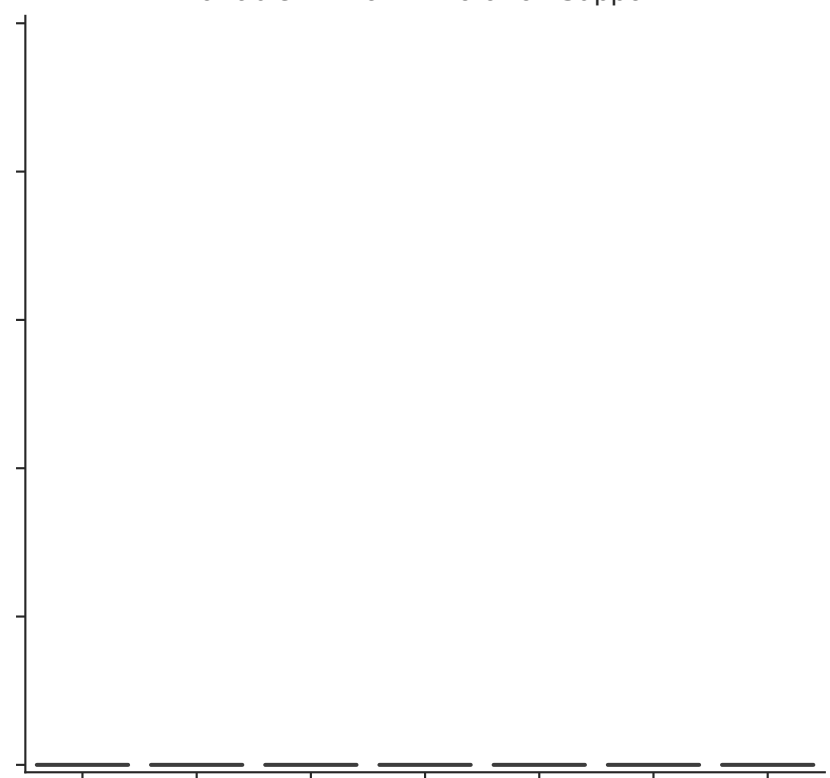
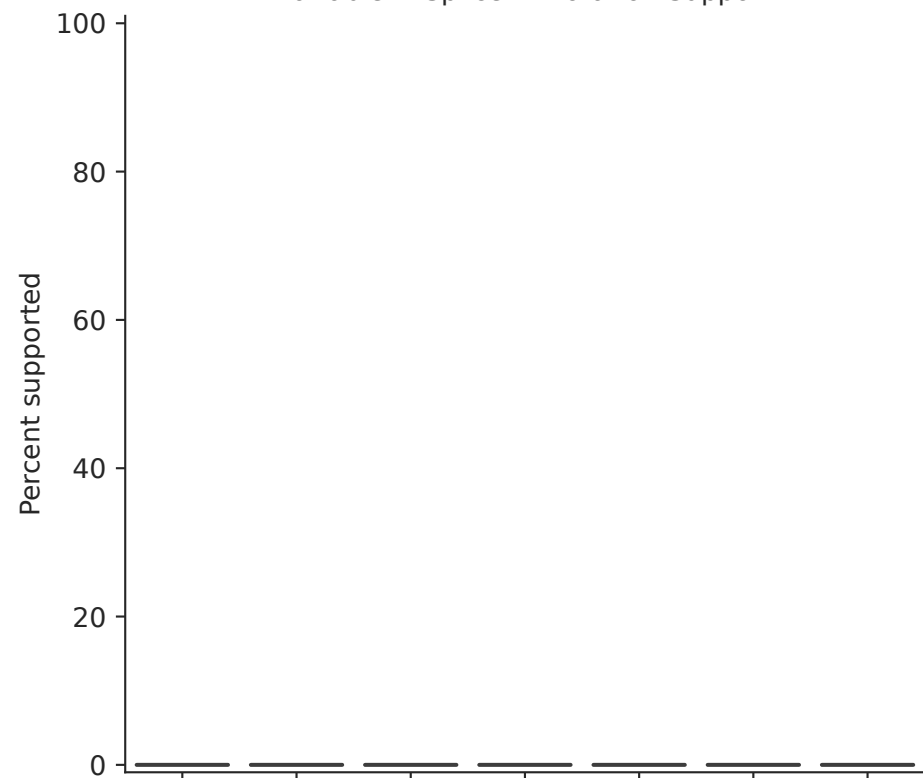


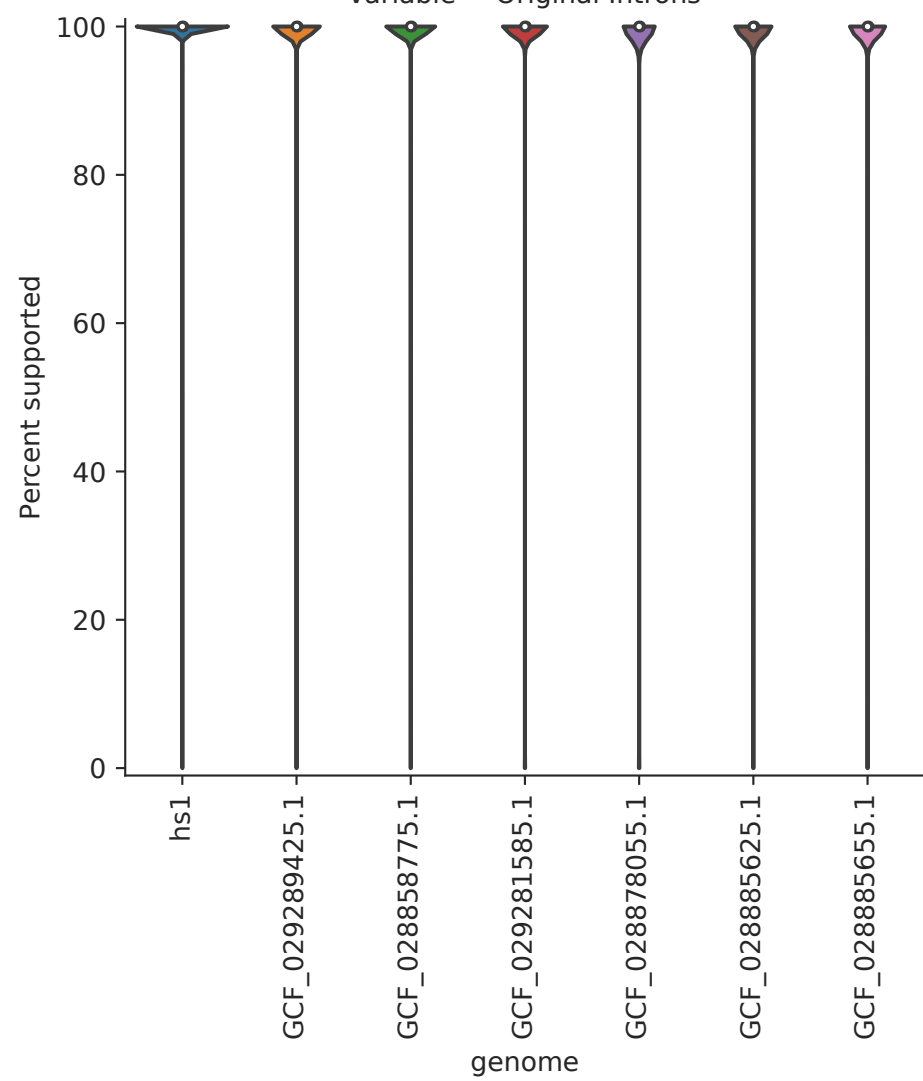
# Reference annotation support

variable = Splice Annotation Support

variable = Exon Annotation Support



variable = Original Introns



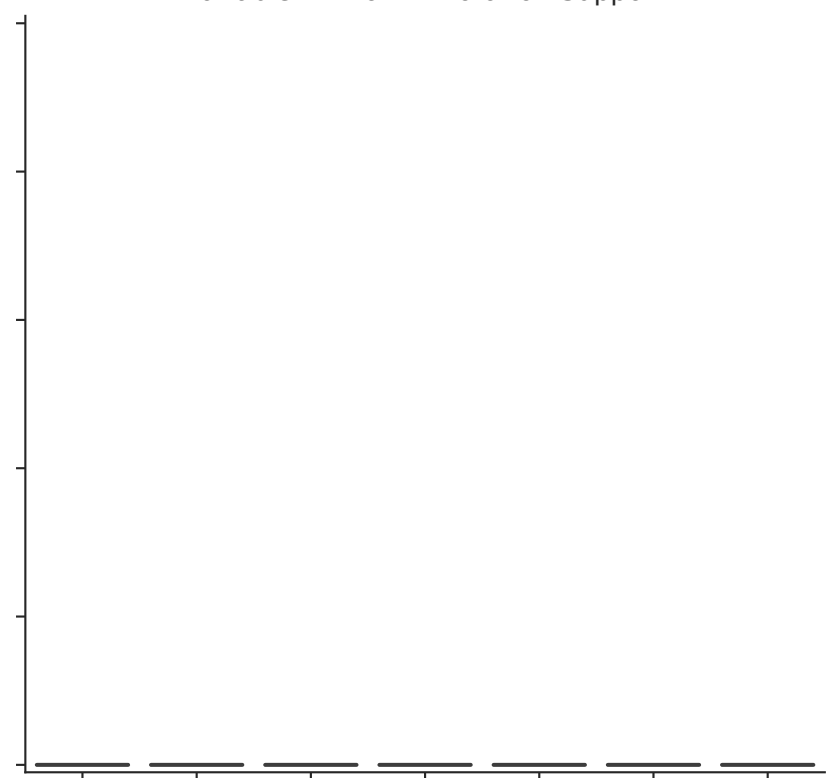
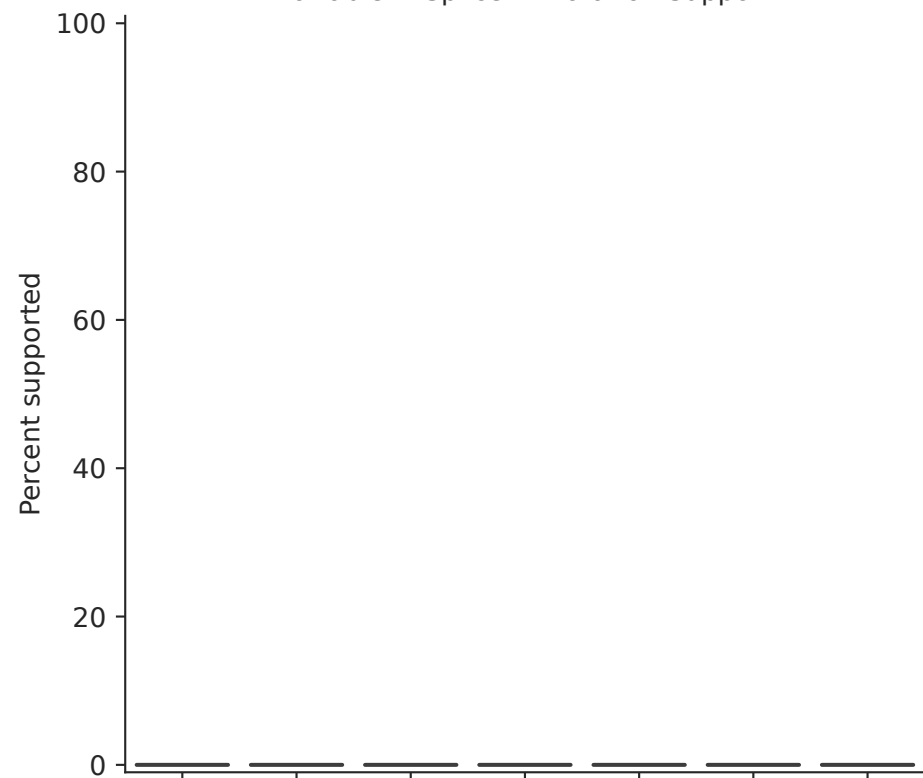
hs1  
GCF\_029289425.1  
GCF\_028858775.1  
GCF\_029281585.1  
GCF\_028878055.1  
GCF\_028885625.1  
GCF\_028885655.1

genome

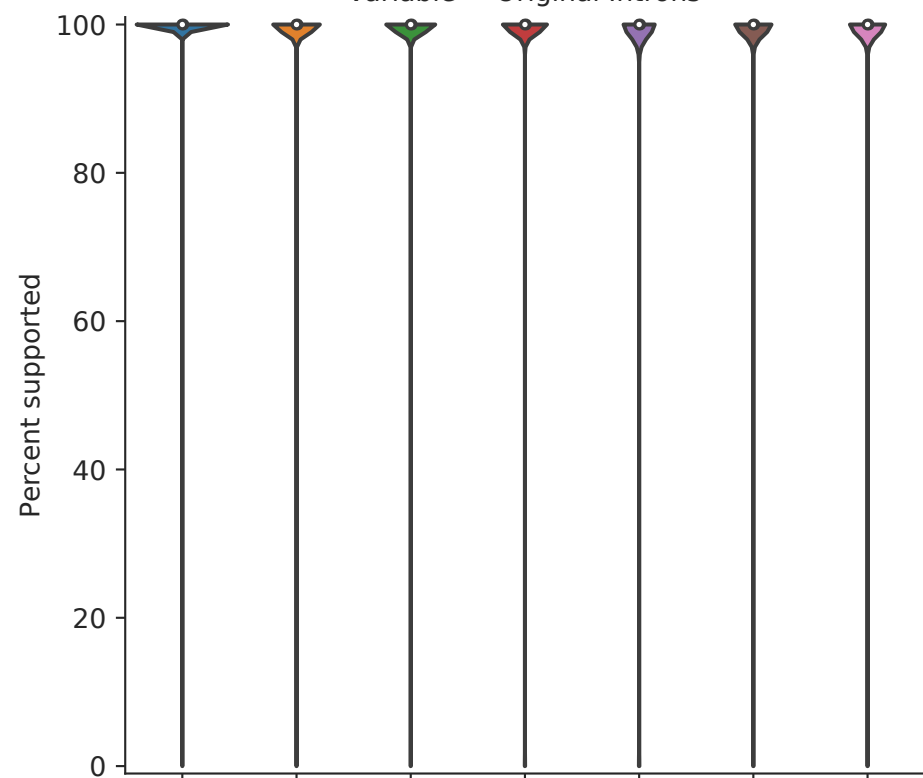
# Reference annotation support for protein\_coding

variable = Splice Annotation Support

variable = Exon Annotation Support



variable = Original Introns



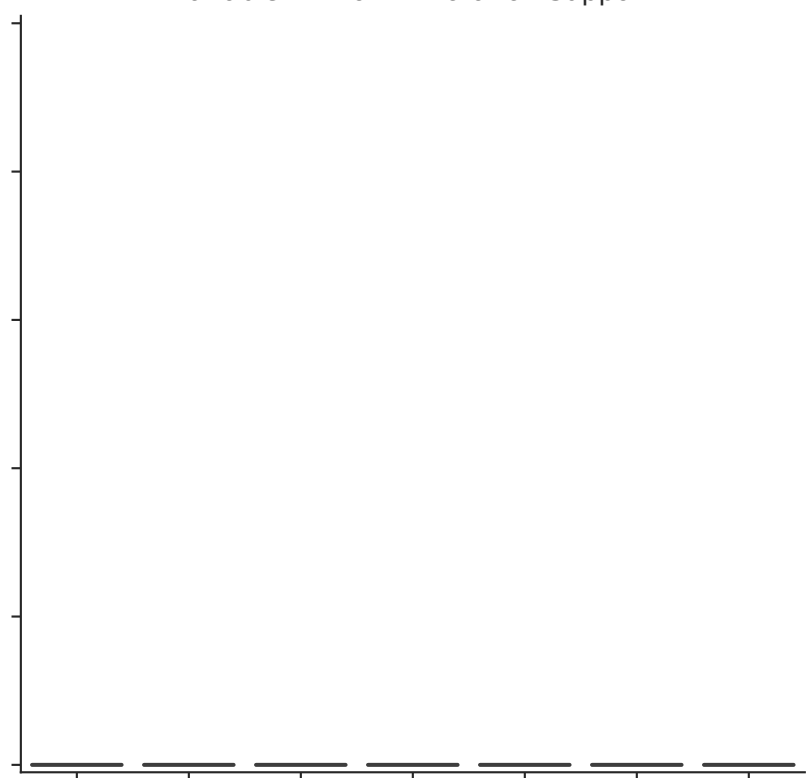
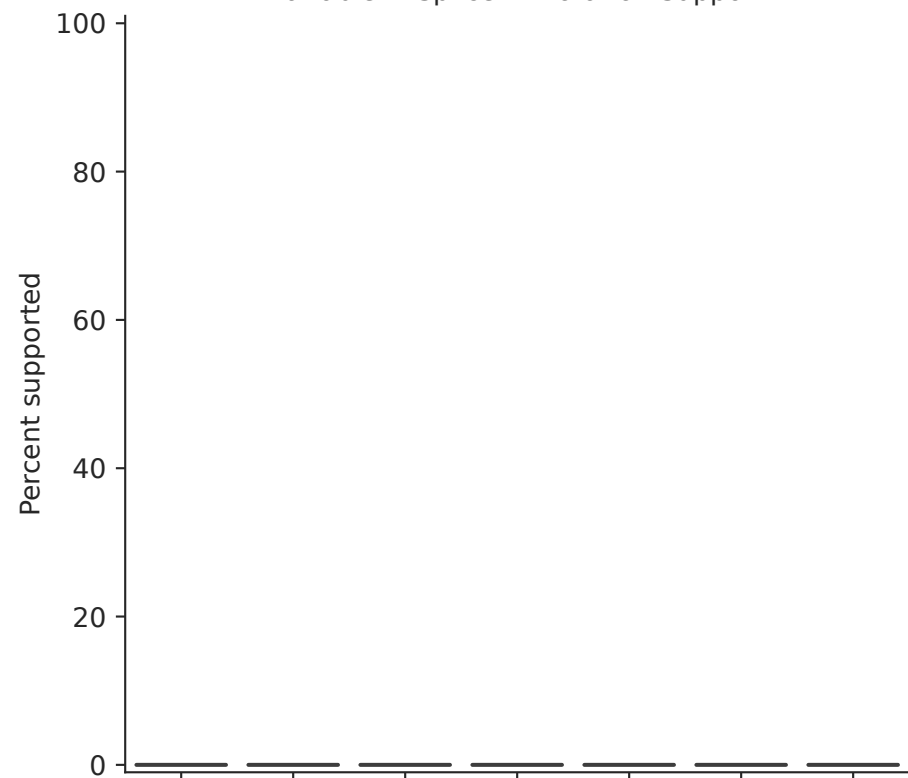
hs1  
GCF\_029289425.1  
GCF\_028858775.1  
GCF\_029281585.1  
genome  
GCF\_028878055.1  
GCF\_028885625.1  
GCF\_028885655.1

genome

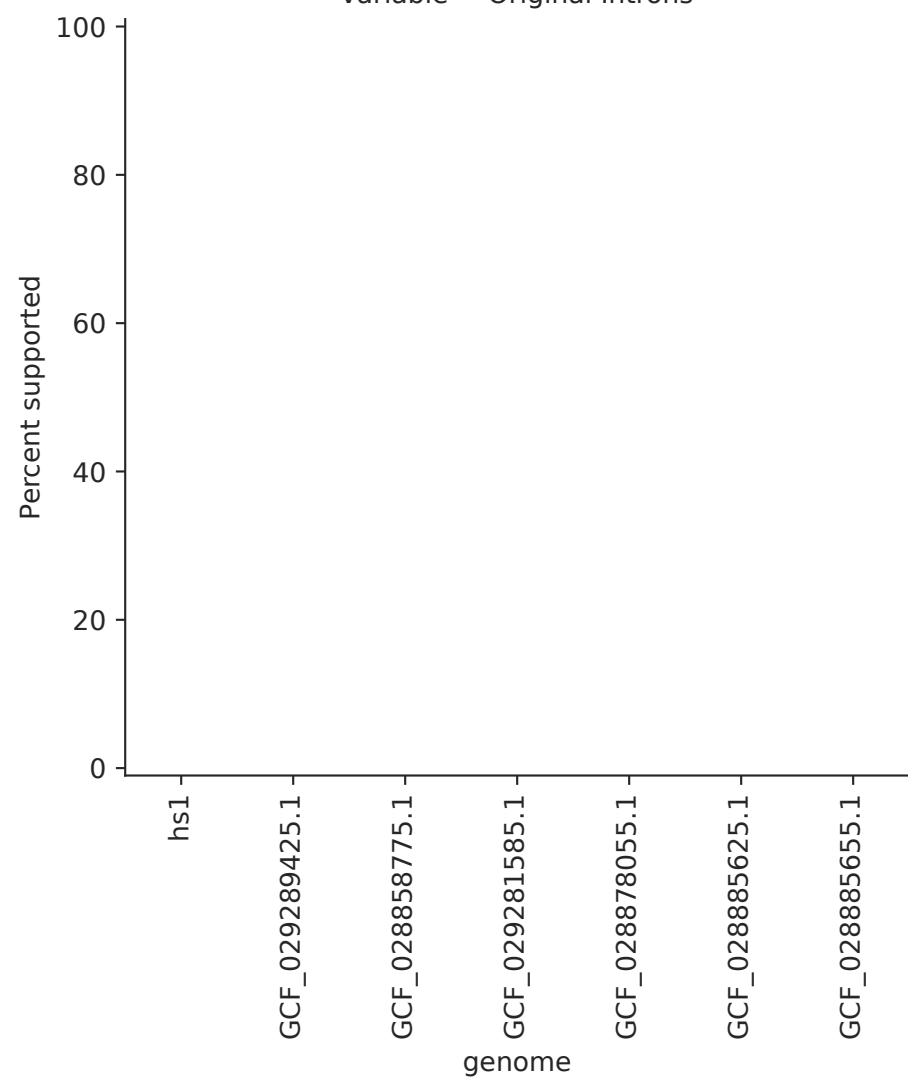
# Reference annotation support for IG\_C\_gene

variable = Splice Annotation Support

variable = Exon Annotation Support



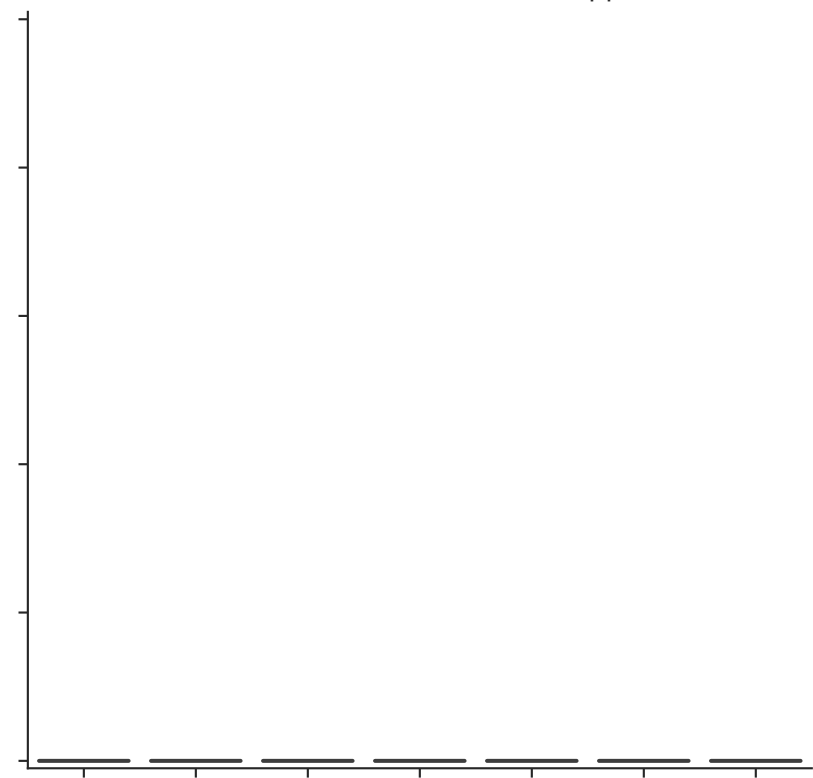
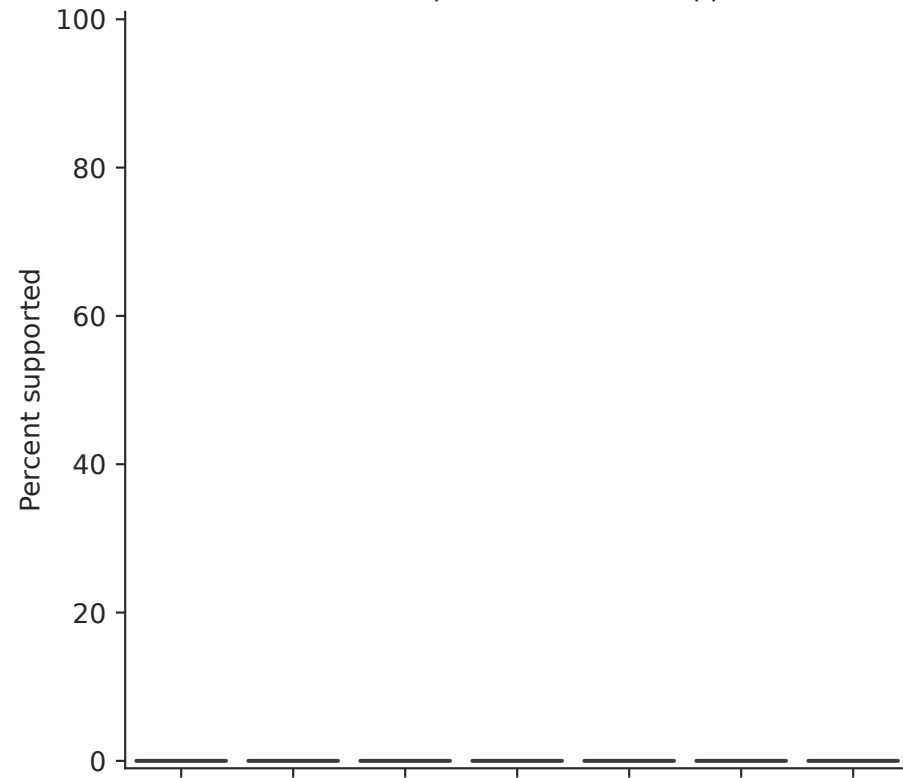
variable = Original Introns



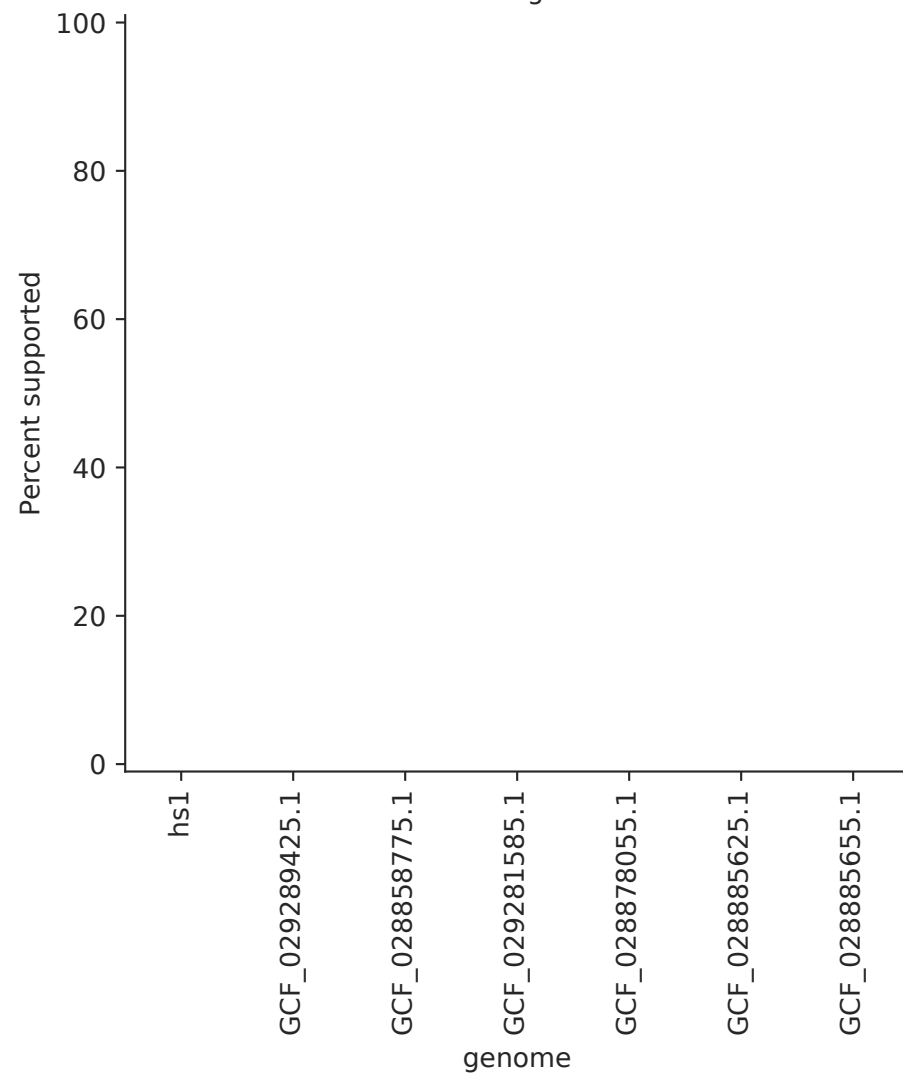
# Reference annotation support for IG\_C\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



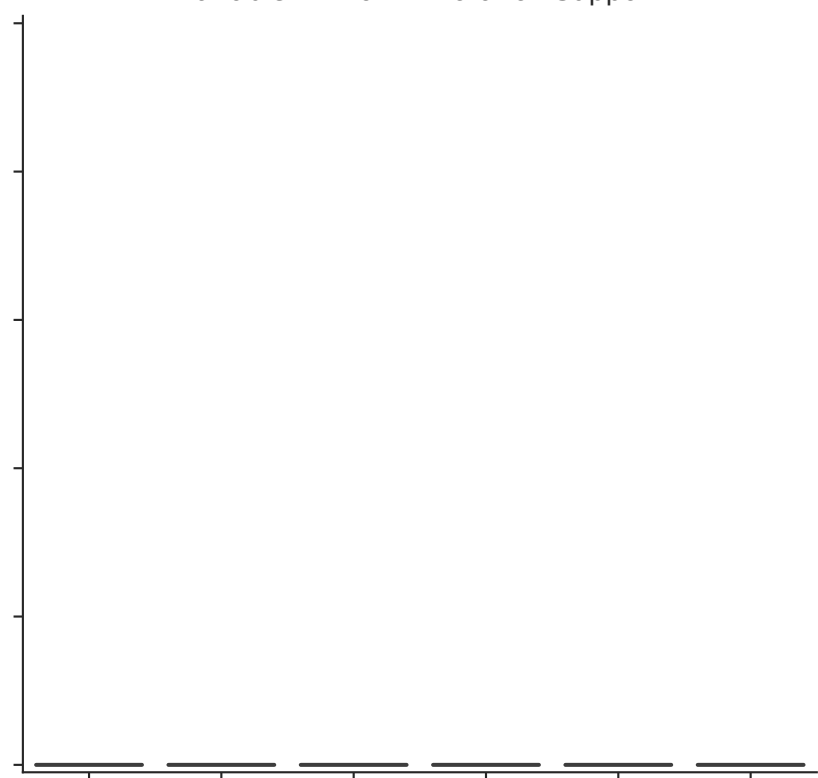
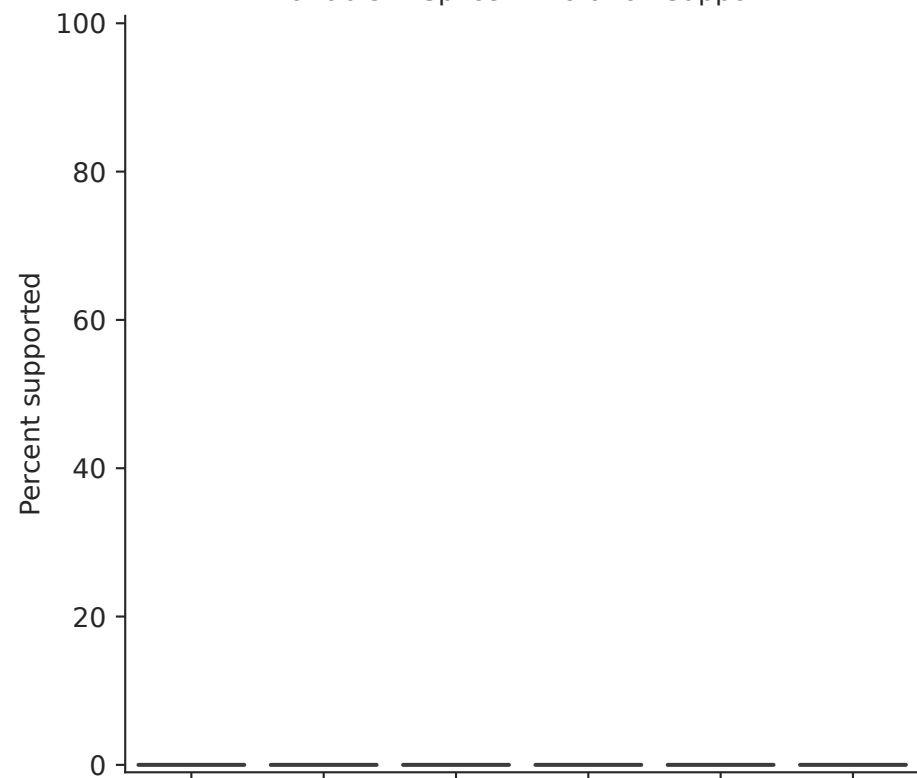
variable = Original Introns



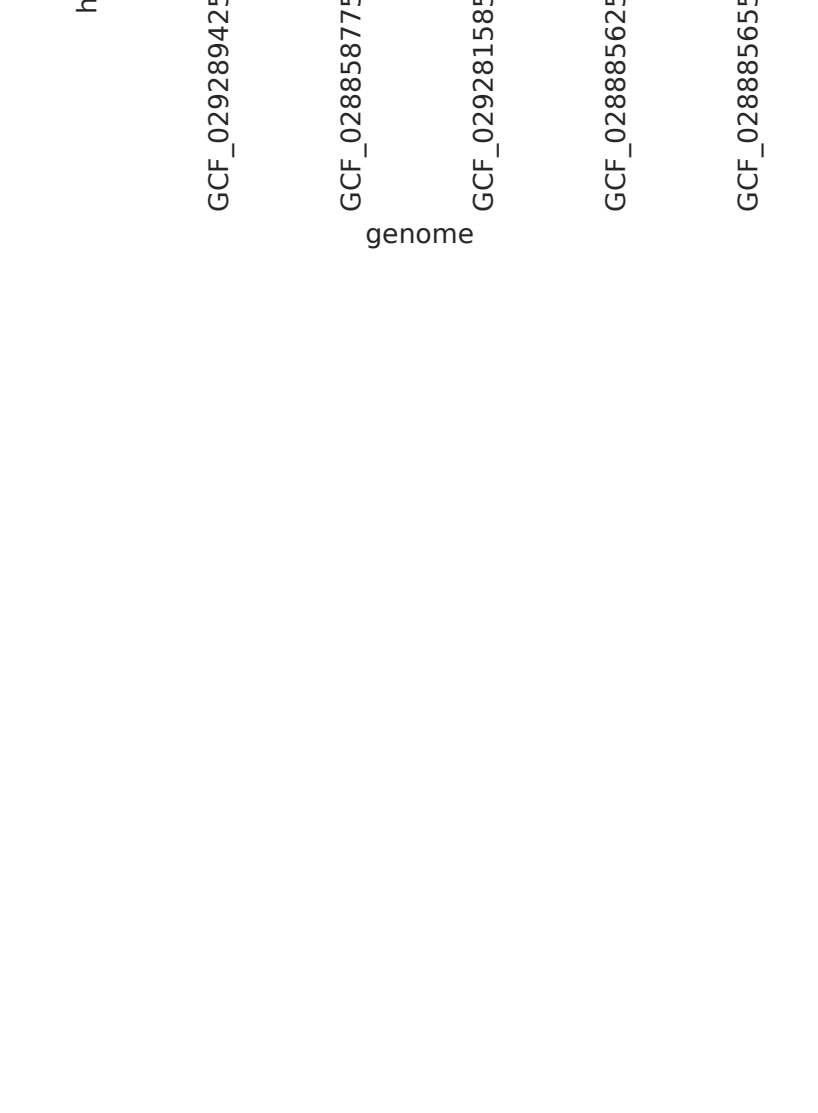
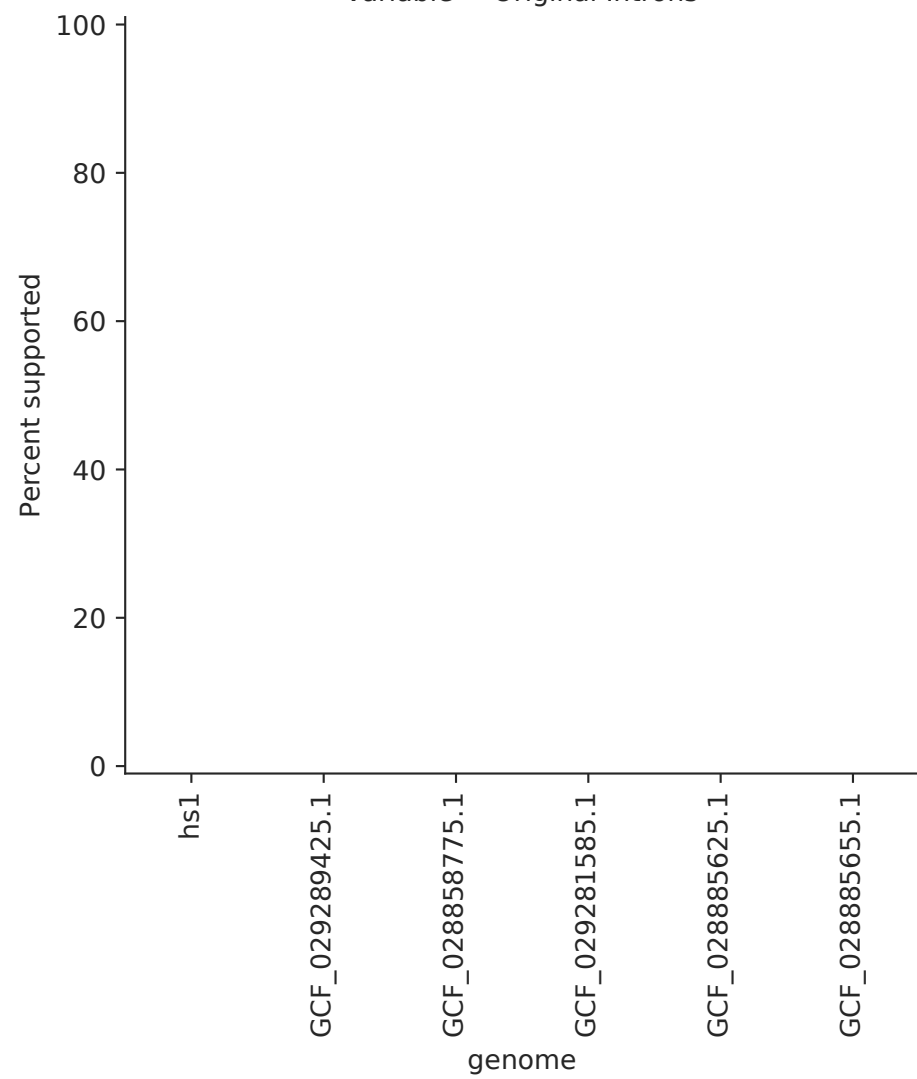
# Reference annotation support for IG\_D\_gene

variable = Splice Annotation Support

variable = Exon Annotation Support



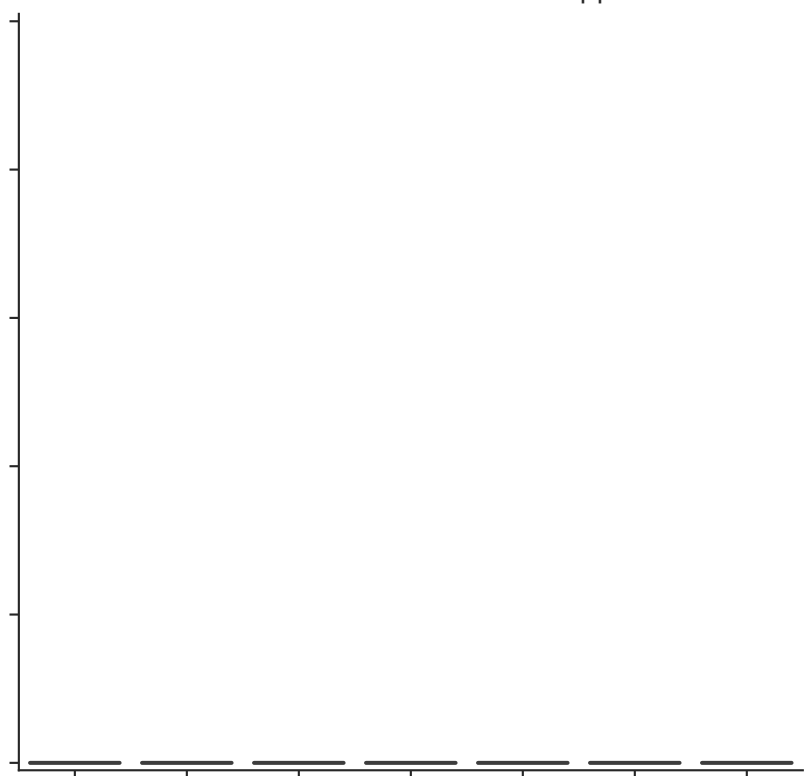
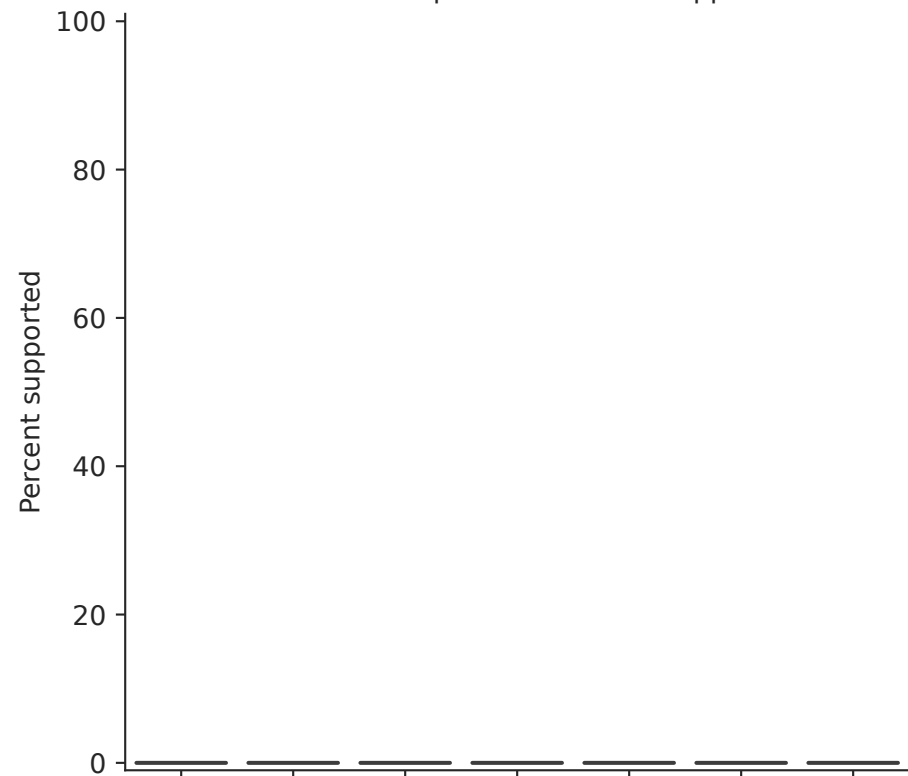
variable = Original Introns



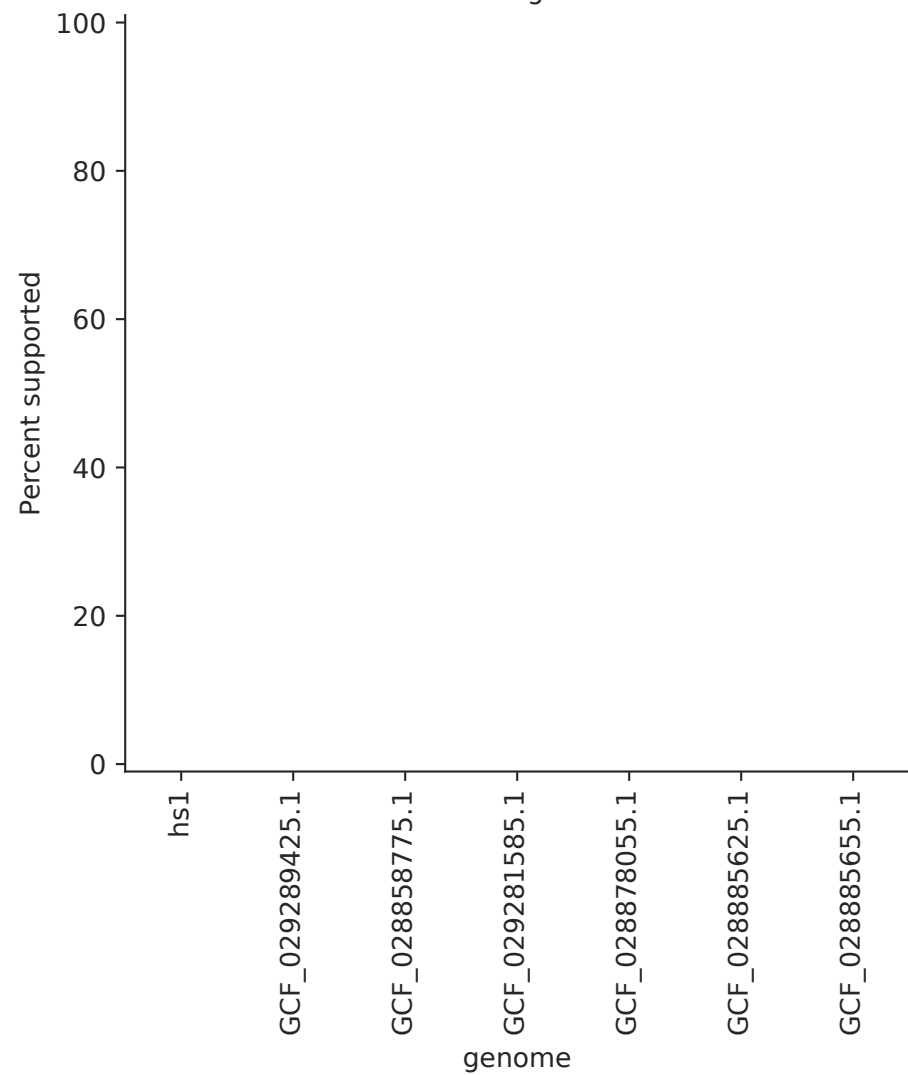
# Reference annotation support for IG\_J\_gene

variable = Splice Annotation Support

variable = Exon Annotation Support



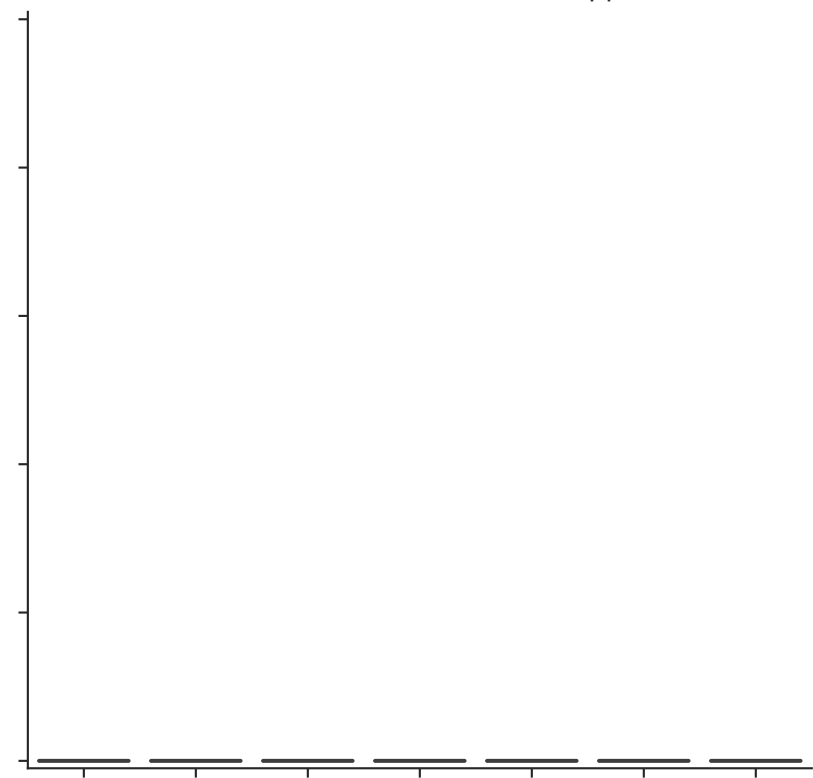
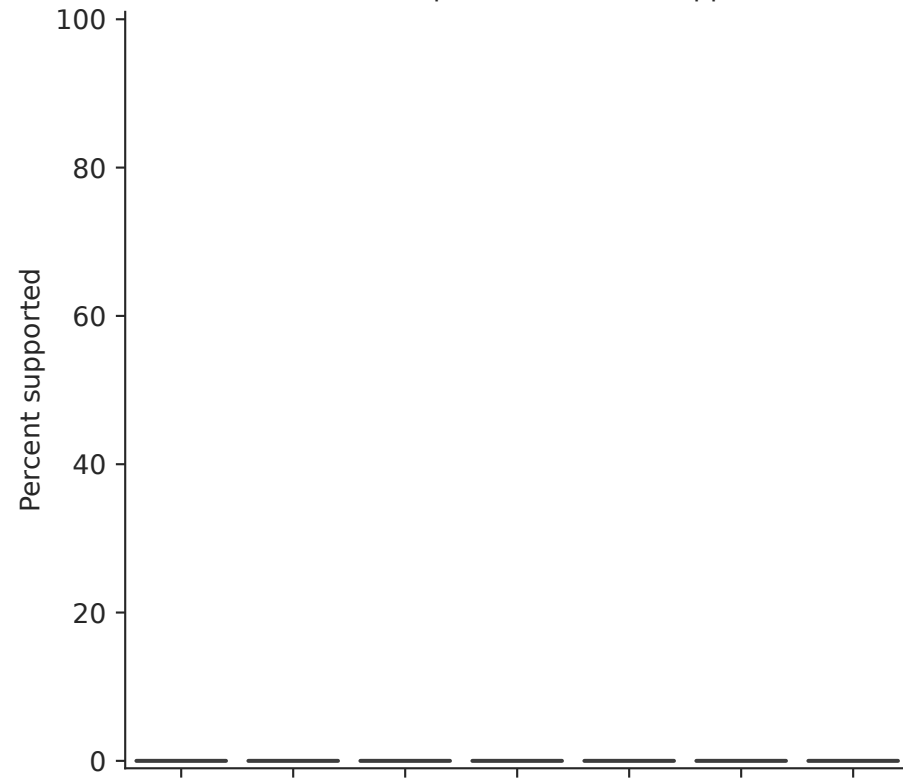
variable = Original Introns



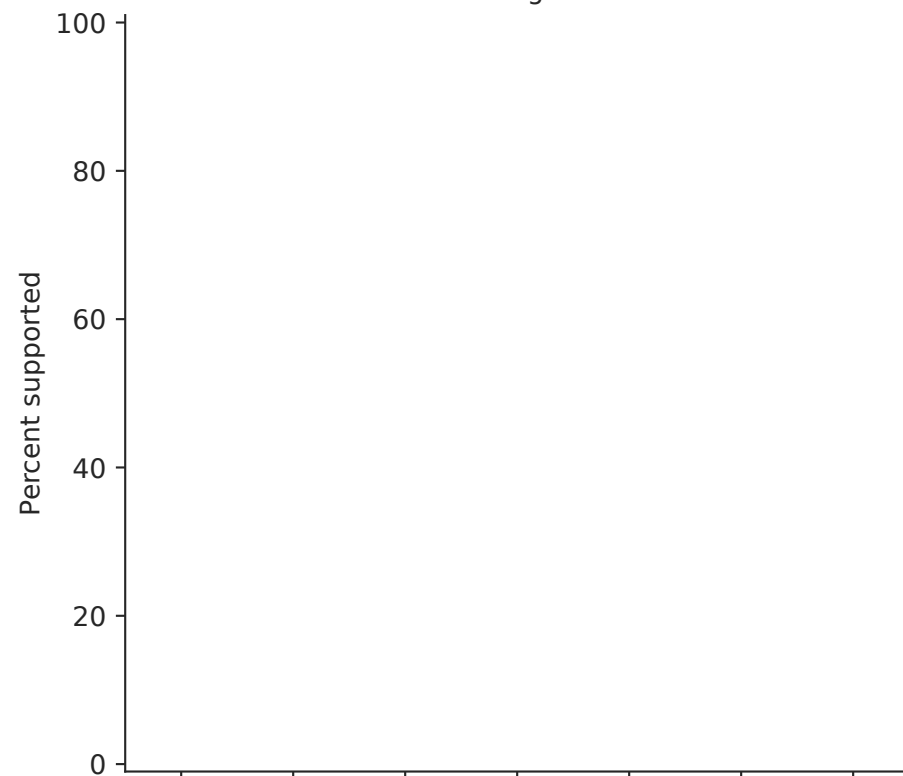
# Reference annotation support for IG\_J\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



variable = Original Introns



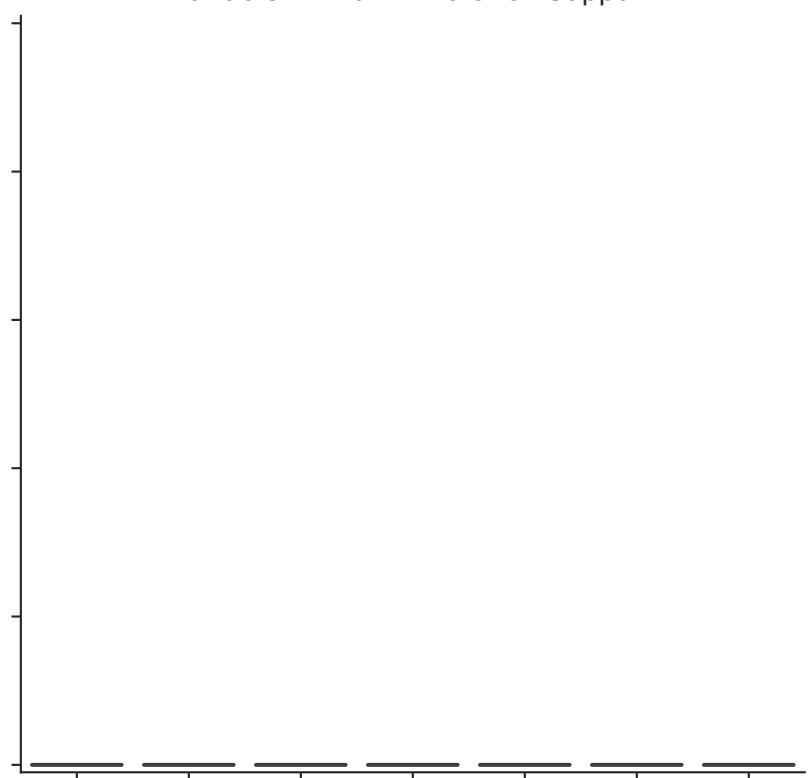
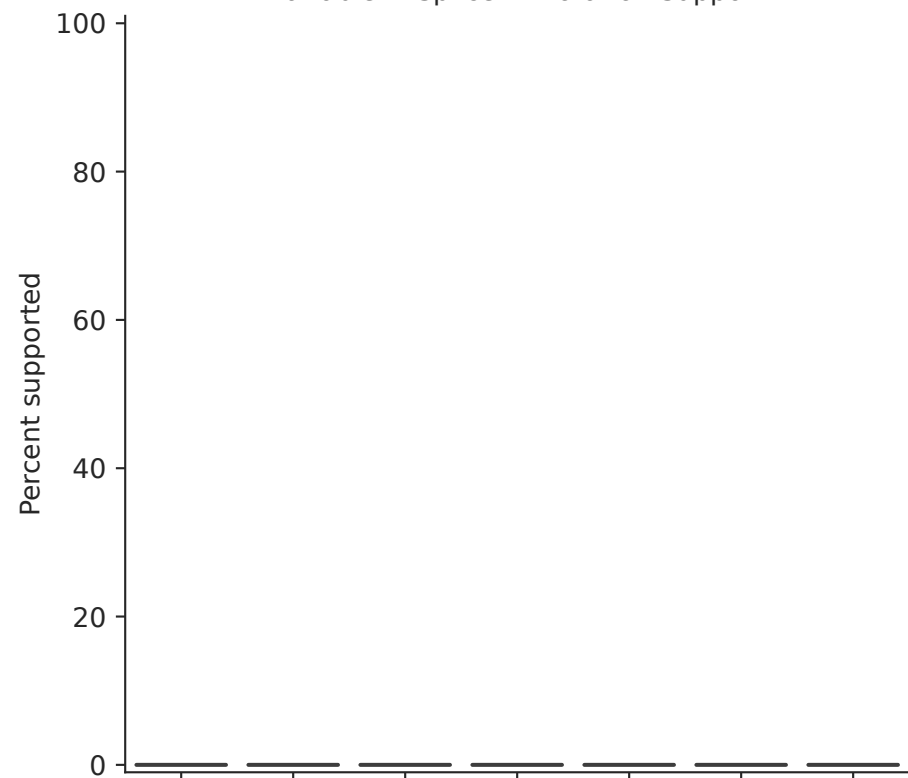
genome

genome

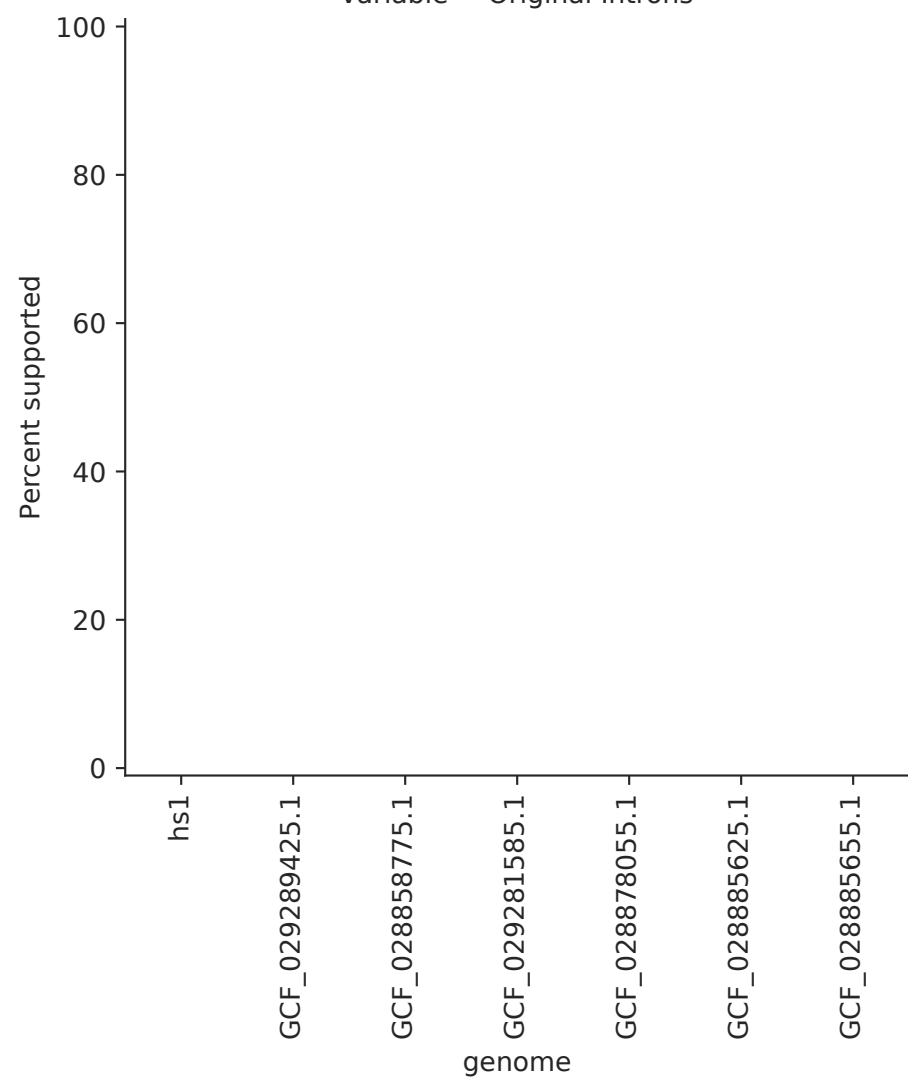
# Reference annotation support for IG\_V\_gene

variable = Splice Annotation Support

variable = Exon Annotation Support



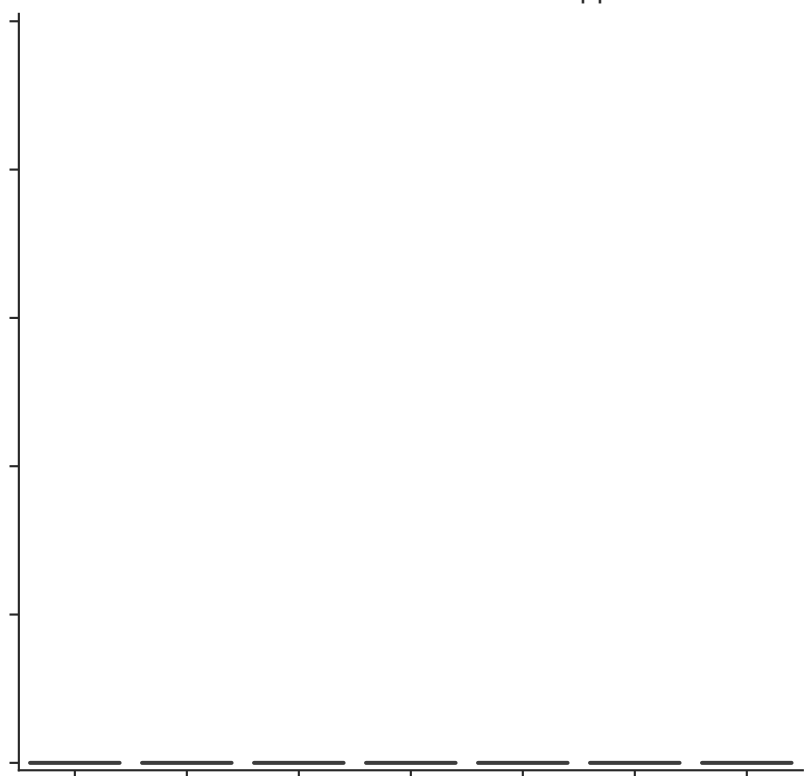
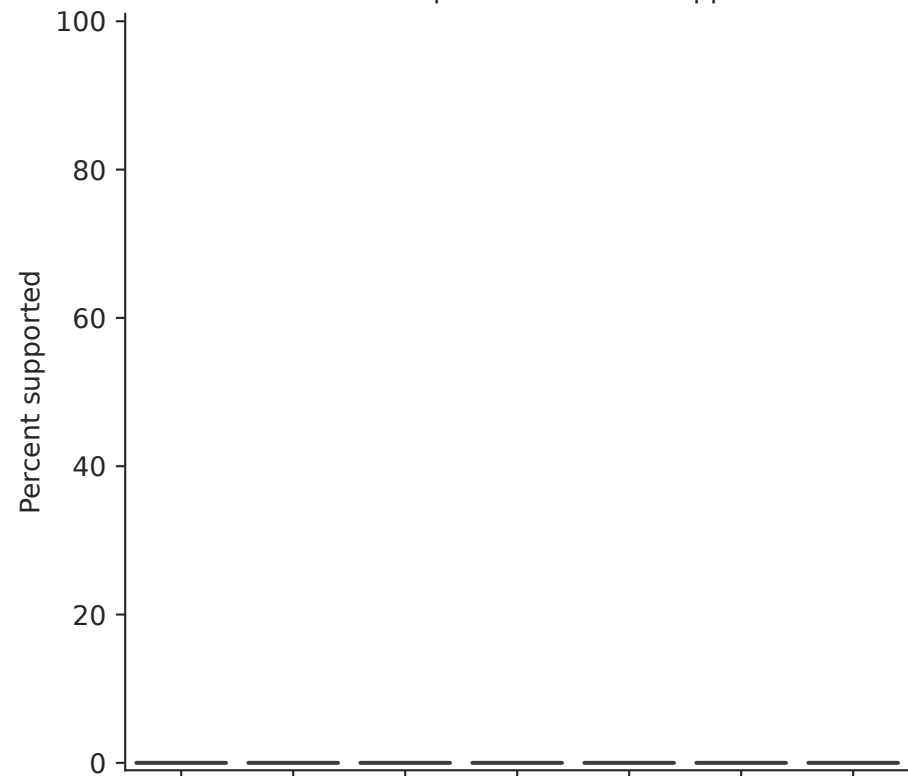
variable = Original Introns



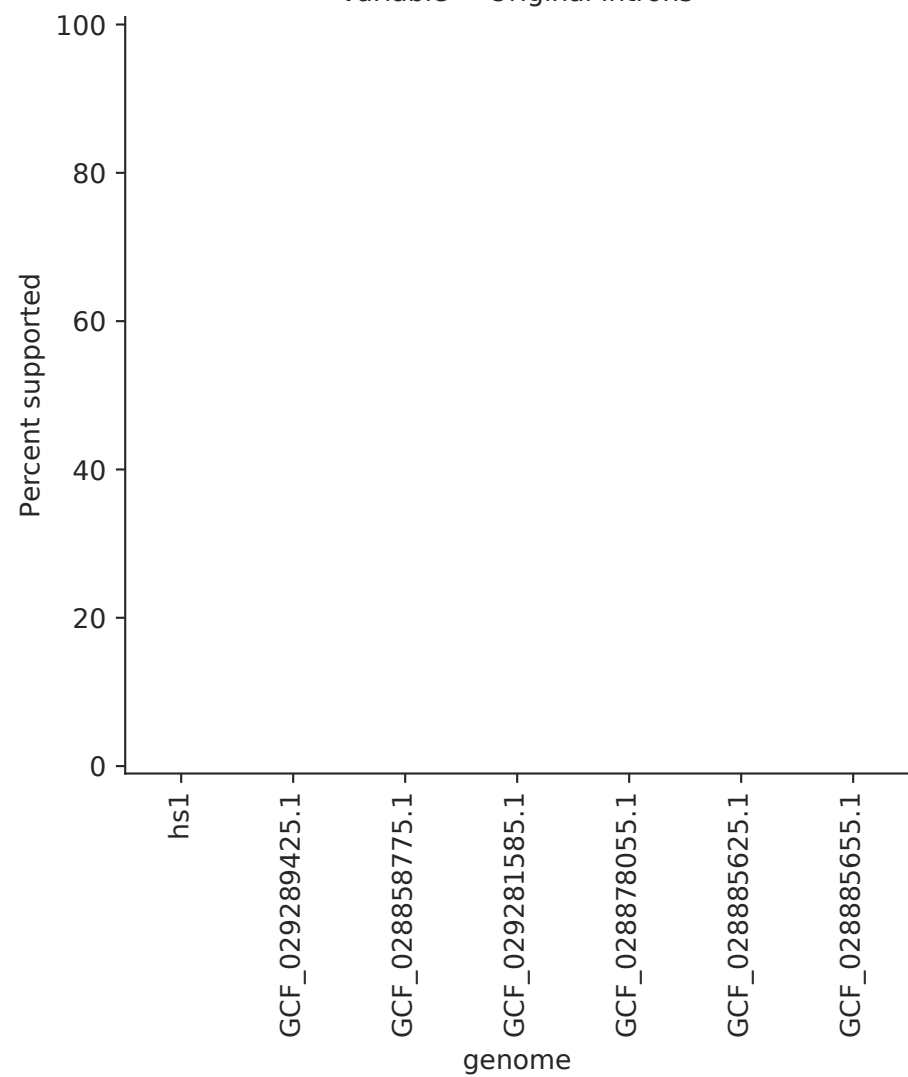
# Reference annotation support for IG\_V\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



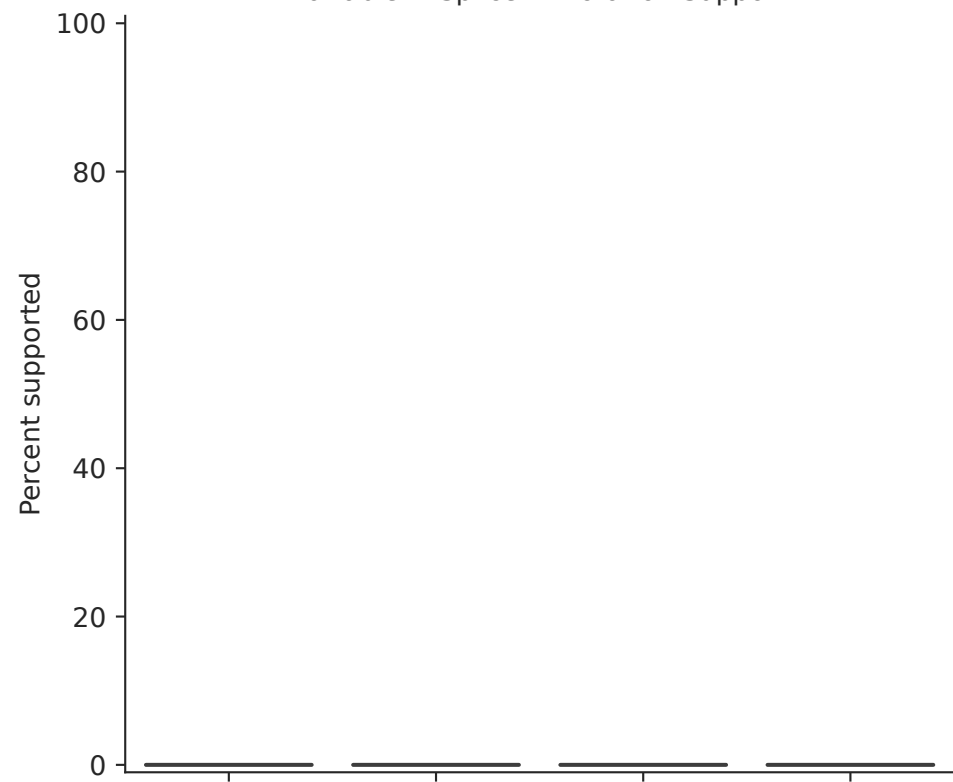
variable = Original Introns



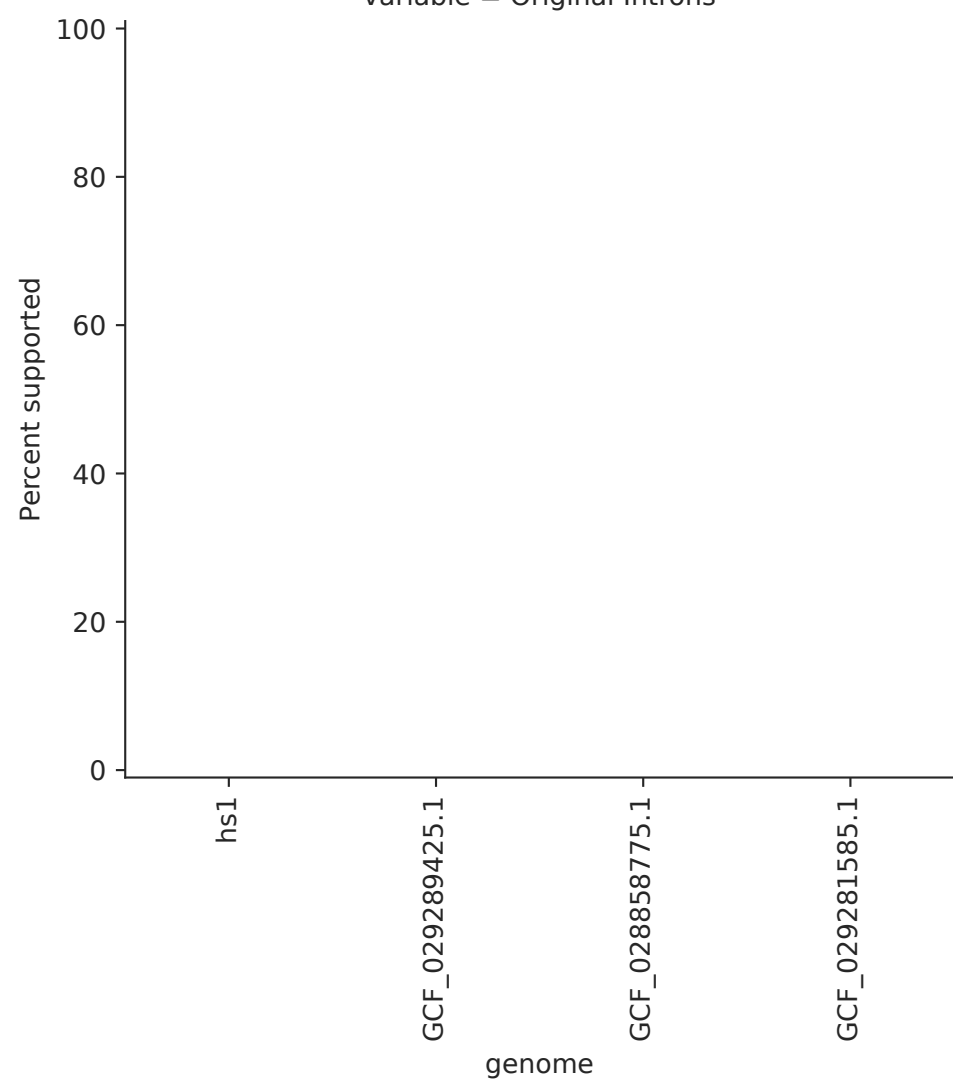
# Reference annotation support for IG\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



variable = Original Introns



genome

hs1

GCF\_029289425.1

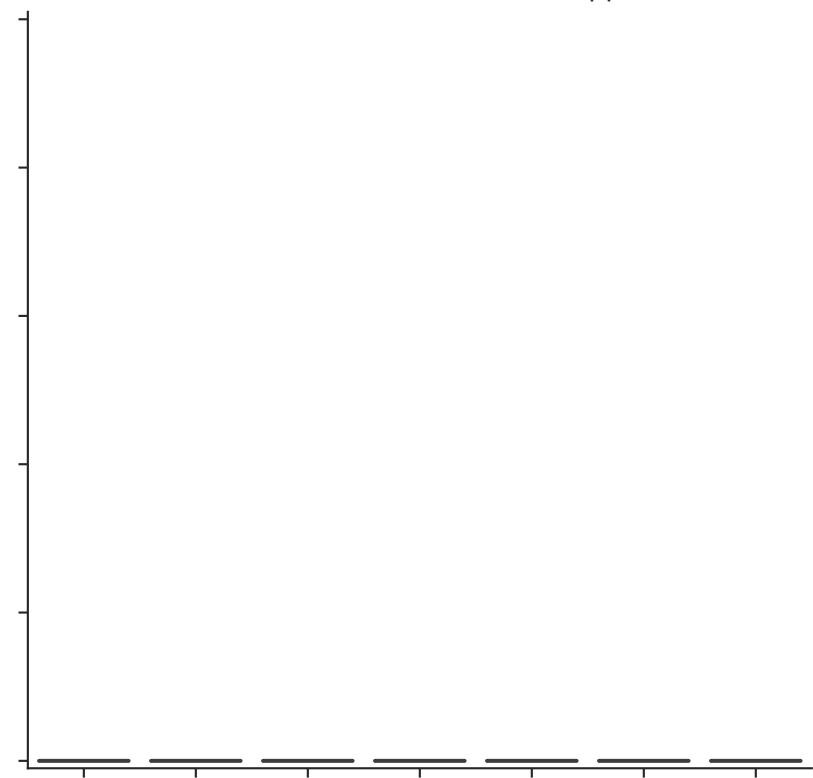
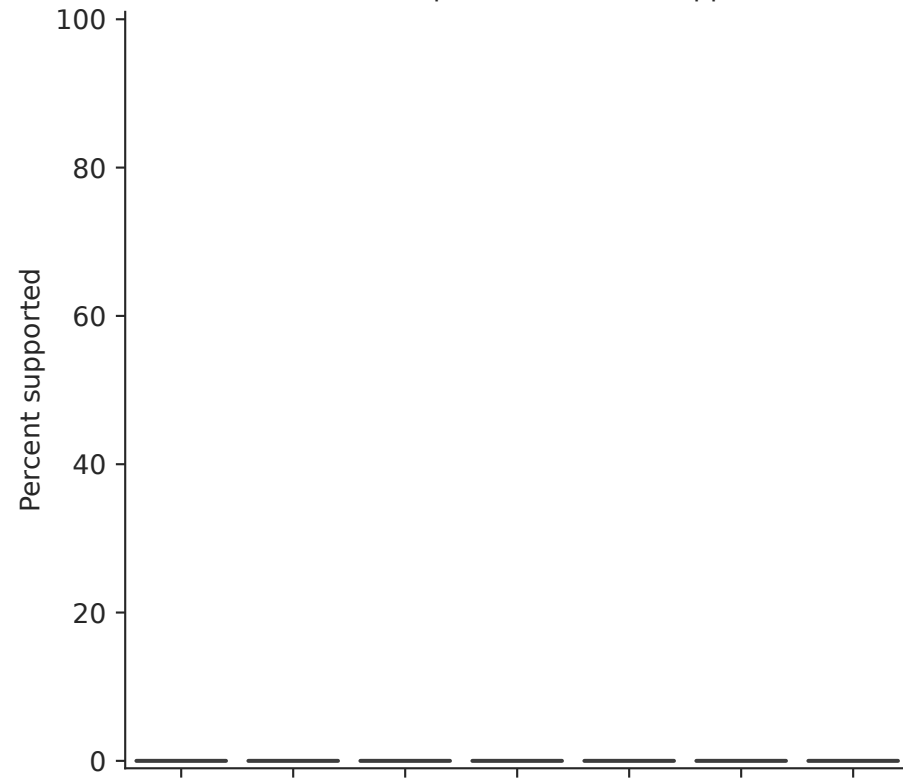
GCF\_028858775.1

GCF\_029281585.1

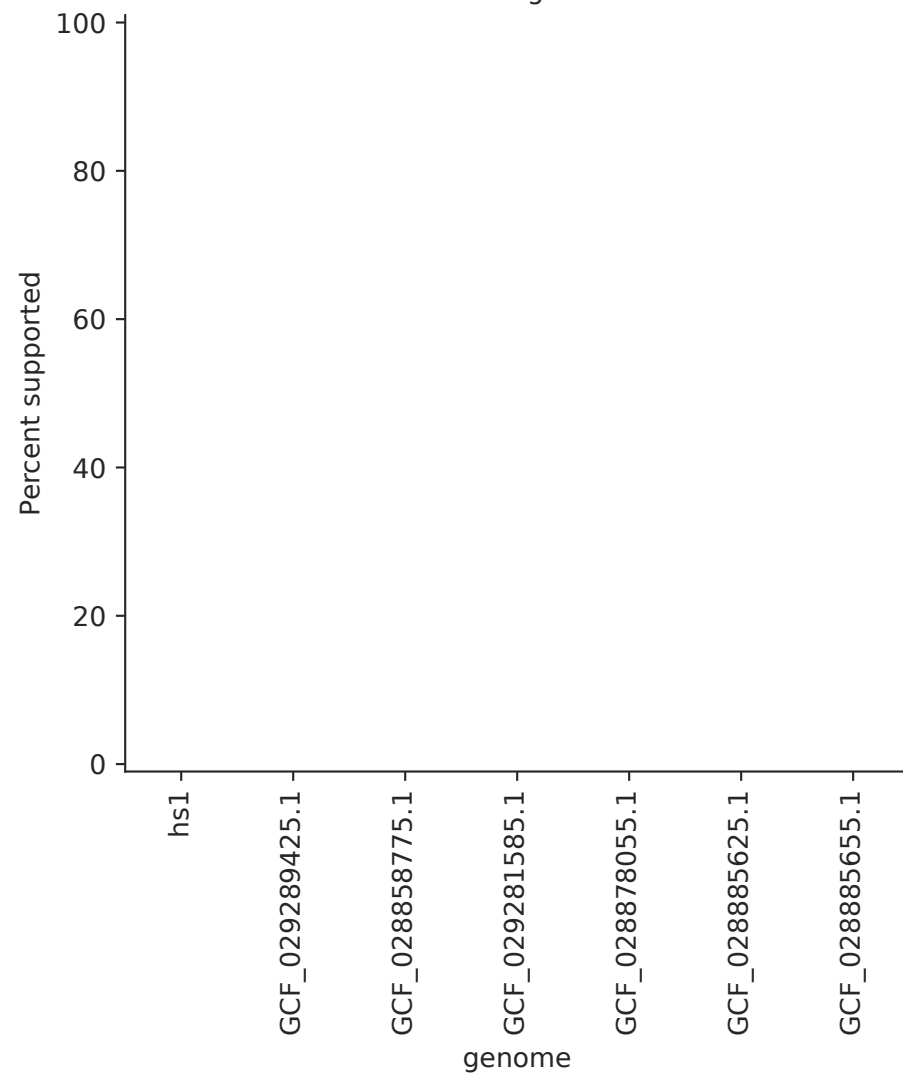
# Reference annotation support for TEC

variable = Splice Annotation Support

variable = Exon Annotation Support



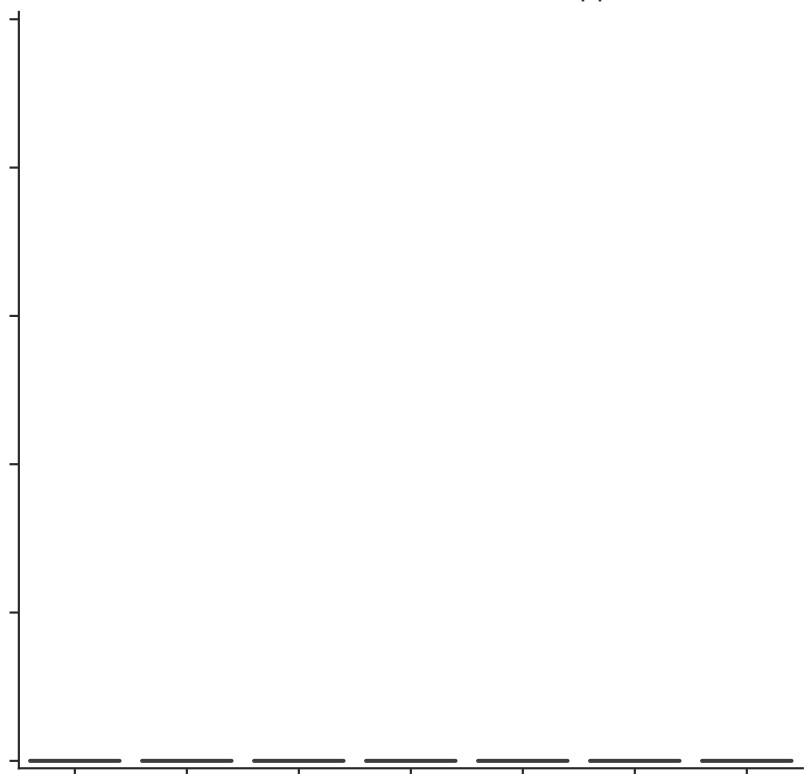
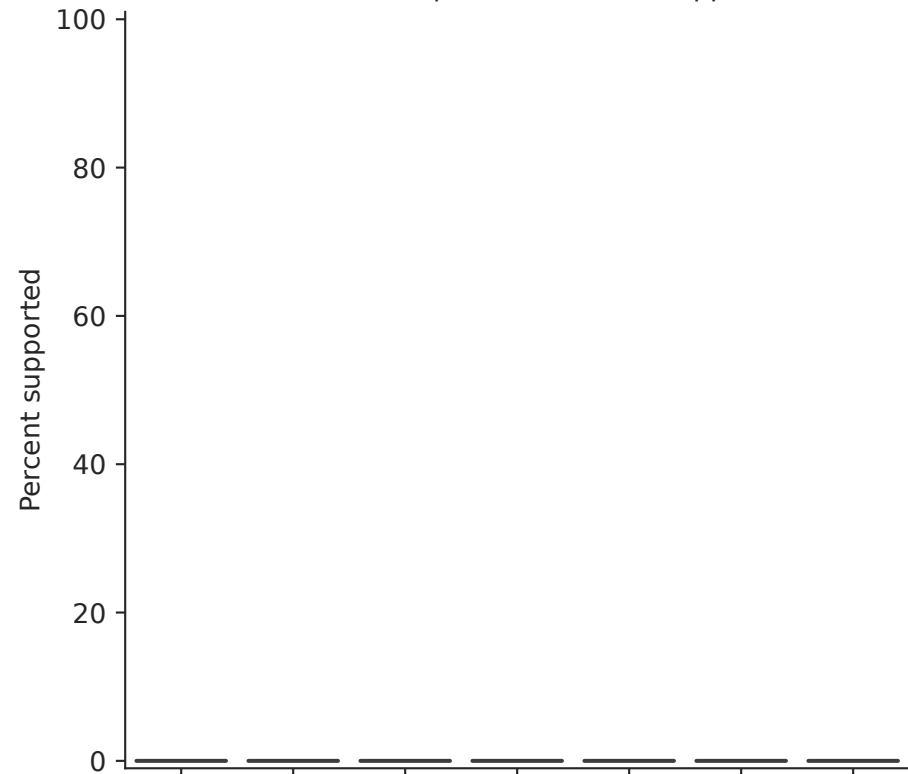
variable = Original Introns



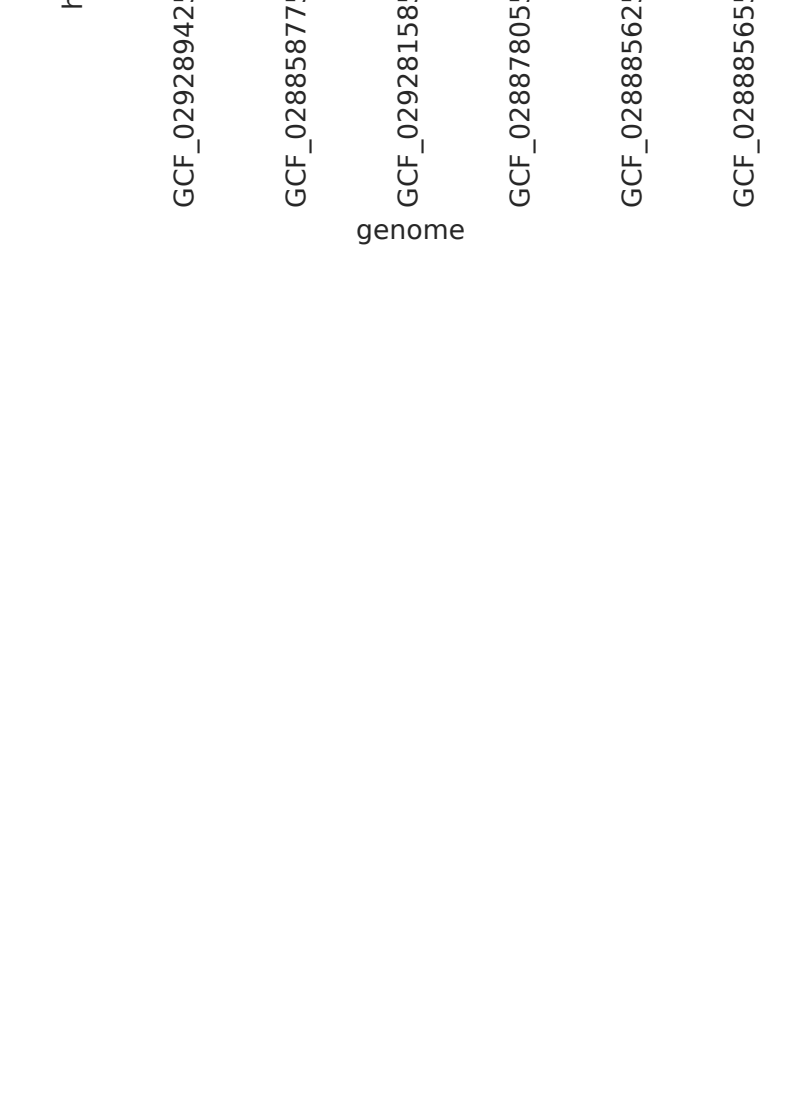
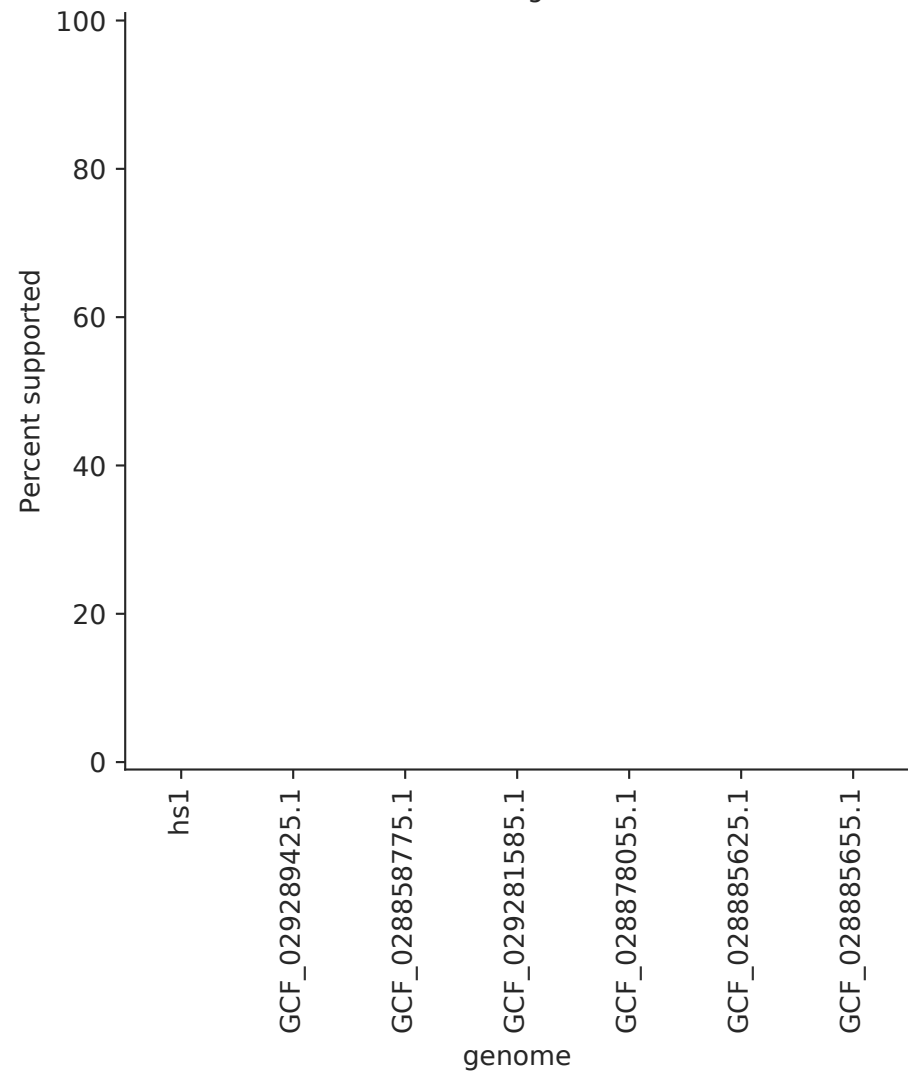
# Reference annotation support for TR\_C\_gene

variable = Splice Annotation Support

variable = Exon Annotation Support



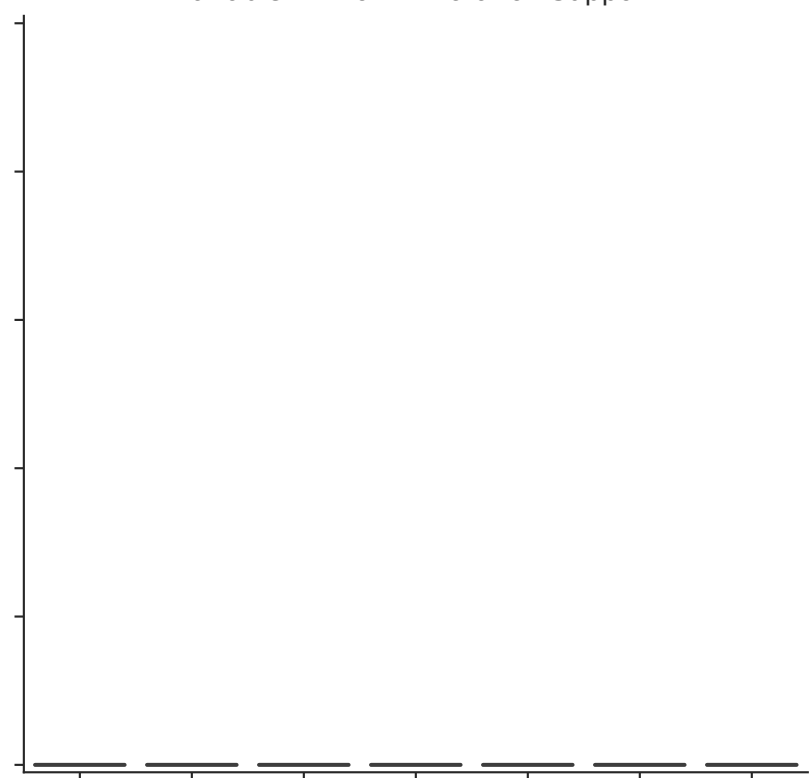
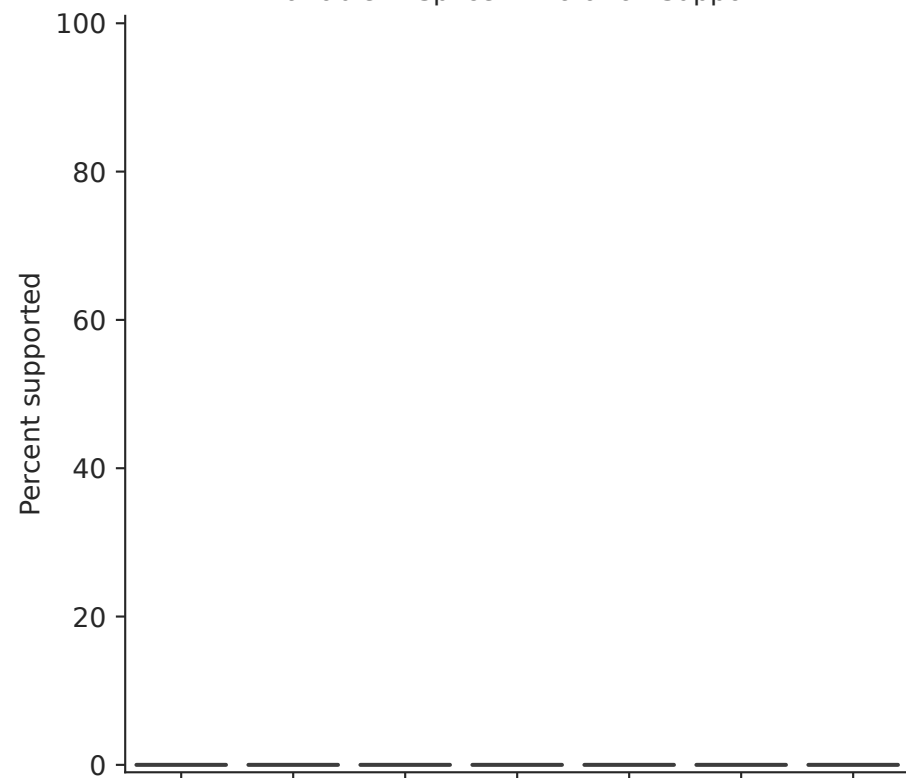
variable = Original Introns



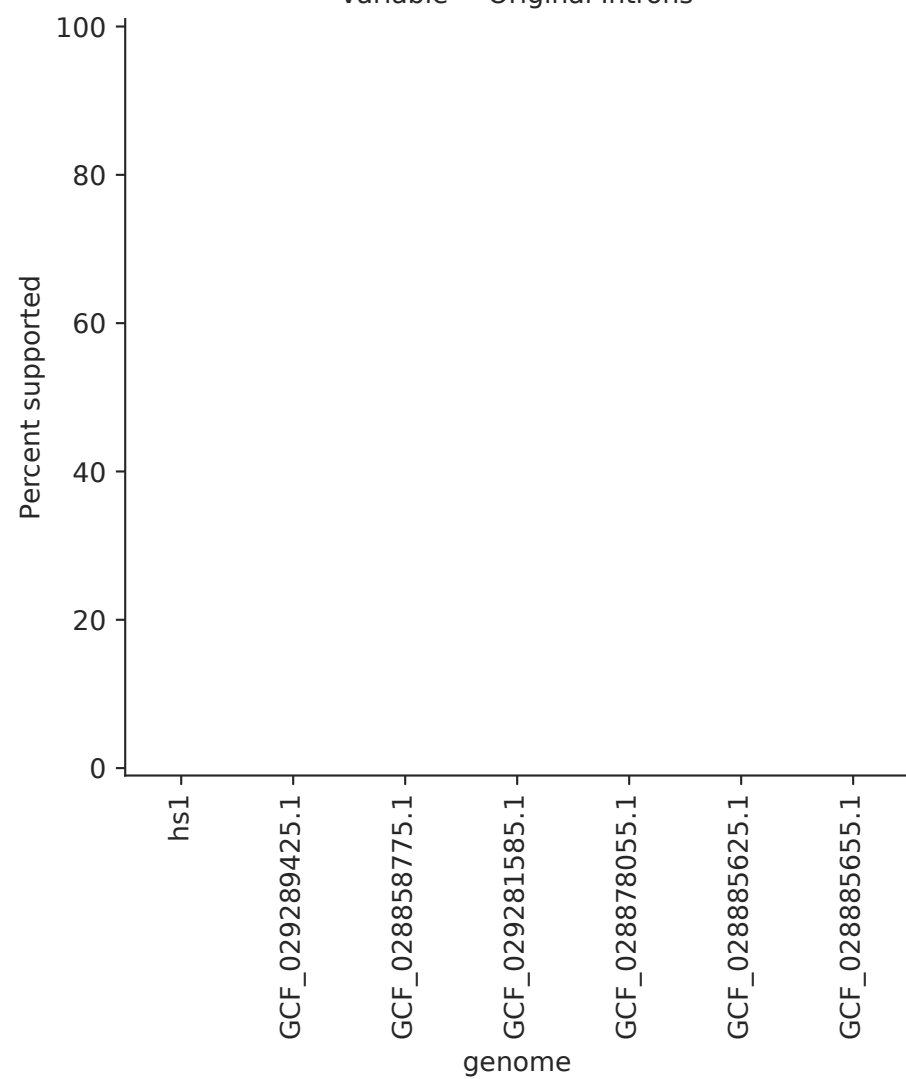
# Reference annotation support for TR\_J\_gene

variable = Splice Annotation Support

variable = Exon Annotation Support



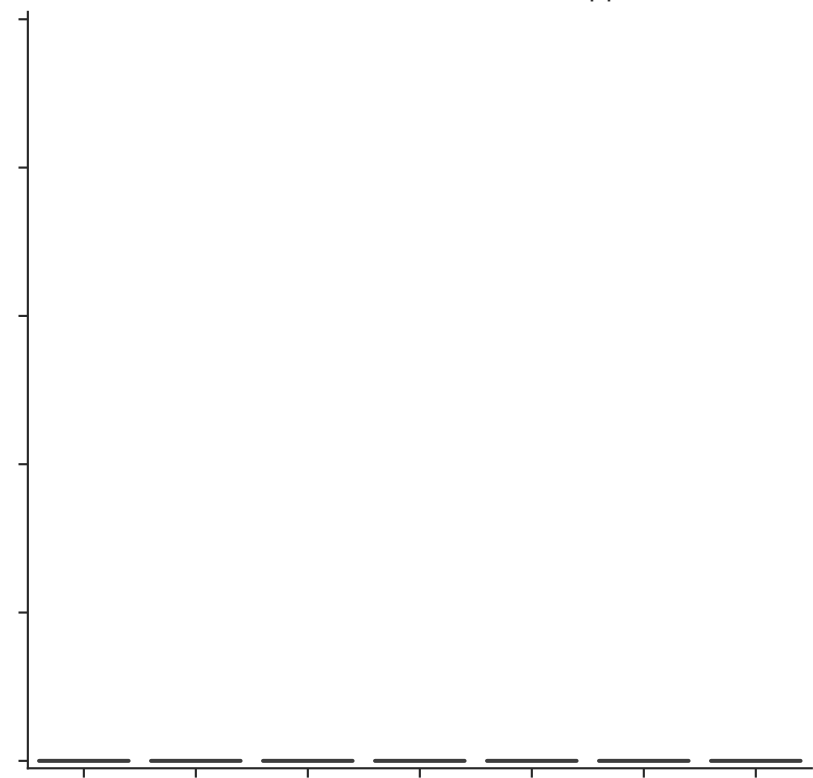
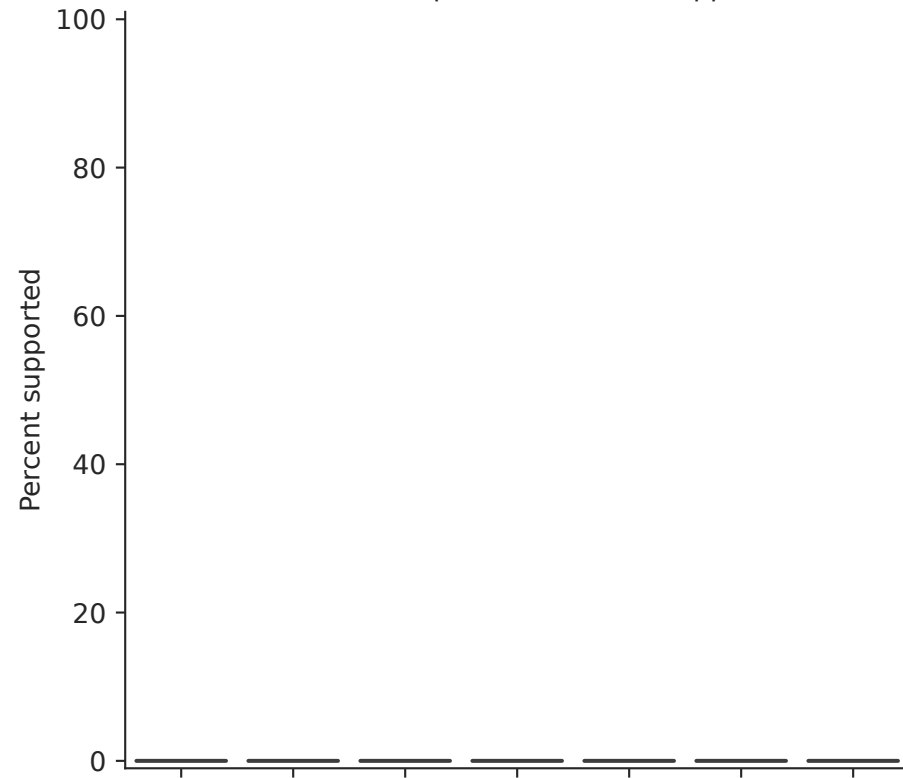
variable = Original Introns



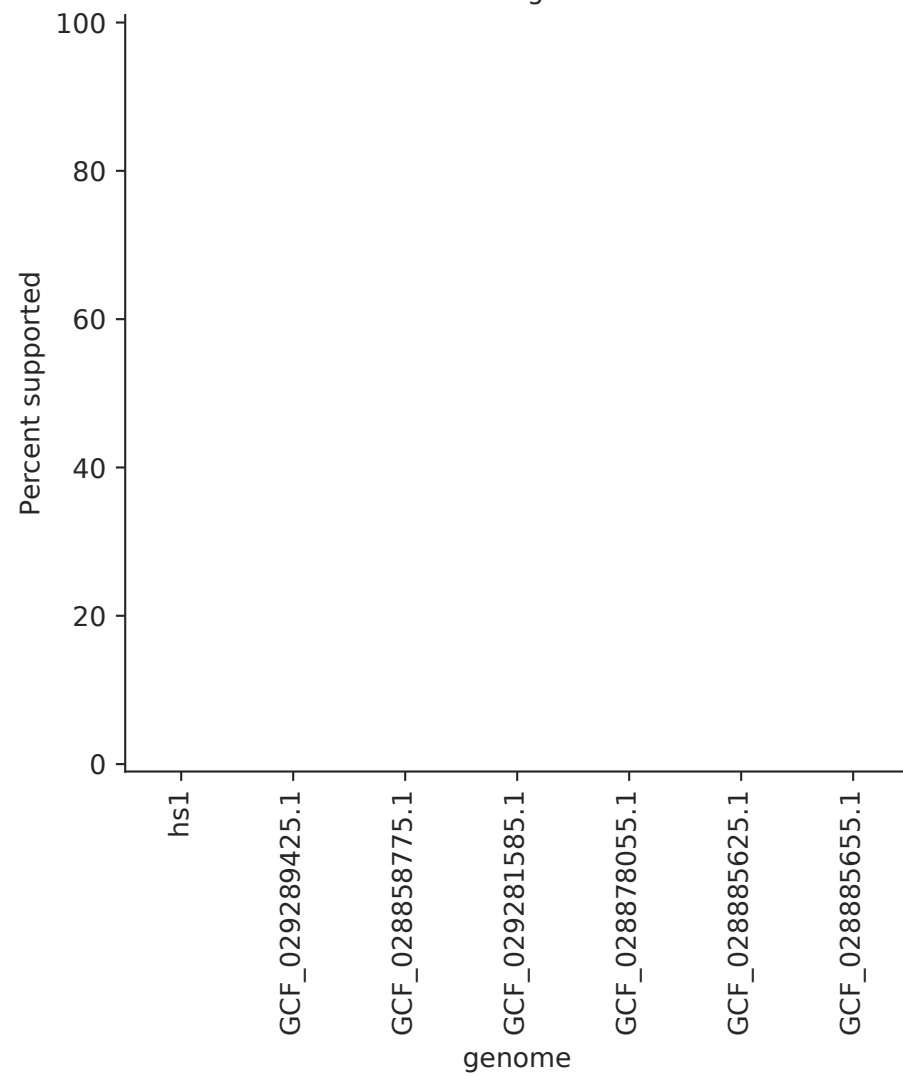
# Reference annotation support for TR\_J\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



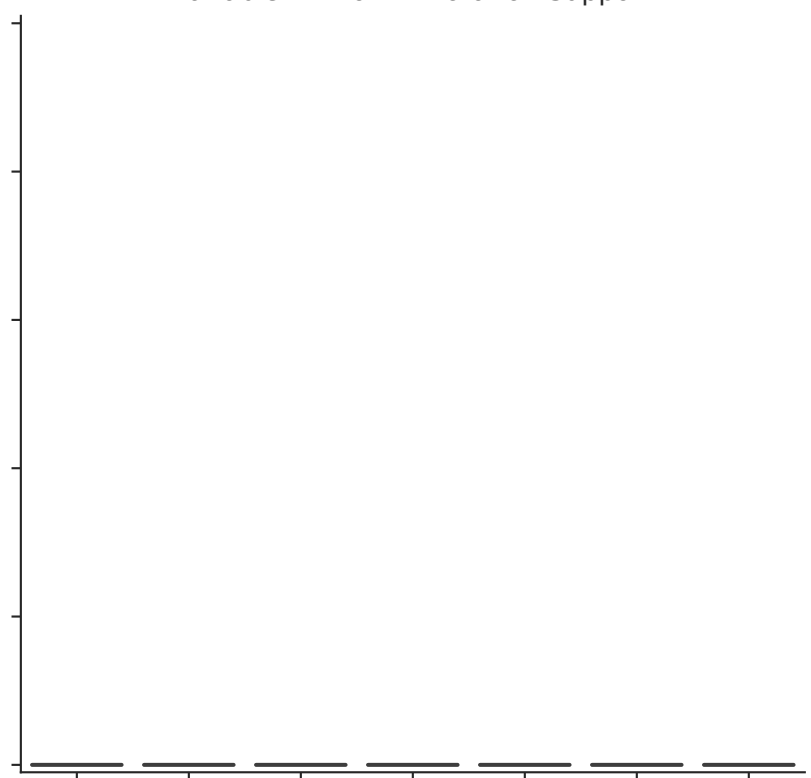
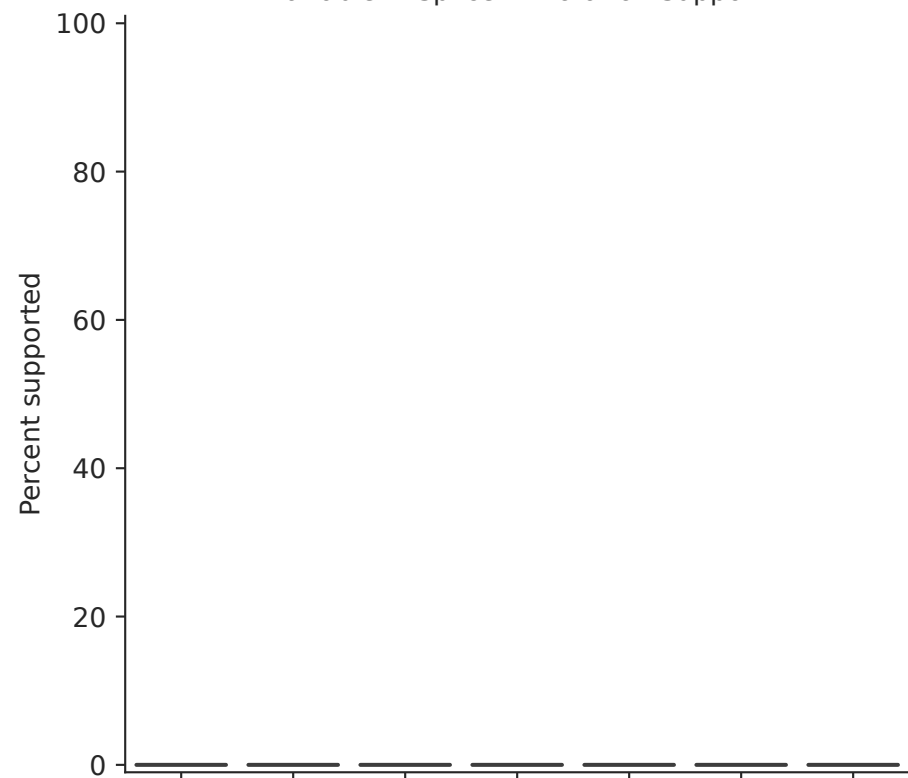
variable = Original Introns



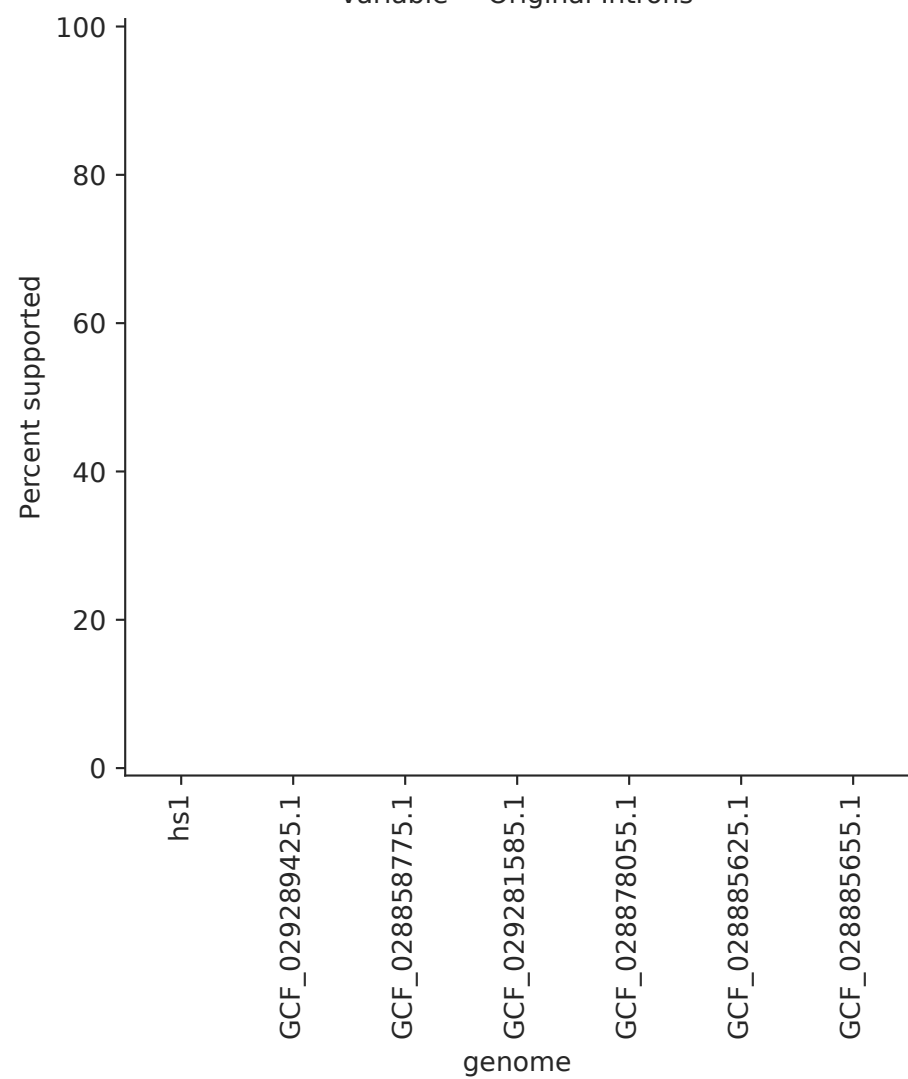
# Reference annotation support for TR\_V\_gene

variable = Splice Annotation Support

variable = Exon Annotation Support



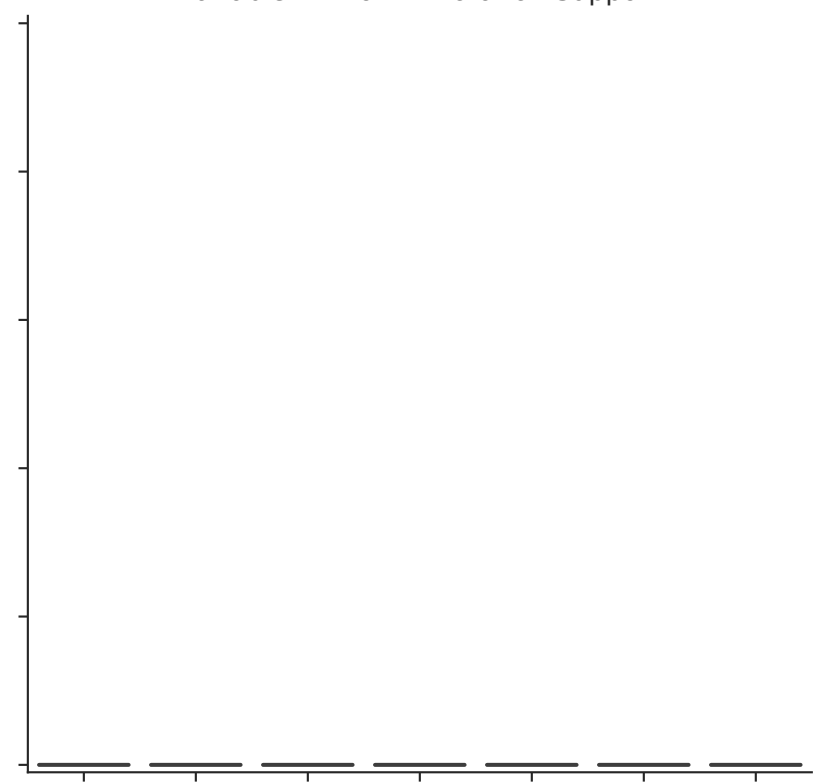
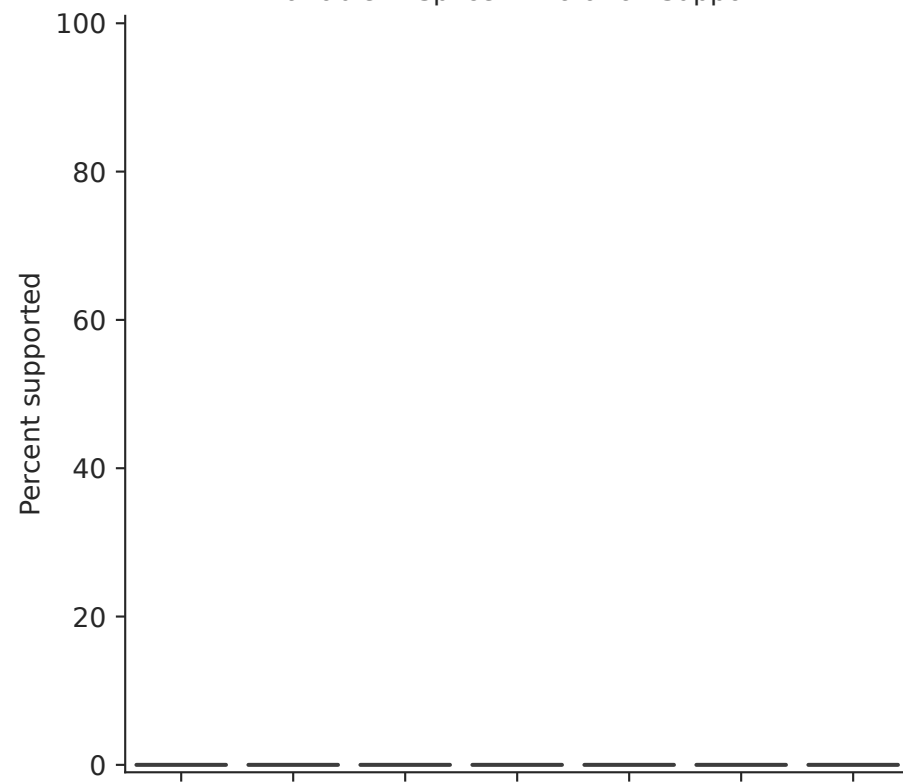
variable = Original Introns



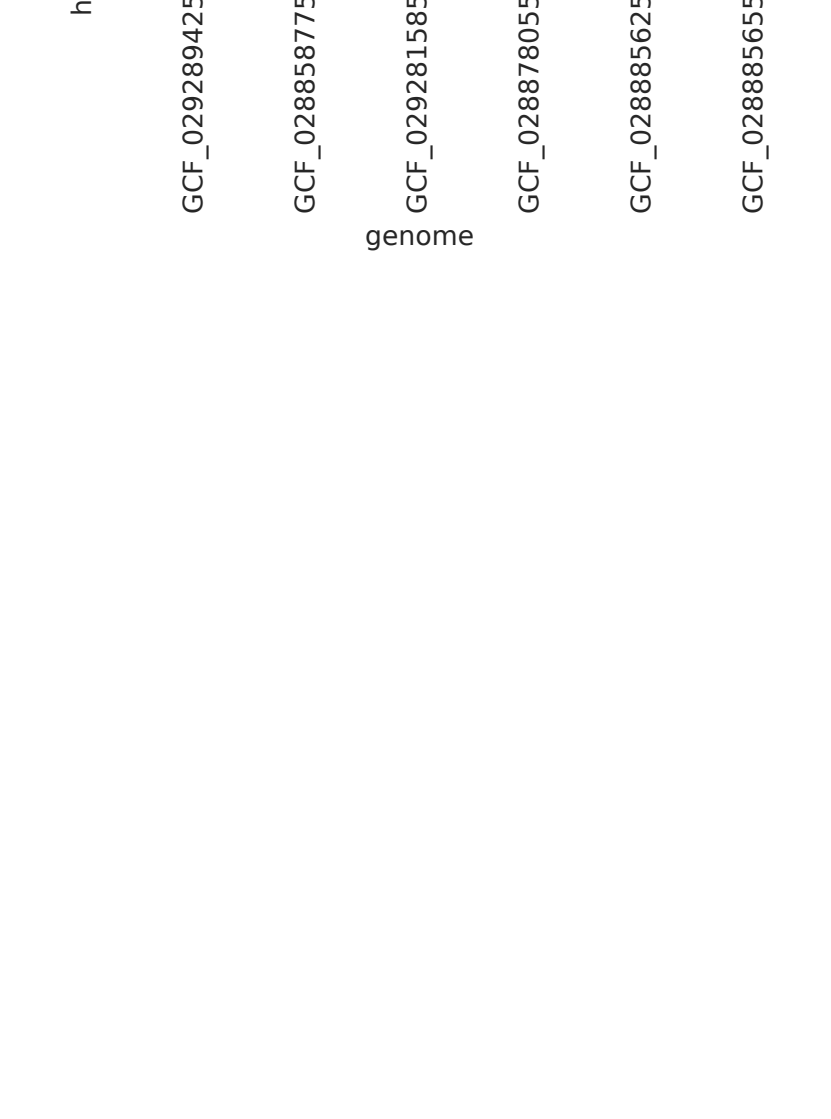
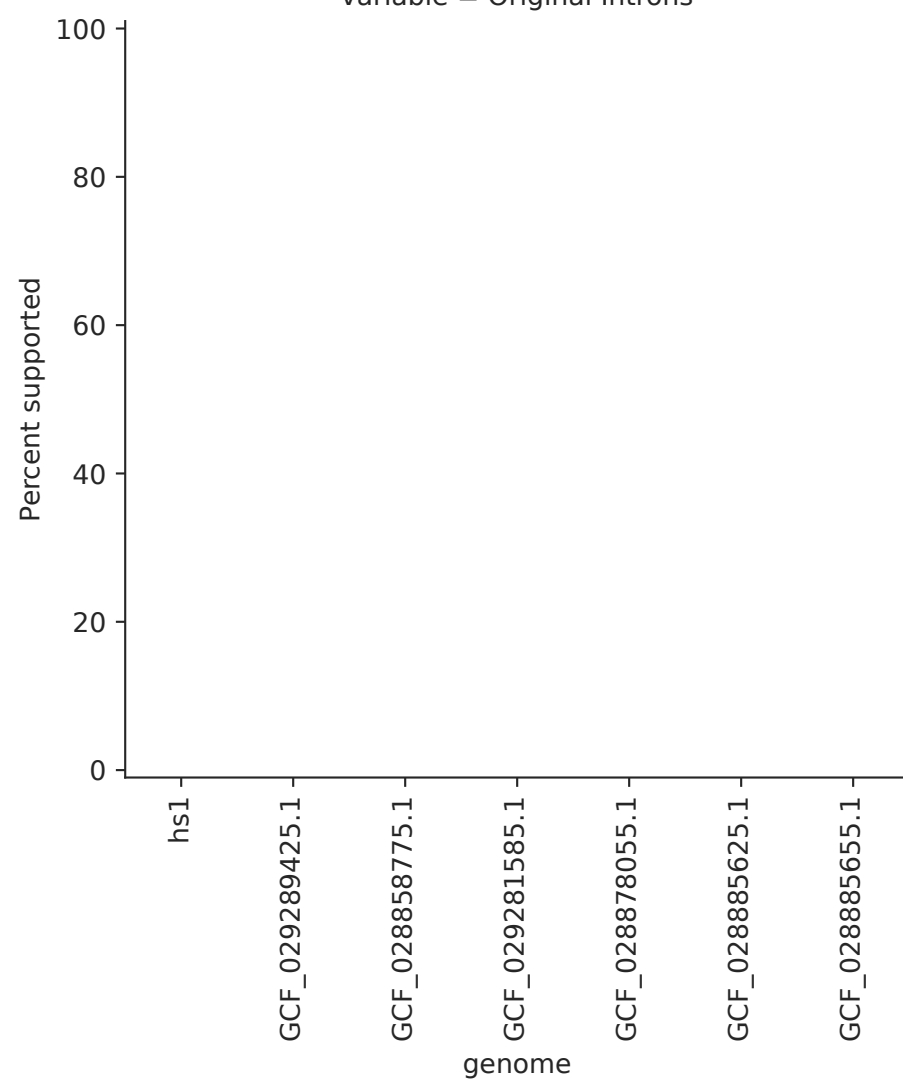
# Reference annotation support for TR\_V\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



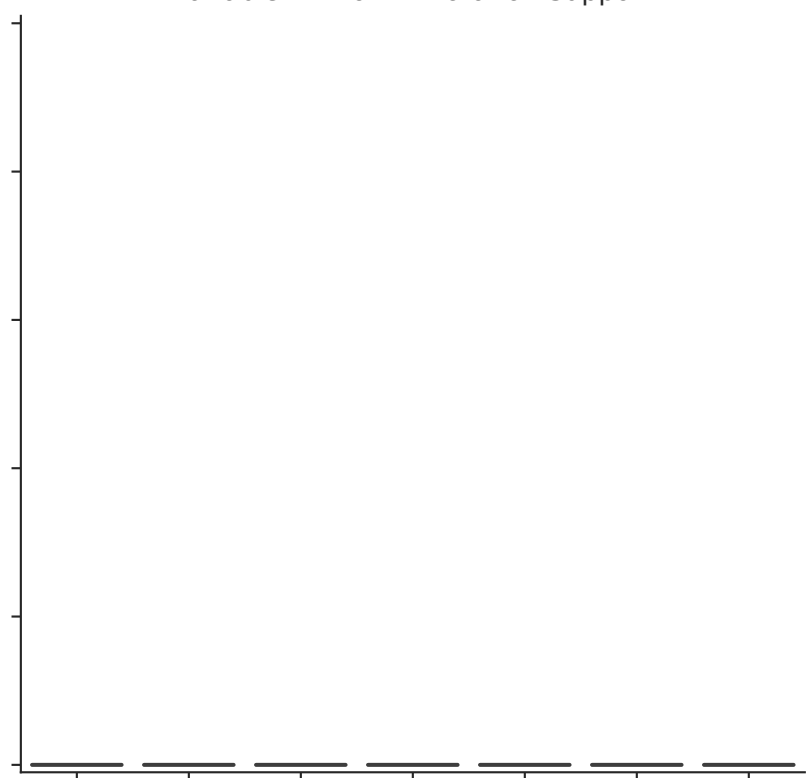
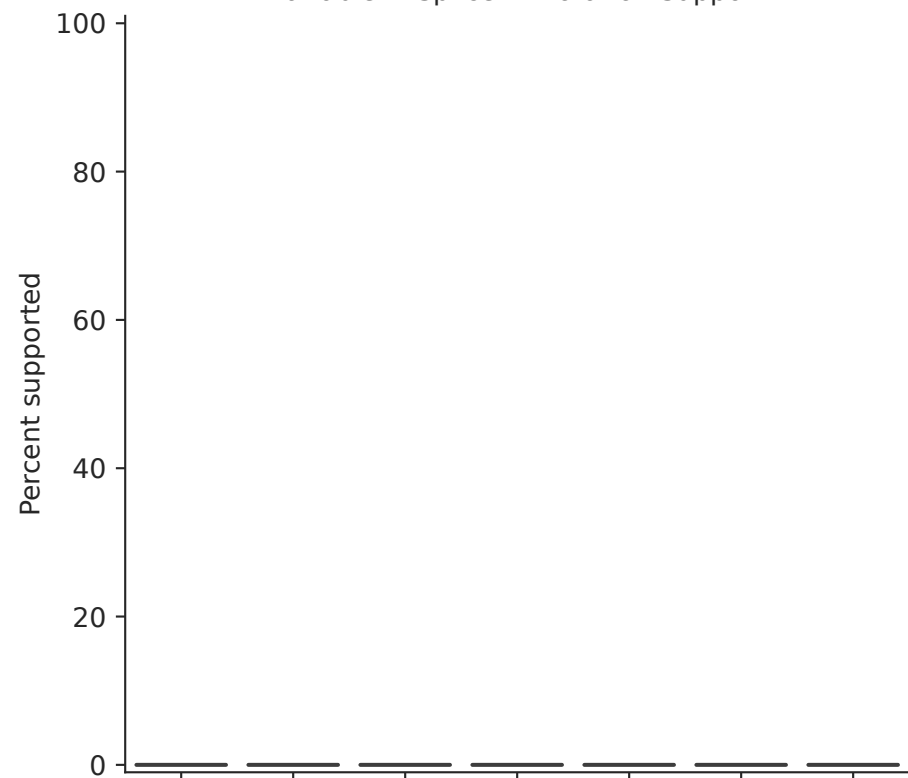
variable = Original Introns



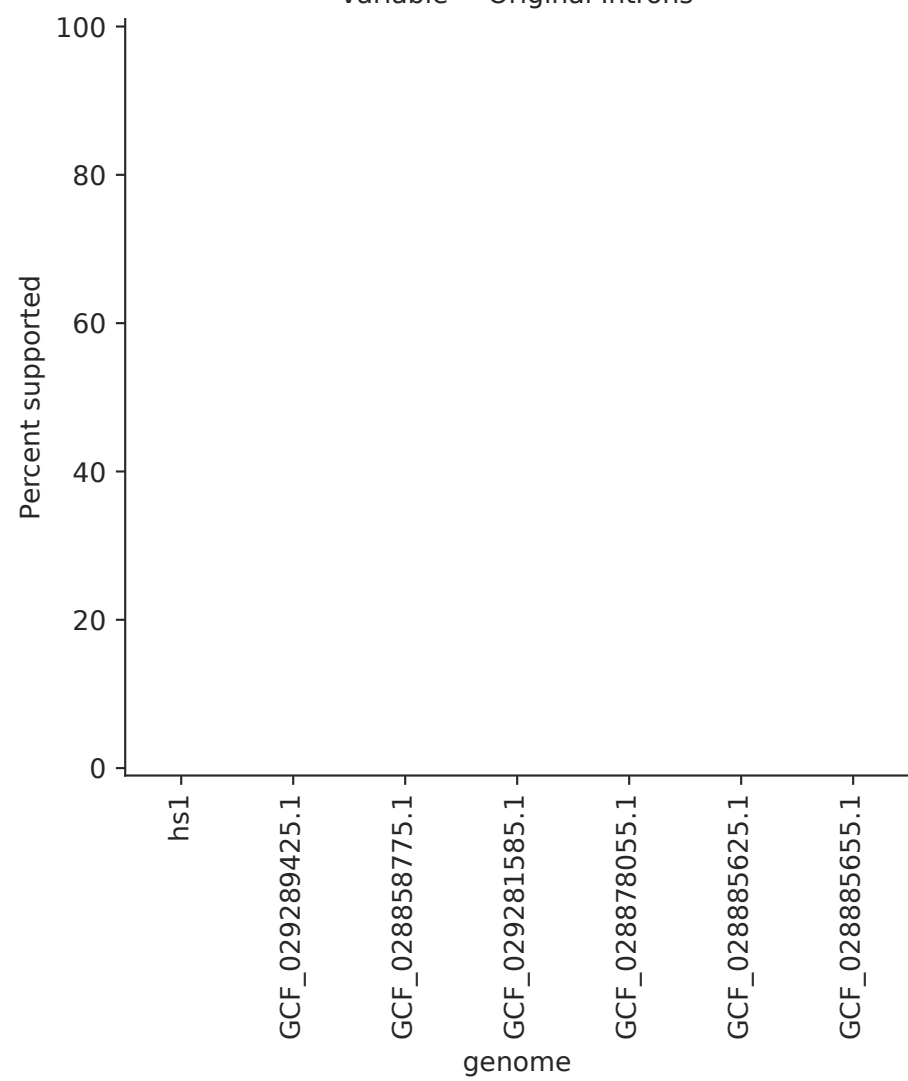
# Reference annotation support for artifact

variable = Splice Annotation Support

variable = Exon Annotation Support



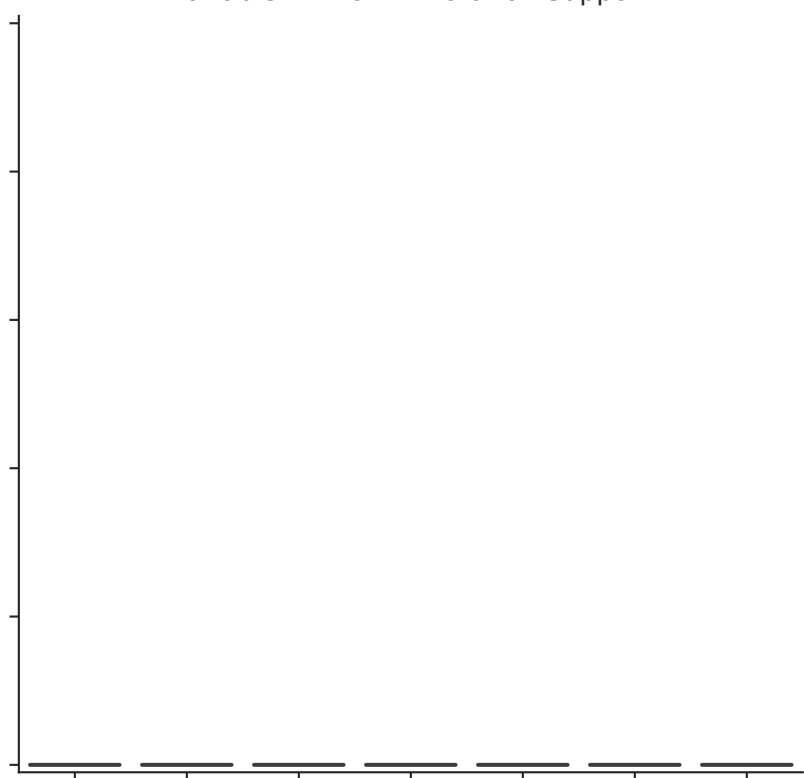
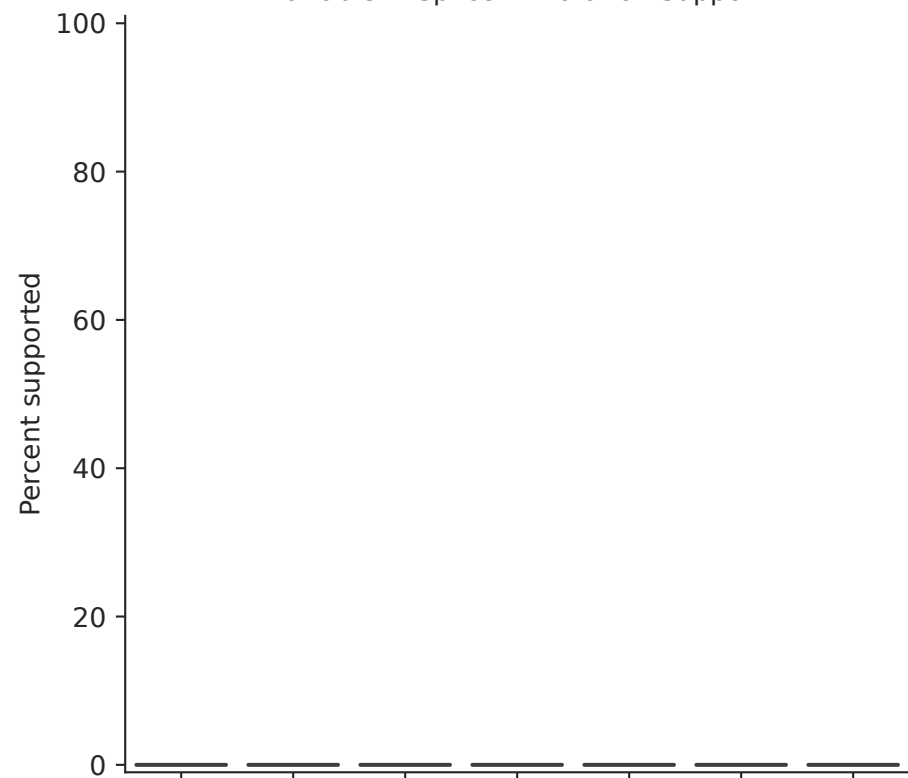
variable = Original Introns



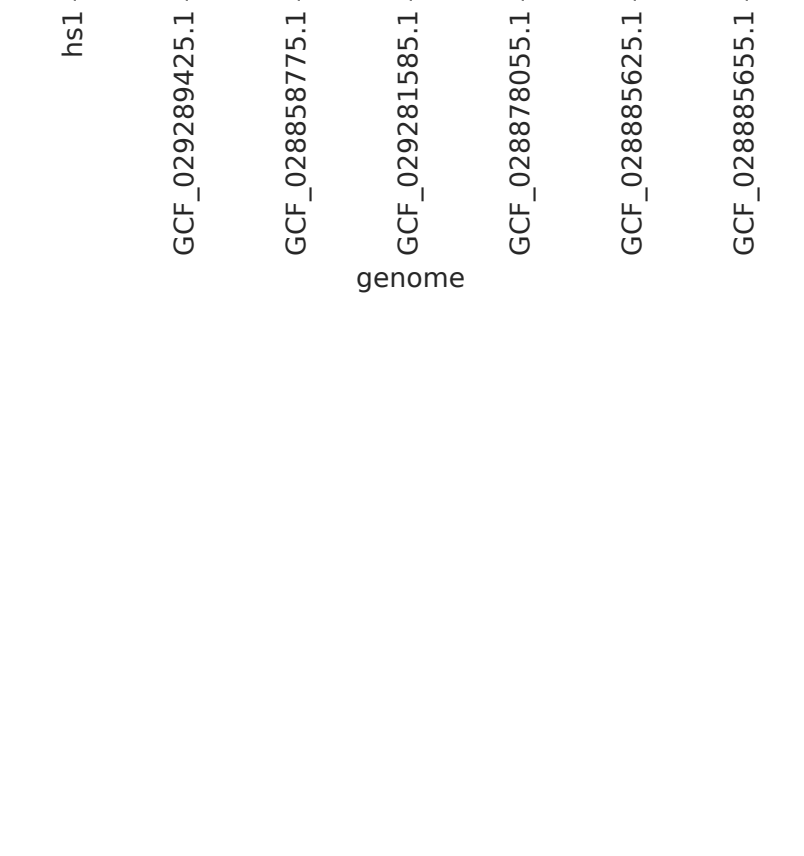
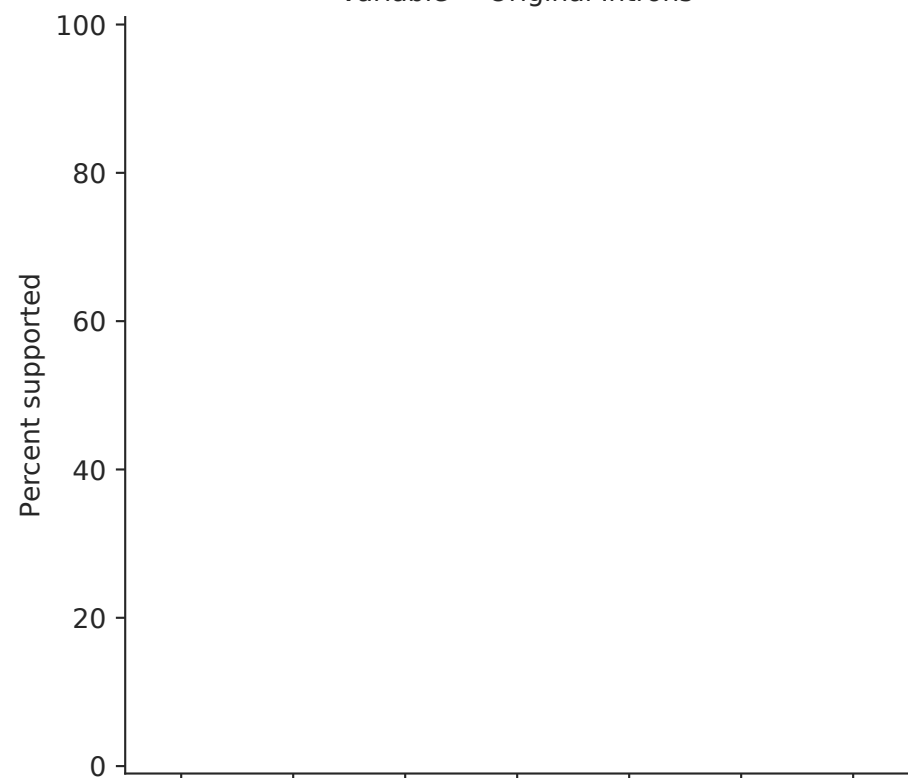
# Reference annotation support for lncRNA

variable = Splice Annotation Support

variable = Exon Annotation Support



variable = Original Introns

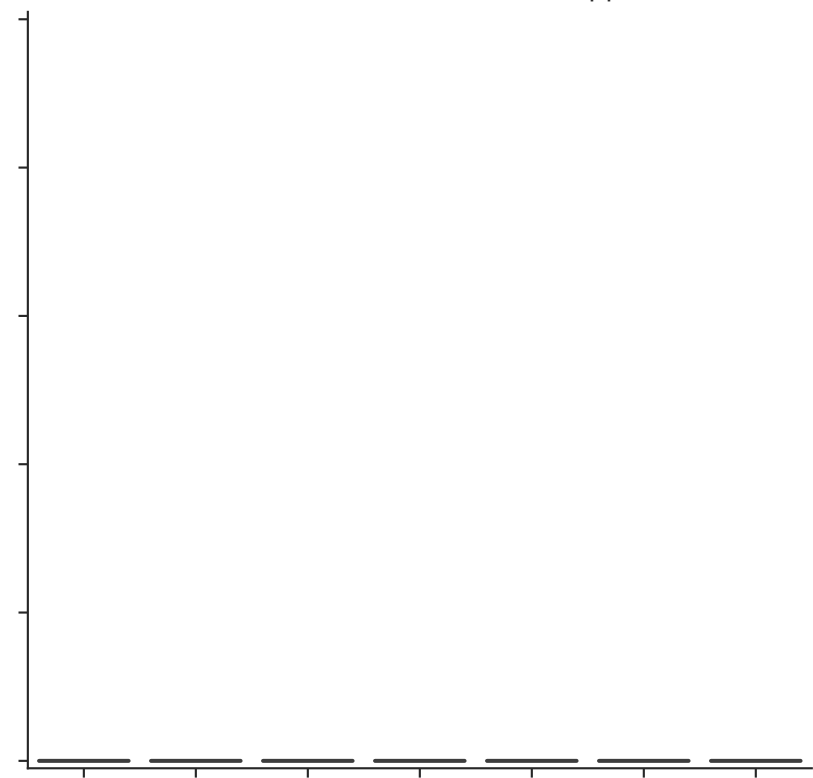
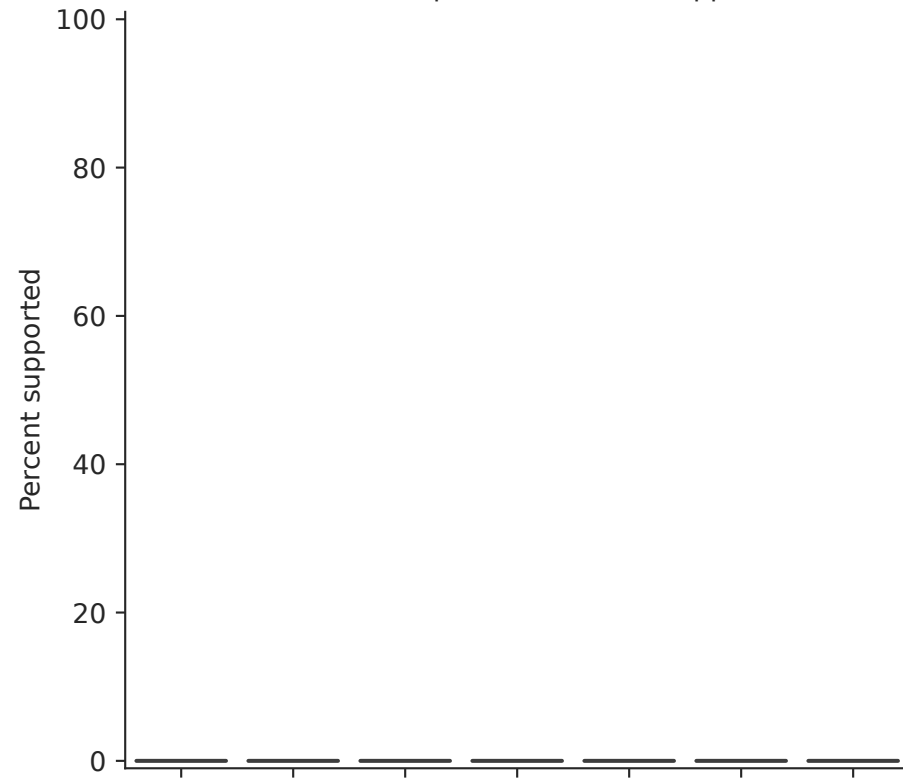


genome

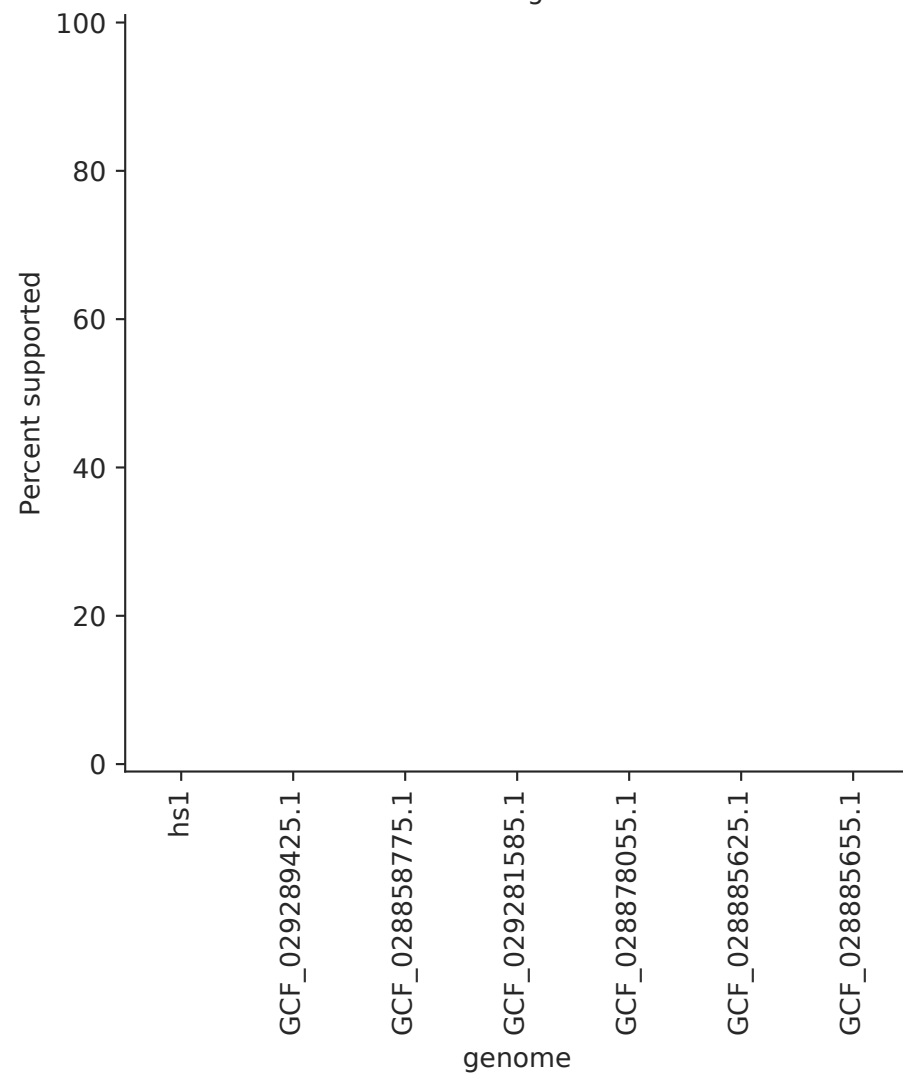
# Reference annotation support for miRNA

variable = Splice Annotation Support

variable = Exon Annotation Support



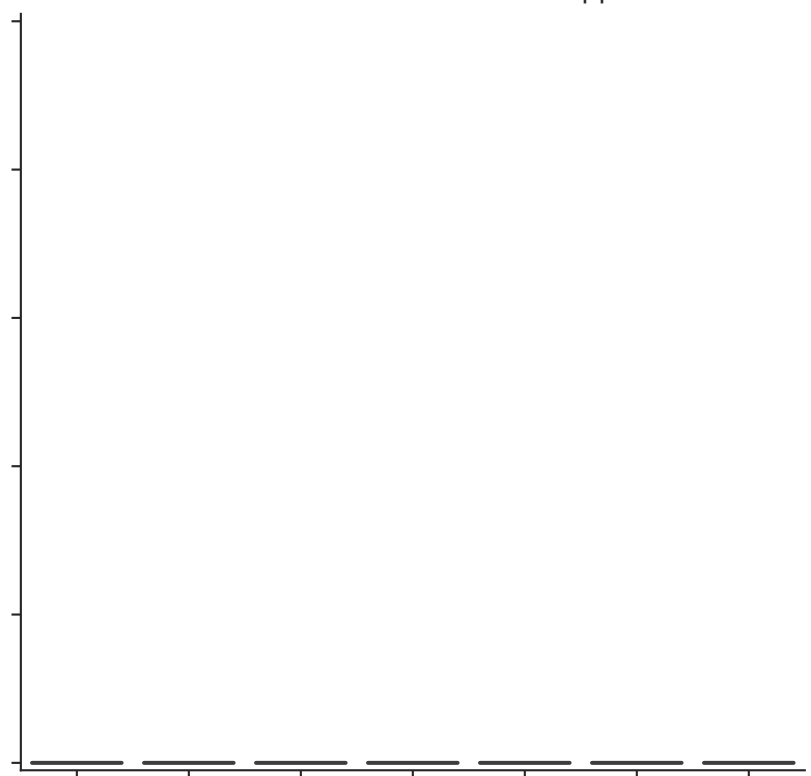
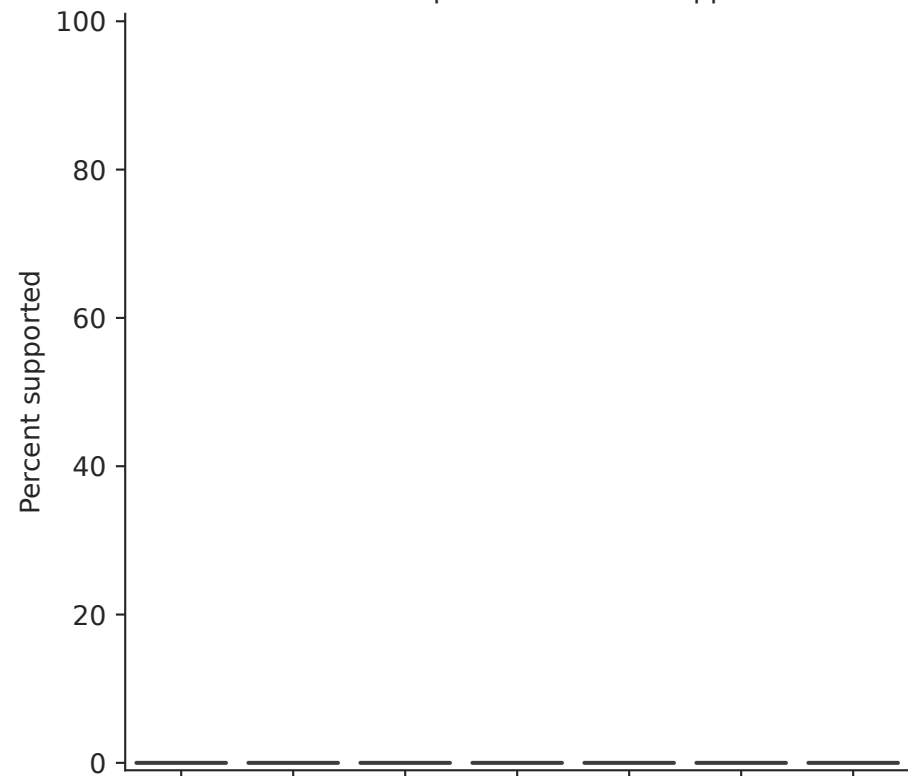
variable = Original Introns



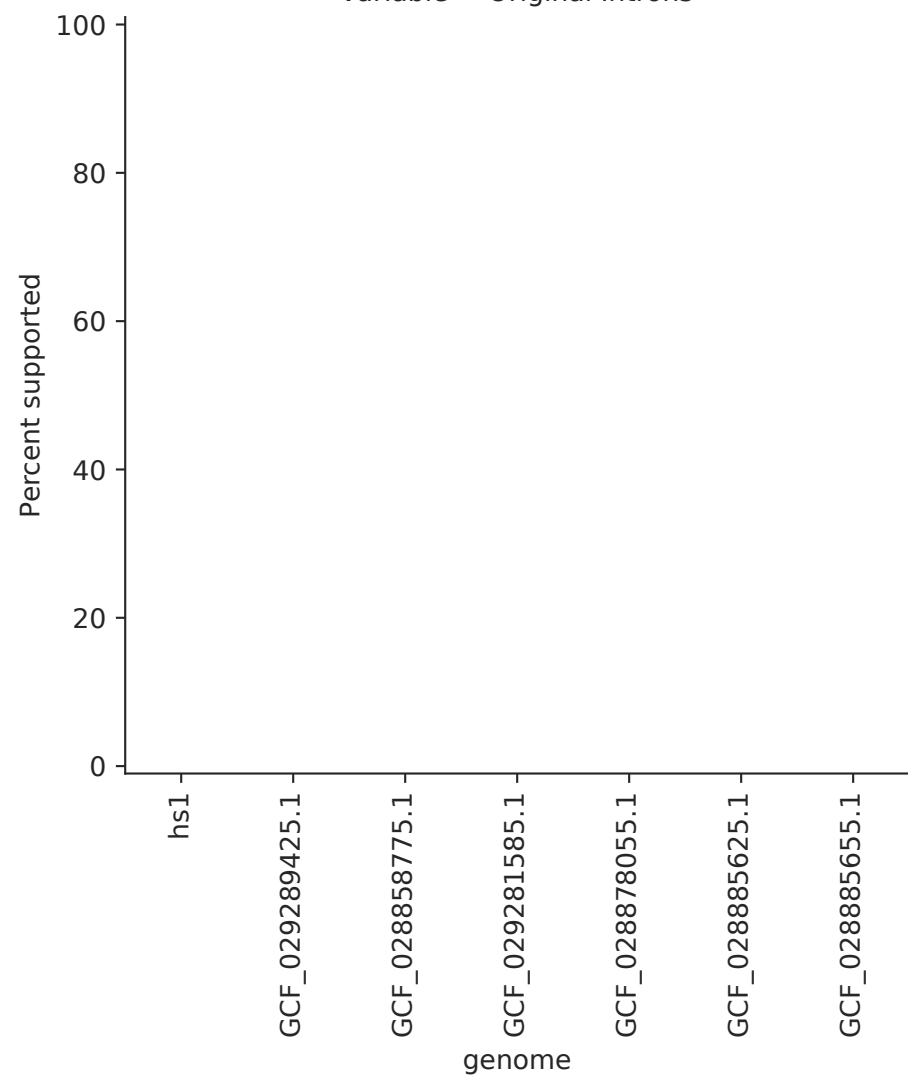
# Reference annotation support for misc\_RNA

variable = Splice Annotation Support

variable = Exon Annotation Support



variable = Original Introns

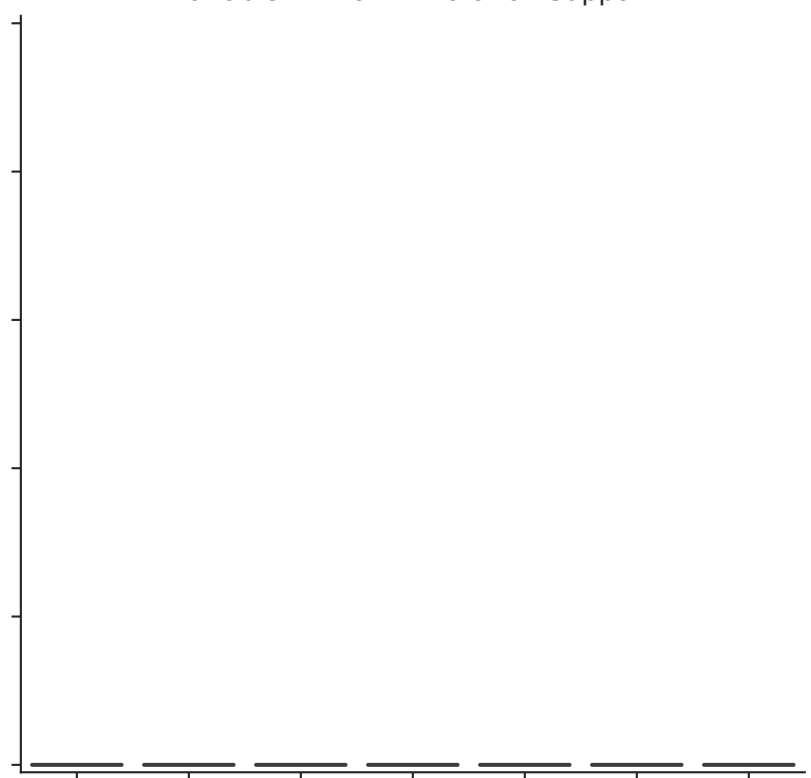
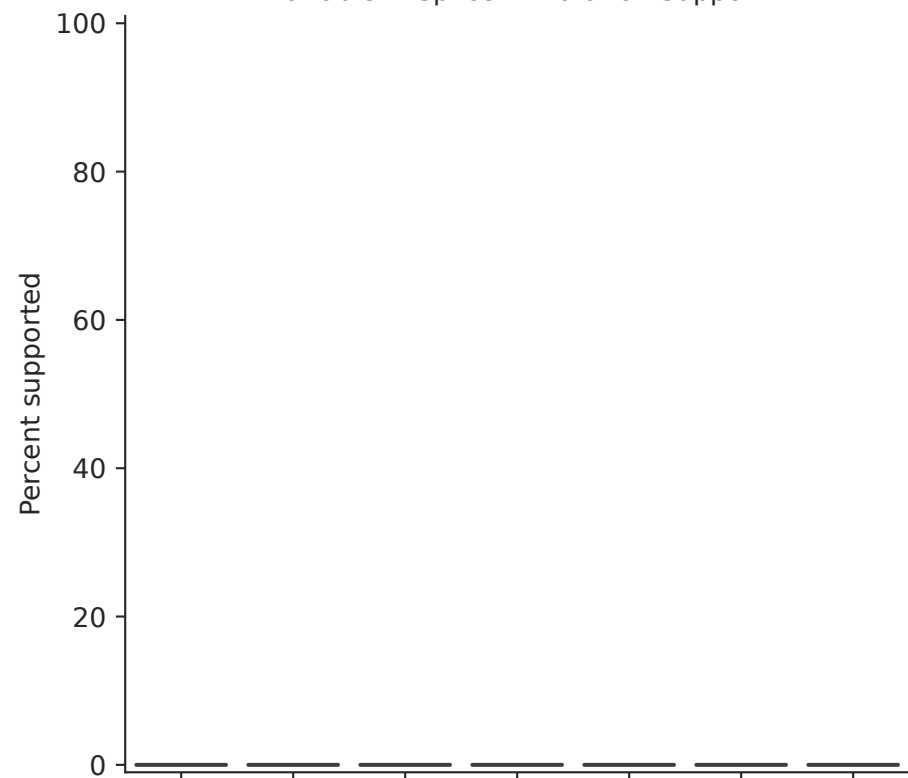


genome

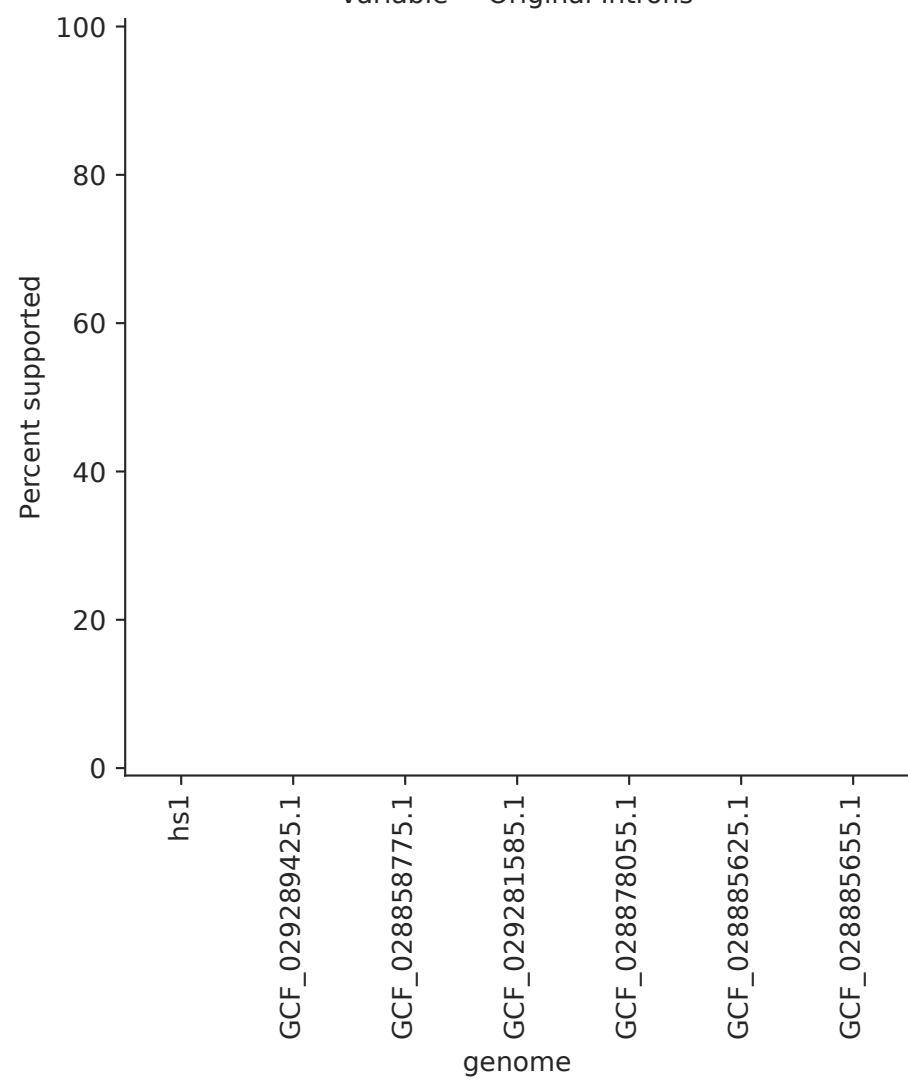
# Reference annotation support for nonsense\_mediated\_decay

variable = Splice Annotation Support

variable = Exon Annotation Support



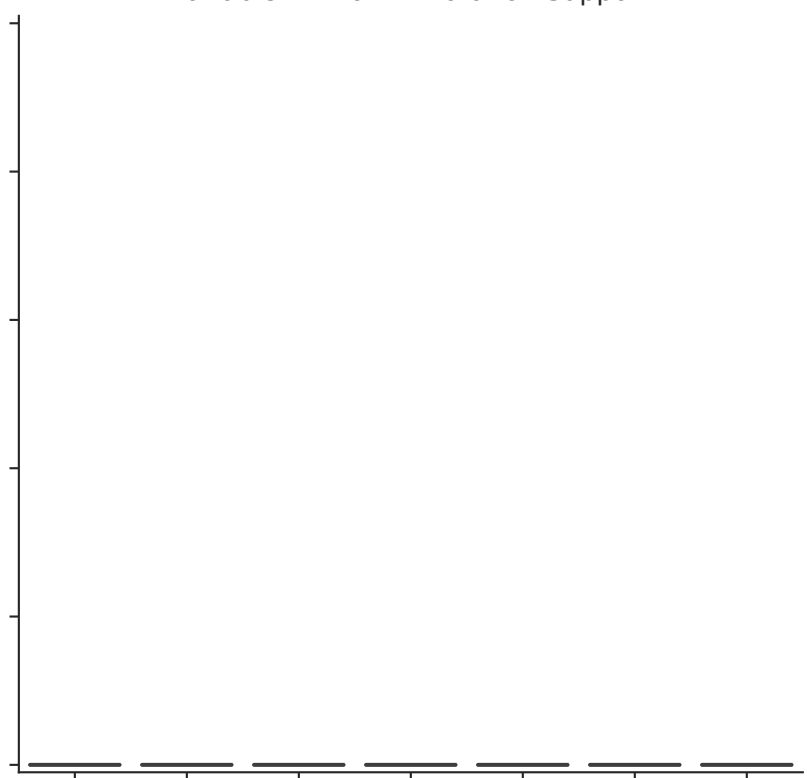
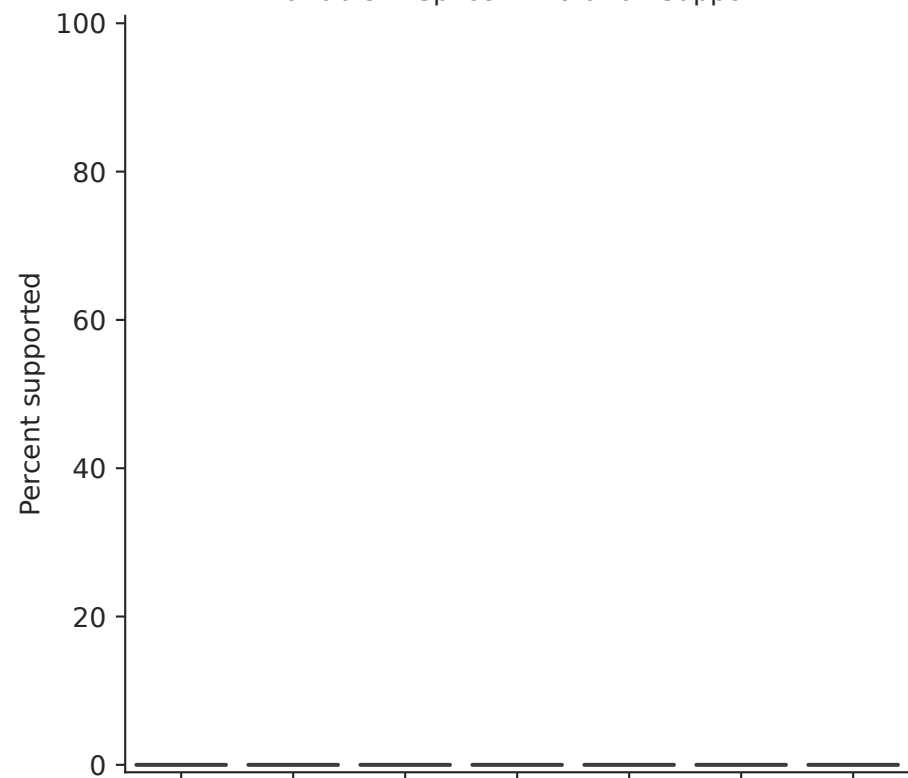
variable = Original Introns



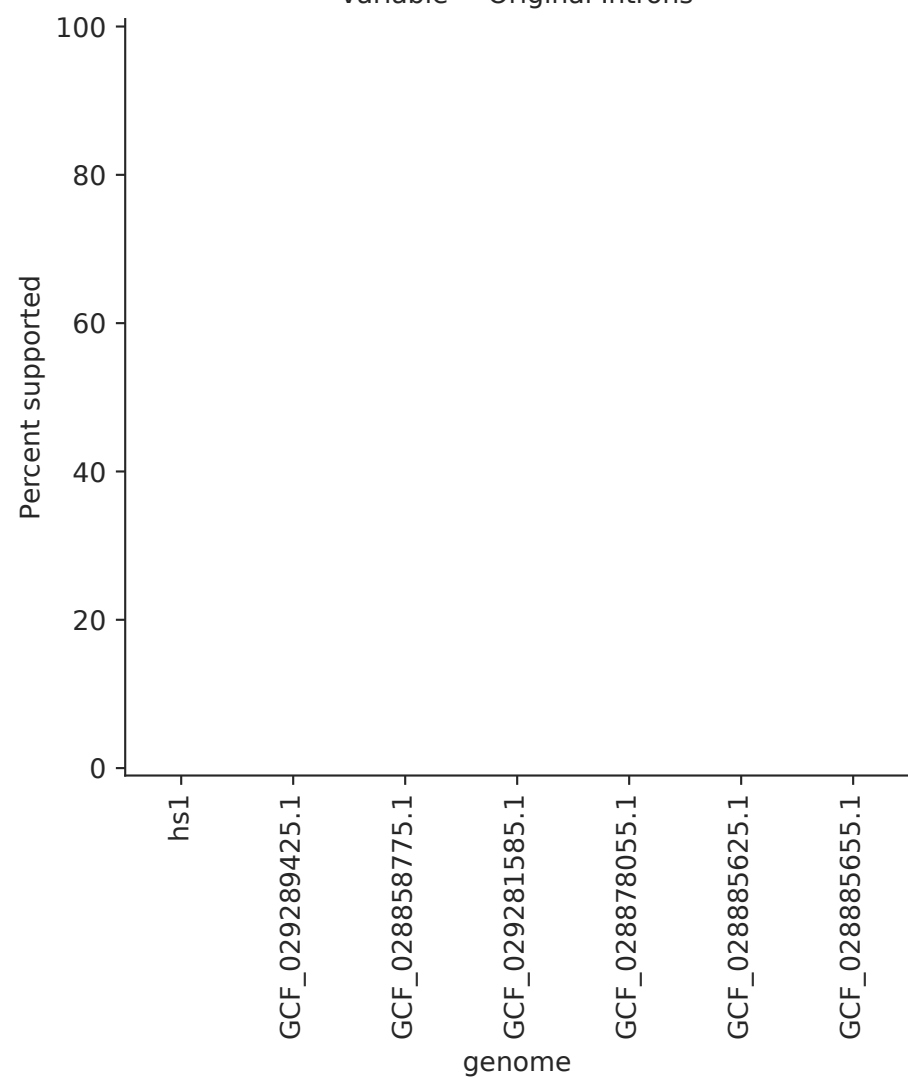
# Reference annotation support for processed\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



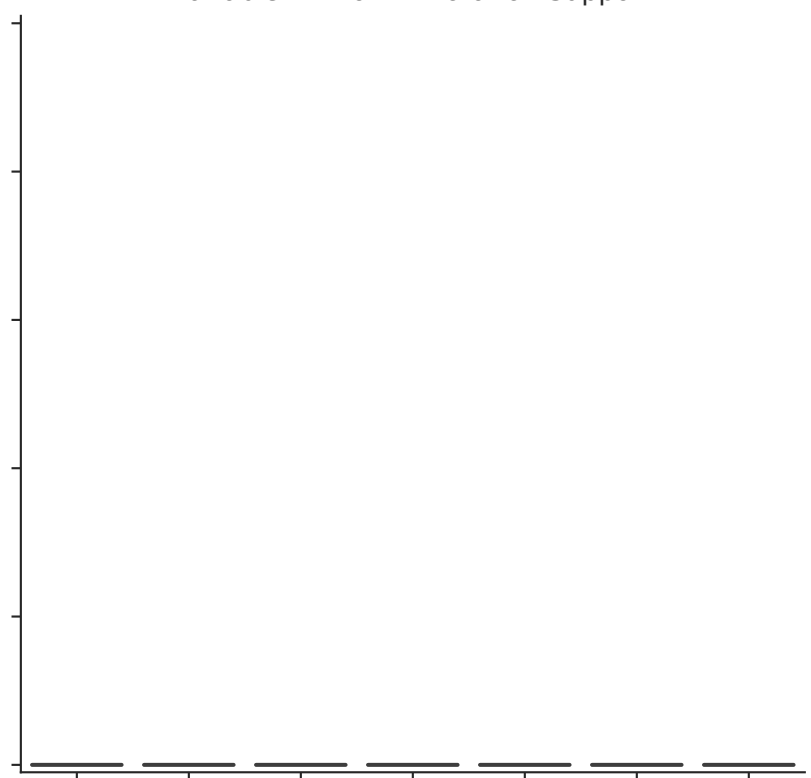
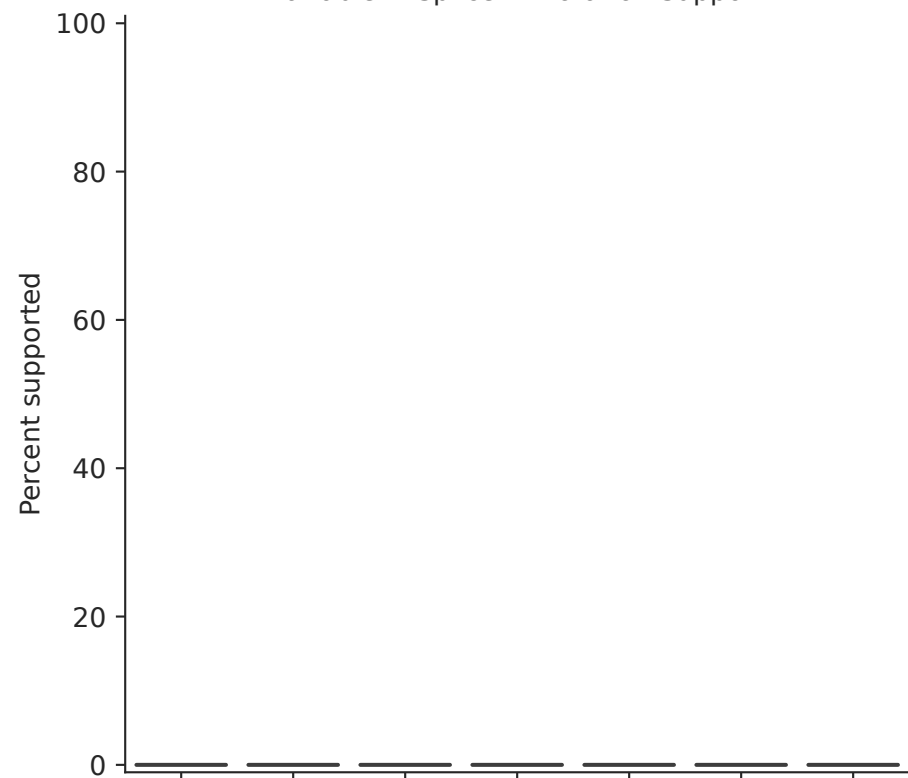
variable = Original Introns



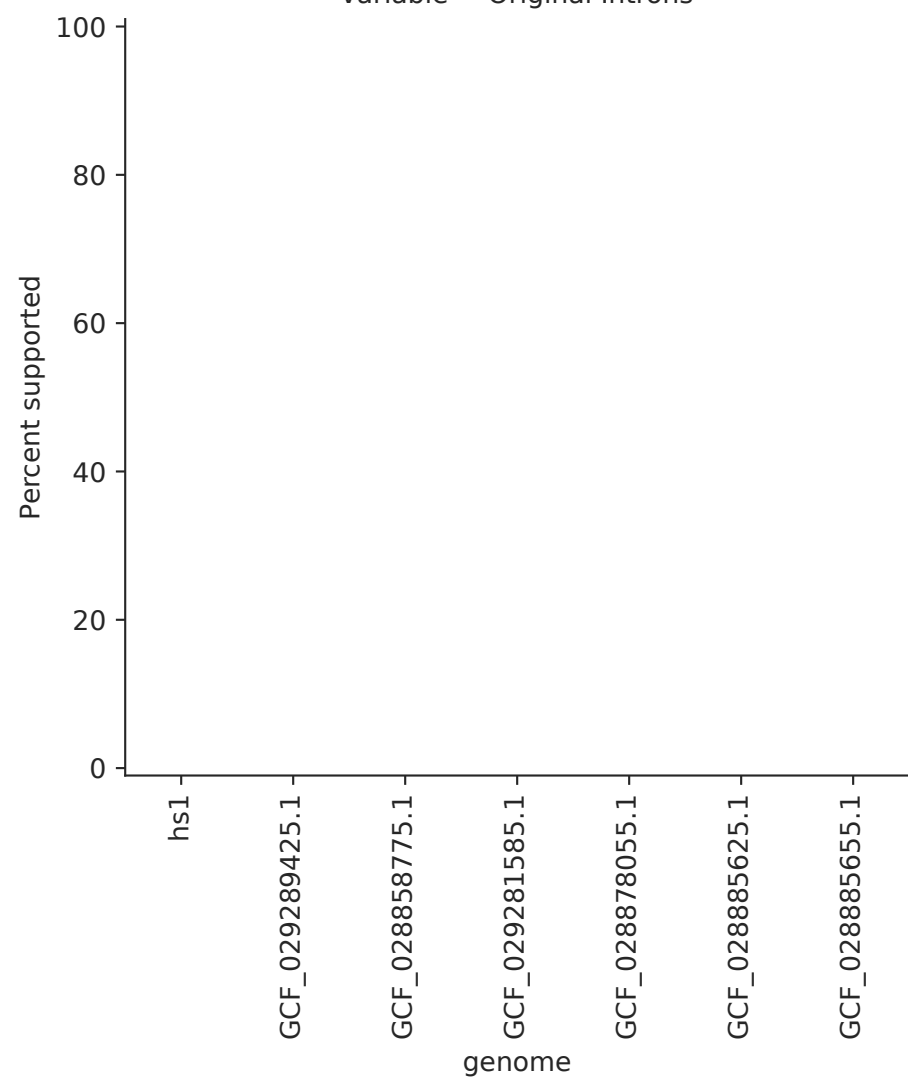
# Reference annotation support for processed\_transcript

variable = Splice Annotation Support

variable = Exon Annotation Support



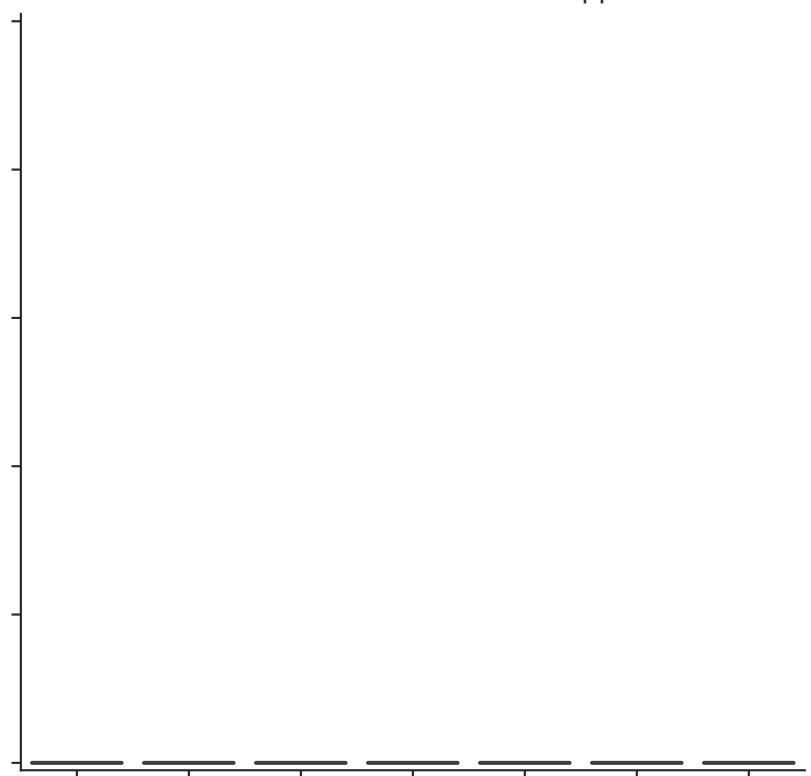
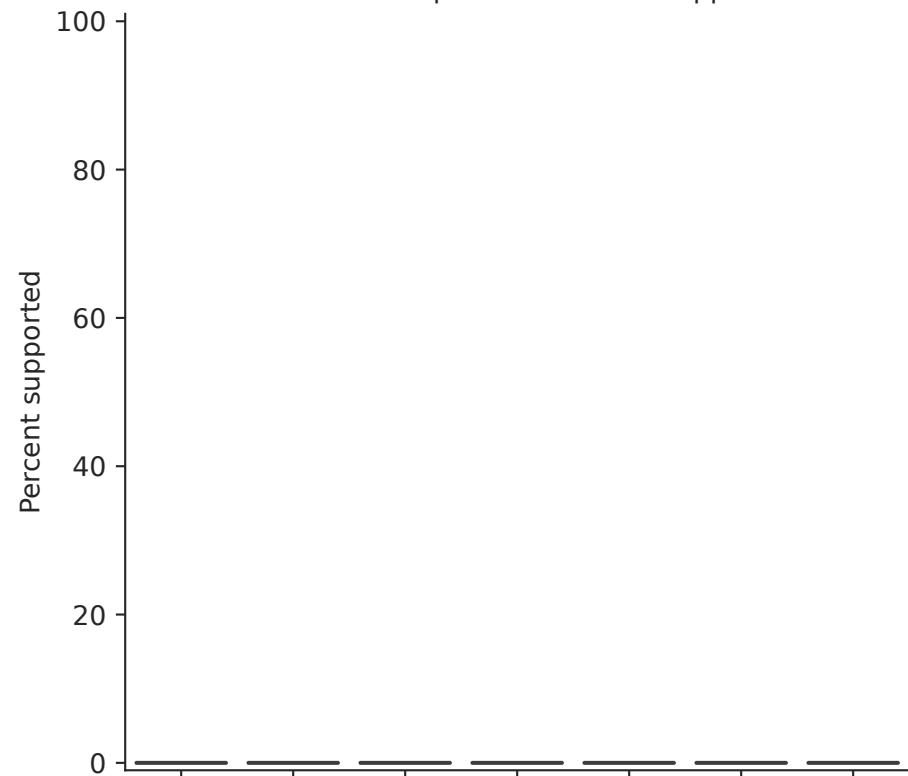
variable = Original Introns



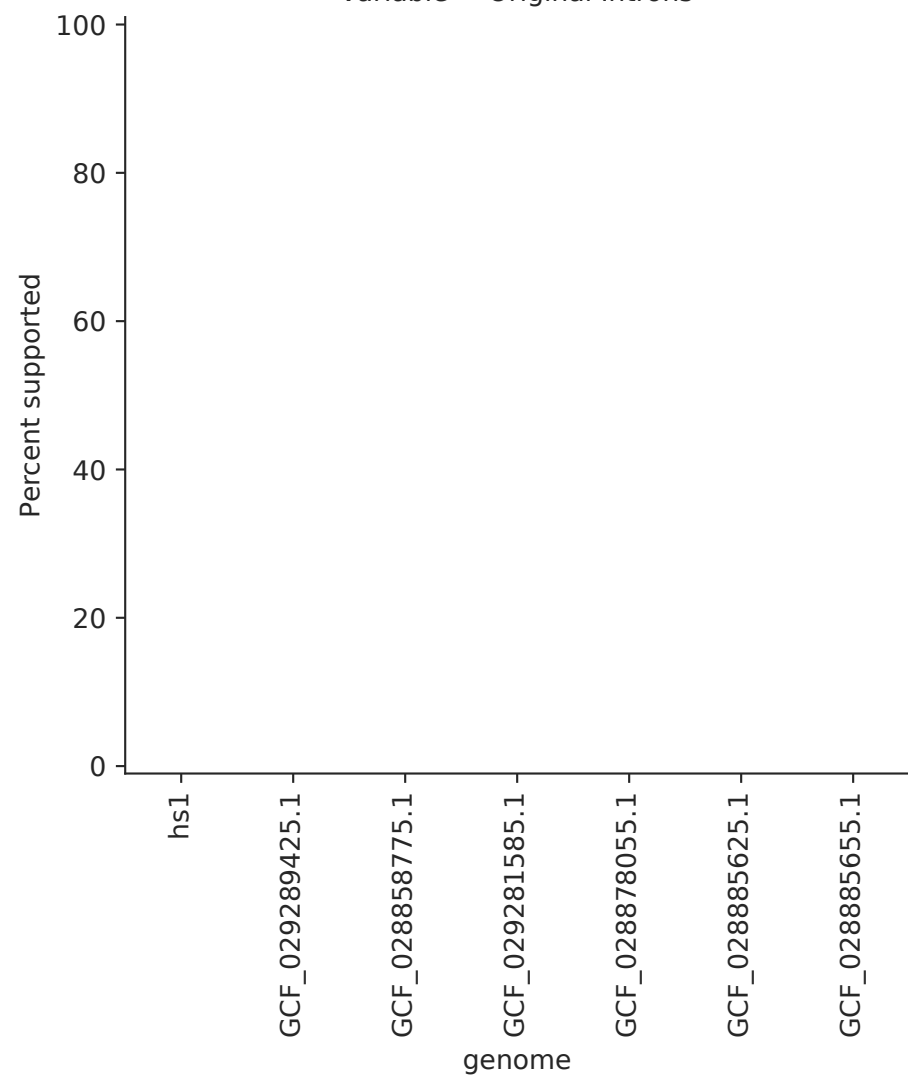
# Reference annotation support for protein\_coding\_LoF

variable = Splice Annotation Support

variable = Exon Annotation Support



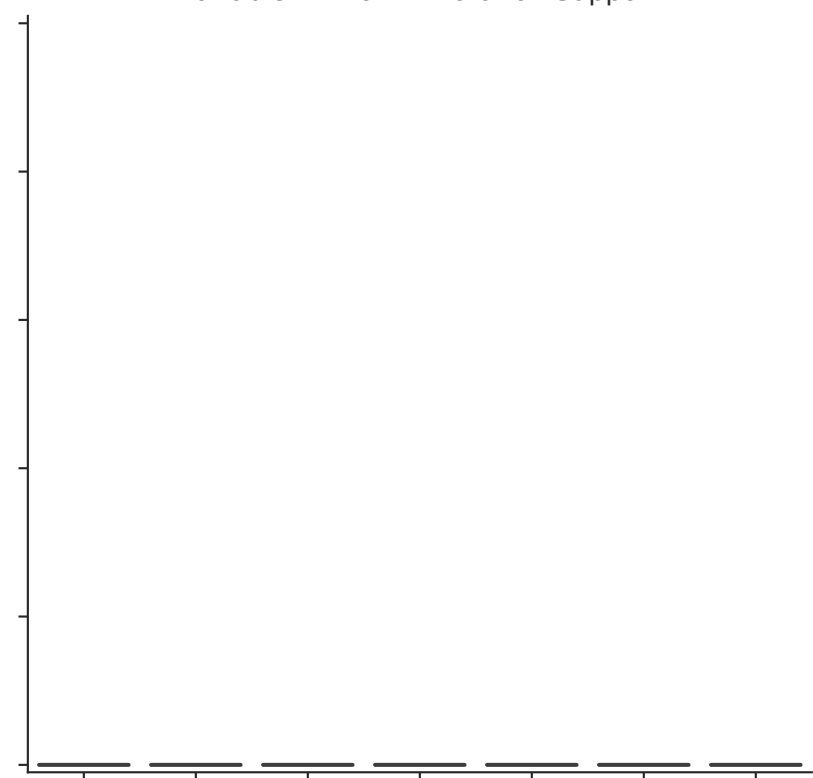
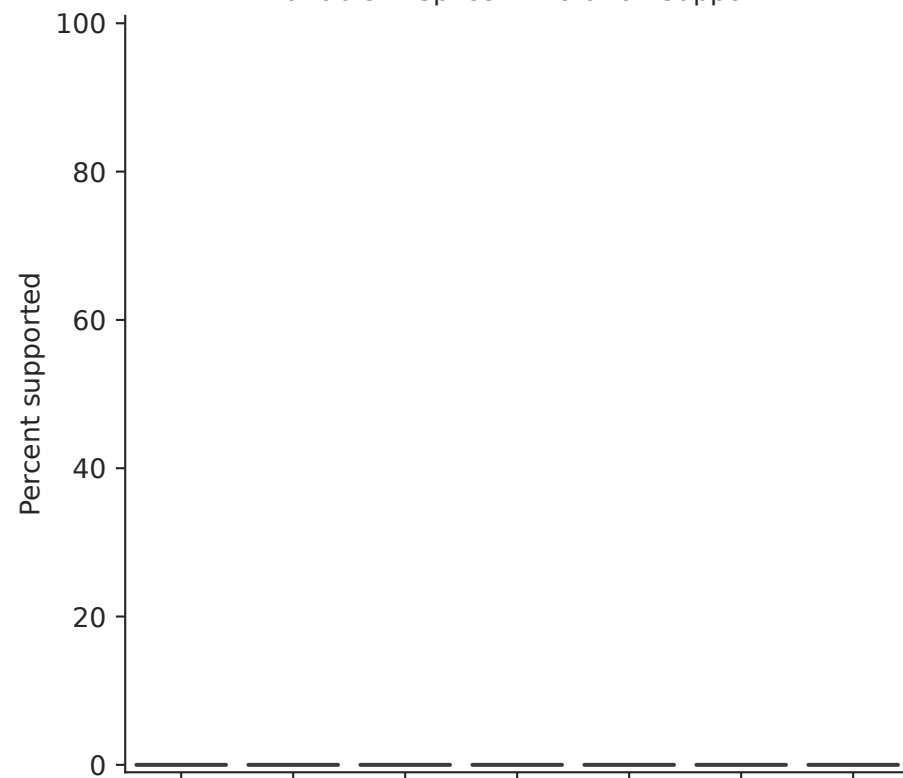
variable = Original Introns



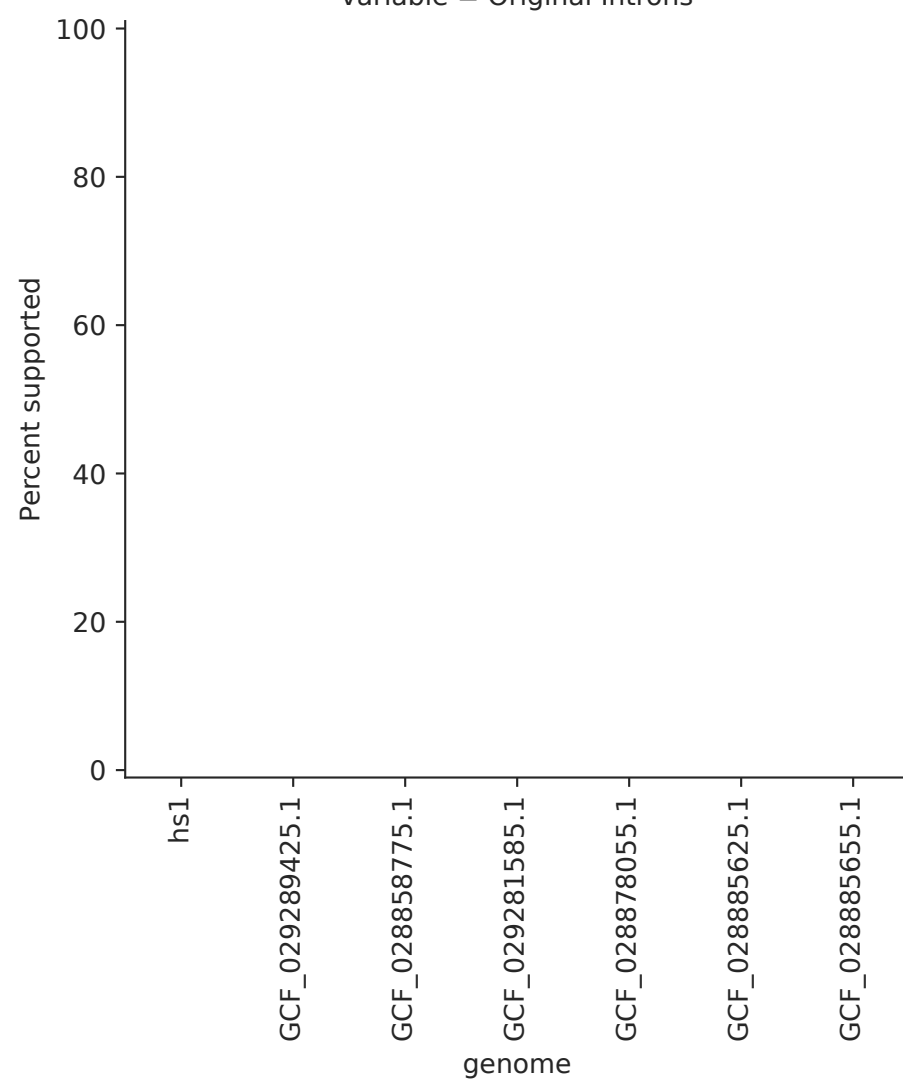
# Reference annotation support for pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support

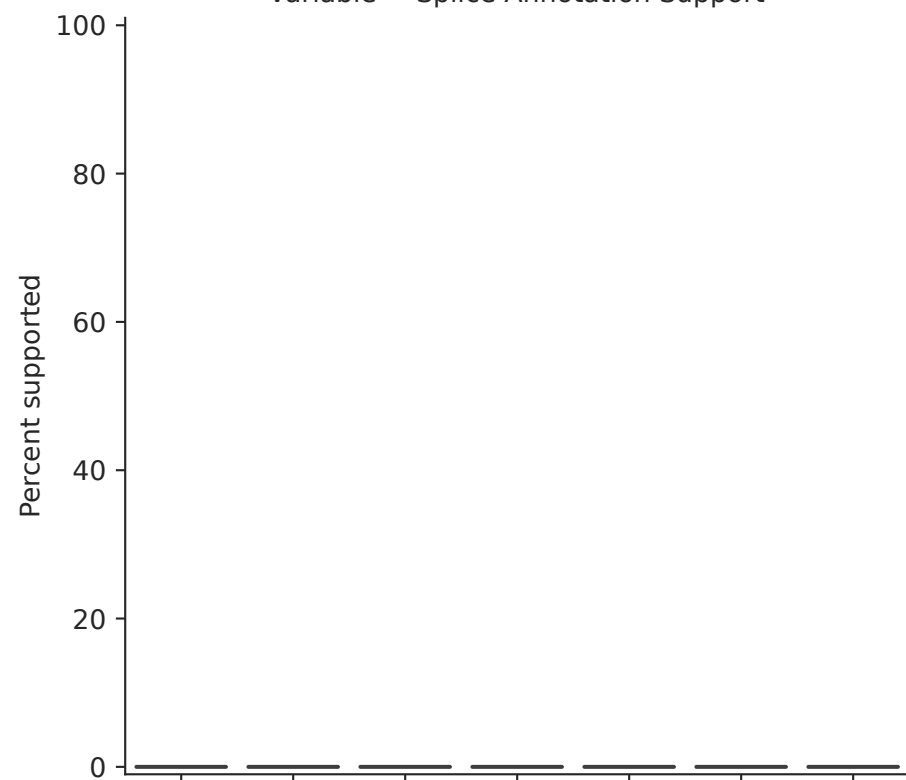


variable = Original Introns

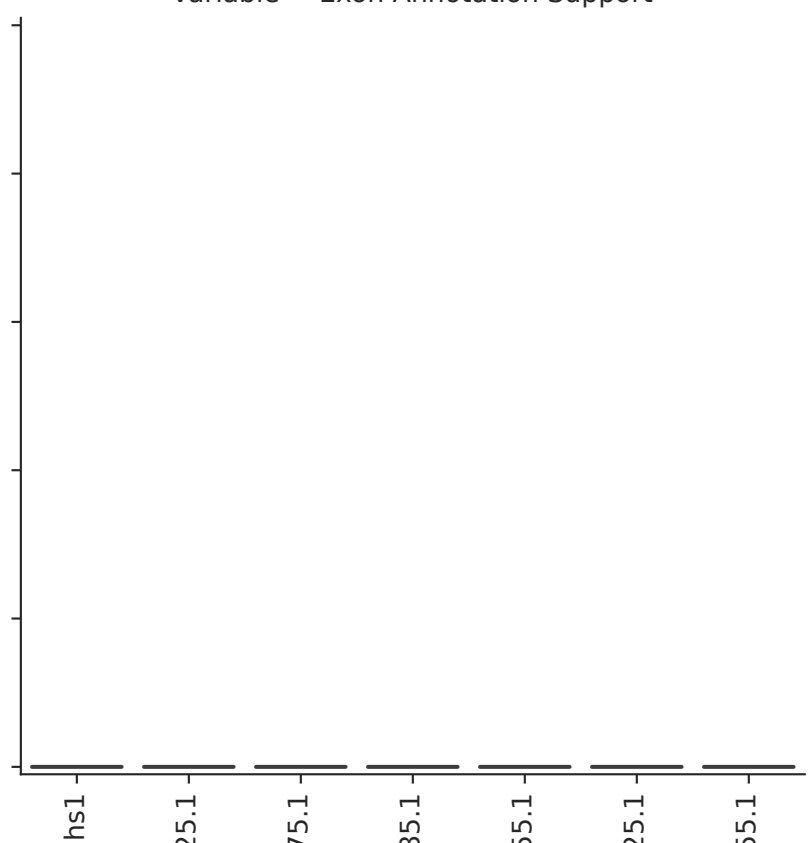


# Reference annotation support for rRNA

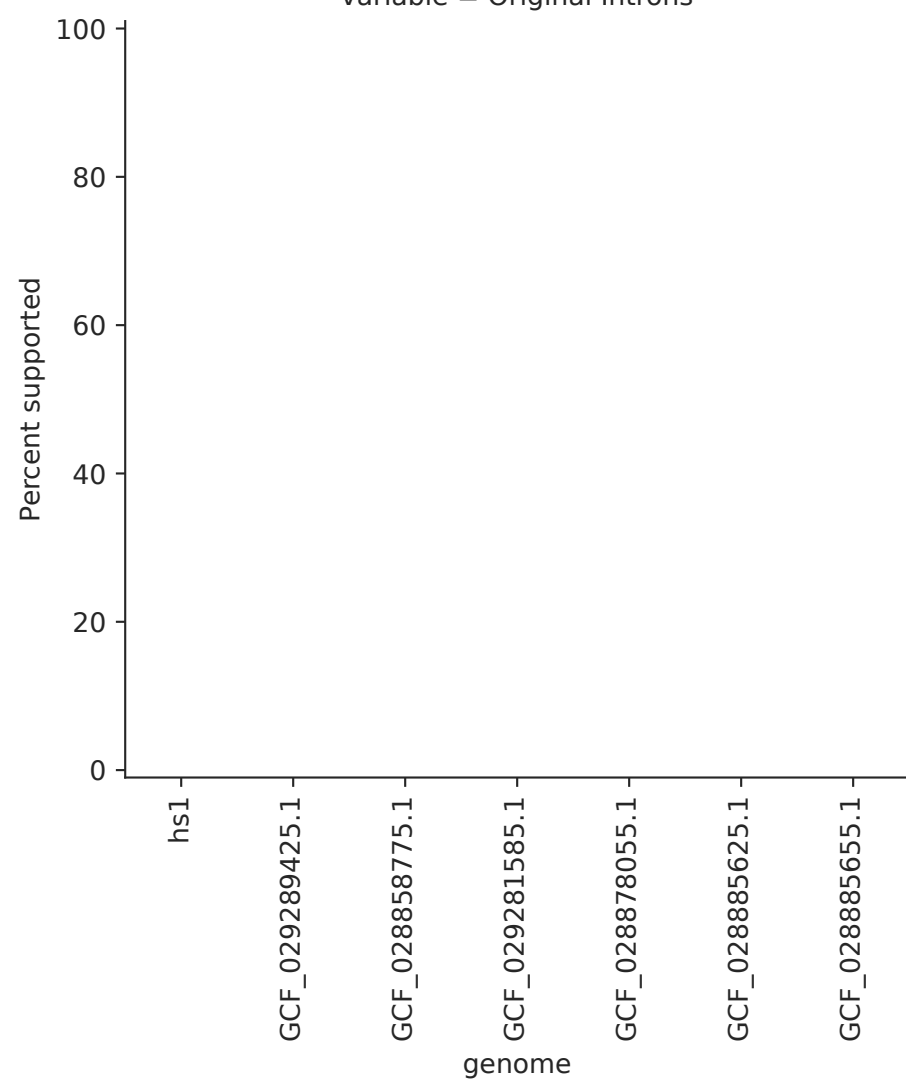
variable = Splice Annotation Support



variable = Exon Annotation Support



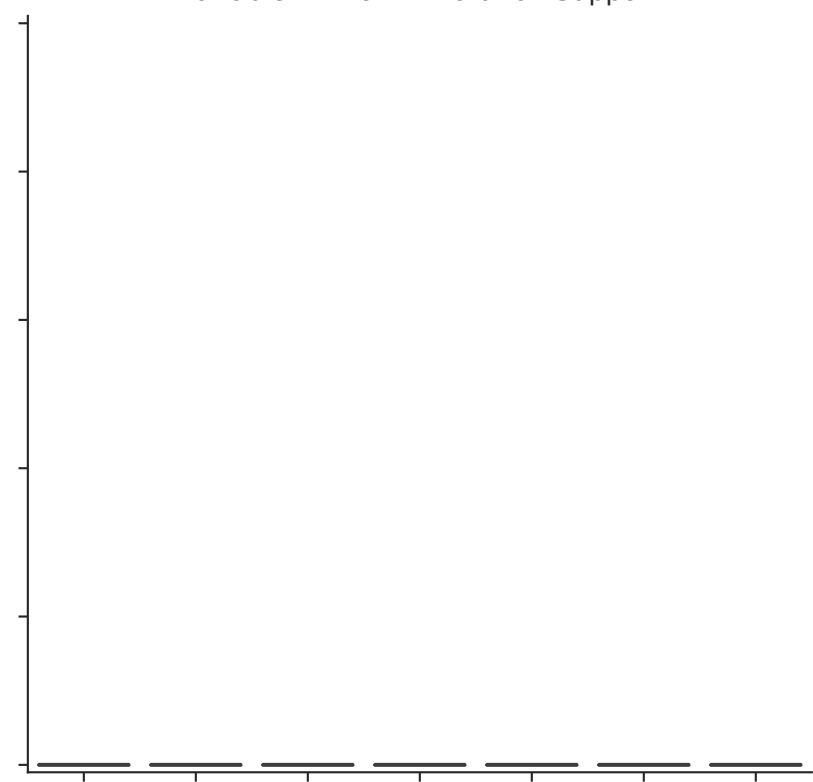
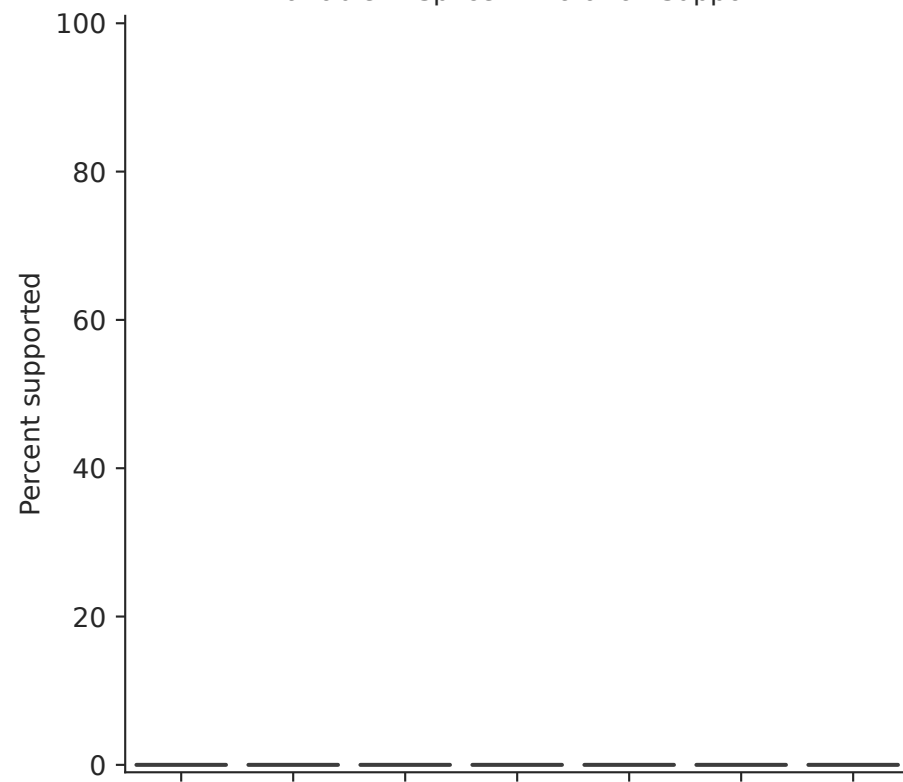
variable = Original Introns



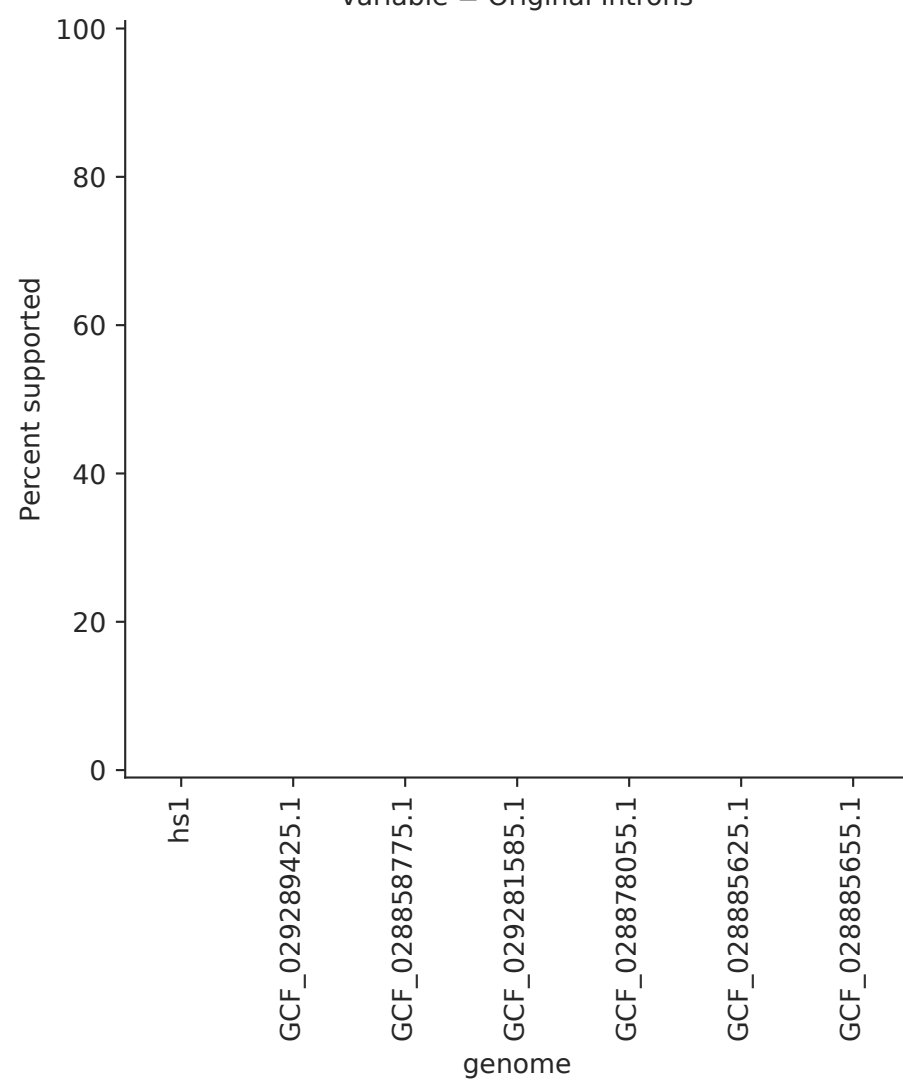
# Reference annotation support for rRNA\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



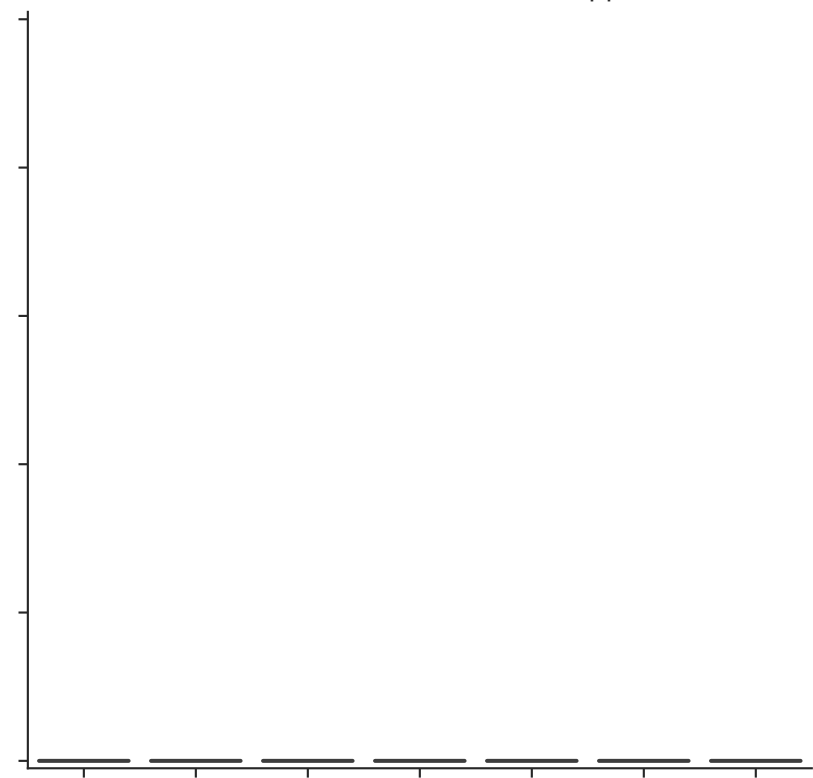
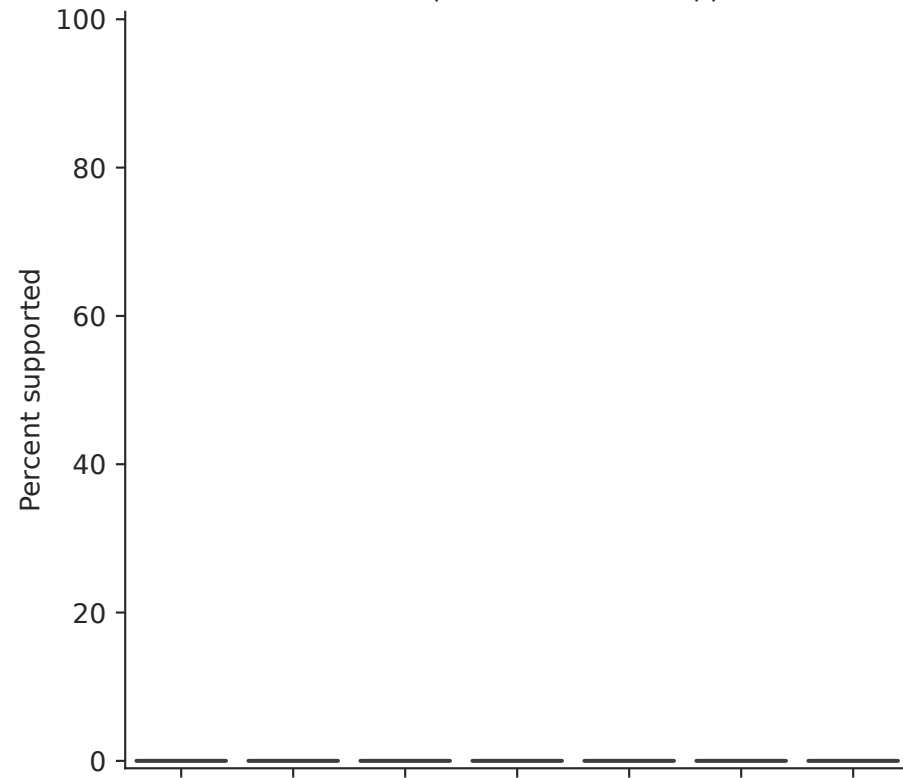
variable = Original Introns



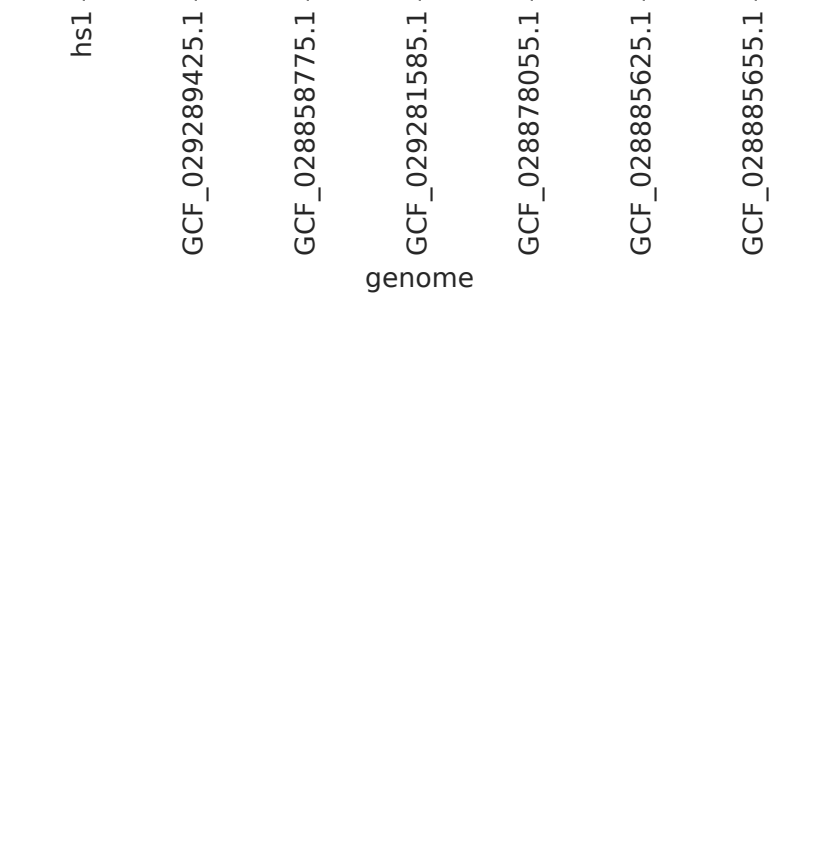
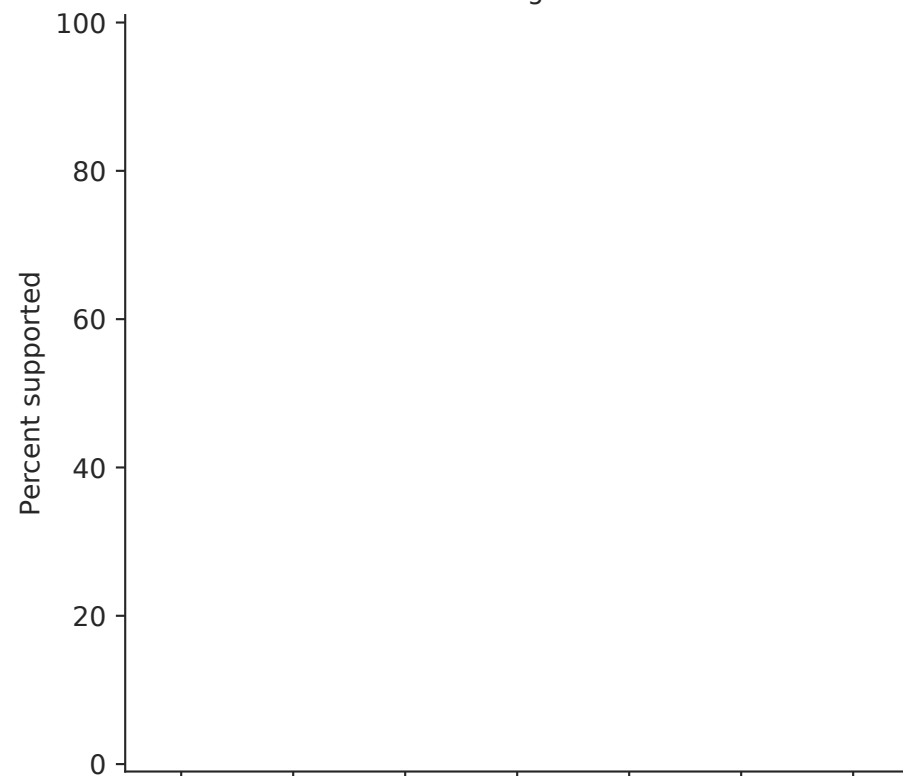
# Reference annotation support for retained\_intron

variable = Splice Annotation Support

variable = Exon Annotation Support



variable = Original Introns



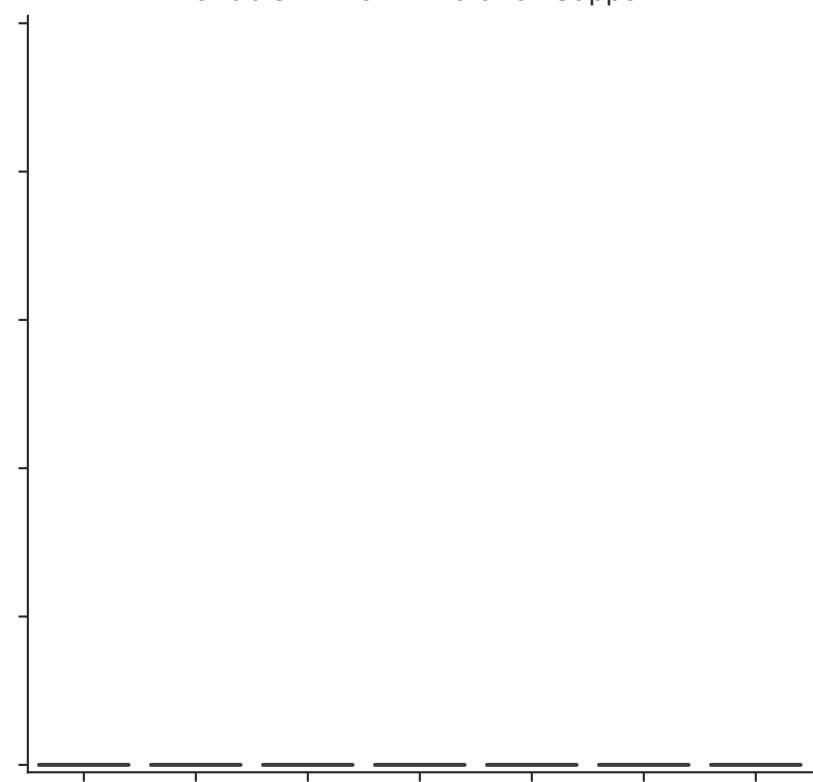
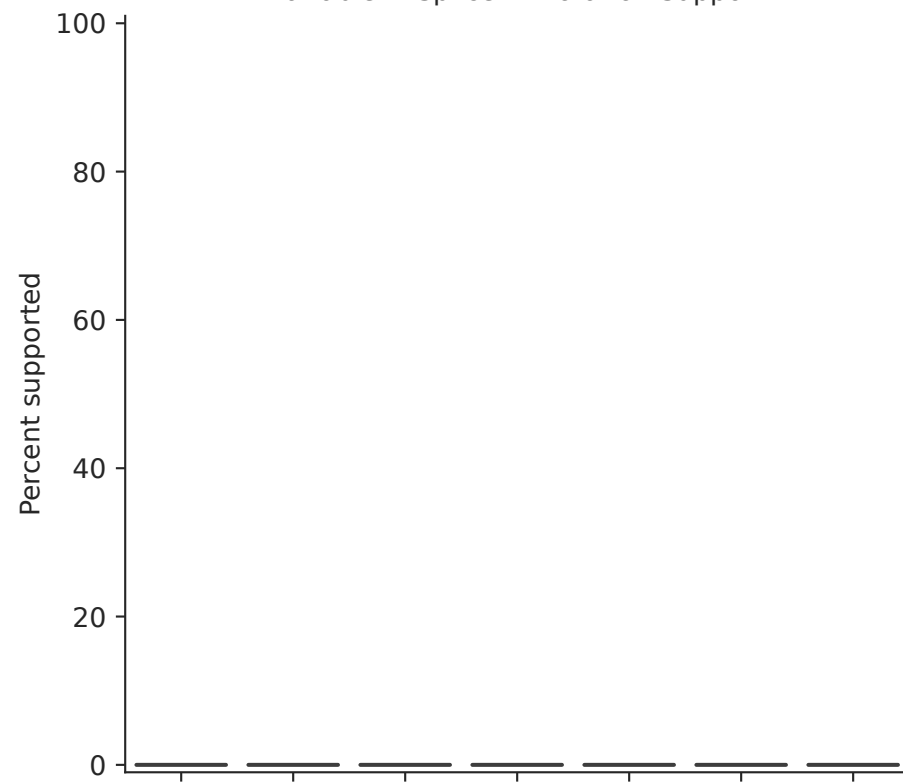
genome

genome

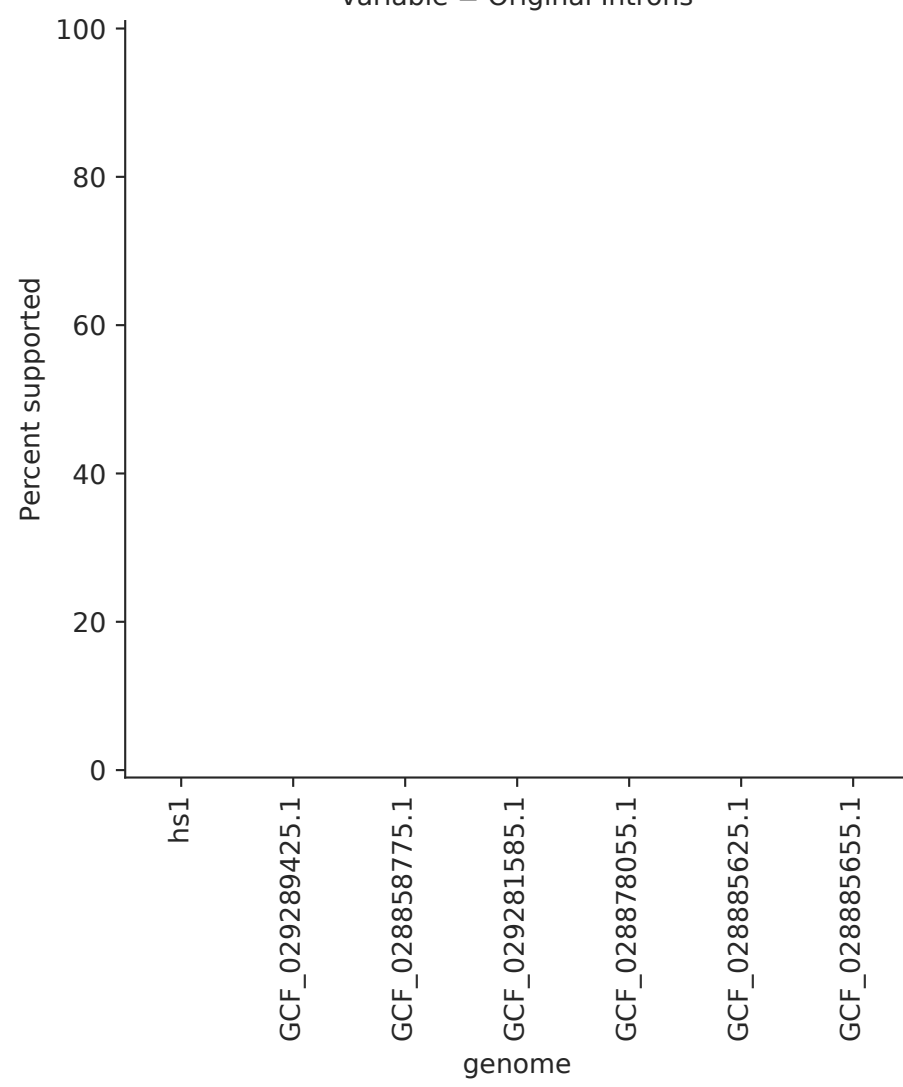
# Reference annotation support for ribozyme

variable = Splice Annotation Support

variable = Exon Annotation Support



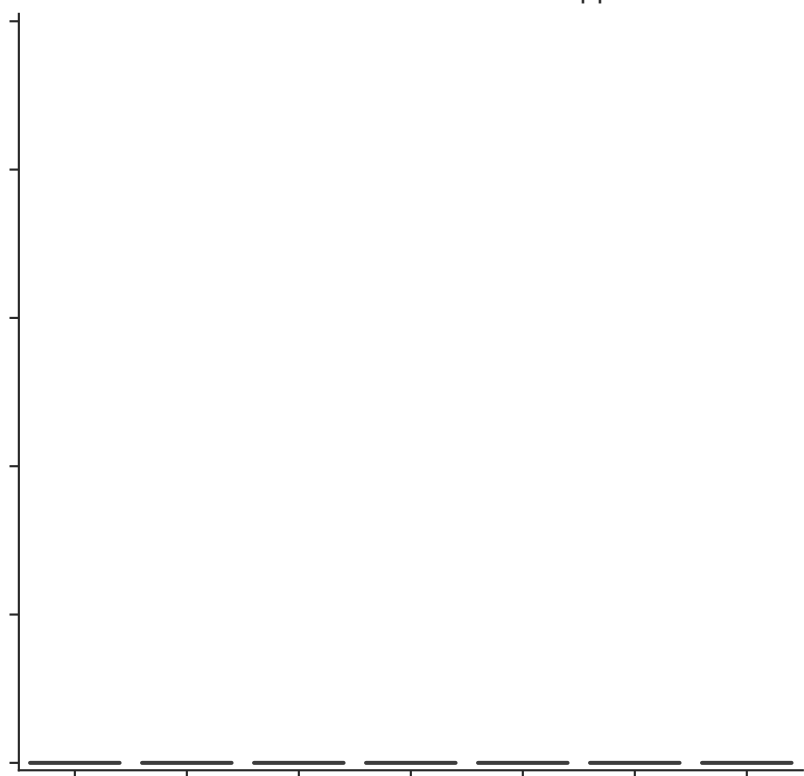
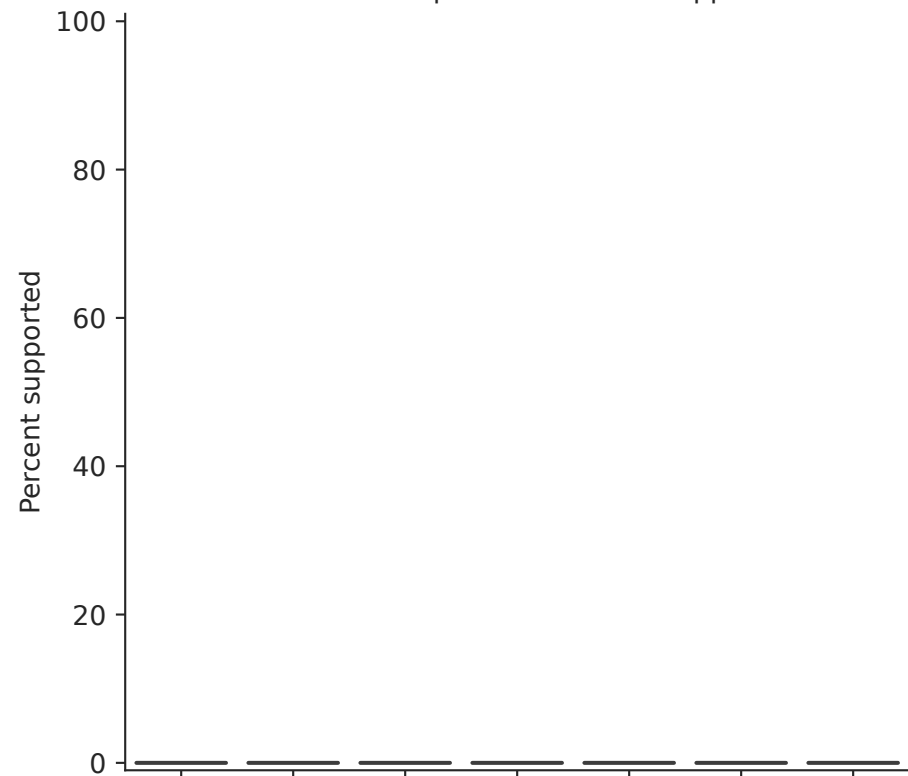
variable = Original Introns



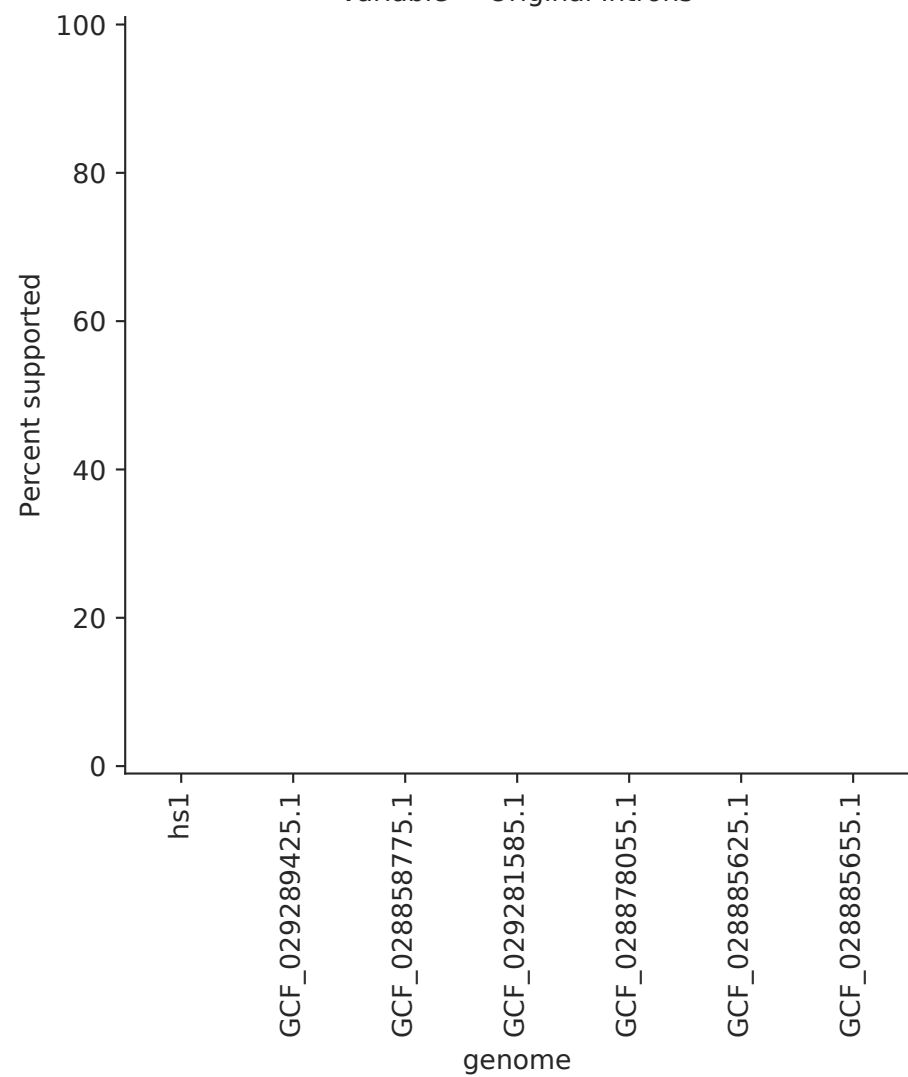
# Reference annotation support for sRNA

variable = Splice Annotation Support

variable = Exon Annotation Support



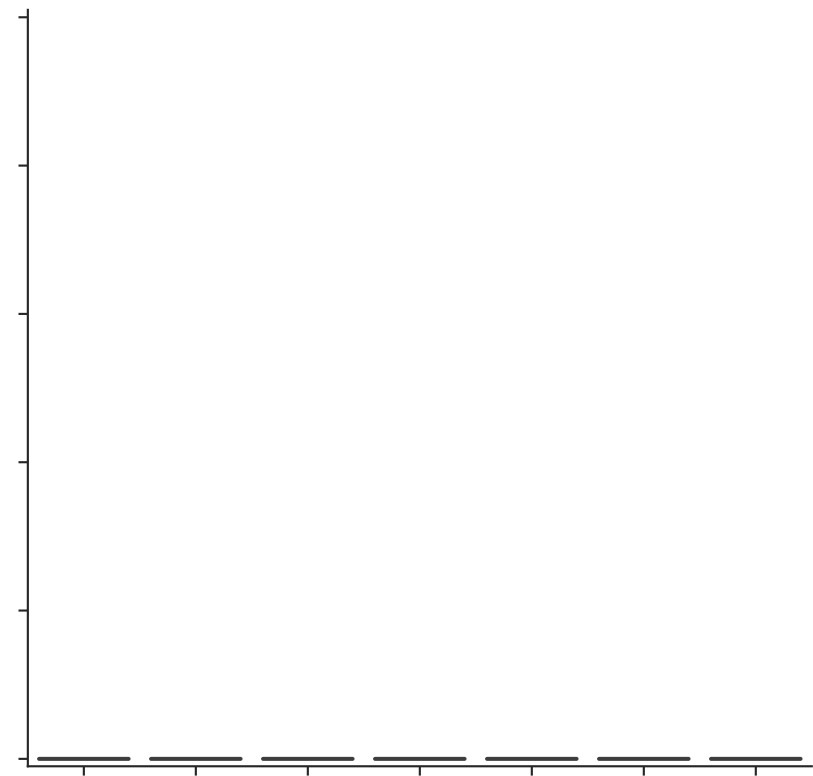
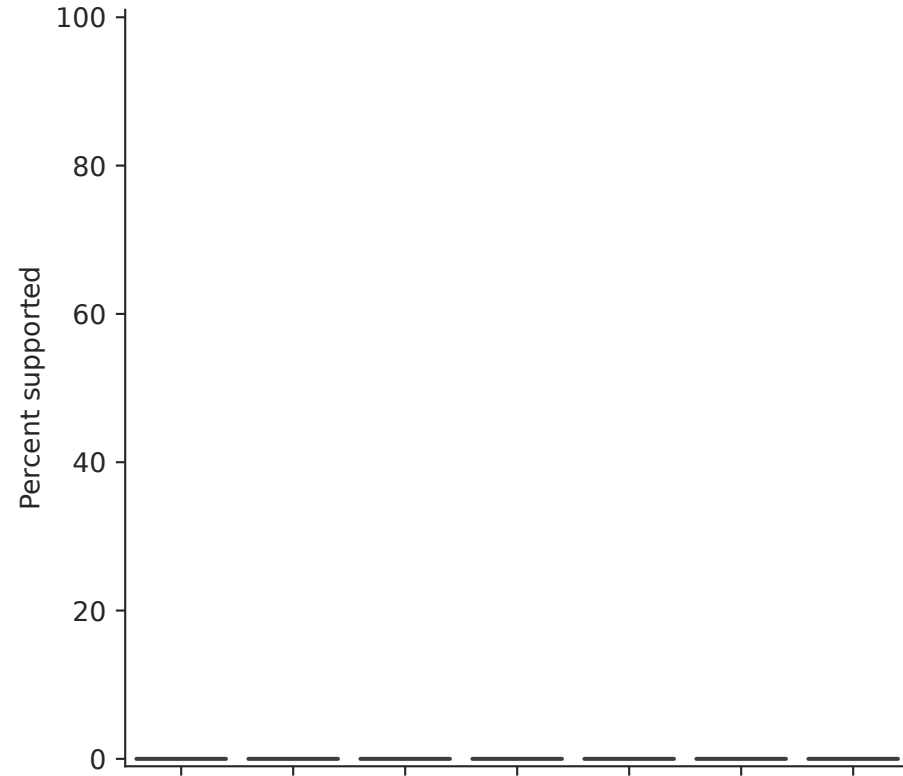
variable = Original Introns



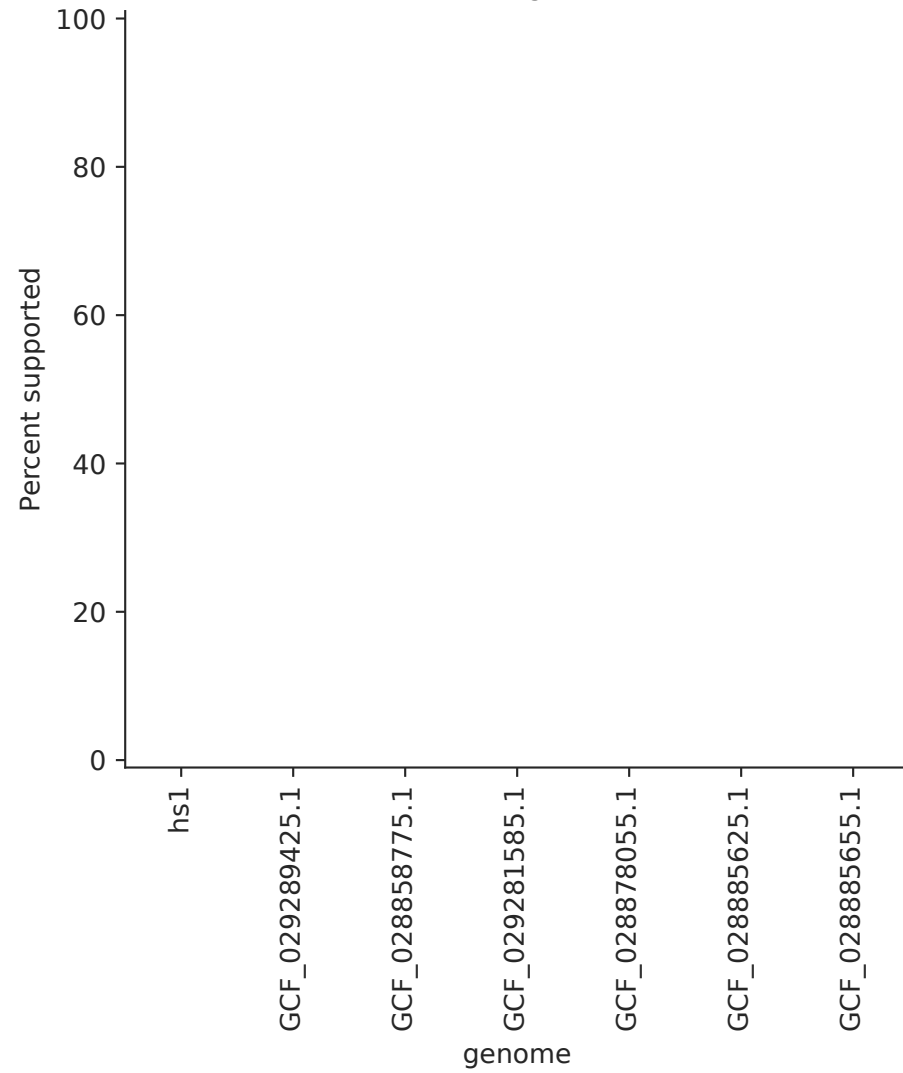
# Reference annotation support for scRNA

variable = Splice Annotation Support

variable = Exon Annotation Support



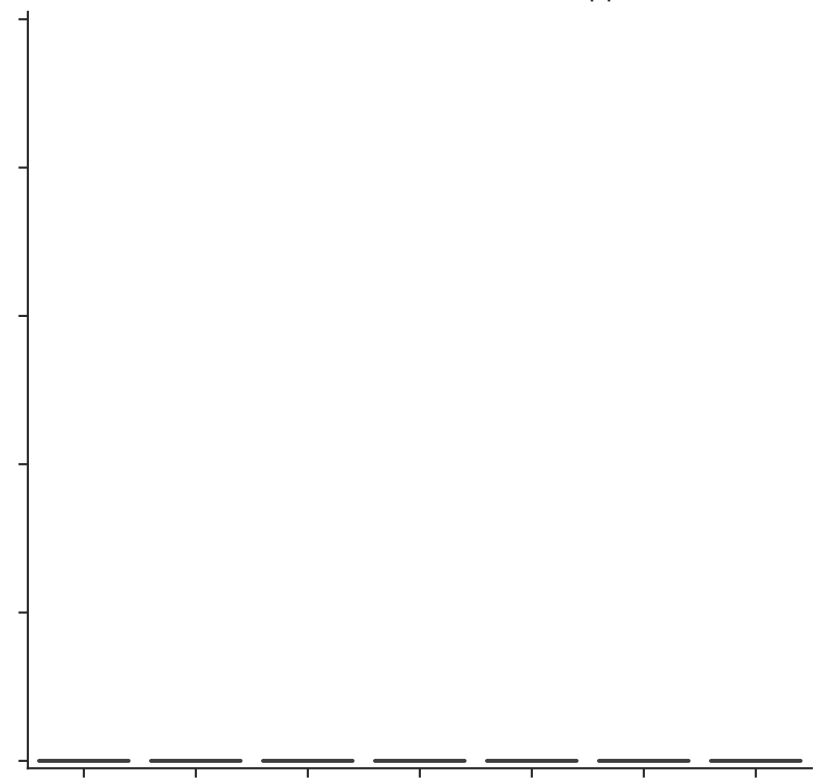
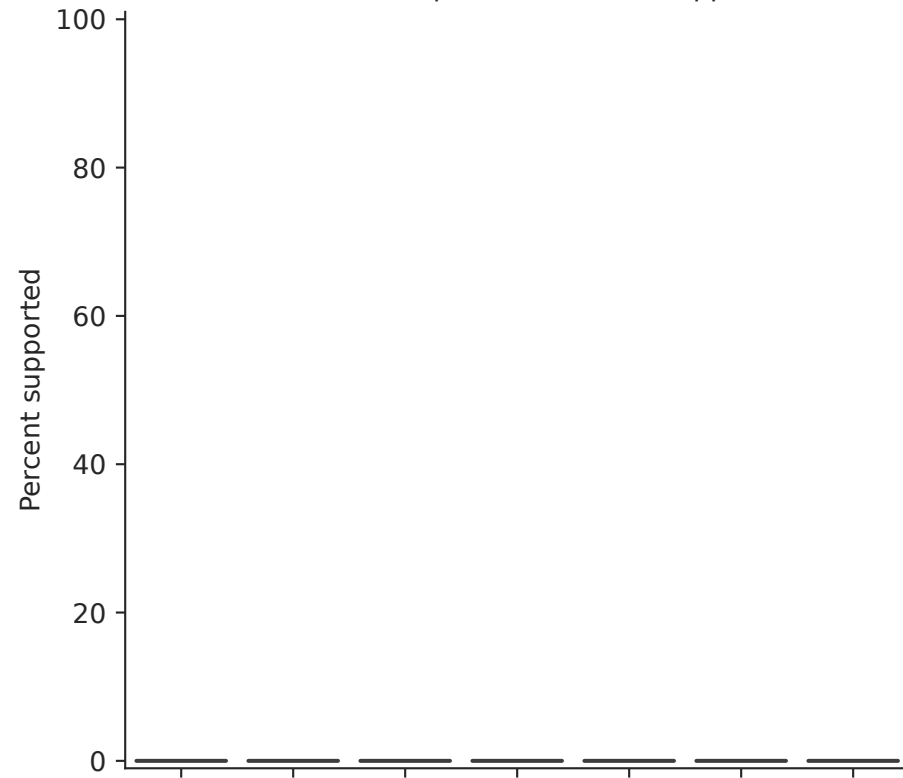
variable = Original Introns



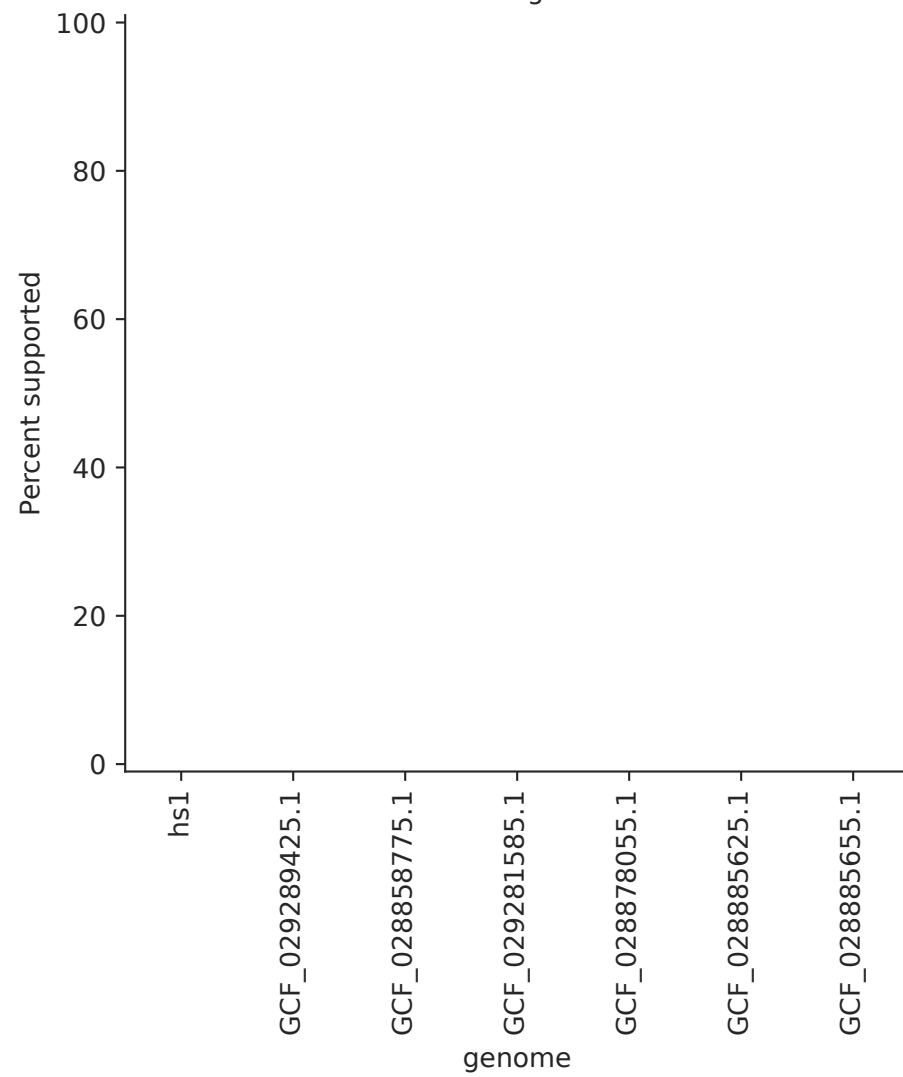
# Reference annotation support for scaRNA

variable = Splice Annotation Support

variable = Exon Annotation Support



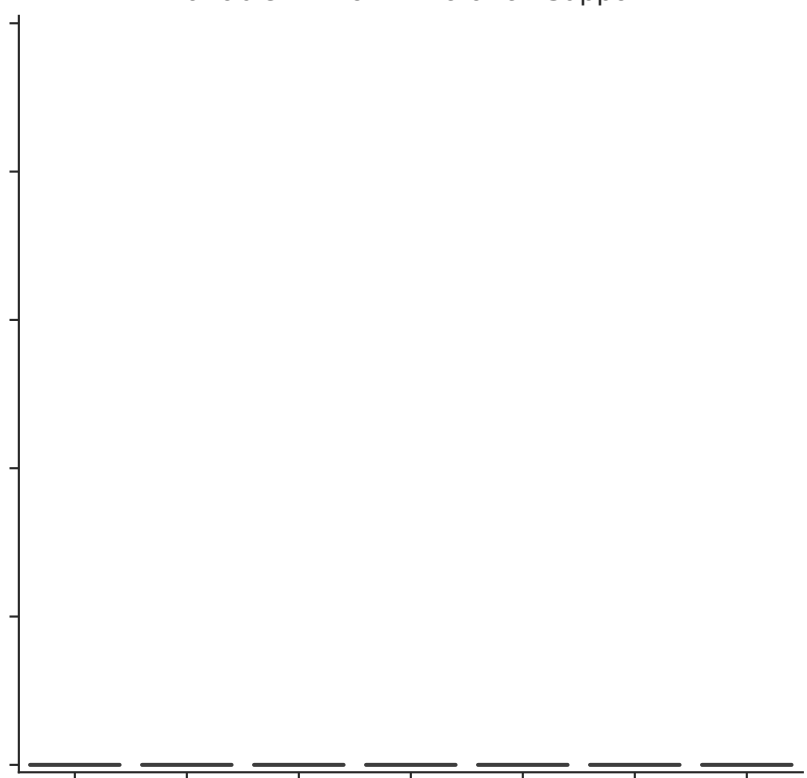
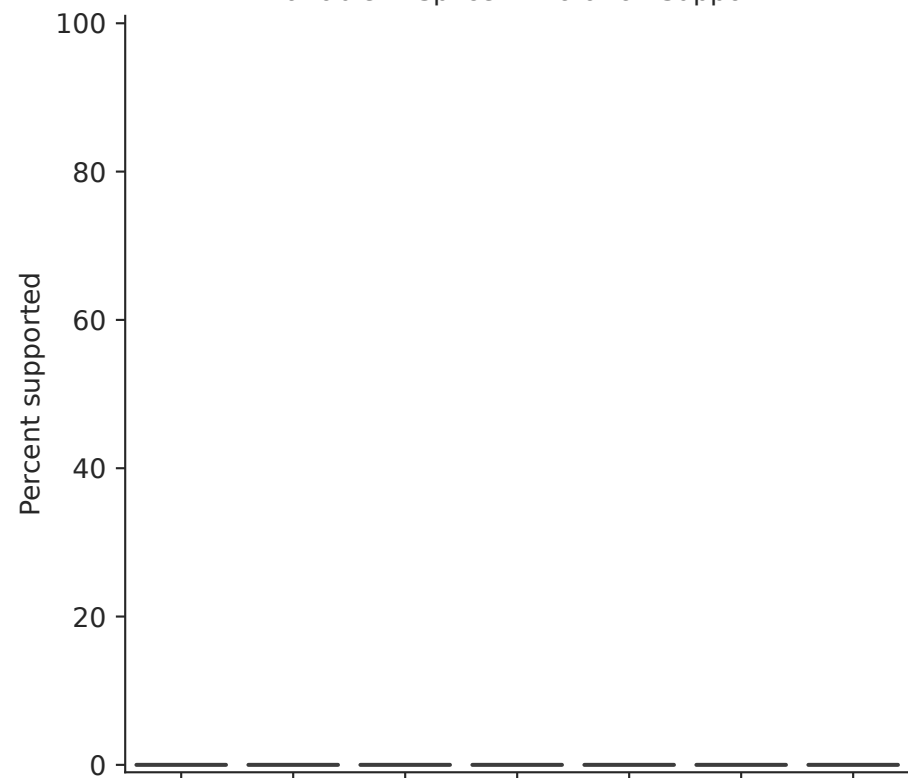
variable = Original Introns



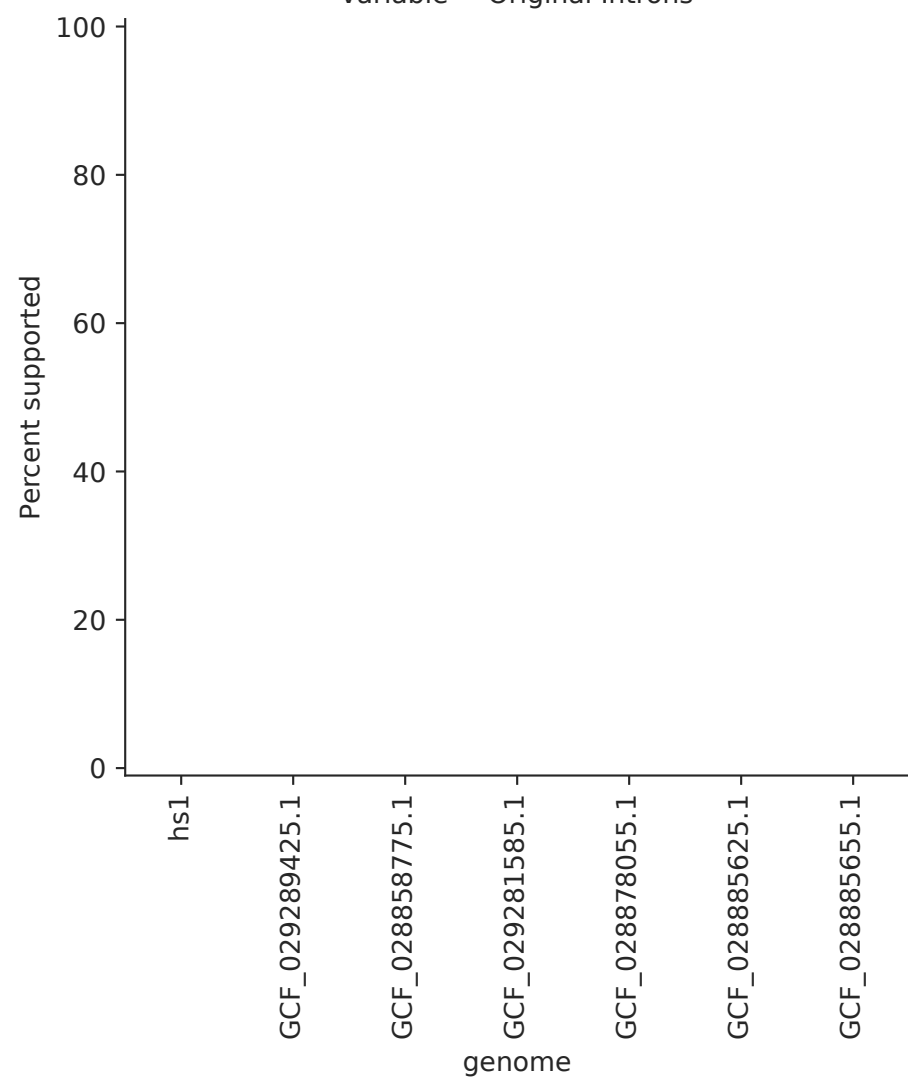
# Reference annotation support for snRNA

variable = Splice Annotation Support

variable = Exon Annotation Support



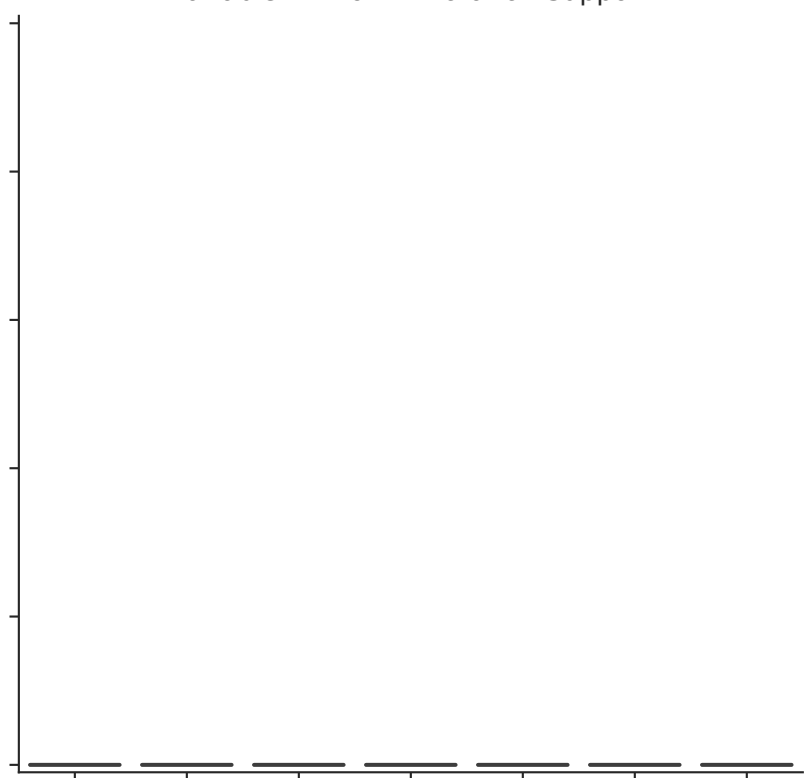
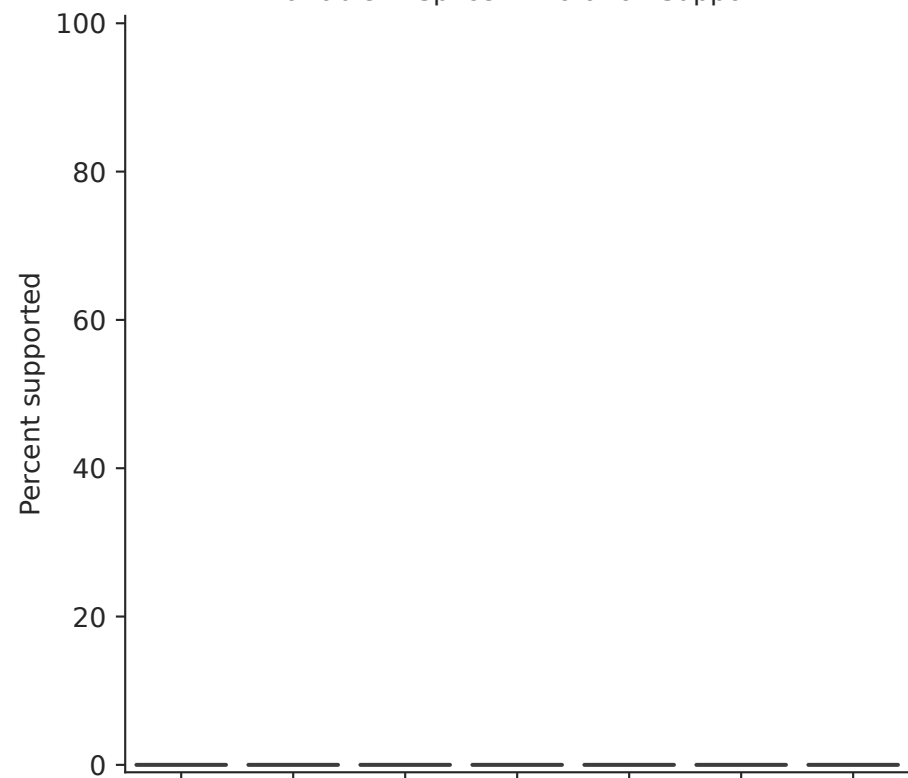
variable = Original Introns



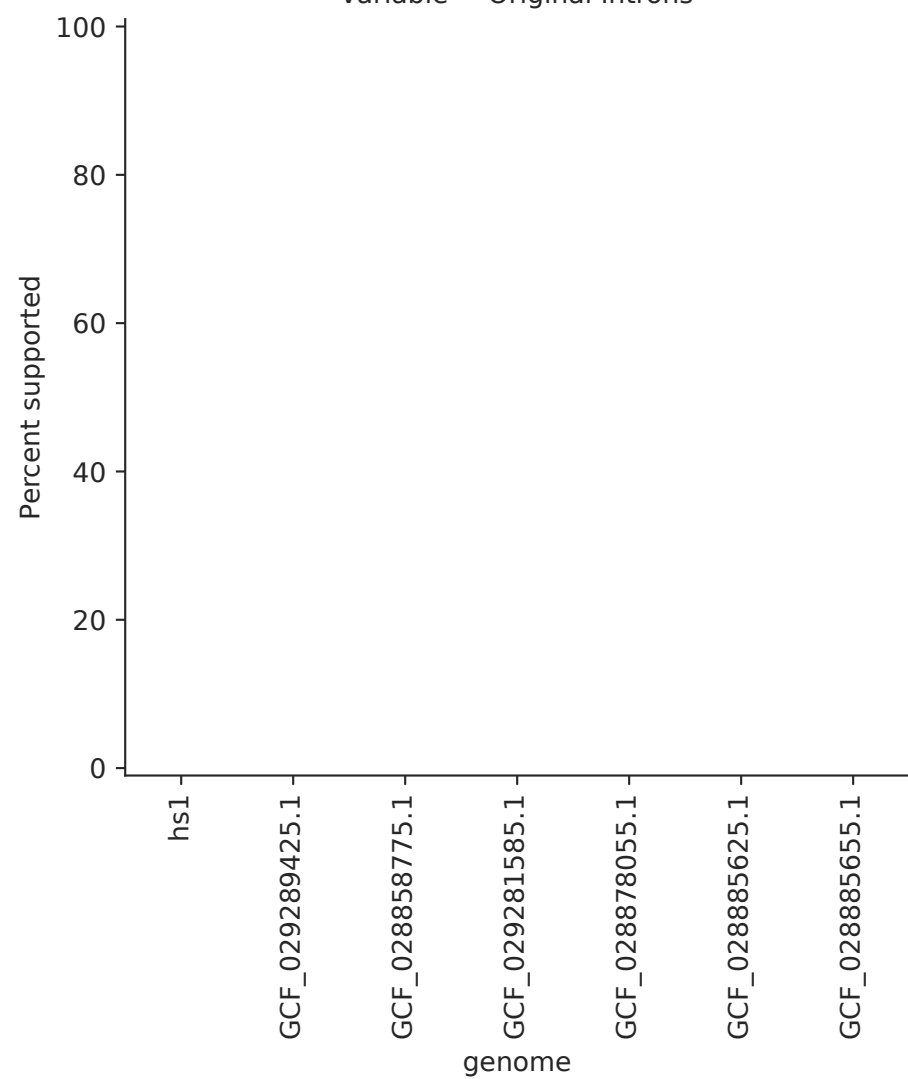
# Reference annotation support for snoRNA

variable = Splice Annotation Support

variable = Exon Annotation Support



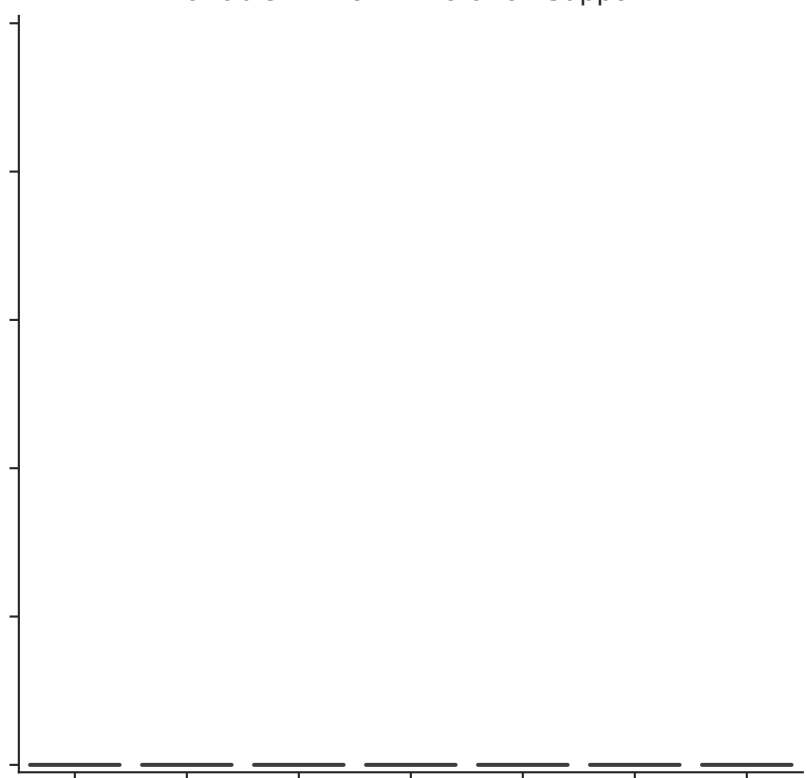
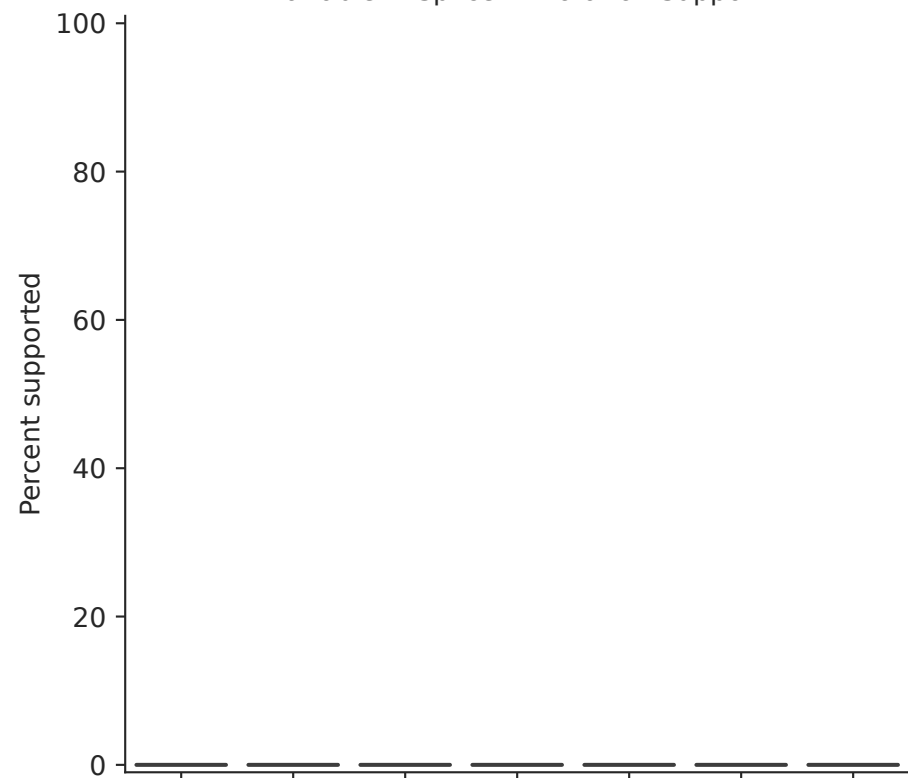
variable = Original Introns



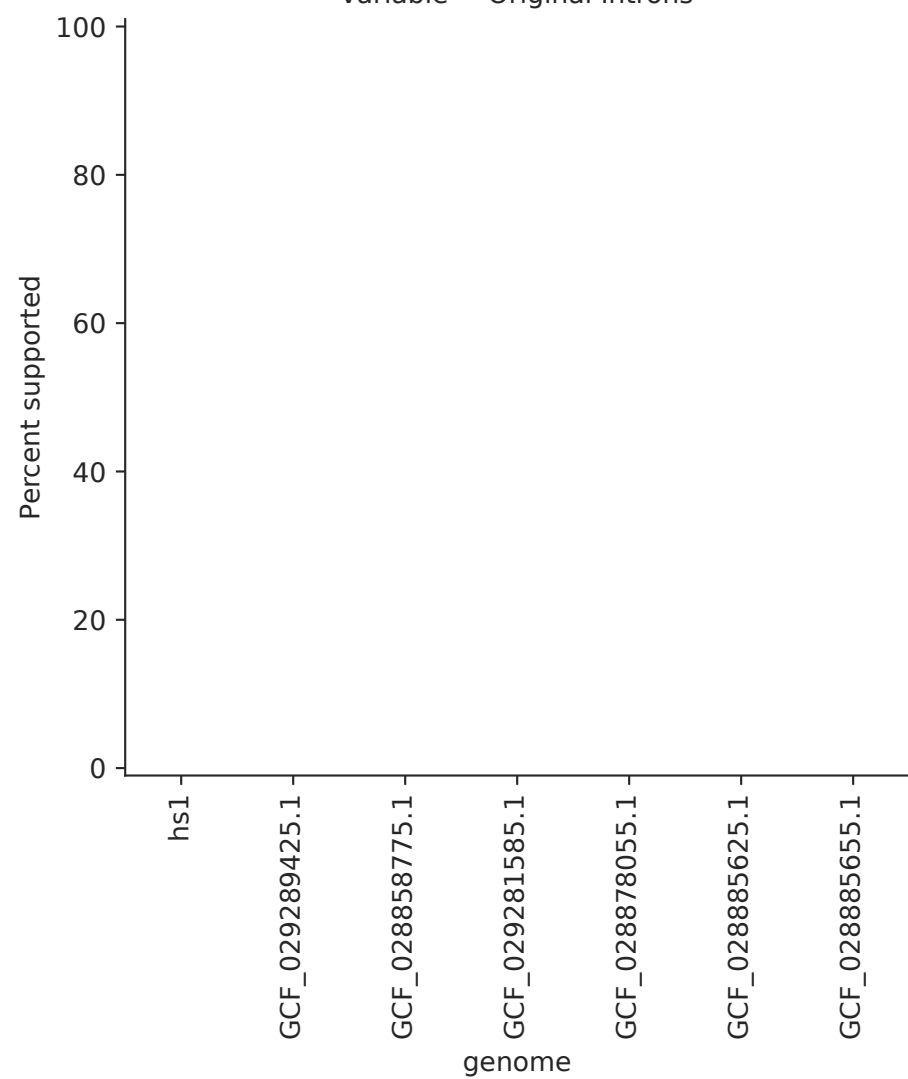
# Reference annotation support for transcribed\_processed\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



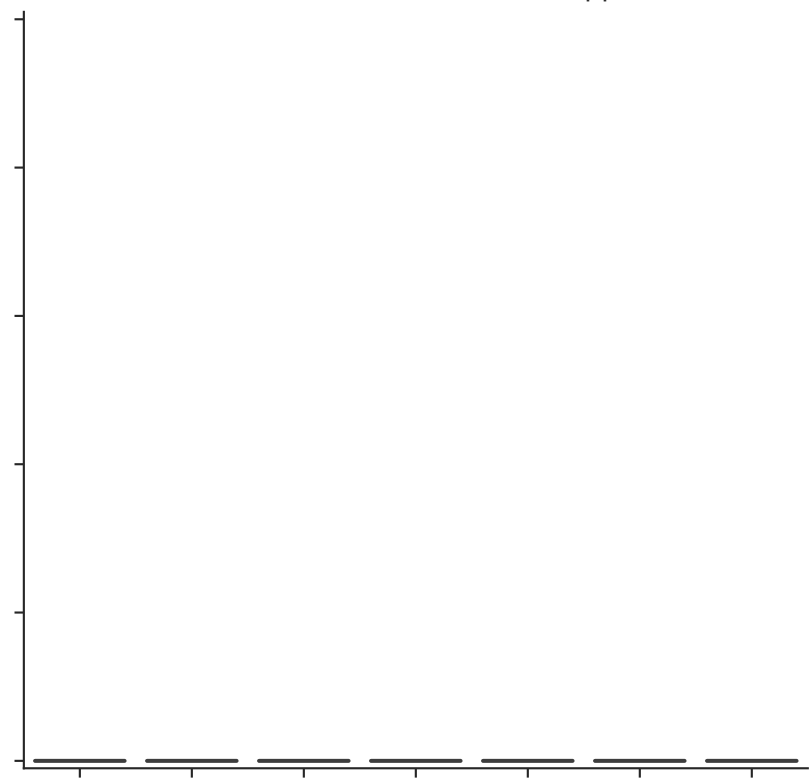
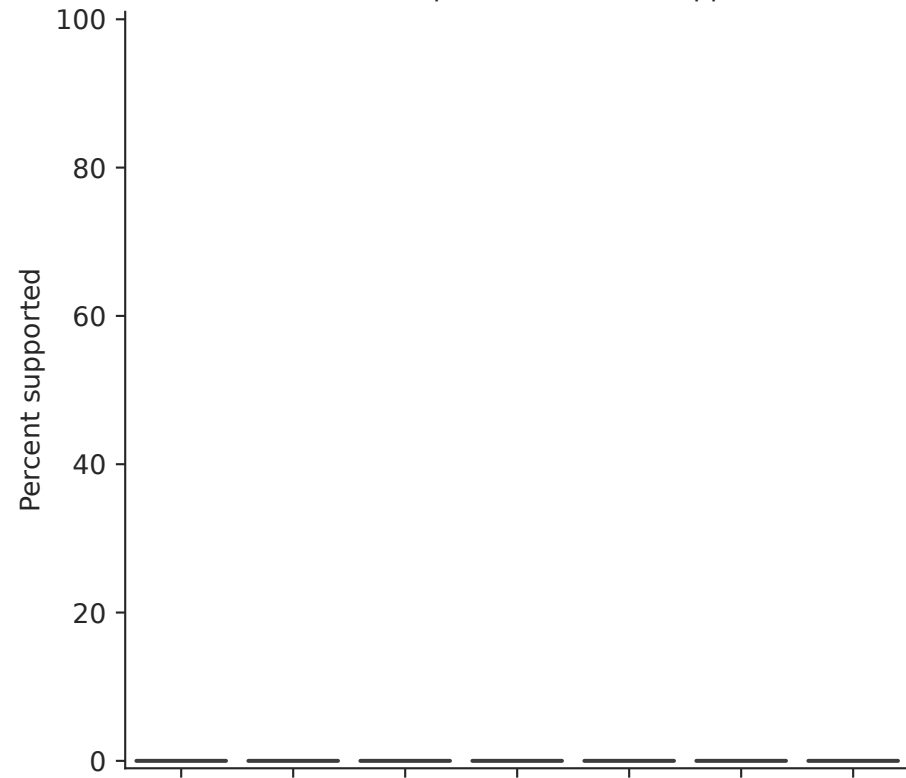
variable = Original Introns



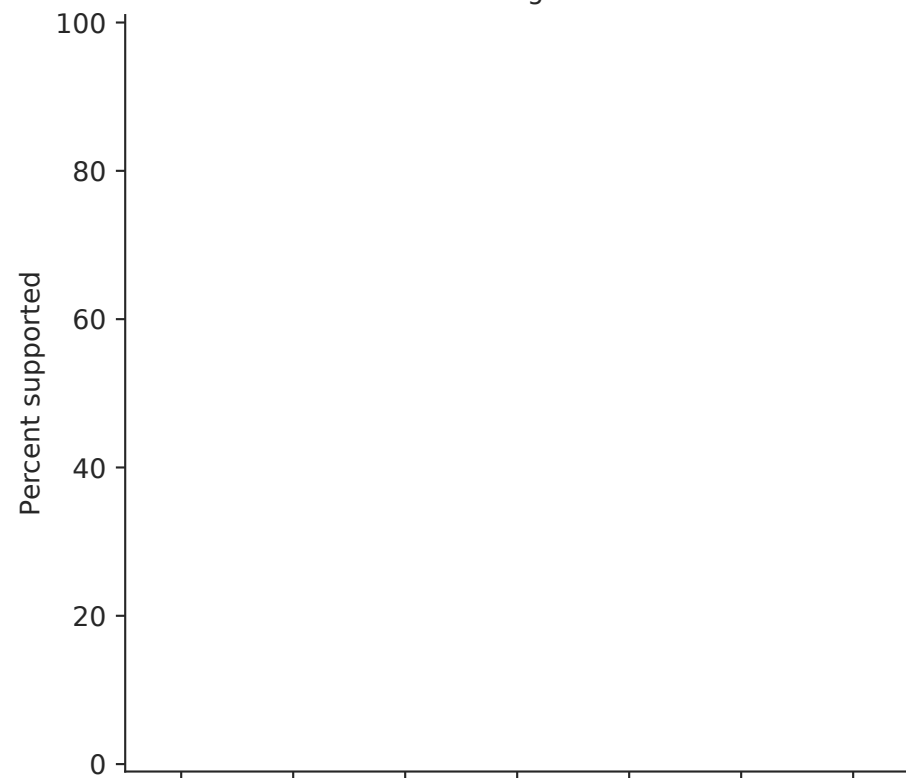
# Reference annotation support for transcribed\_unitary\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



variable = Original Introns



hs1

GCF\_029289425.1

GCF\_028858775.1

GCF\_029281585.1

GCF\_028878055.1

GCF\_028885625.1

GCF\_028885655.1

genome

hs1

GCF\_029289425.1

GCF\_028858775.1

GCF\_029281585.1

GCF\_028878055.1

GCF\_028885625.1

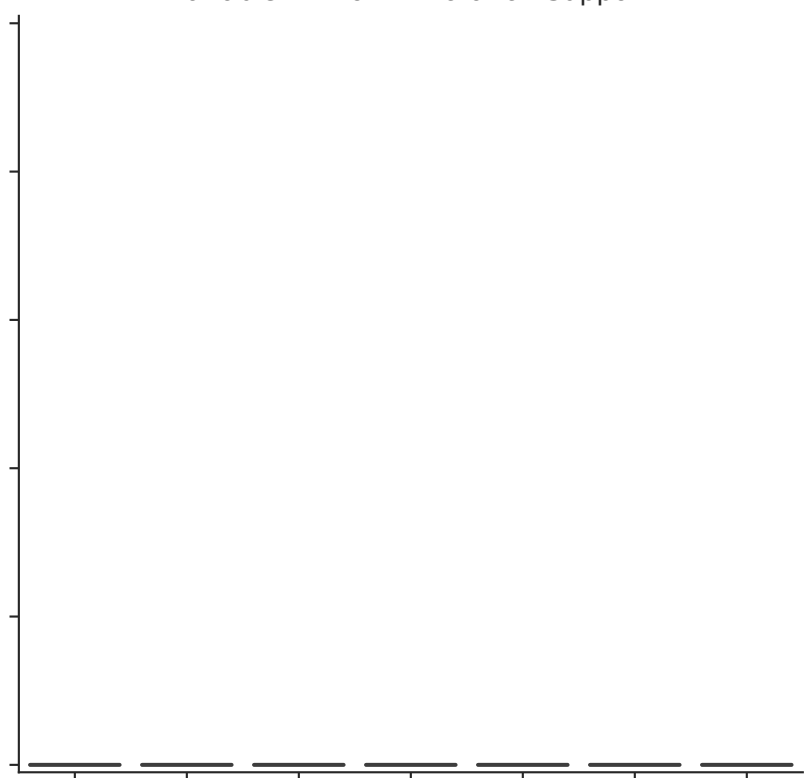
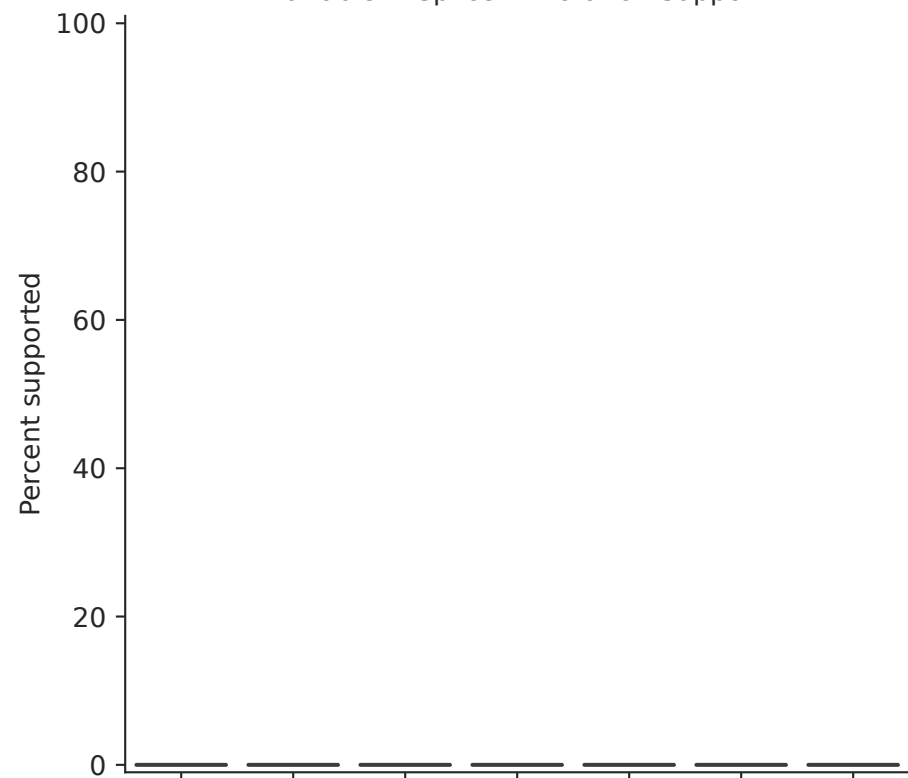
GCF\_028885655.1

genome

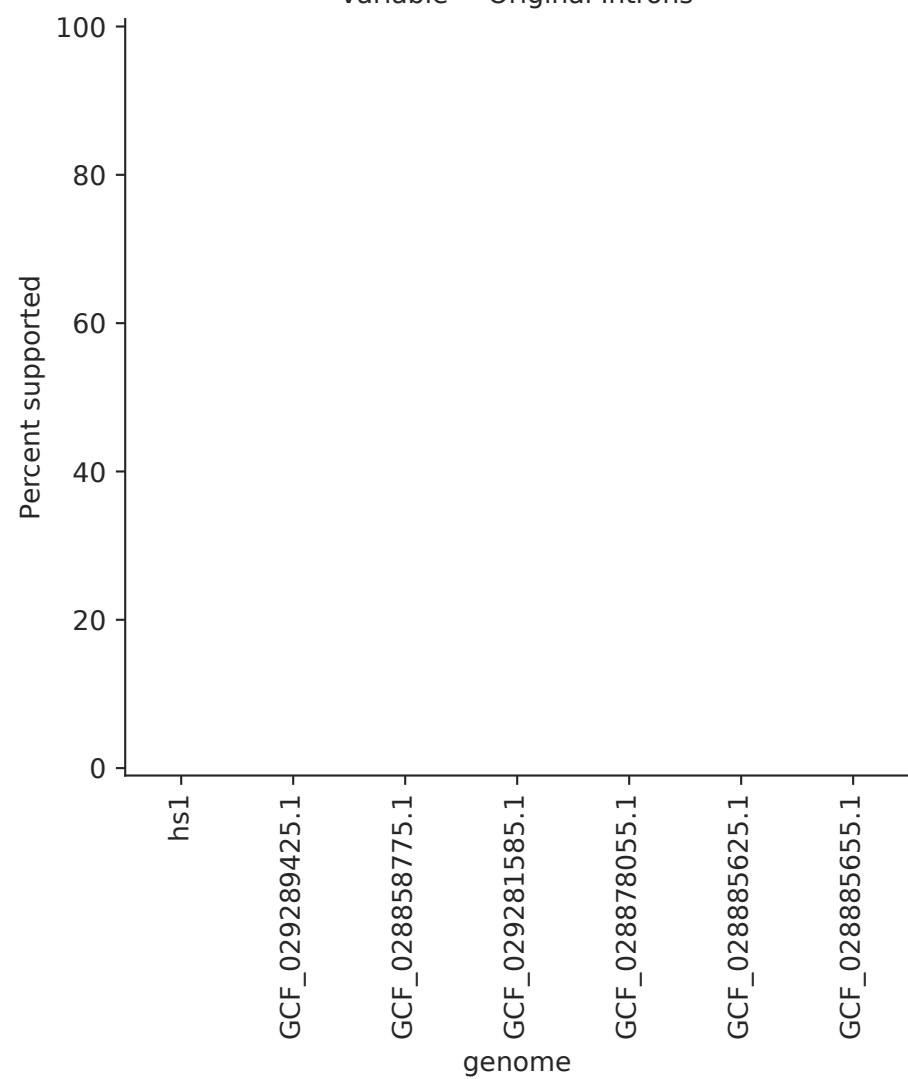
# Reference annotation support for transcribed\_unprocessed\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



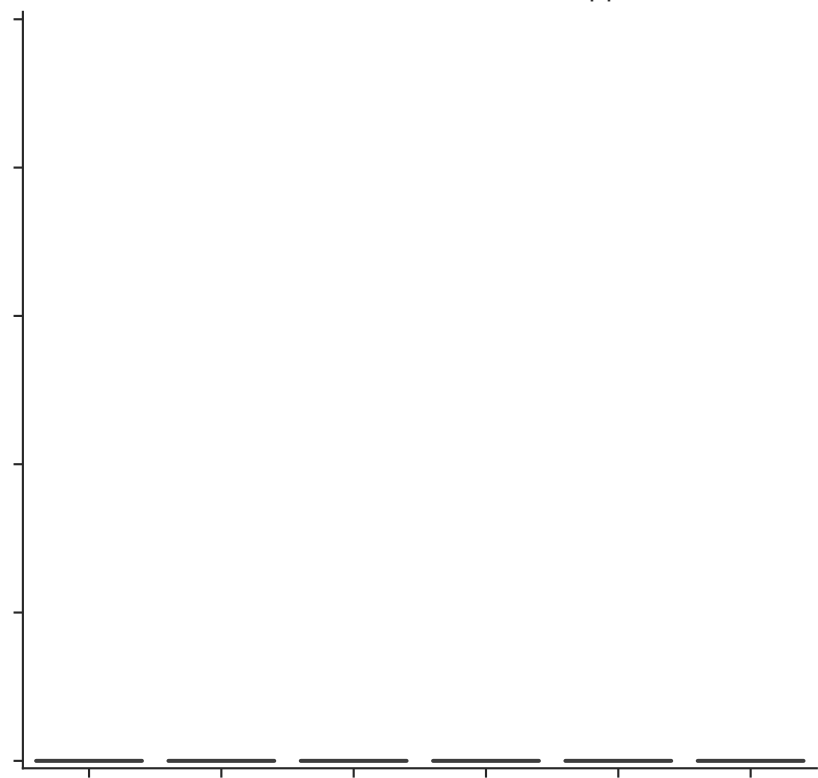
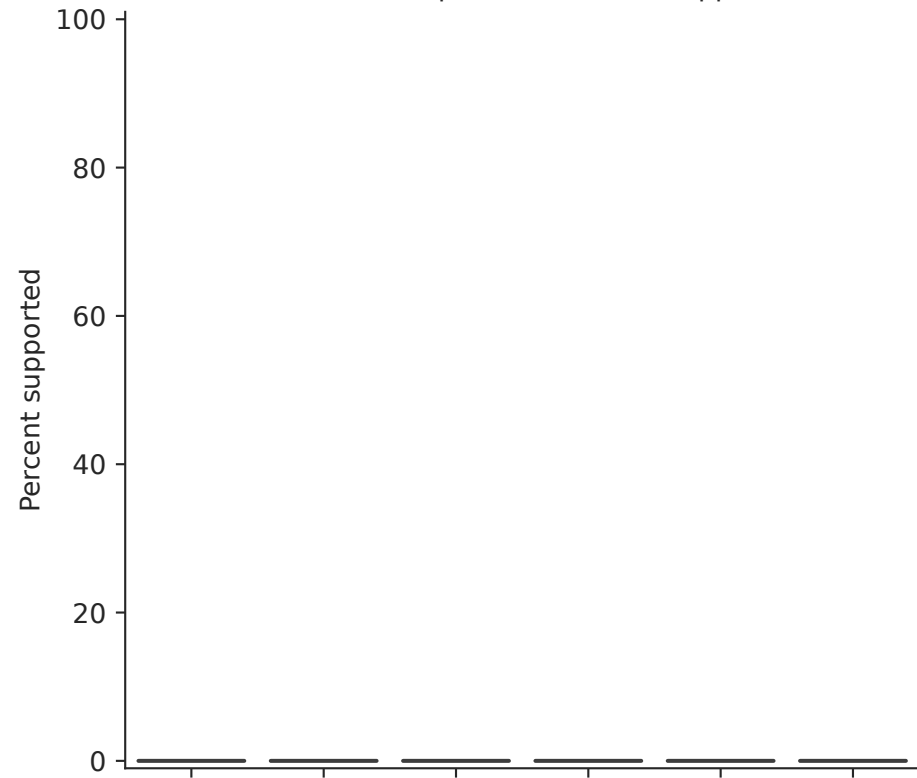
variable = Original Introns



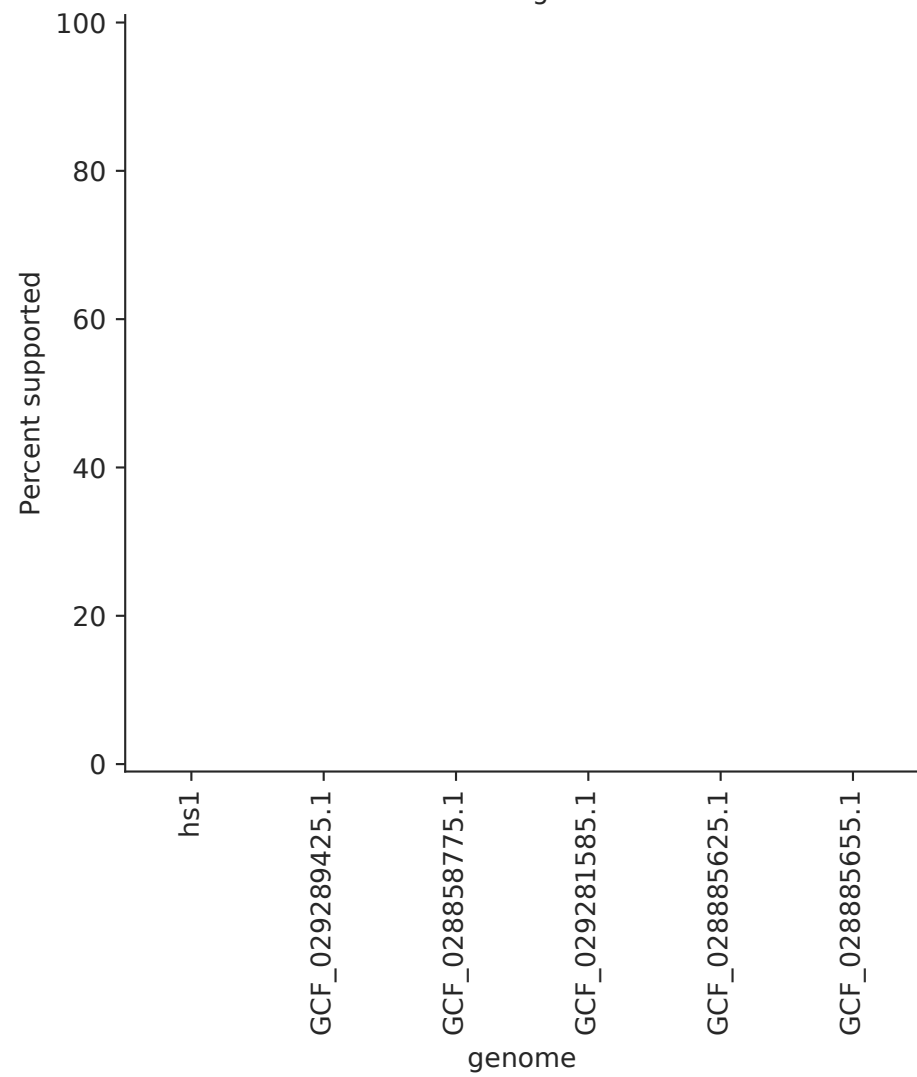
# Reference annotation support for translated\_processed\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



variable = Original Introns

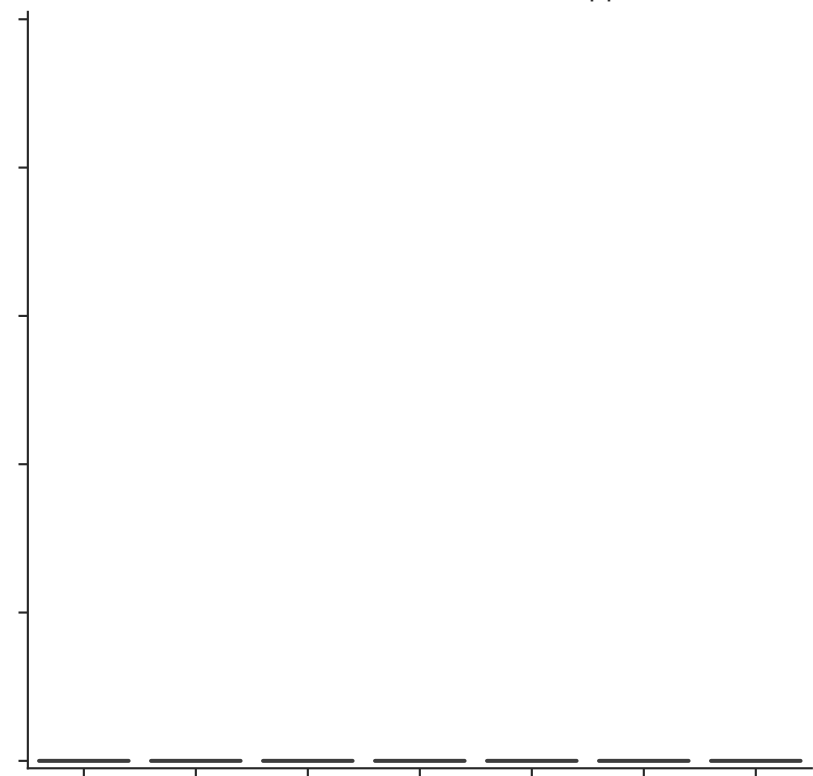
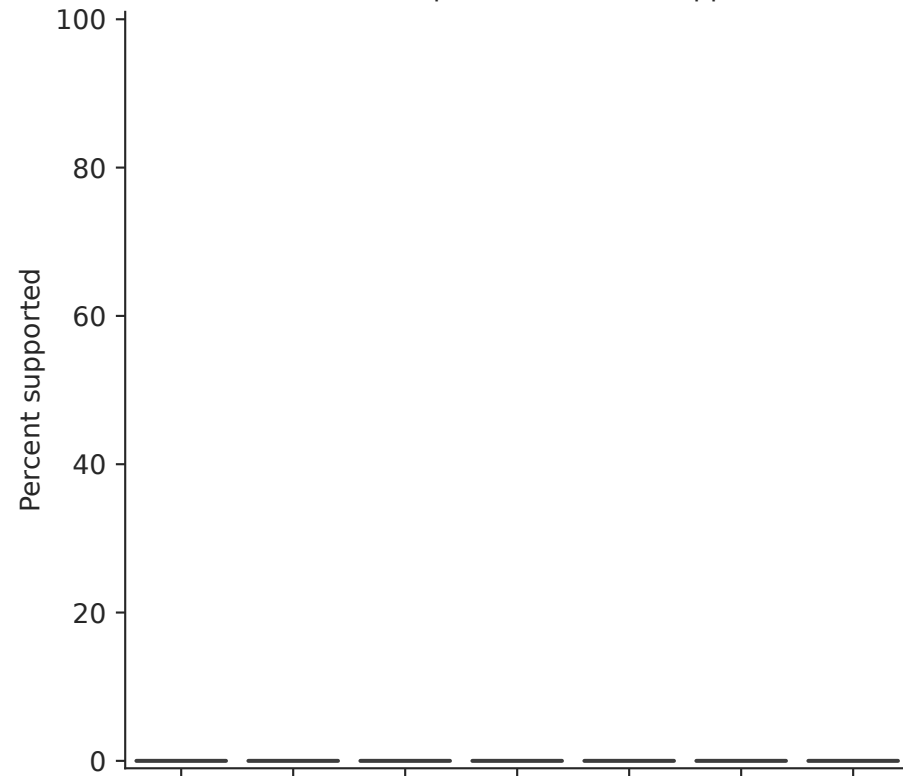


hs1  
GCF\_029289425.1  
GCF\_028858775.1  
GCF\_029281585.1  
GCF\_028885625.1  
GCF\_028885655.1  
genome

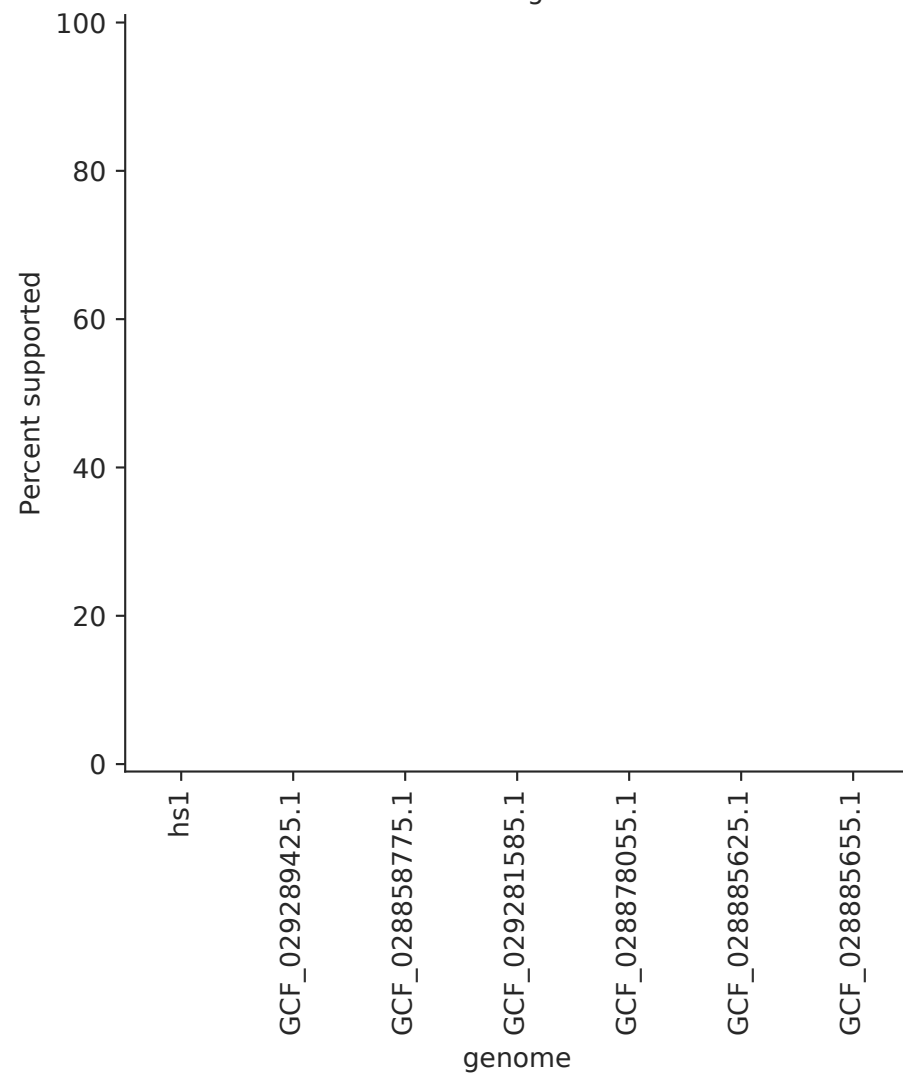
# Reference annotation support for translated\_unprocessed\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



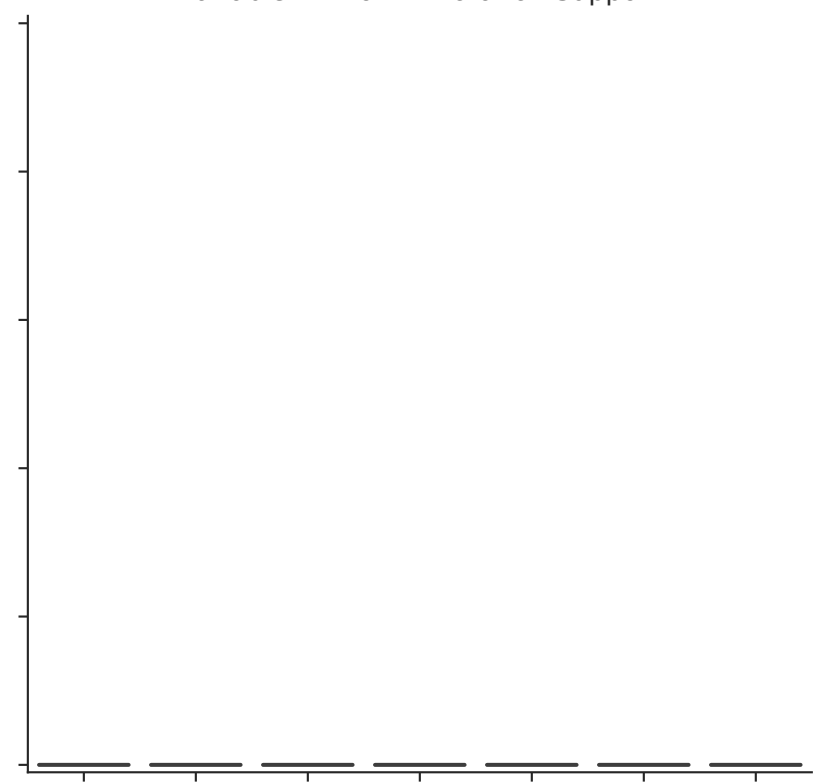
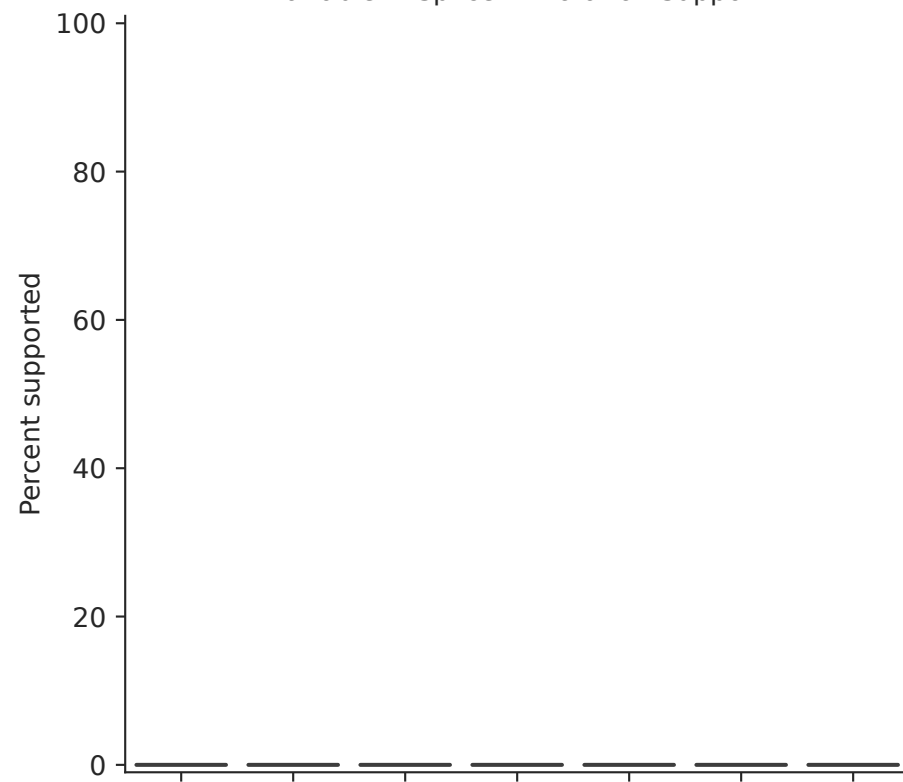
variable = Original Introns



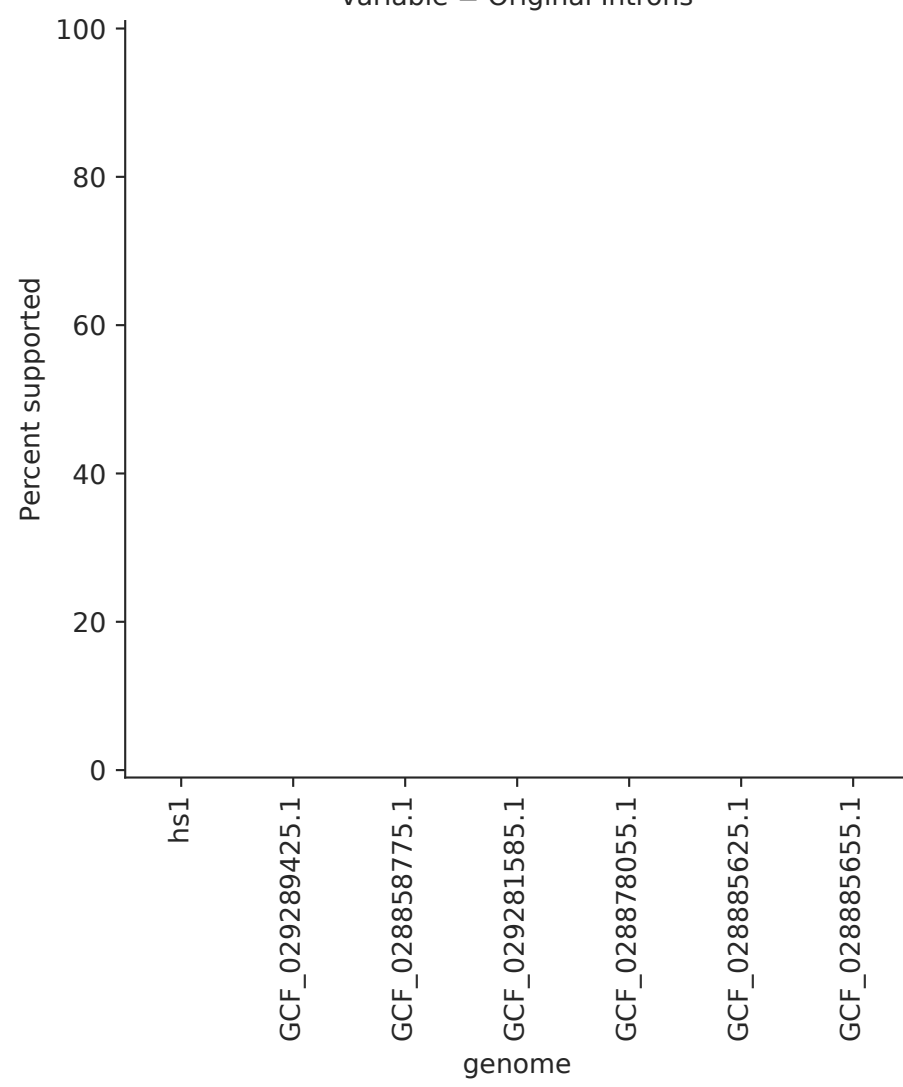
# Reference annotation support for unitary\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



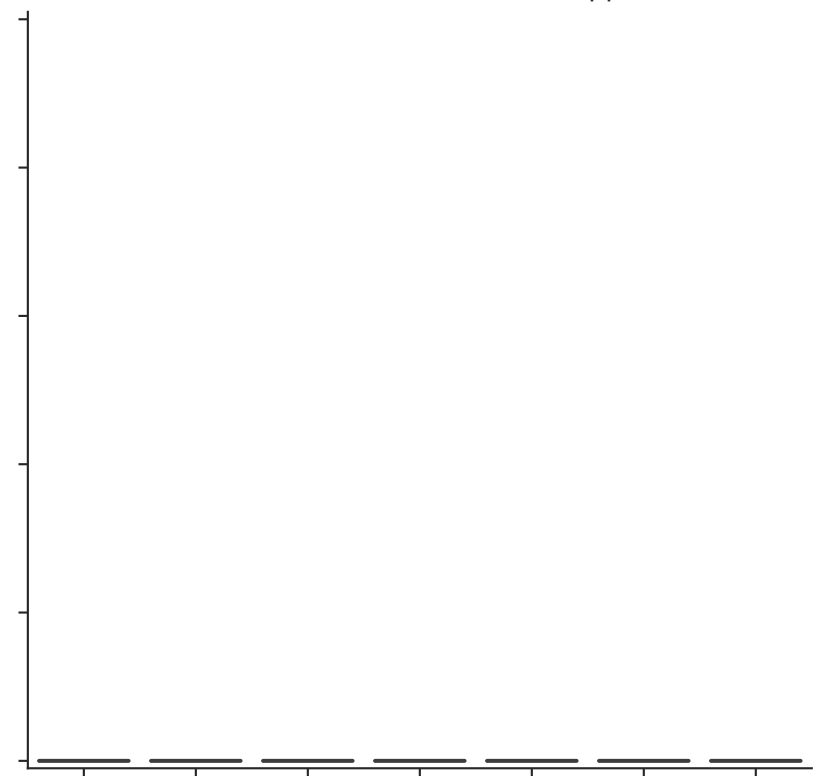
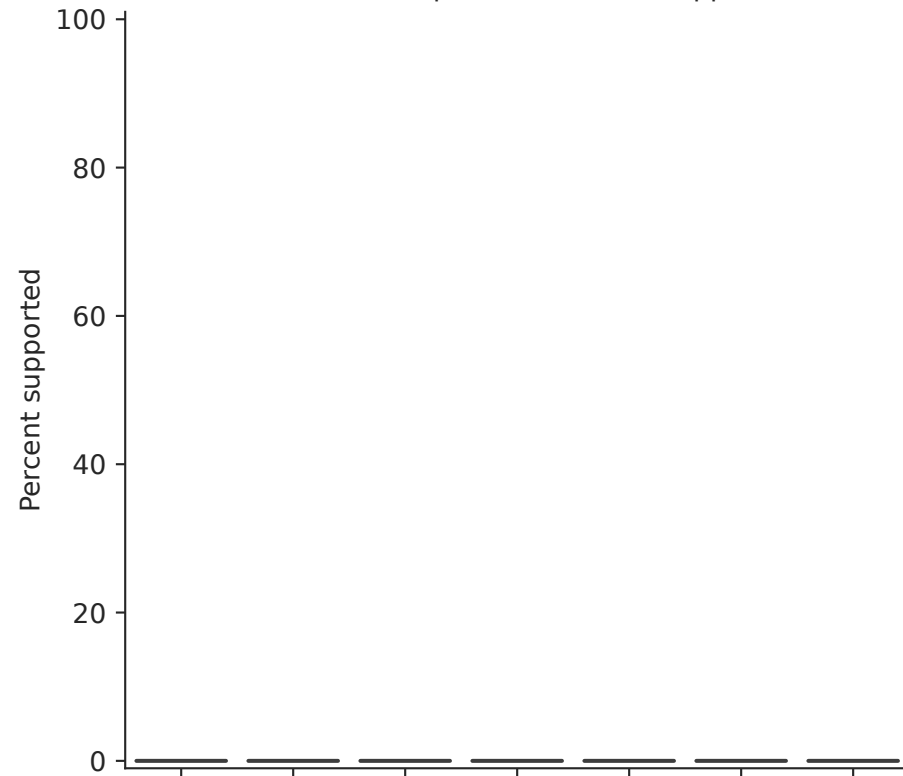
variable = Original Introns



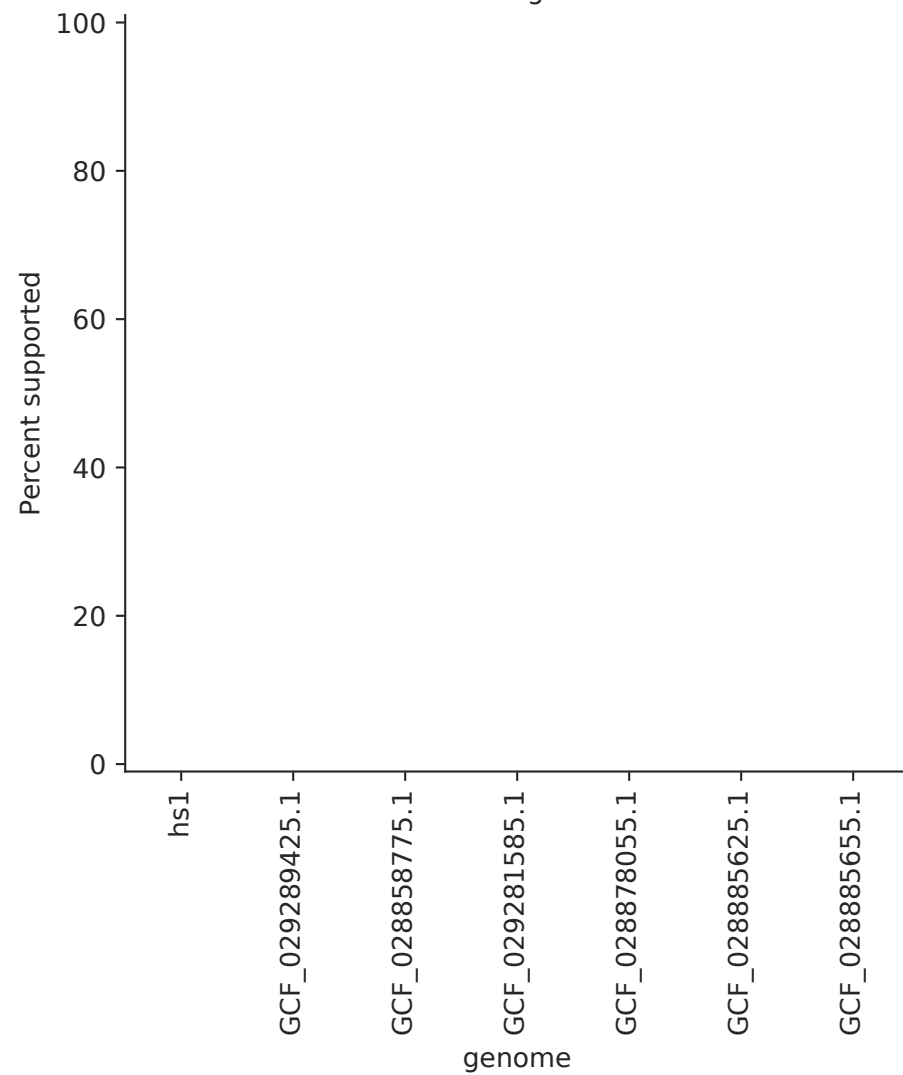
# Reference annotation support for unprocessed\_pseudogene

variable = Splice Annotation Support

variable = Exon Annotation Support



variable = Original Introns



hs1

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GCF\_028858775.1

GCF\_029281585.1

GCF\_028878055.1

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GCF\_028885655.1

genome

hs1

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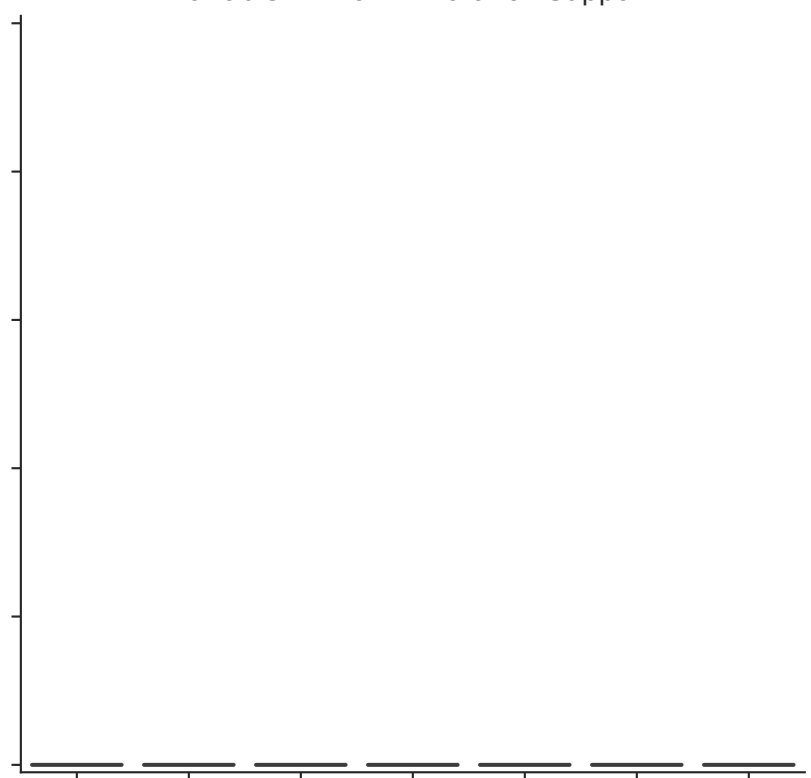
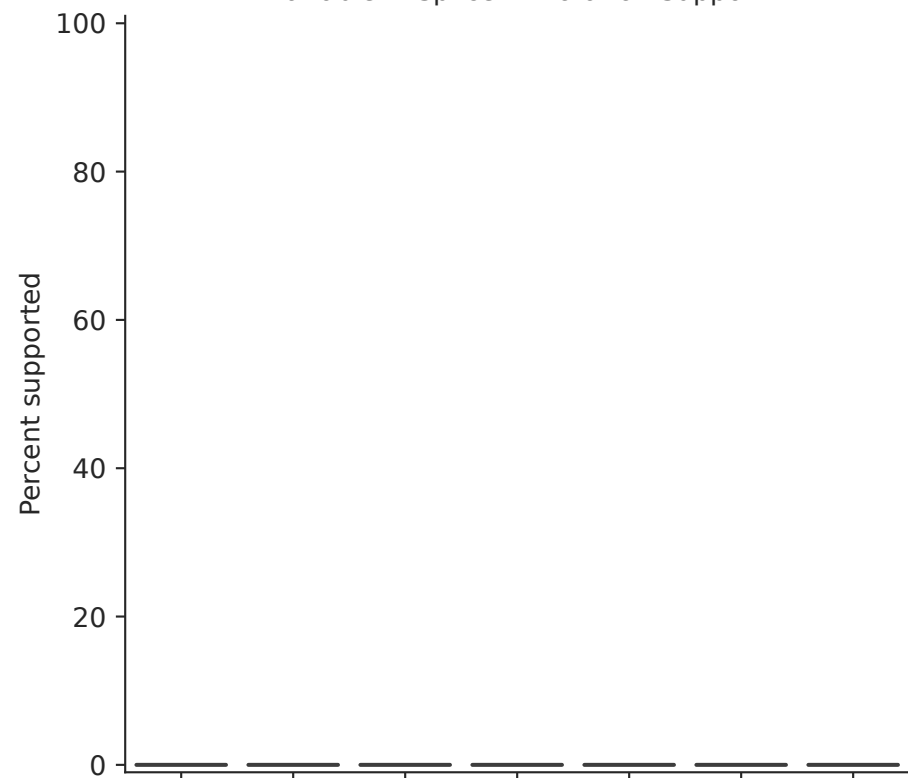
GCF\_028885655.1

genome

# Reference annotation support for vault\_RNA

variable = Splice Annotation Support

variable = Exon Annotation Support



variable = Original Introns

