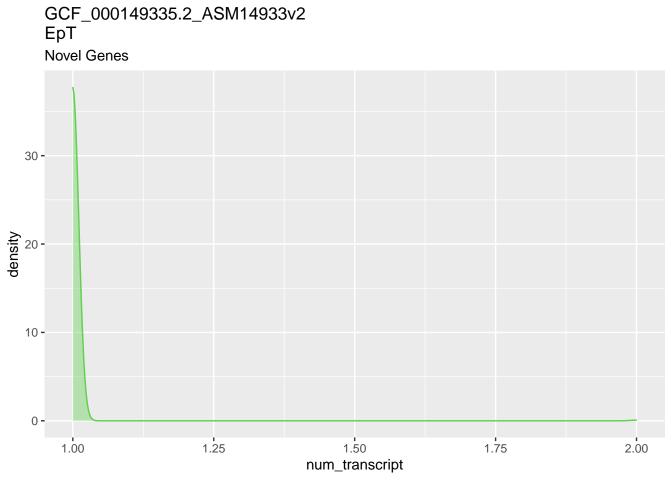
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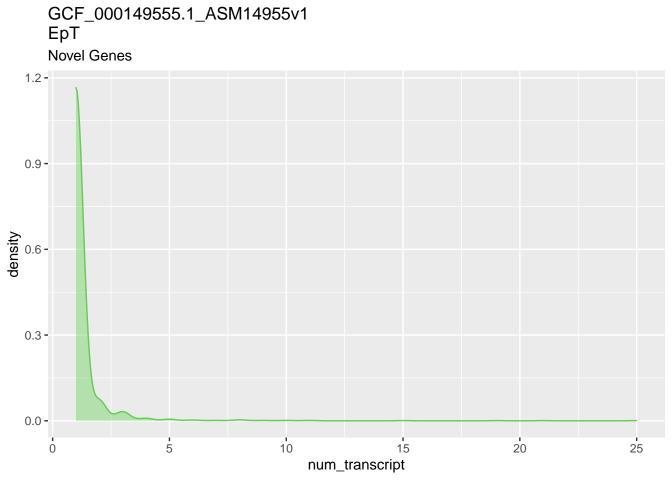




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	Novel Genes
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density	
ŏ	
	num transcript

num_transcript

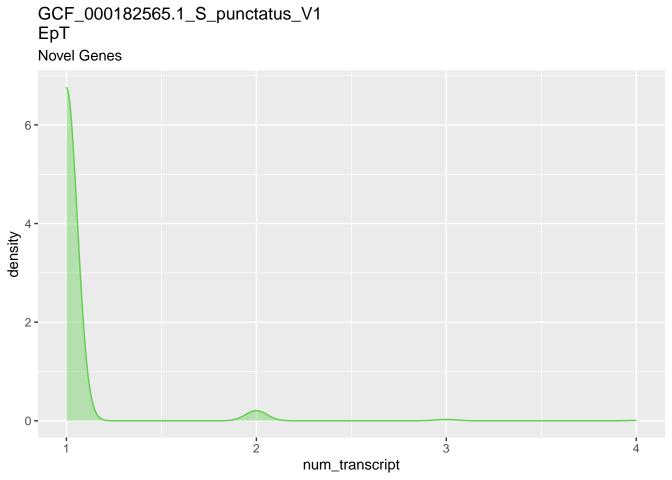




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GCF_000150705.2_Paracocci_br_Pb01_V2 EpT	
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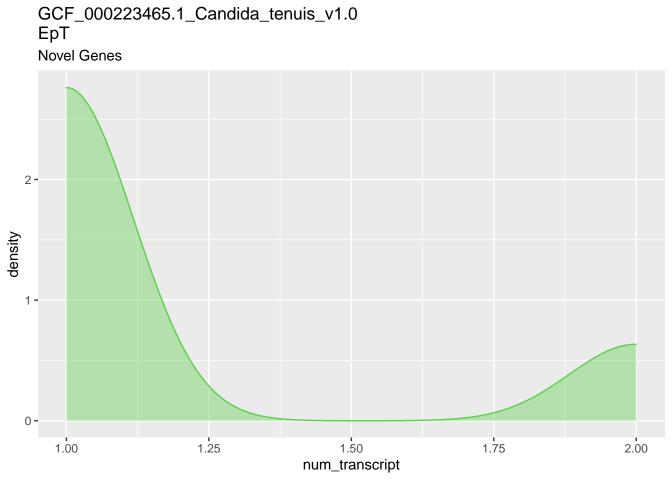


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GCF_000230375.1 EpT Novel Genes	_ASM23037v1	

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density	
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num_transcript

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GCF_000271645.1_Treme1 EpT Novel Genes

	GCF_000275845.1_Dichomitus_squalens_v1.0 EpT
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density	
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	num_transcript

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ЕрТ
Novel Genes

GCF_000300595.1_Phanerochaete_carnosa_HHB-10118-Sp_v1.0 EpT	
Novel Genes	

GCF_000313525.1_ASM31352v1 EpT **Novel Genes**

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density	
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num_transcript

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GCF_000497045.1_PSEUBRA1 EpT Novel Genes

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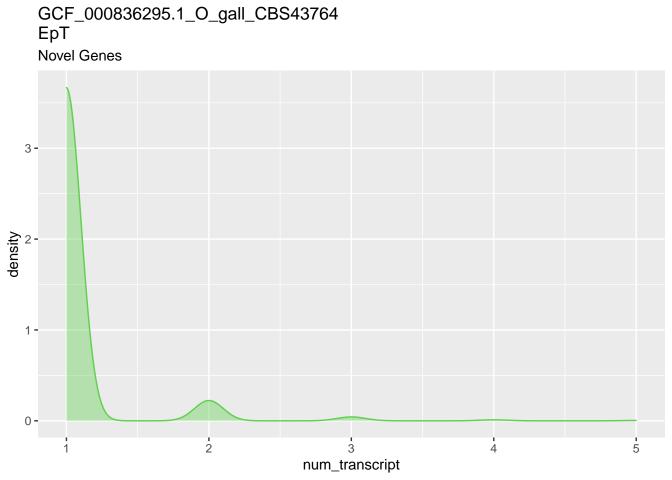
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num_transcript

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	Novel Genes
	
density	
Ü	
	num_transcript

	GCF_000835555.1_Rhin_mack_CBS_650_93_V1 EpT
	Novel Genes
density	
	num_transcript



GCF_000938715.1_LALA0 EpT Novel Genes

GCF_001027345.1_Triol1 EpT Novel Genes

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GCF_001329695.1_Rhoba1_1 EpT Novel Genes

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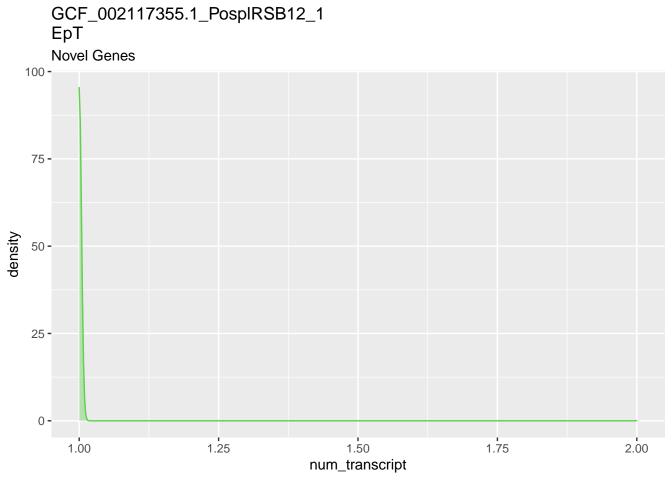
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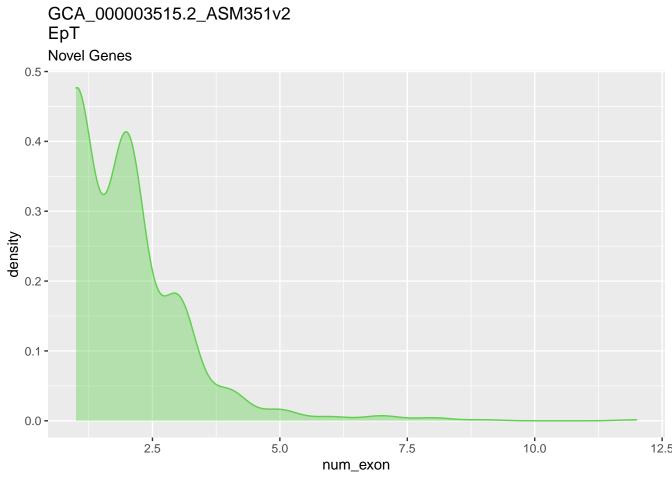
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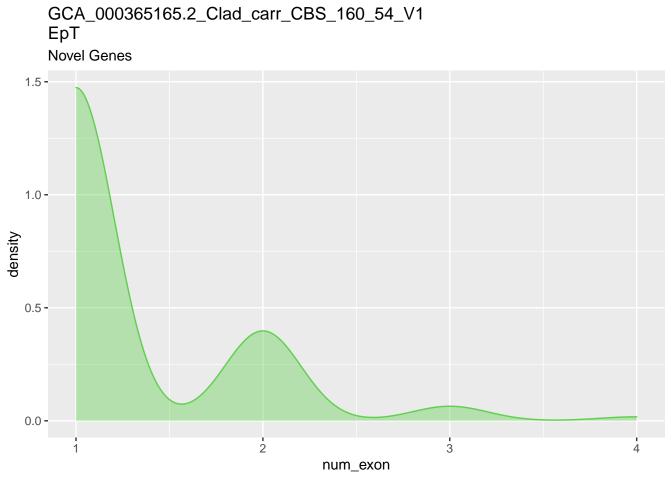
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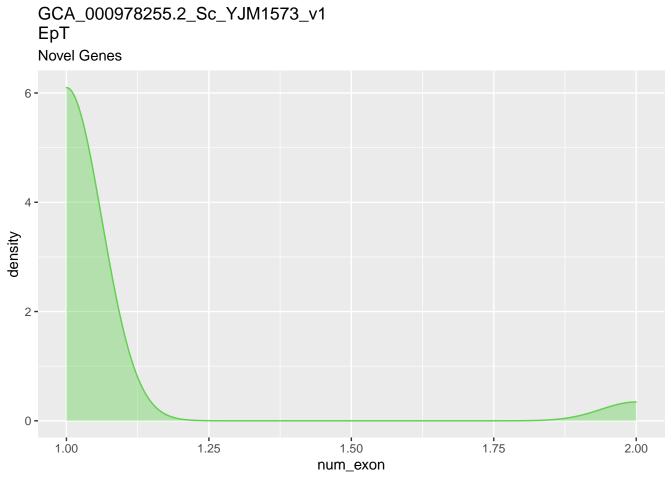
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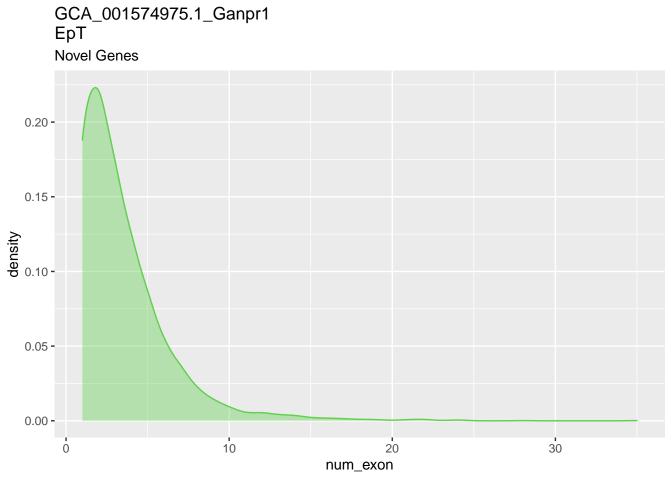


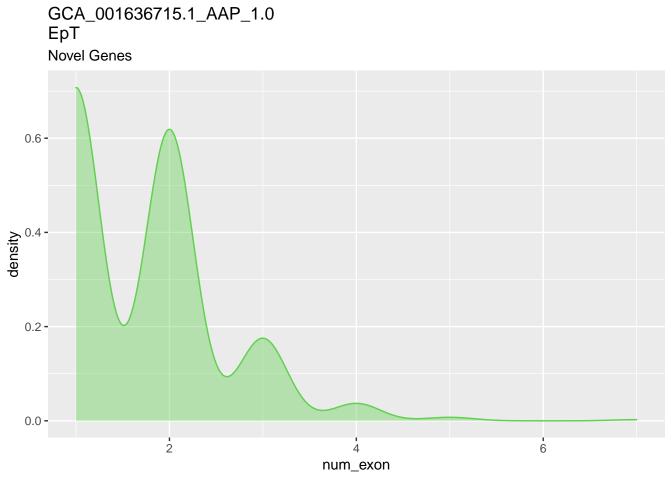
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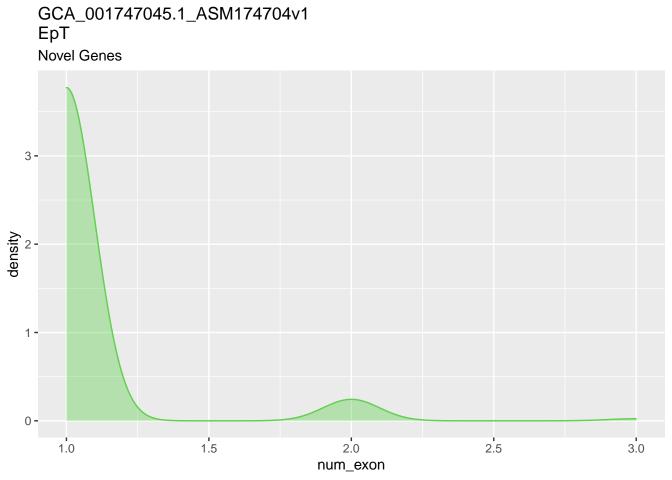


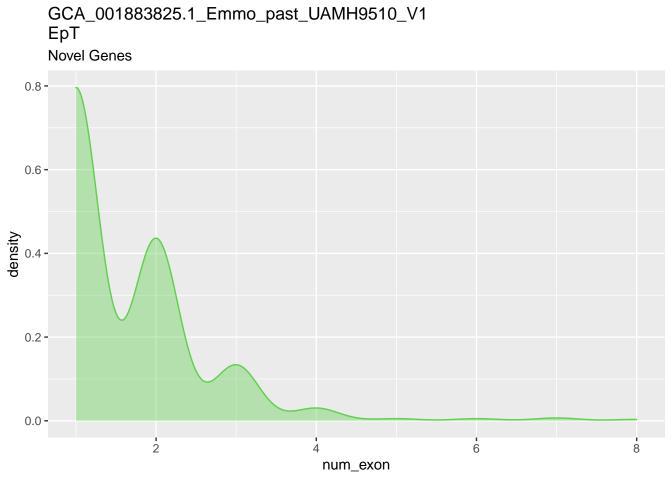


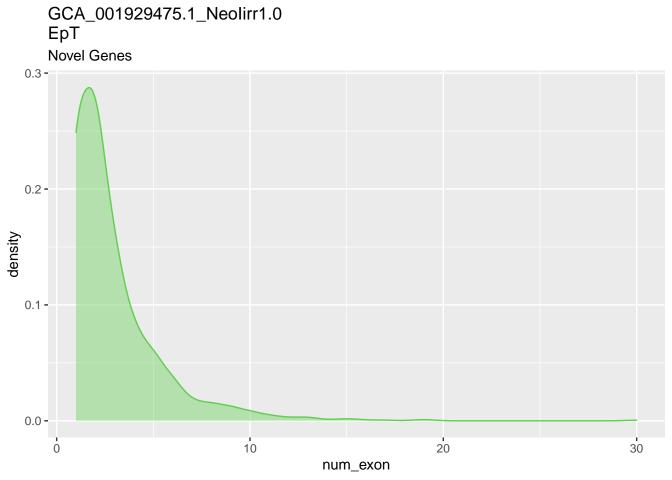


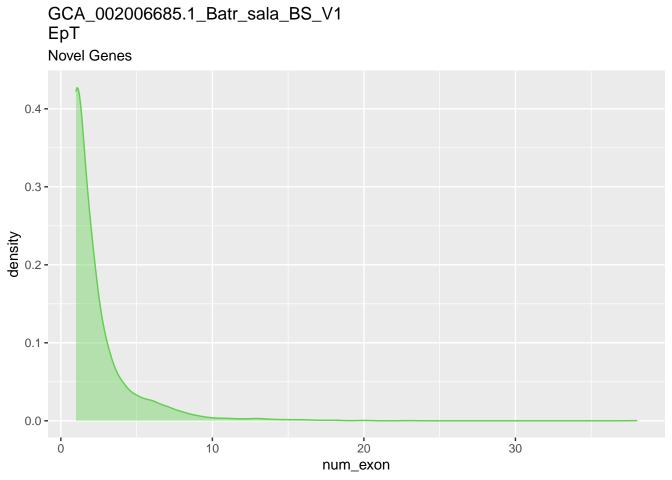


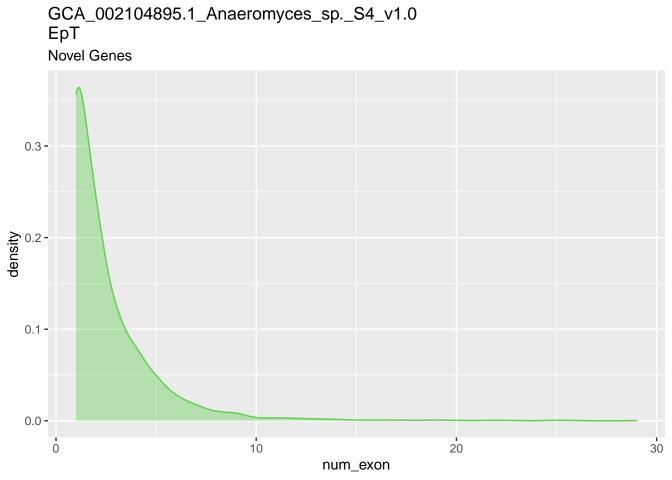


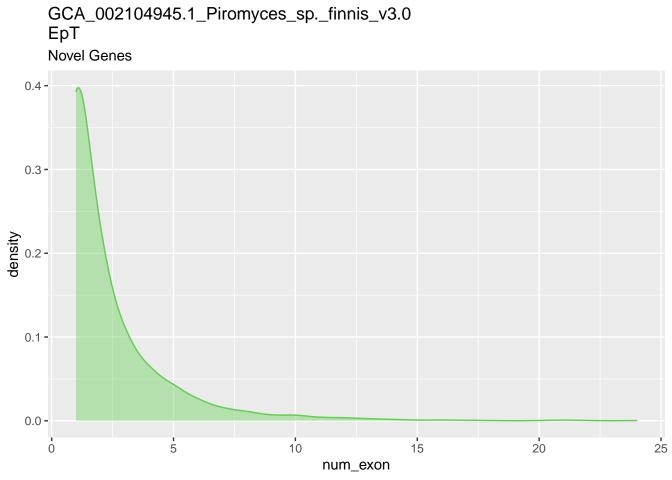


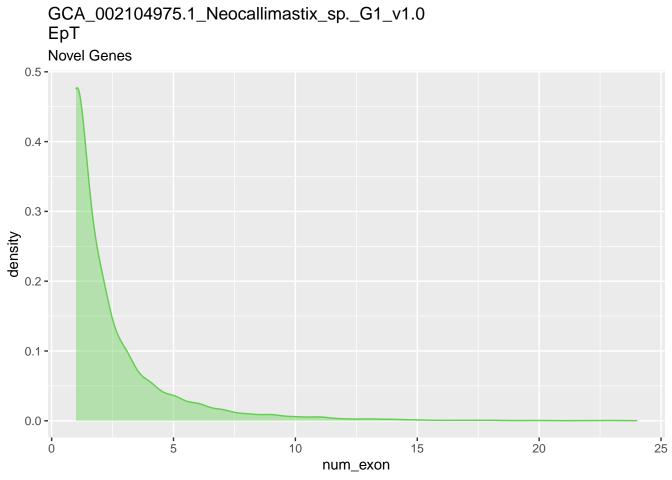


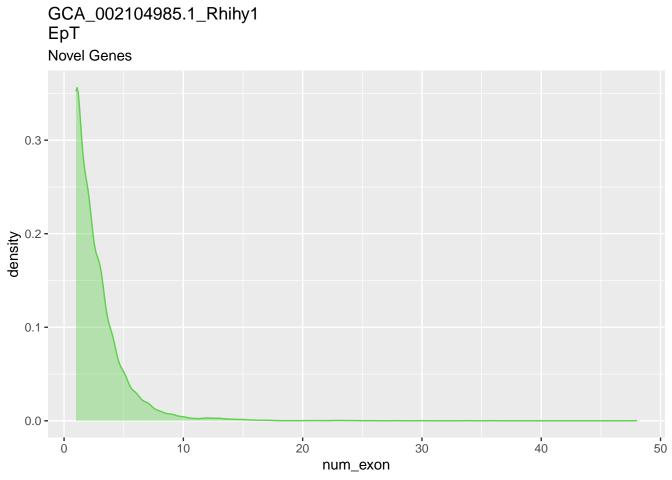


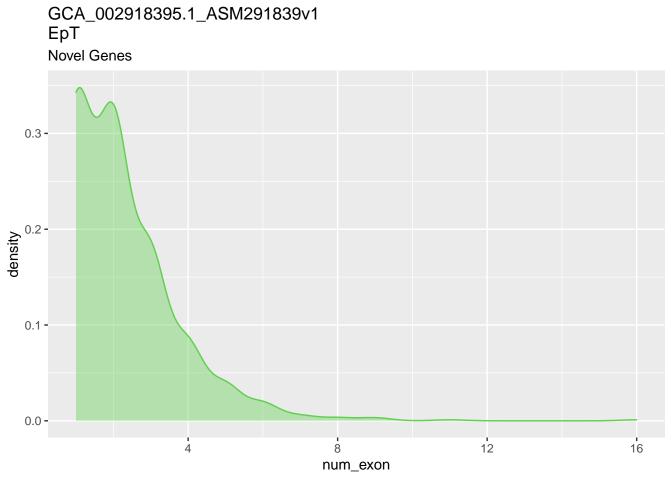


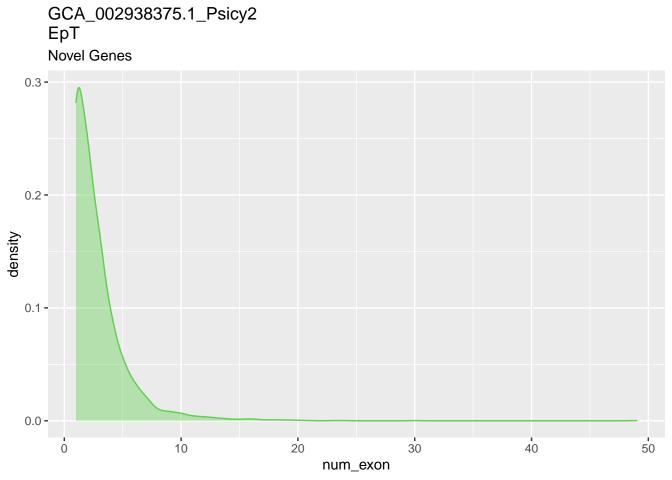


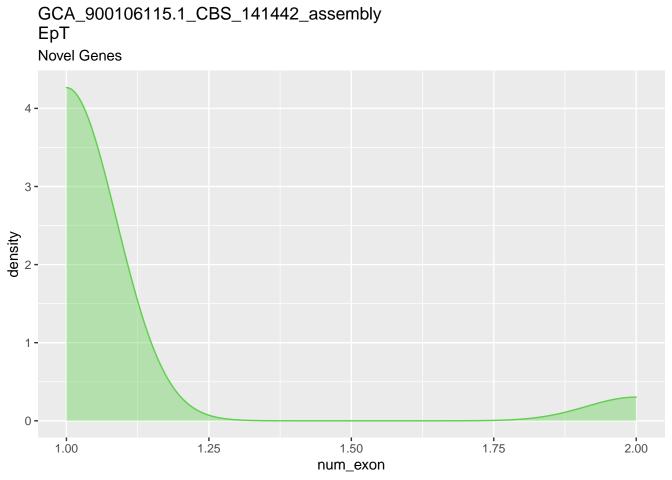


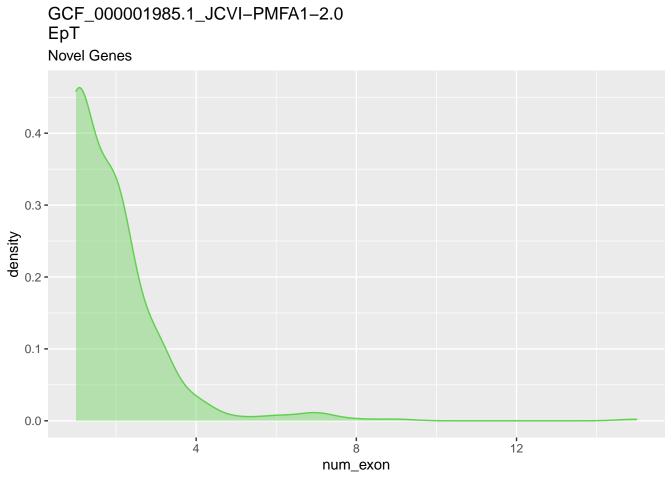


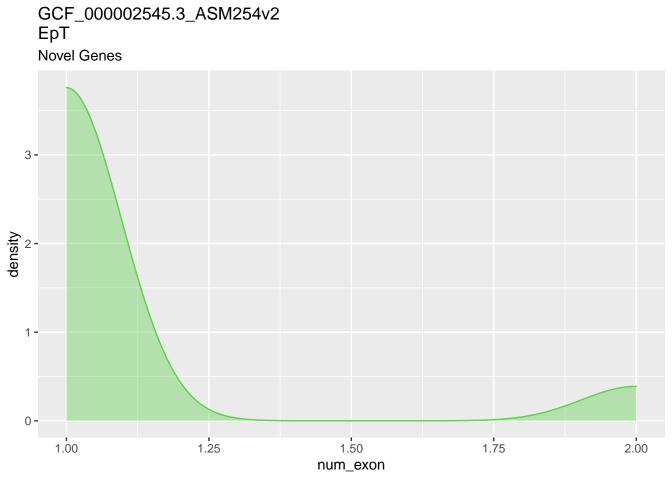


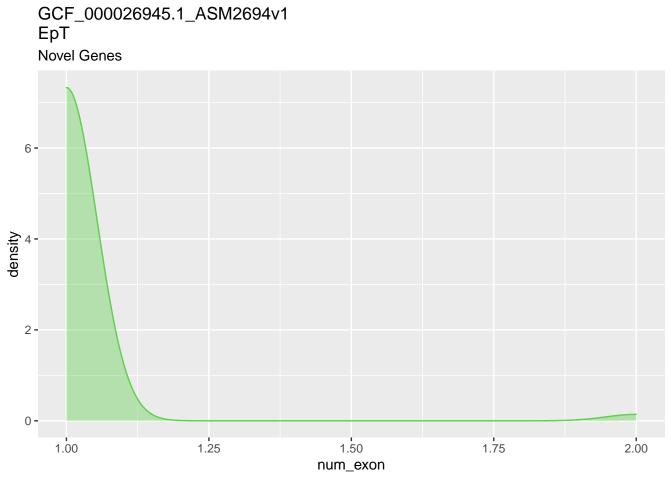


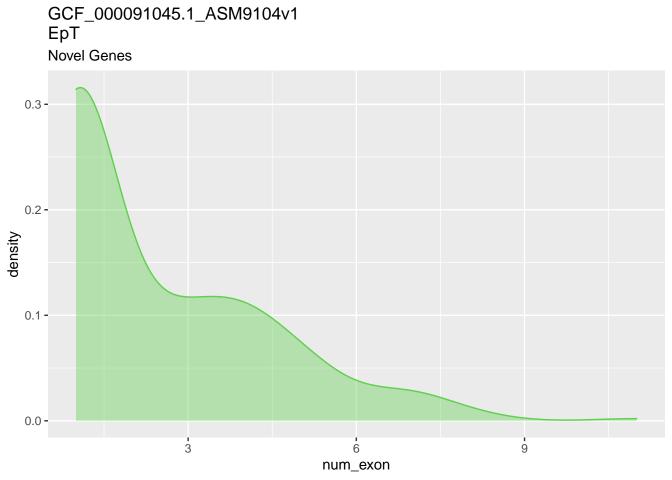


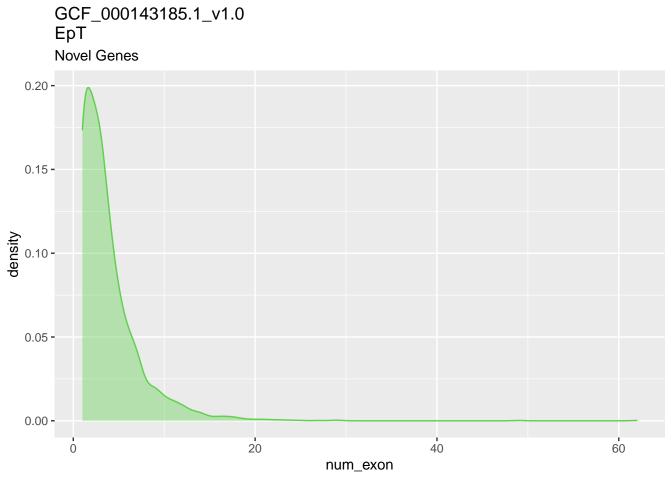


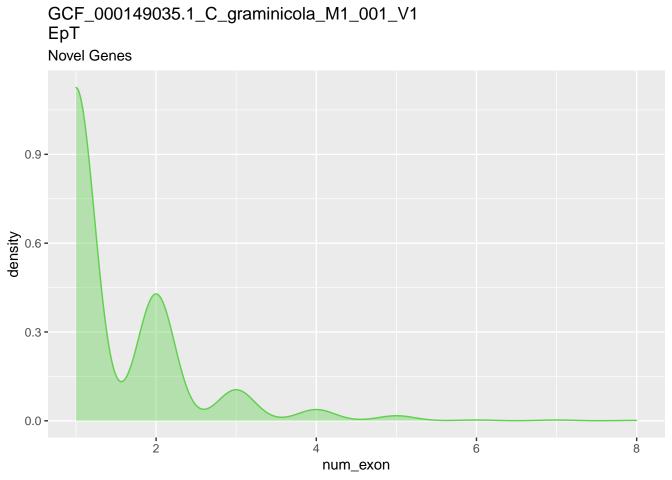




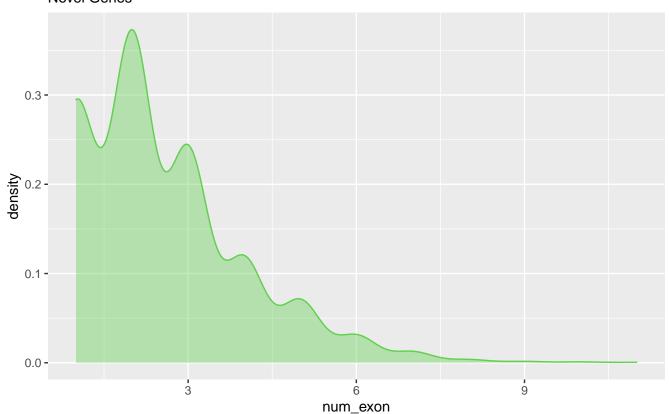


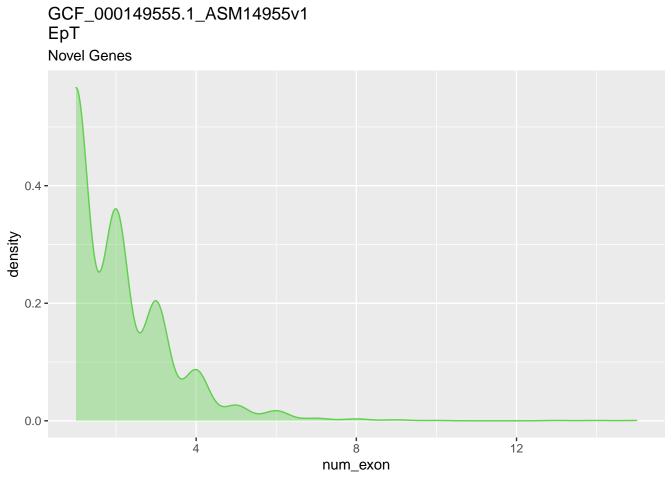


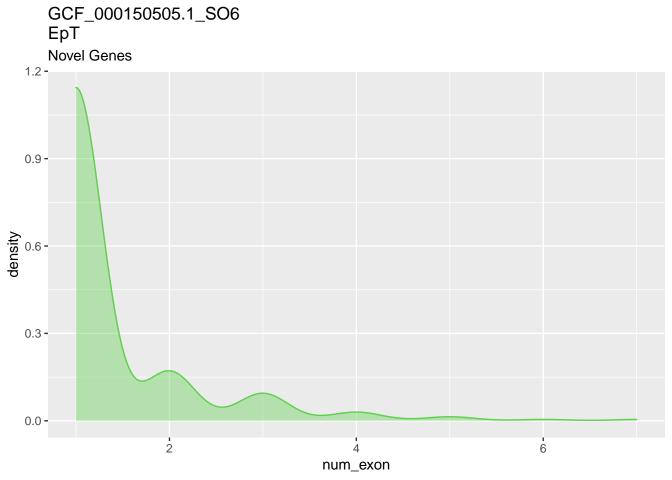


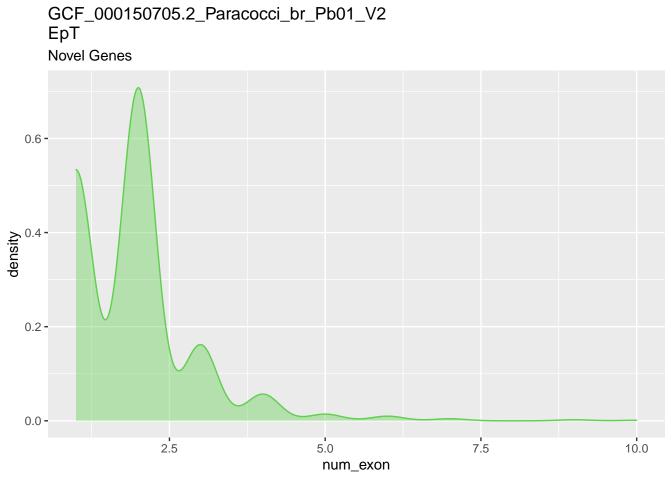


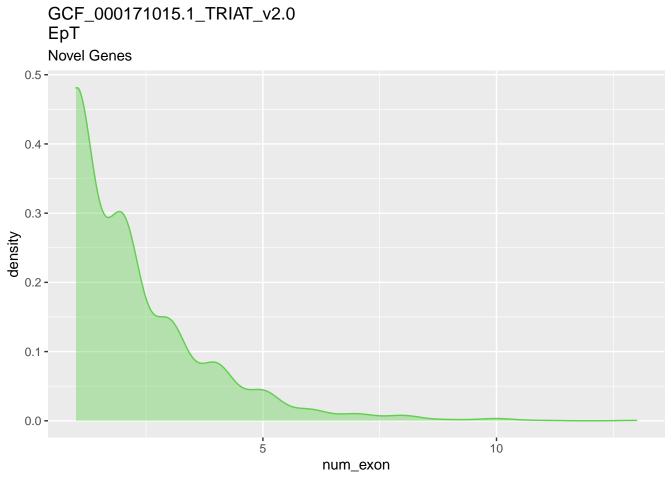
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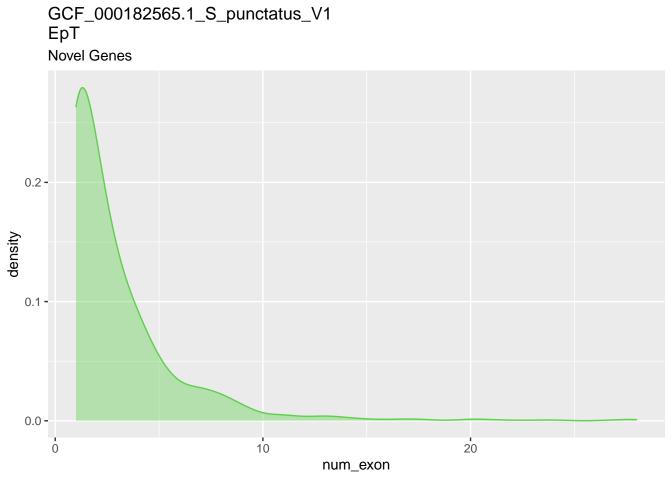


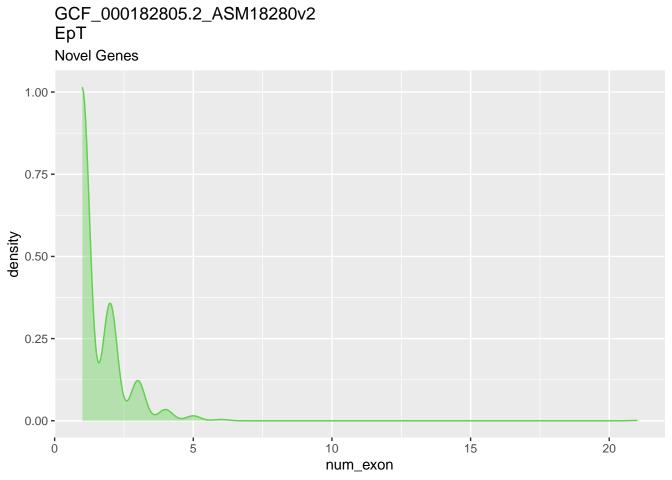


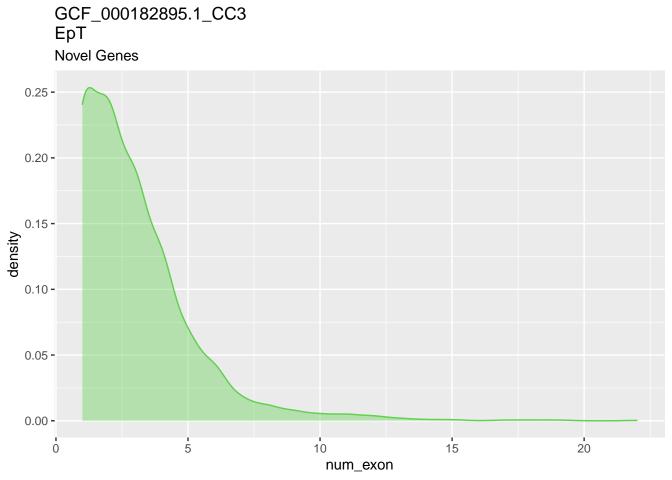


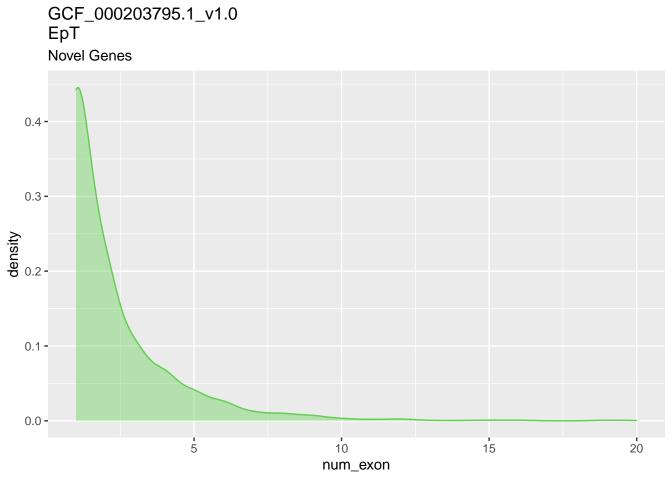


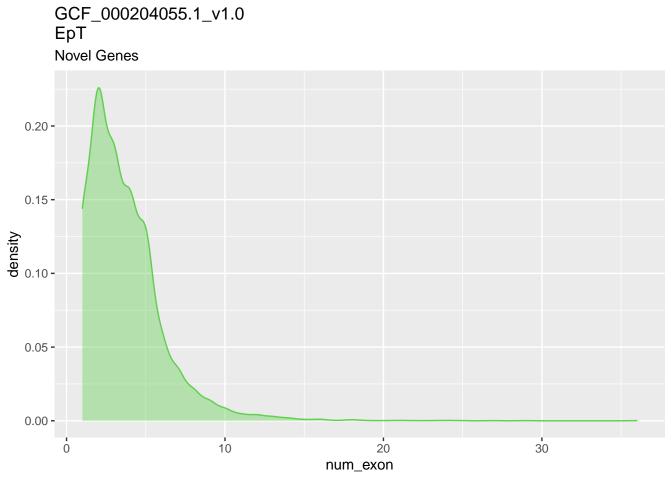


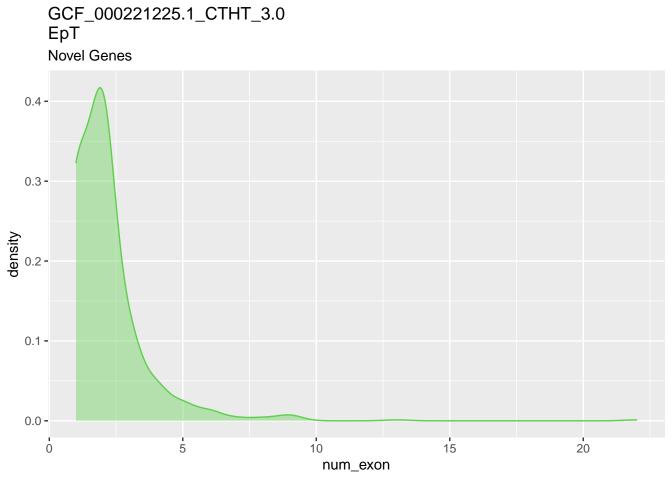


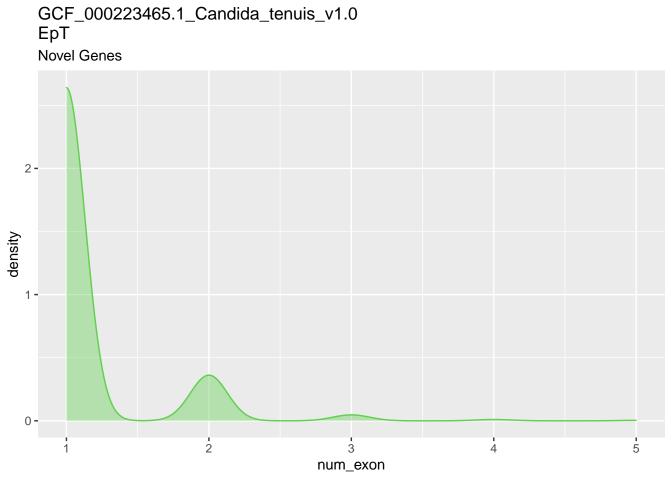


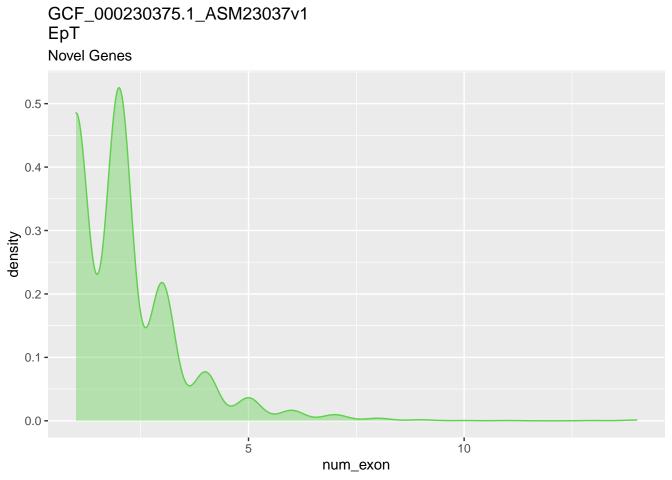


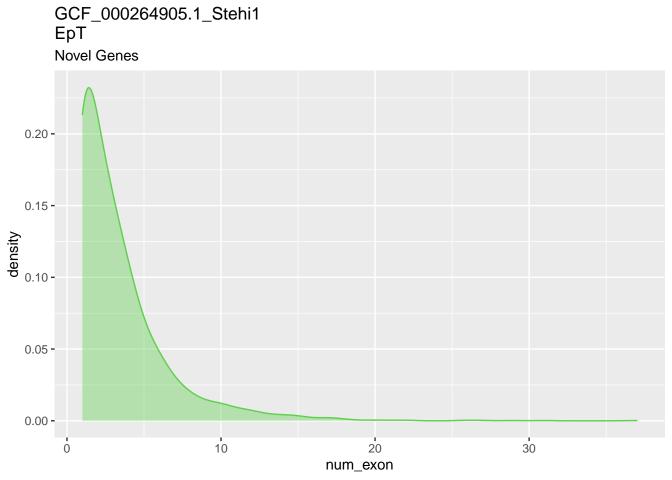


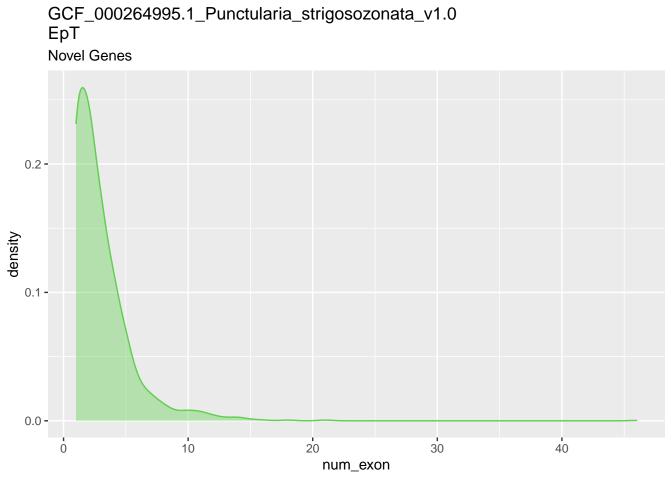


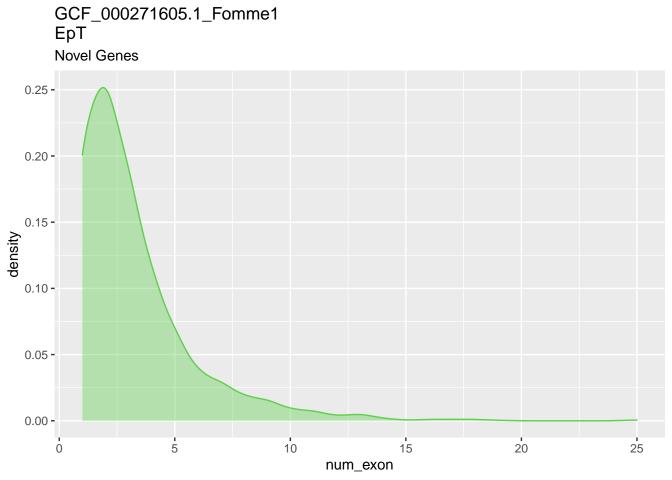


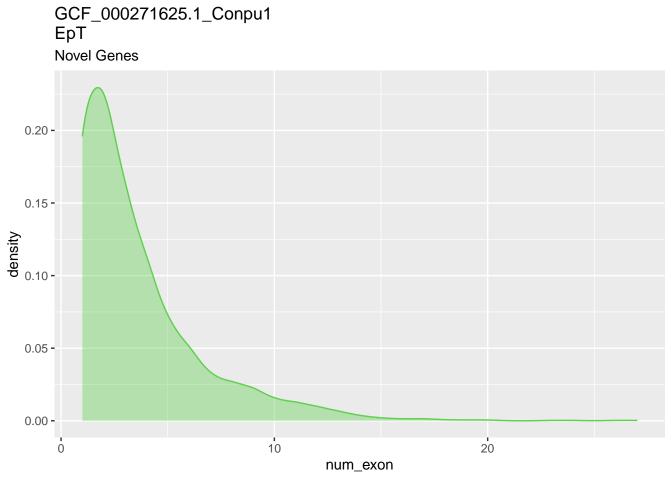


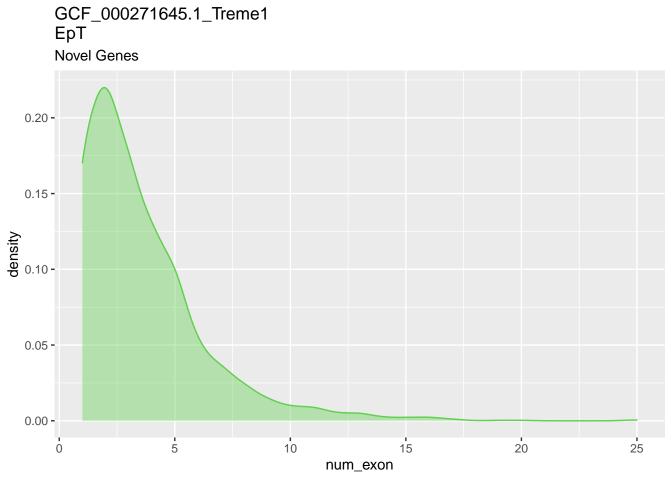


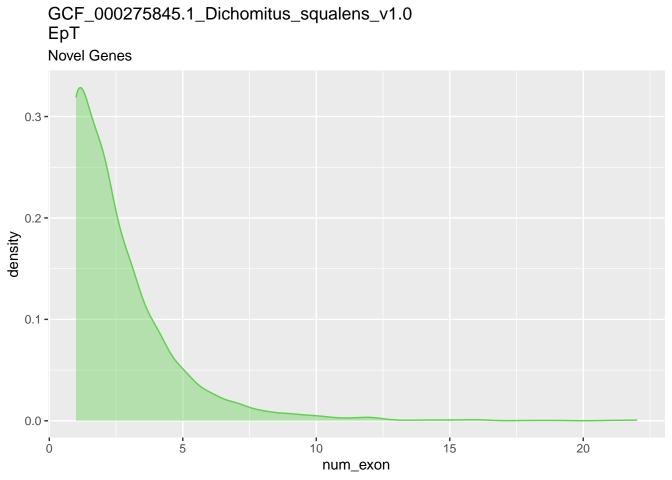


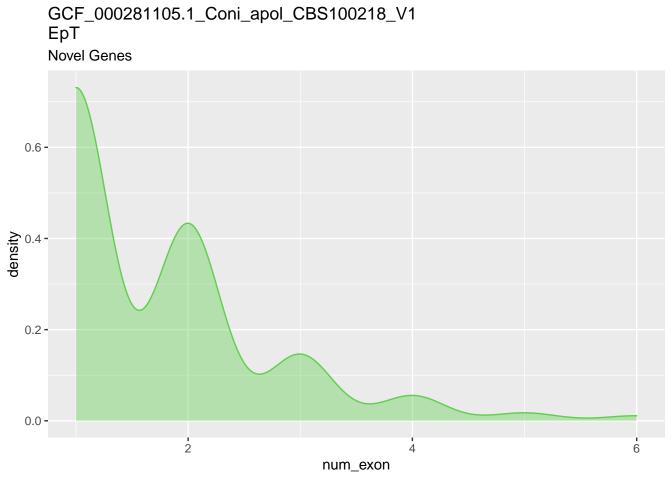


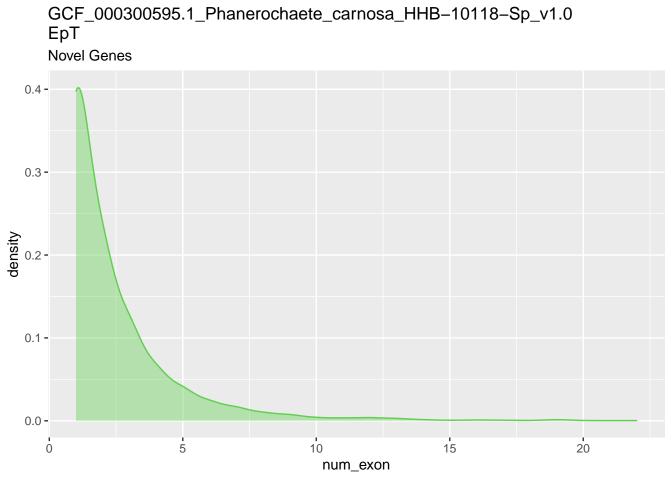


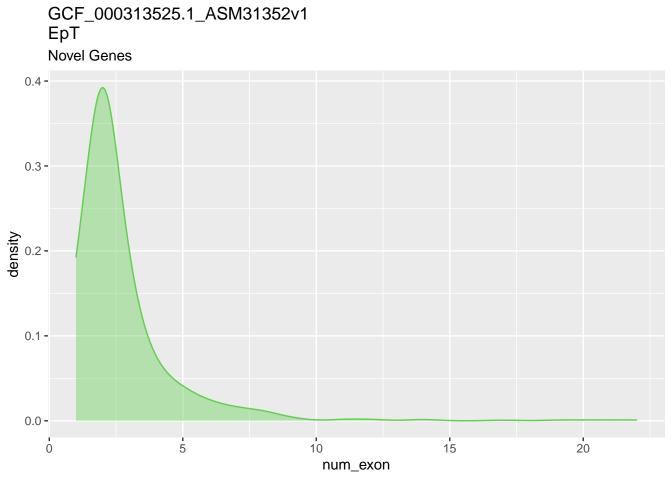


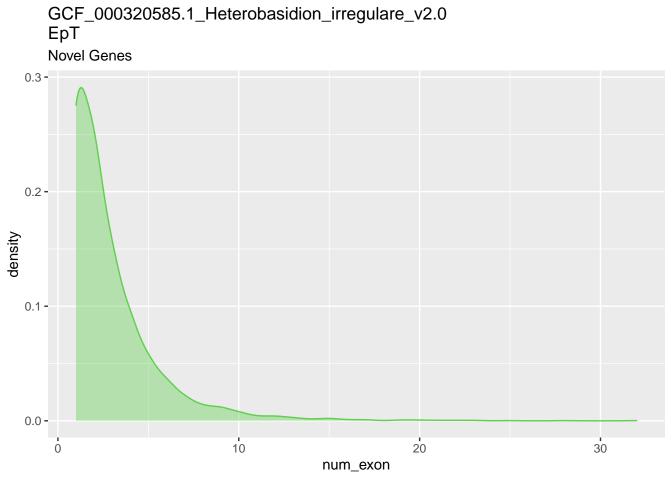


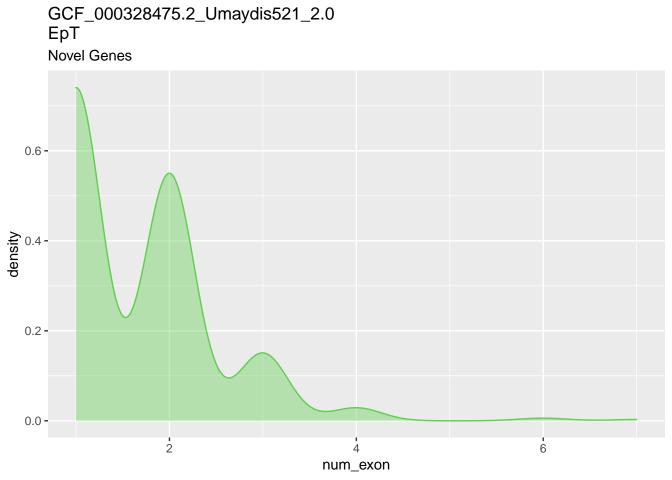


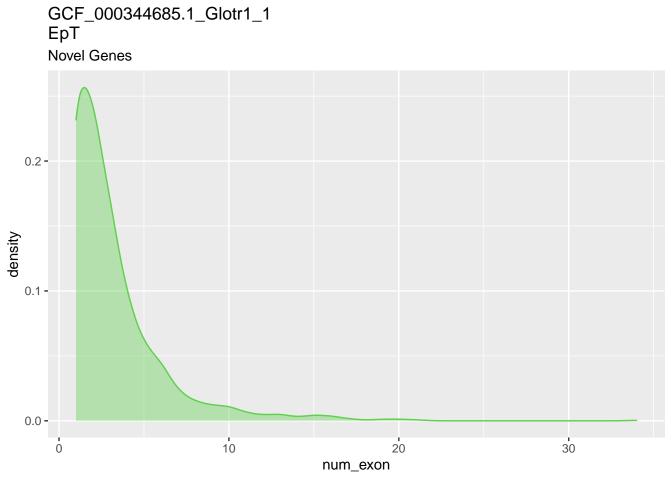


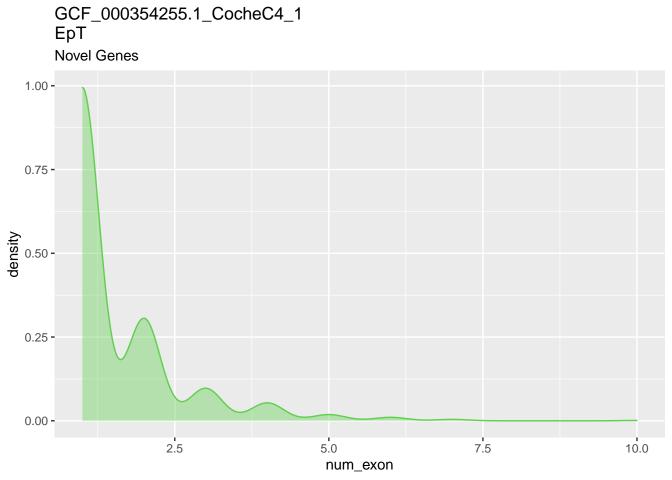


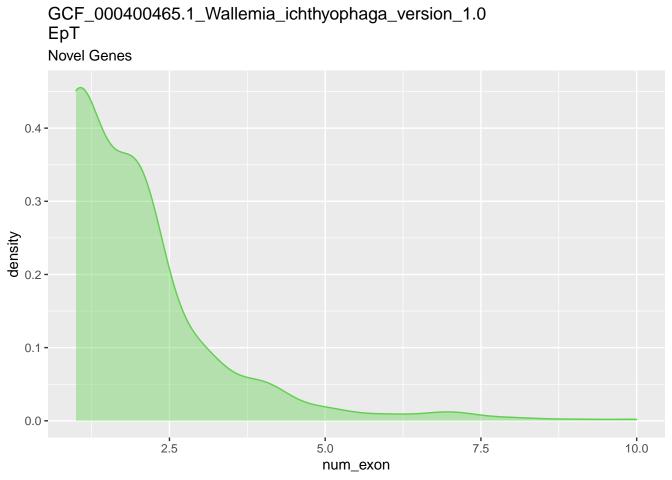


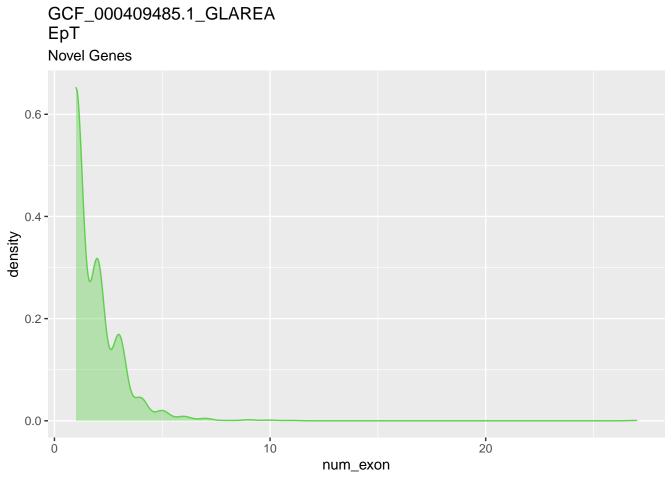


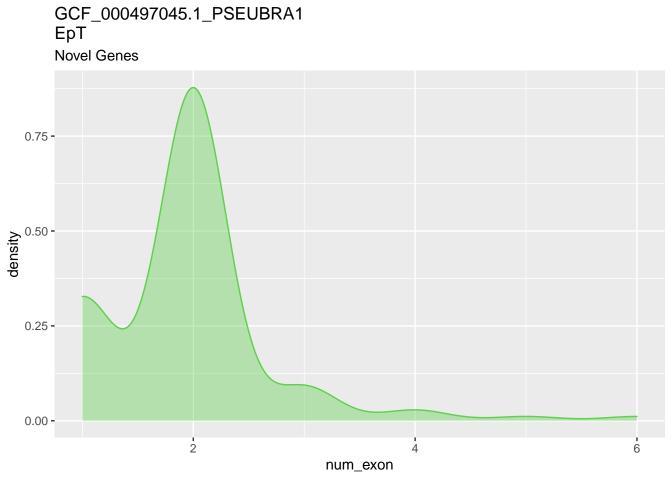


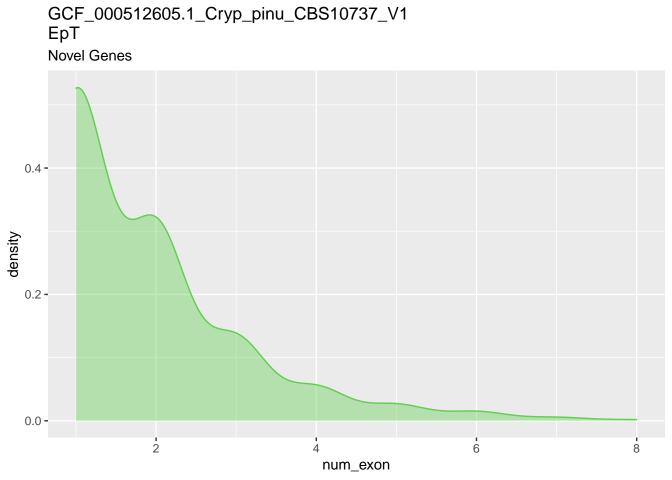


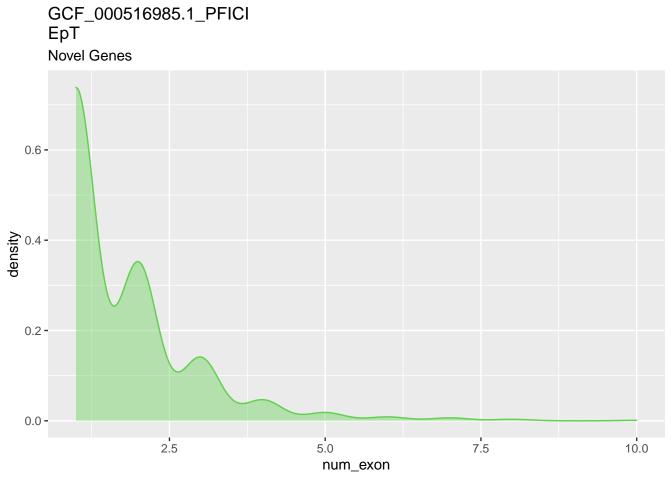


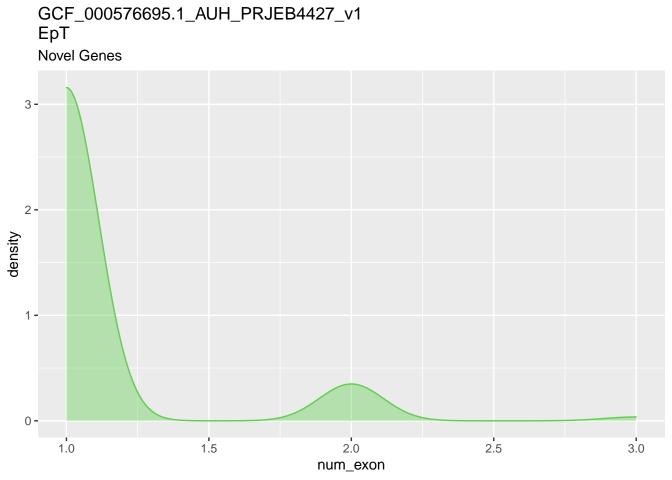


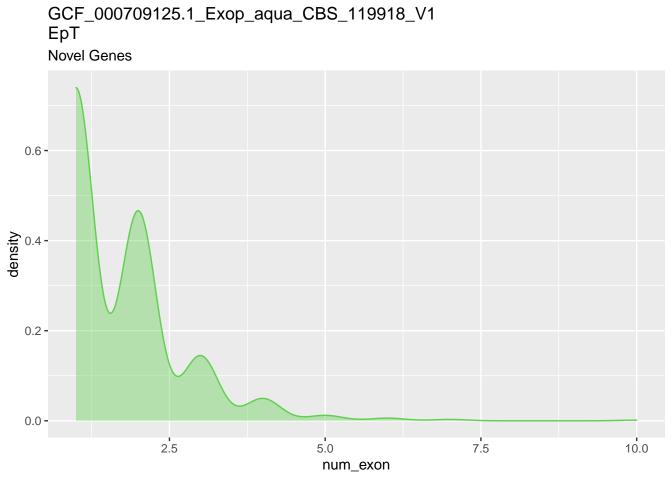


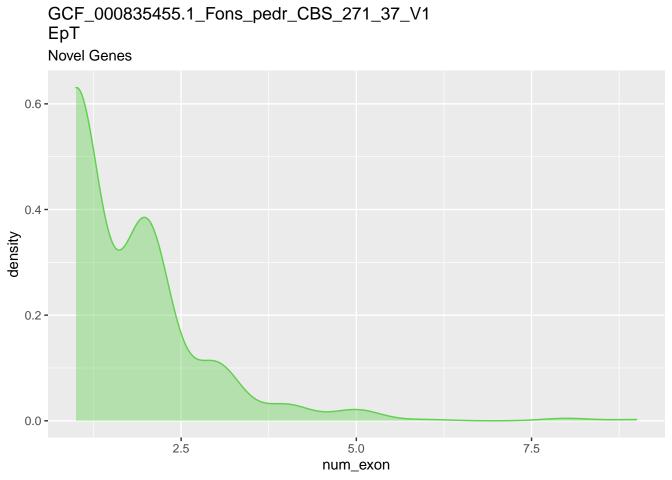


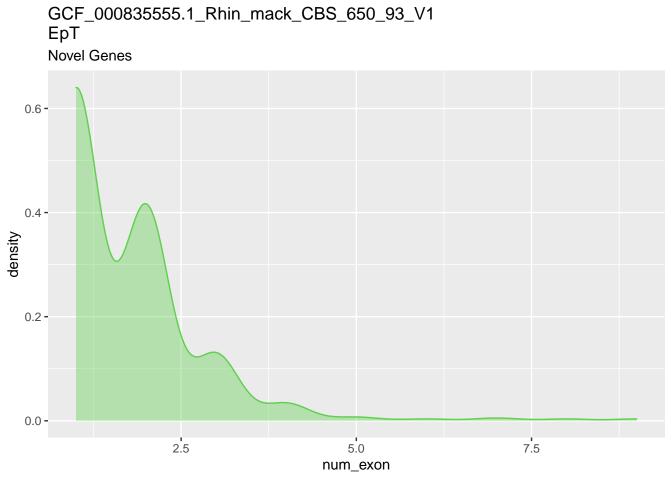


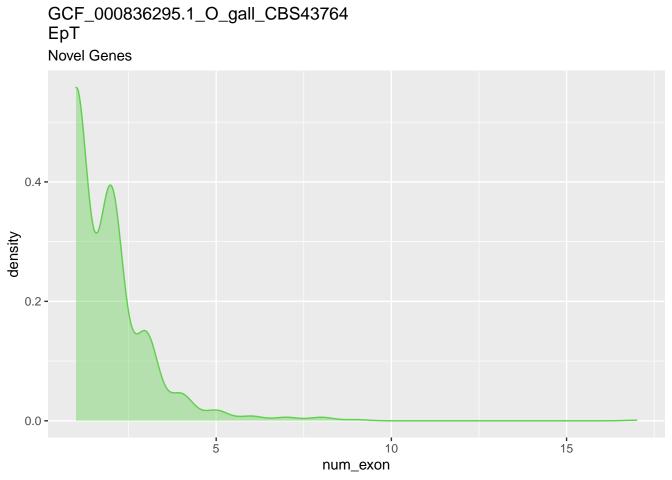


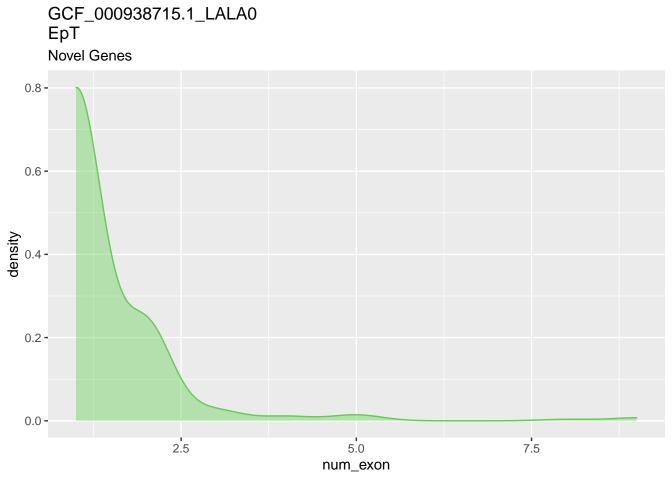


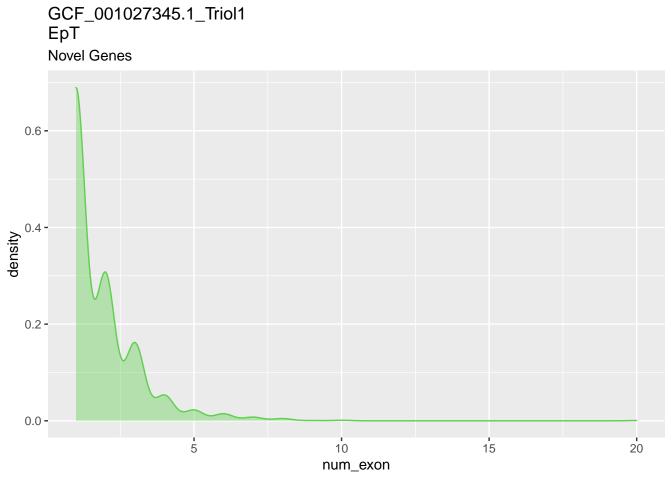


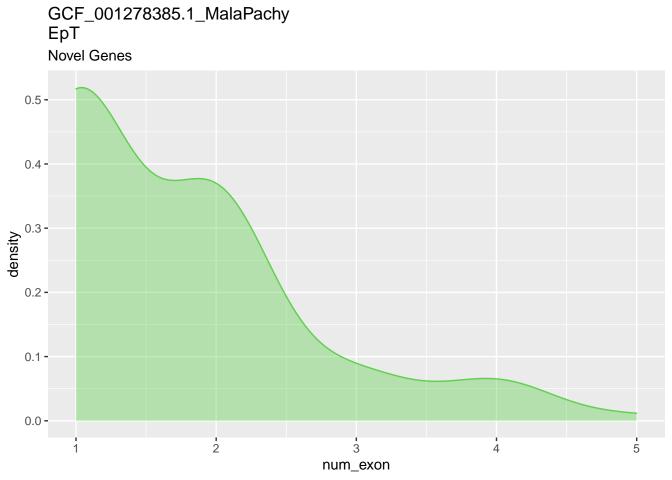


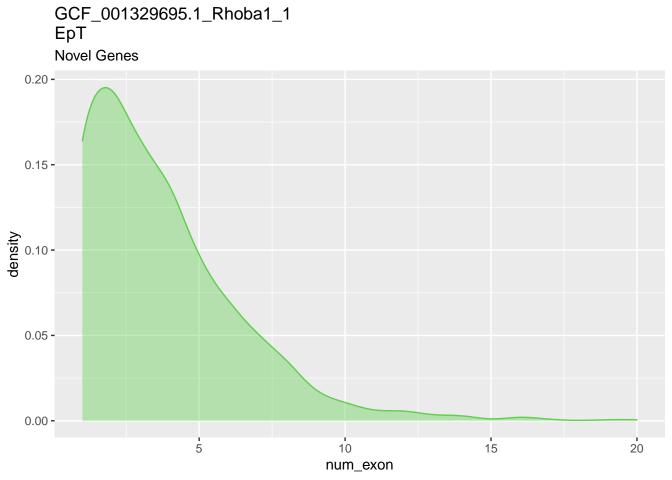


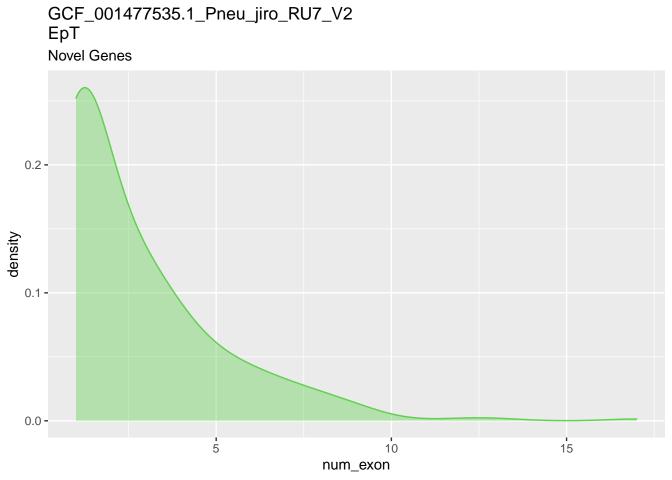


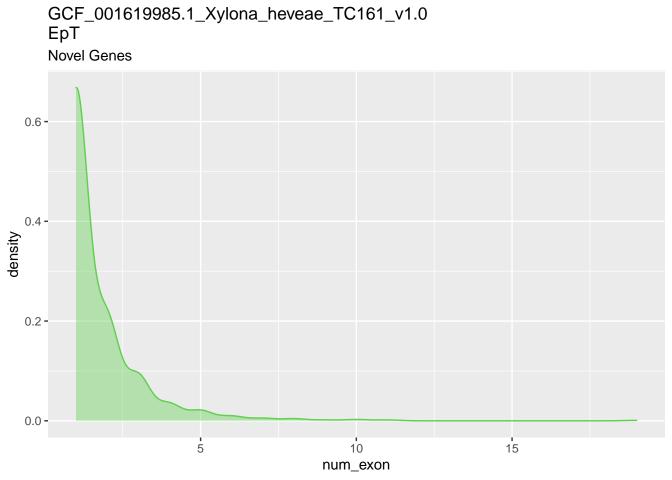


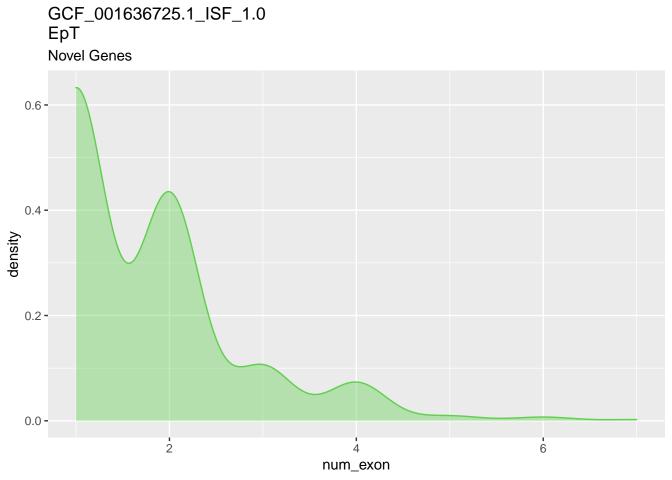


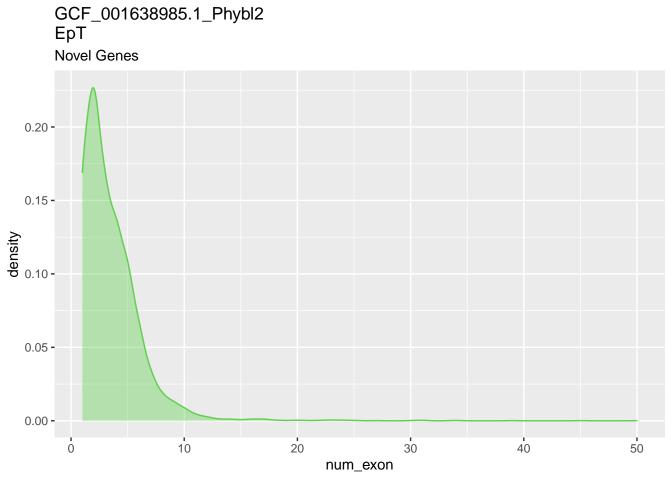


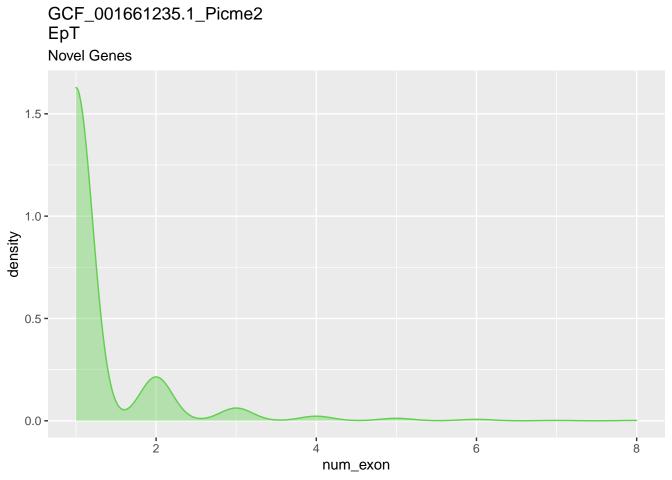


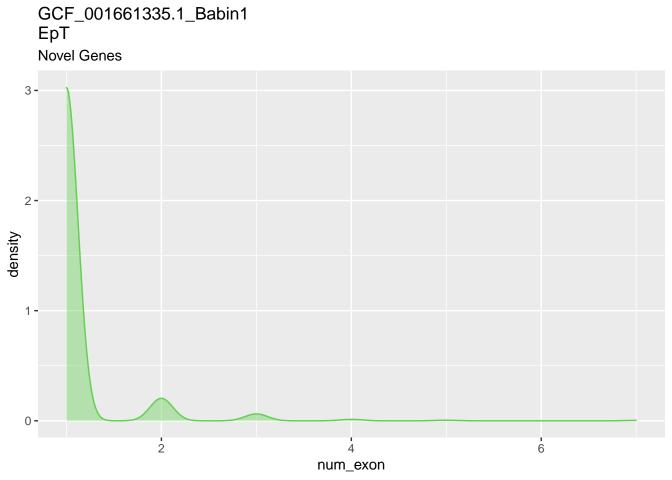


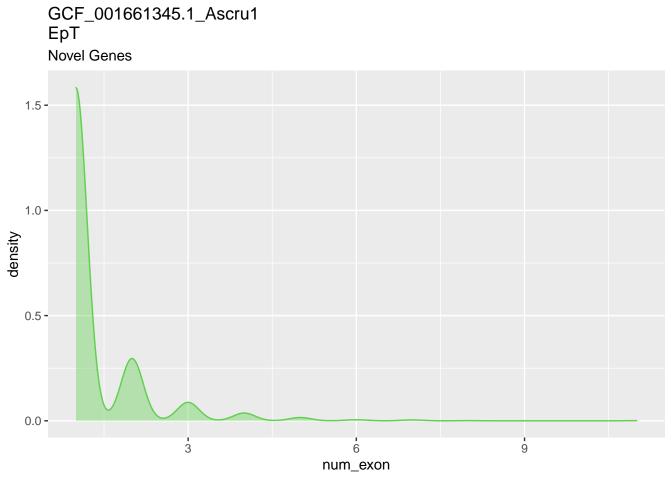


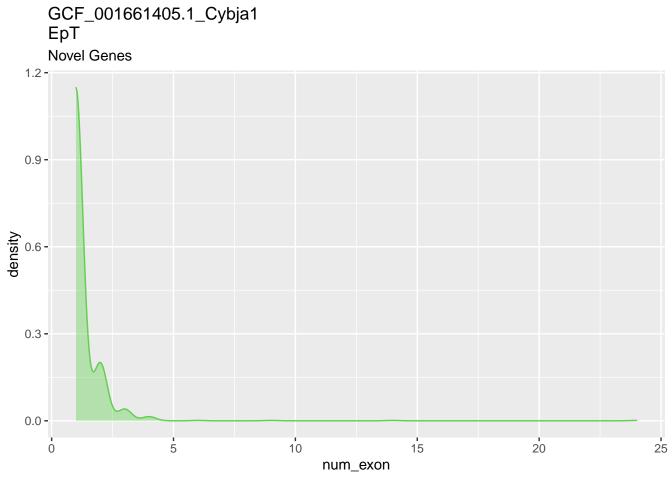


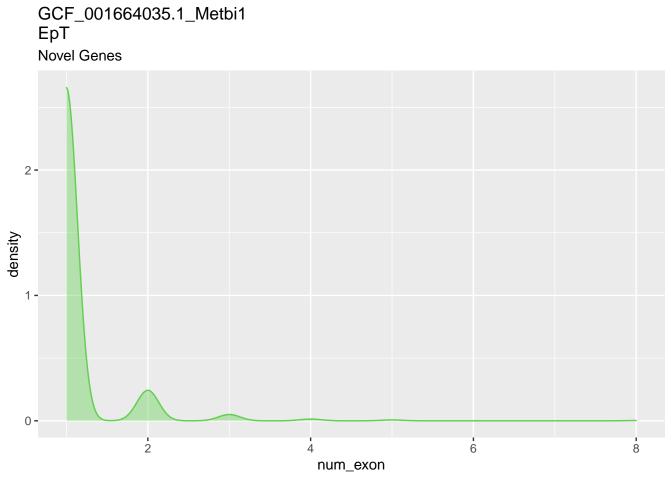


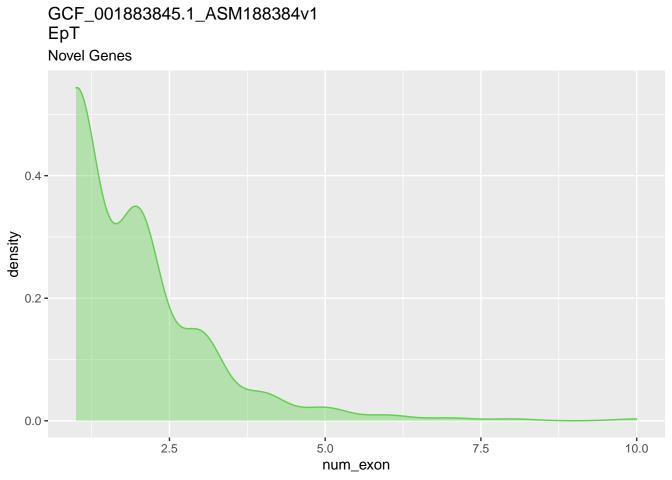


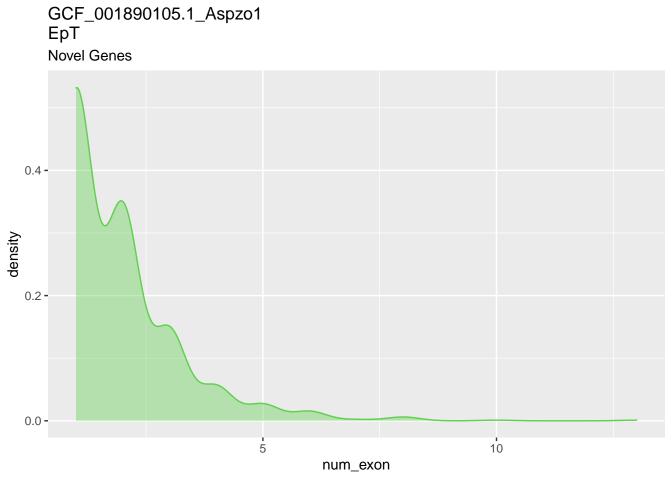


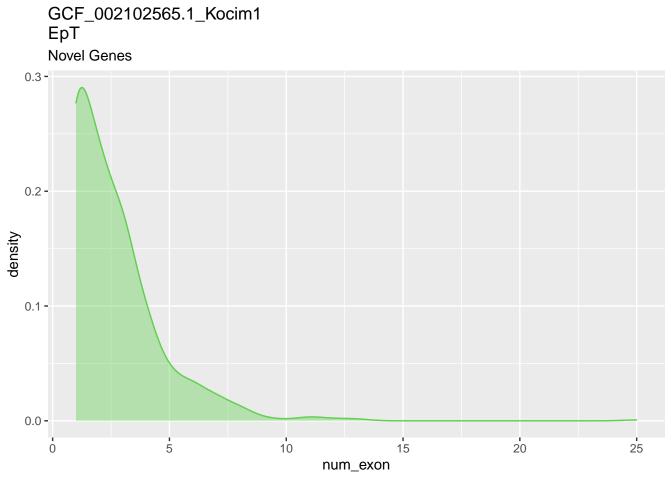


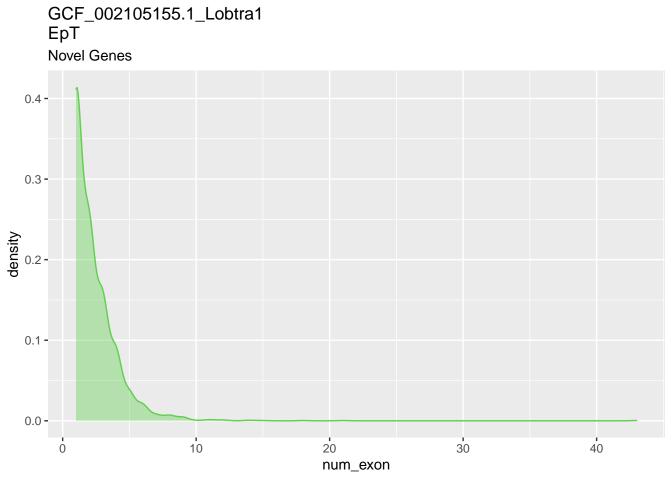


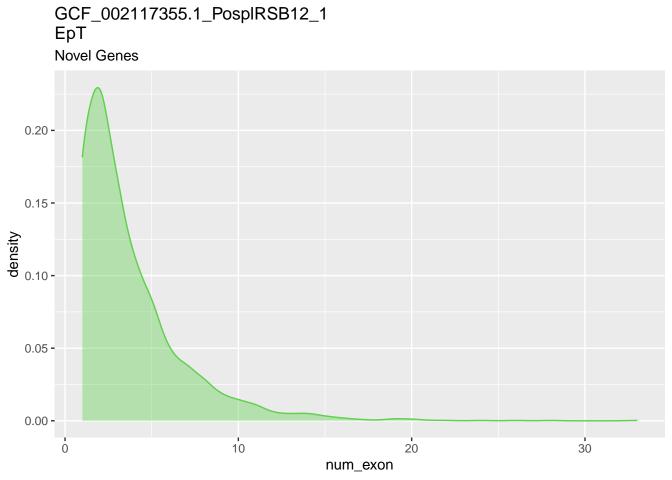


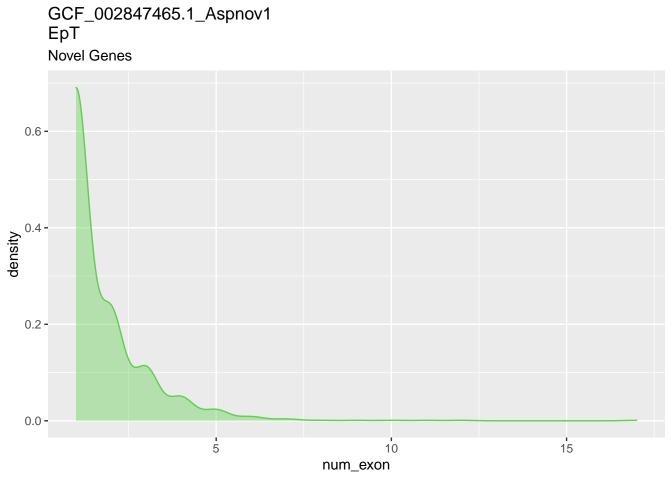


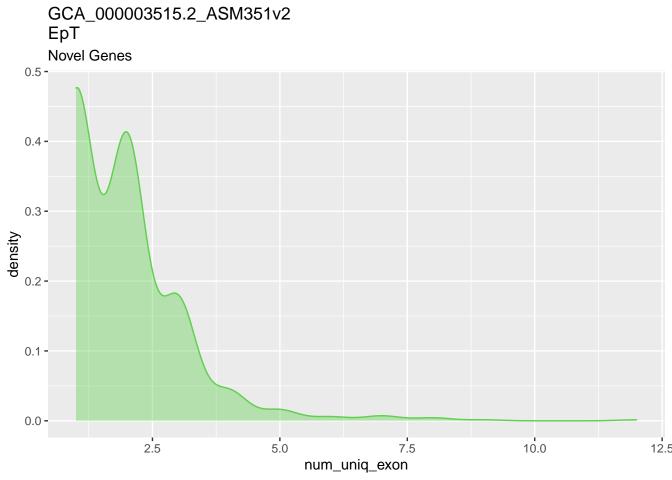


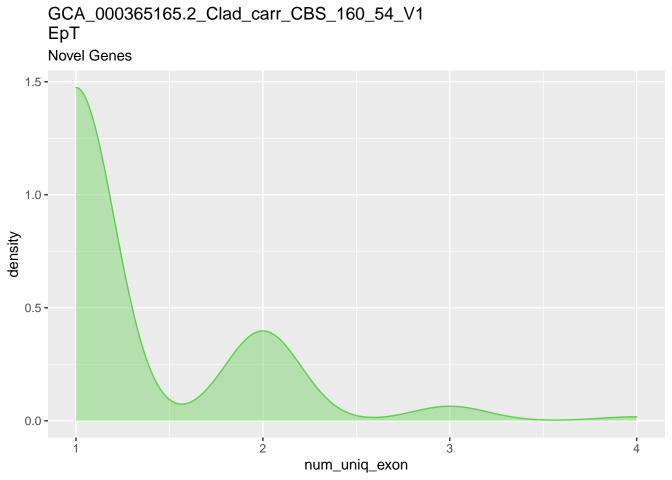


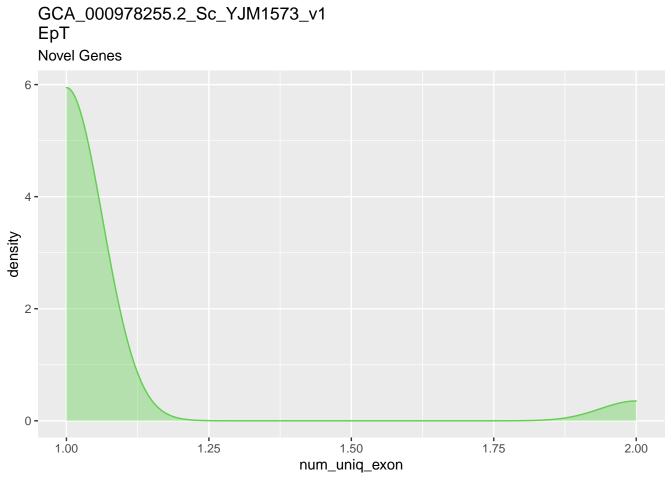


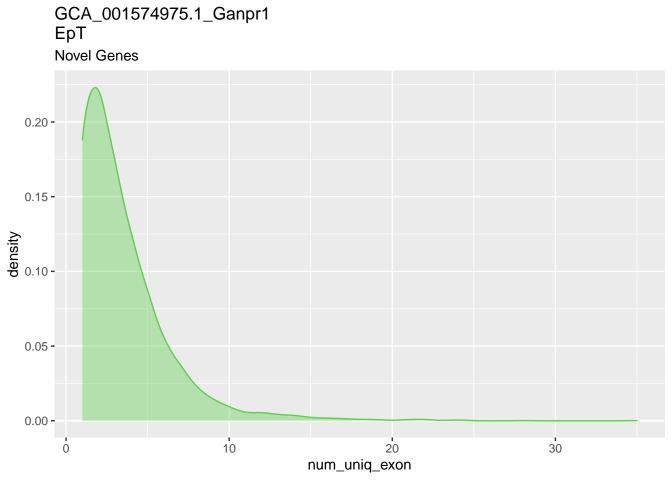


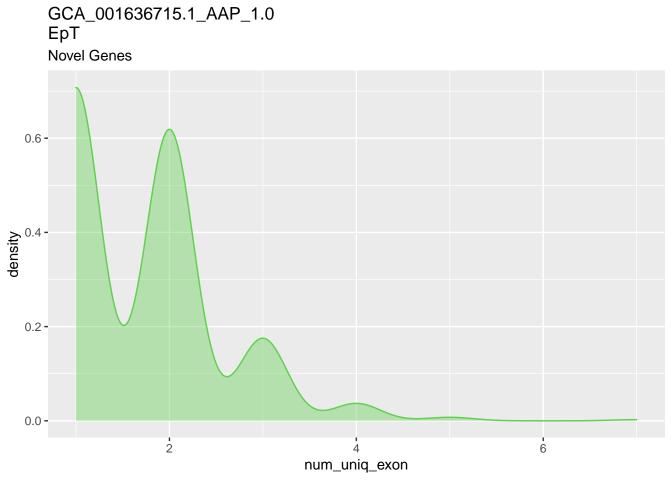


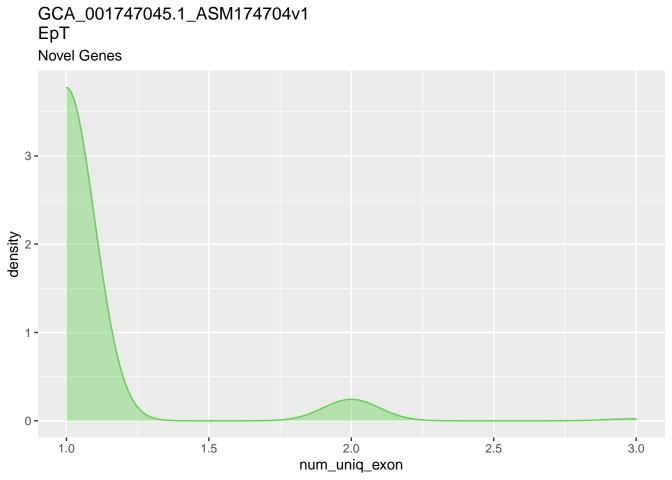


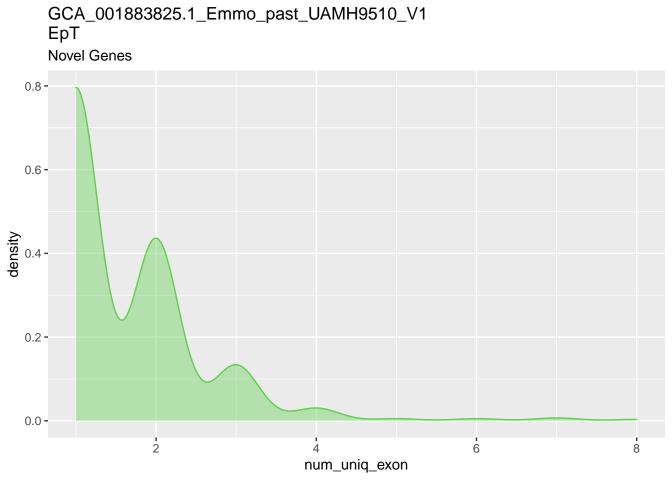


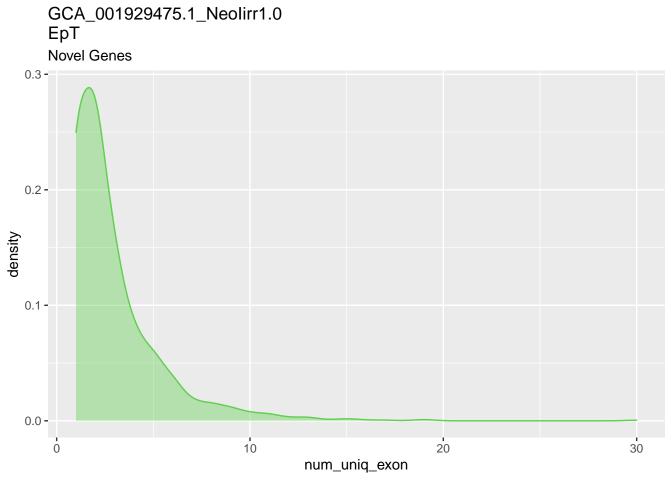


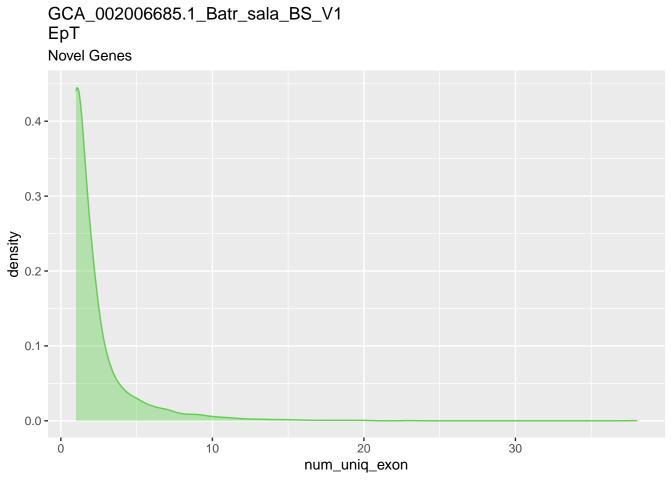


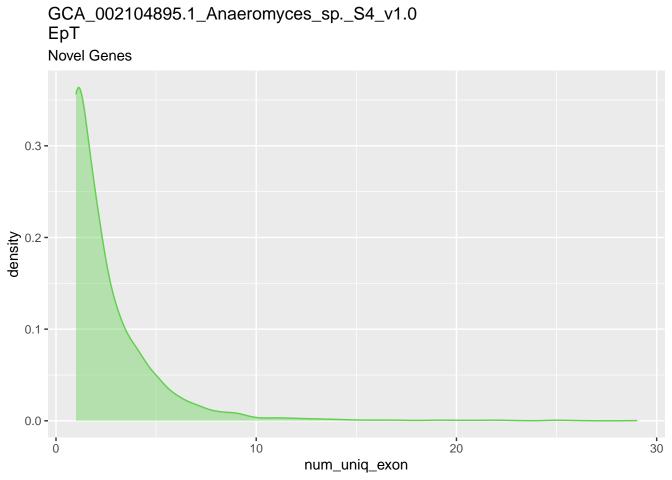


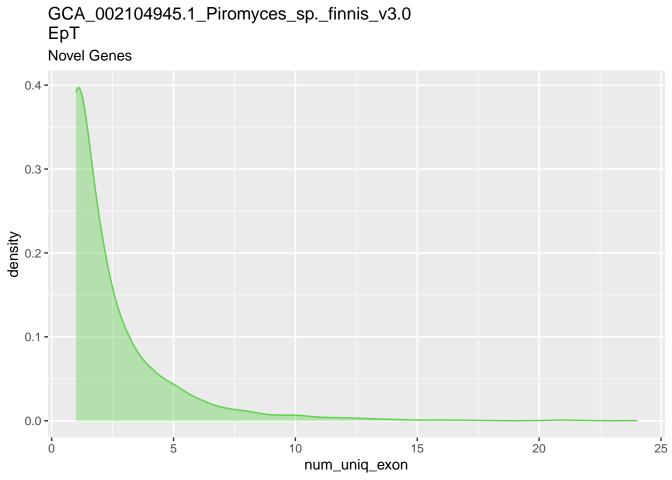


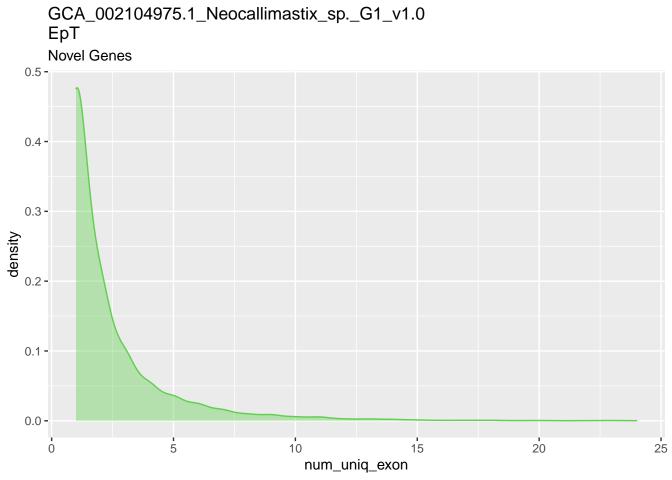


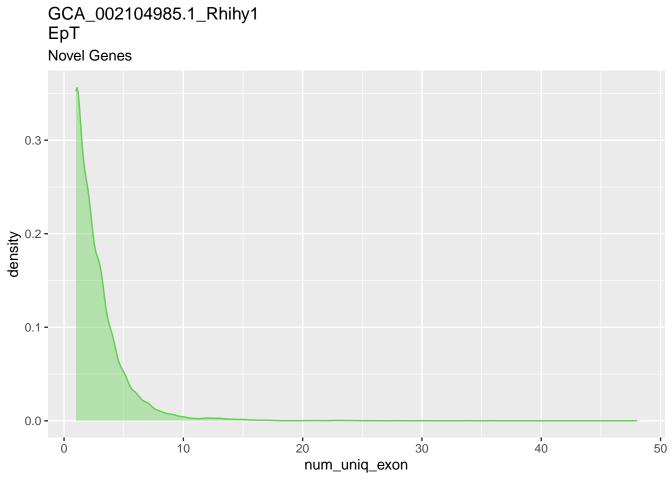


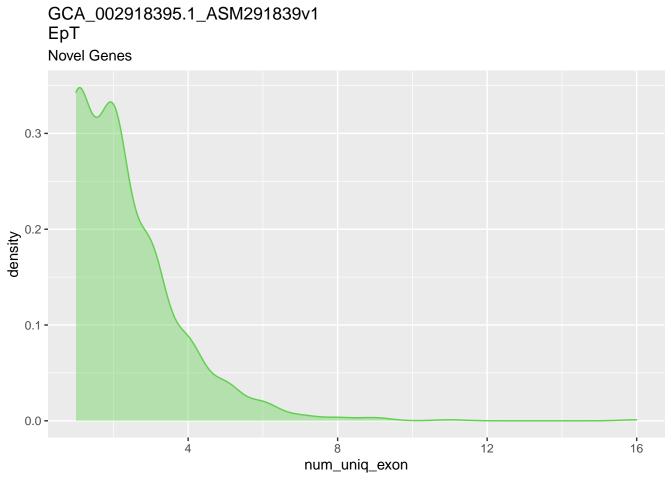


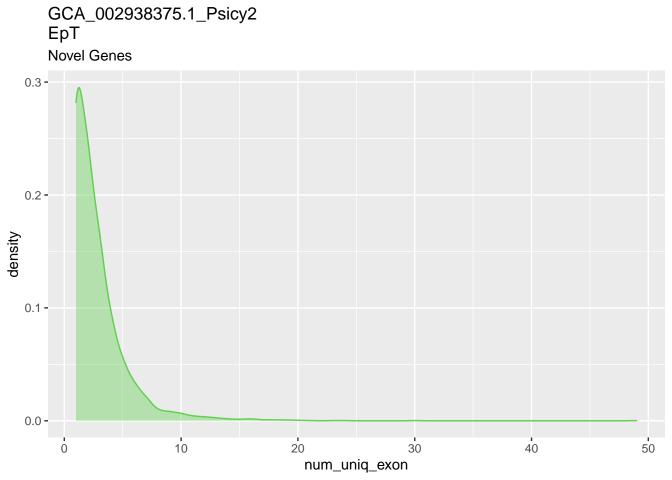


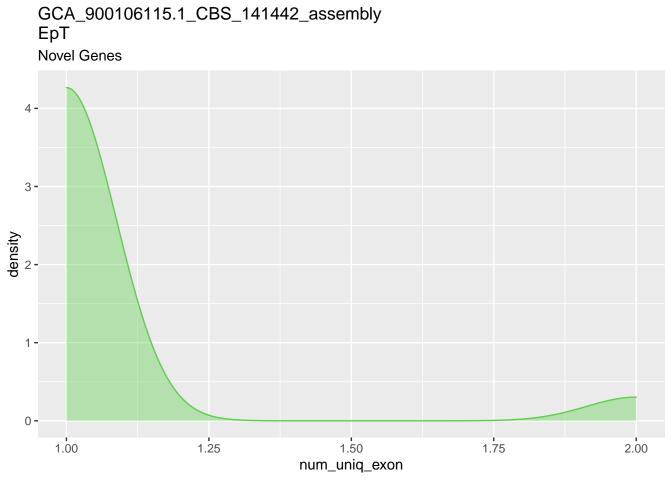


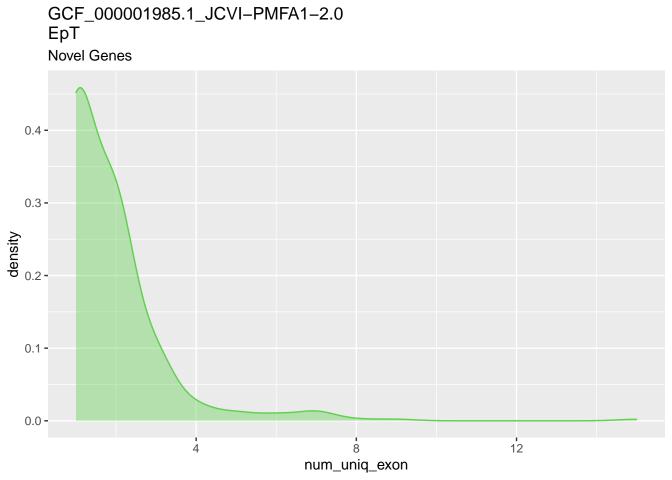


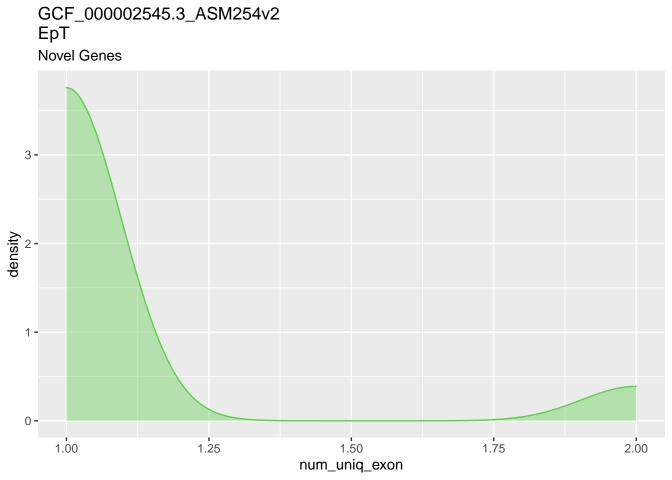


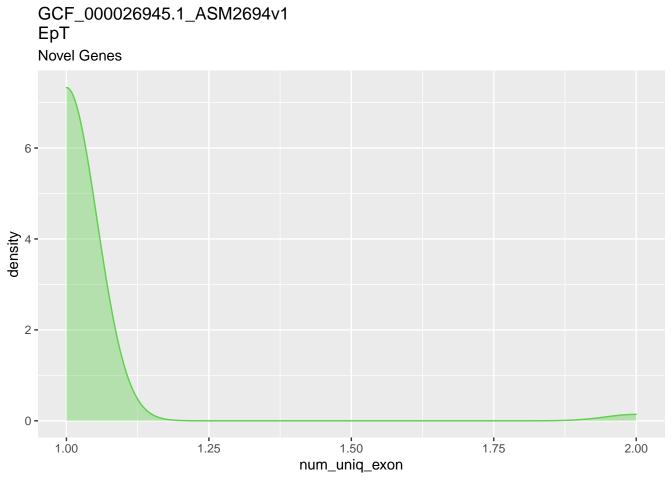


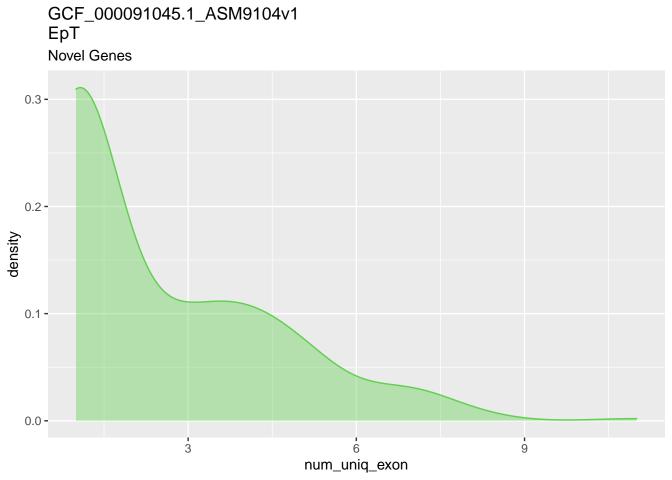


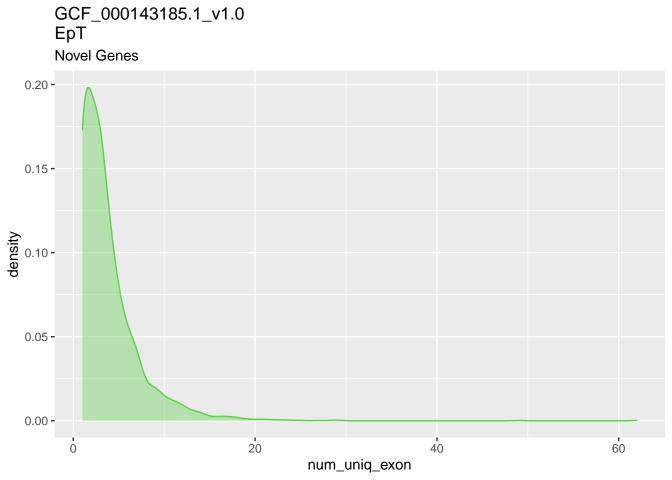


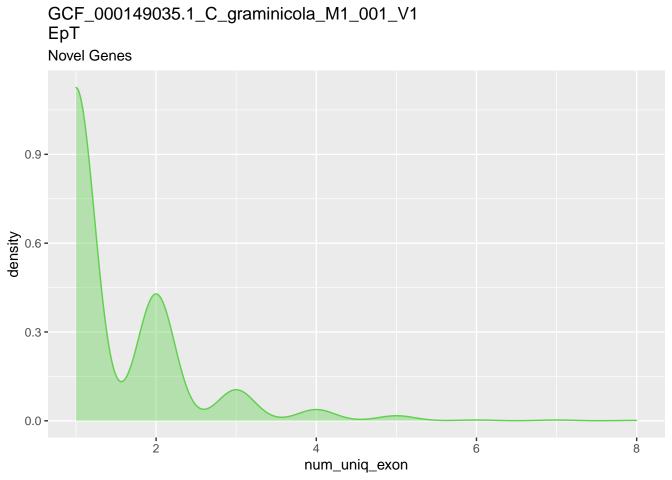












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