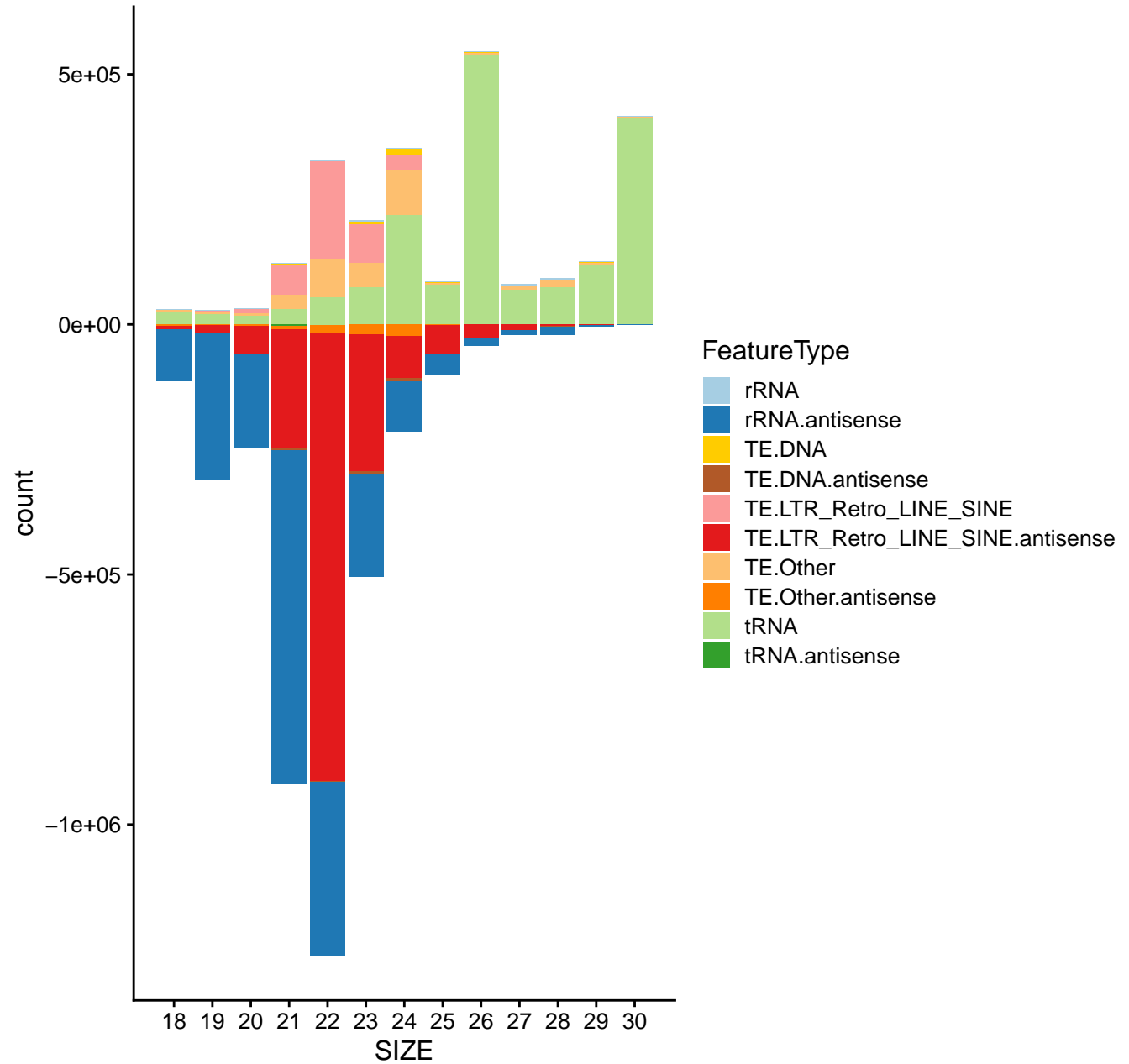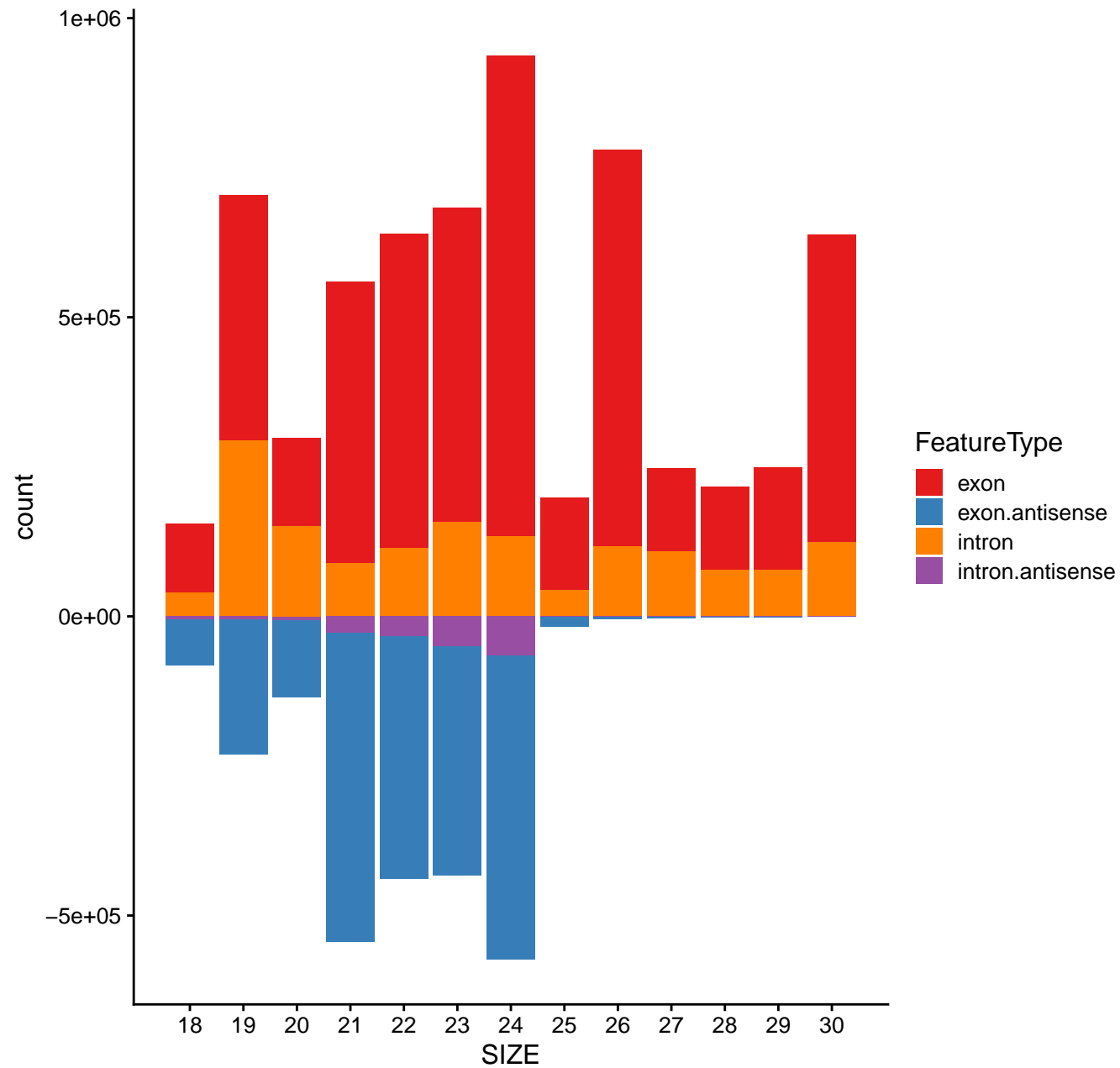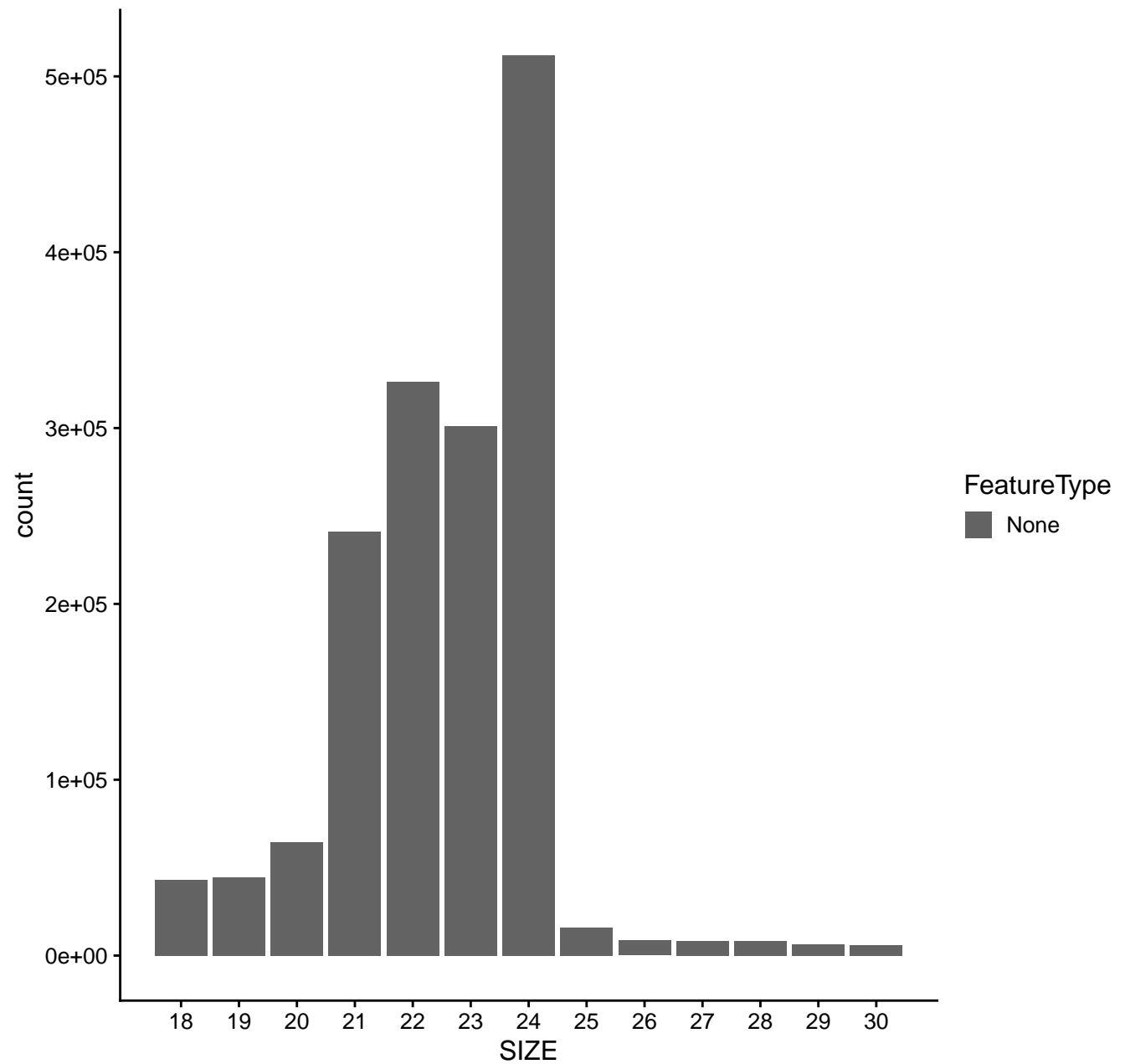**Combined all Libraries sRNA profile NonCoding Features for Uniqu...**
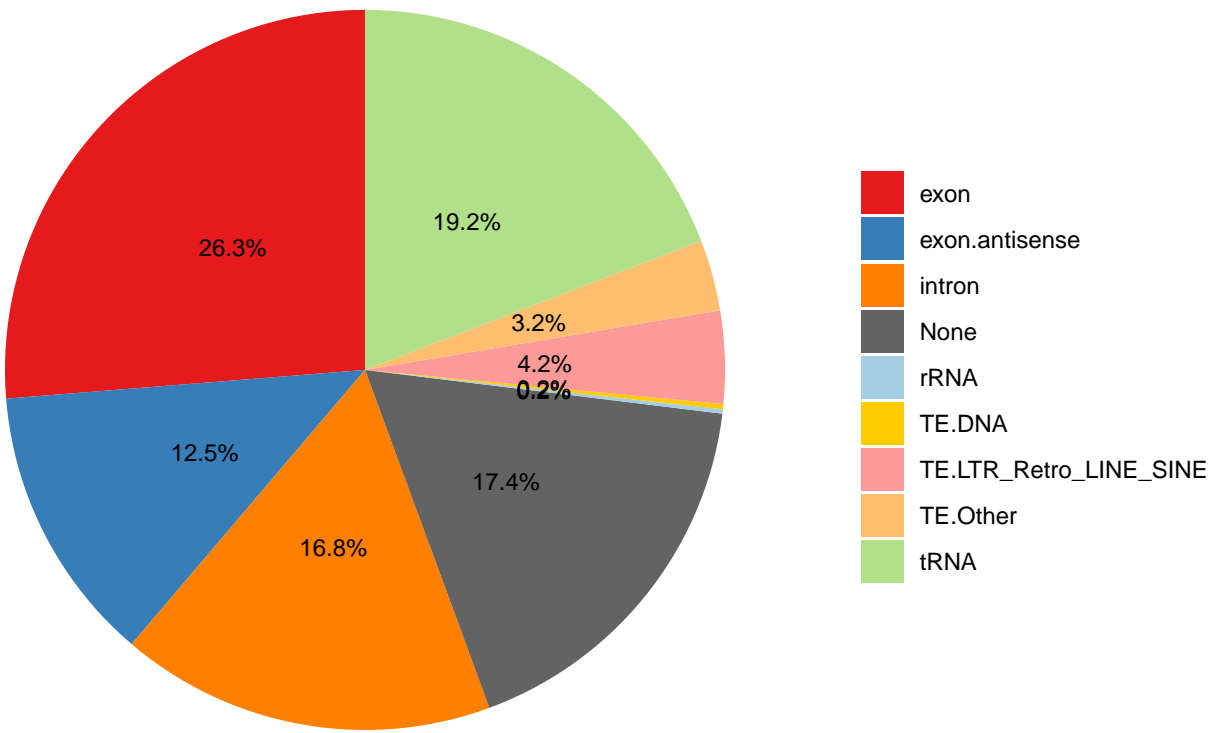
Combined all Libraries sRNA profile Exon,Intron Features for Uniqu

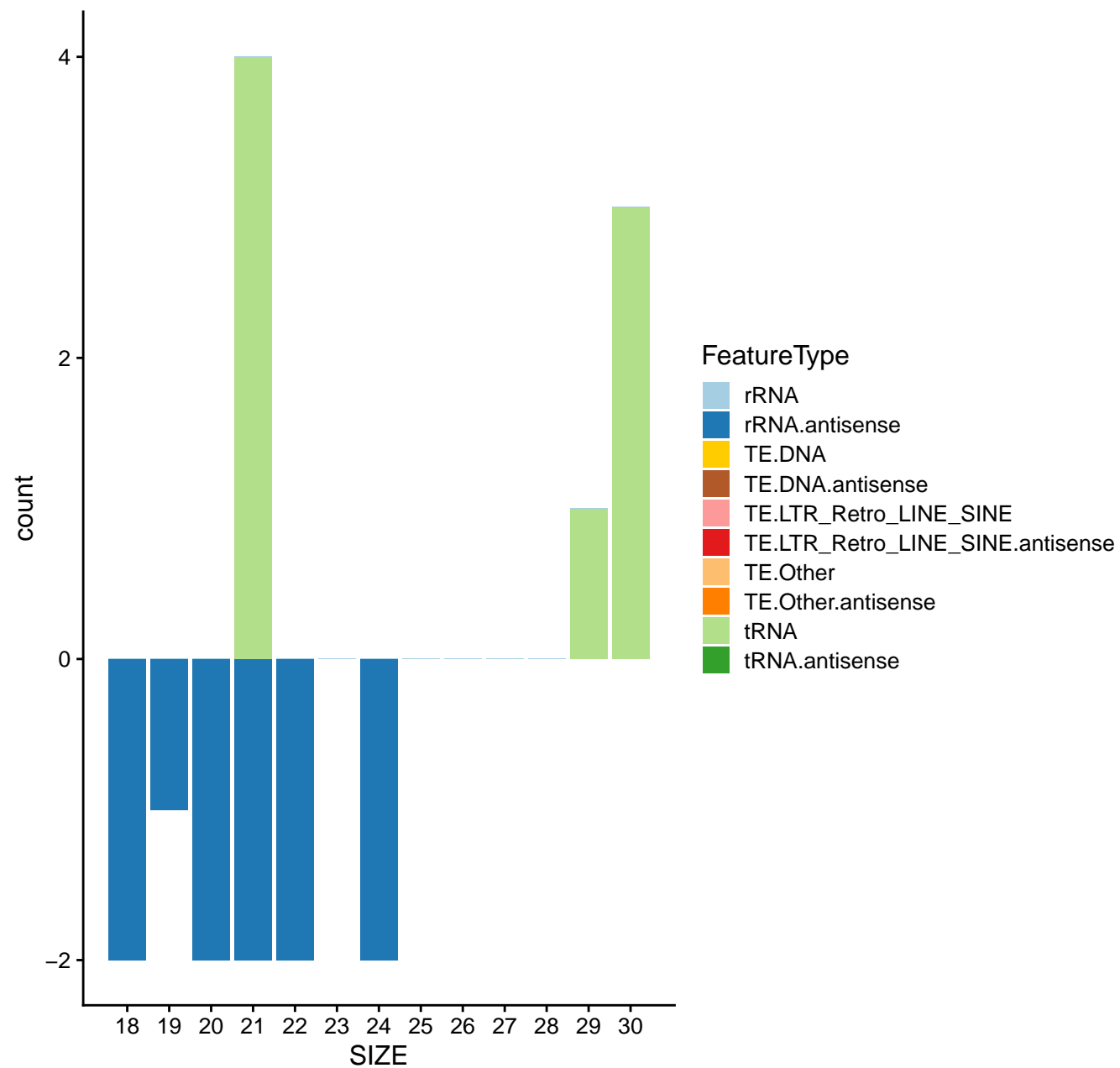# Combined all Libraries sRNA profile for Non–Genic location for Uni

Total for all Libraries sRNA profile for Unique Mapping

🟥	exon
🟦	exon.antisense
🟧	intron
⬛	None
🟦	rRNA
🟨	TE.DNA
🟥	TE.LTR_Retro_LINE_SINE
🟧	TE.Other
🟩	tRNA

Pie chart values: 26.3%, 12.5%, 16.8%, 17.4%, 0.2%, 4.2%, 3.2%, 19.2%

Botrytis Mycelium sRNA profile NonCoding Feature for Unique Mappin

Botrytis Mycelium Libraries sRNA profile Exon,Intron Features for Uniq

Botrytis Mycelium Libraries sRNA profile for Nongenic location for U

FeatureType
None
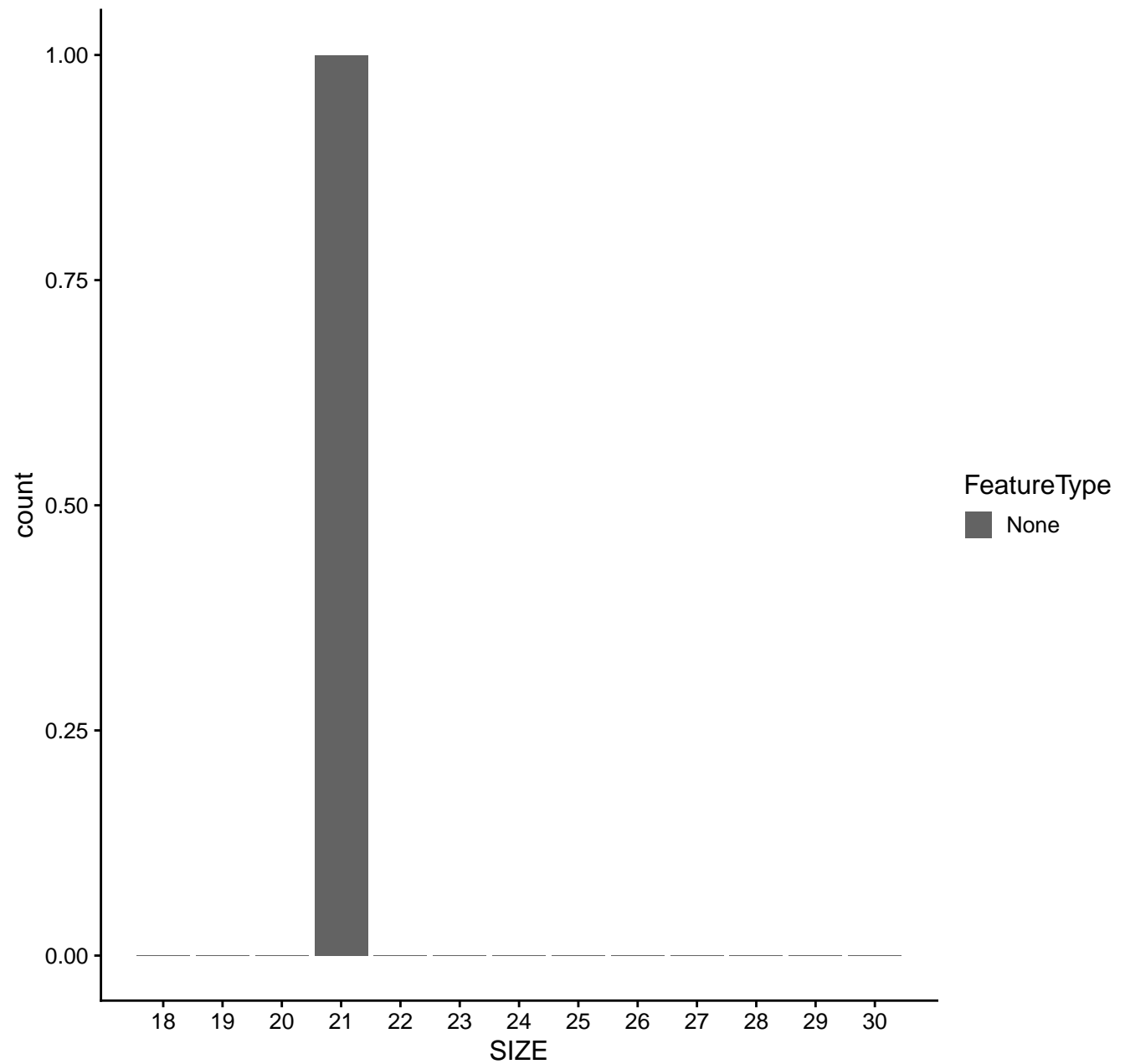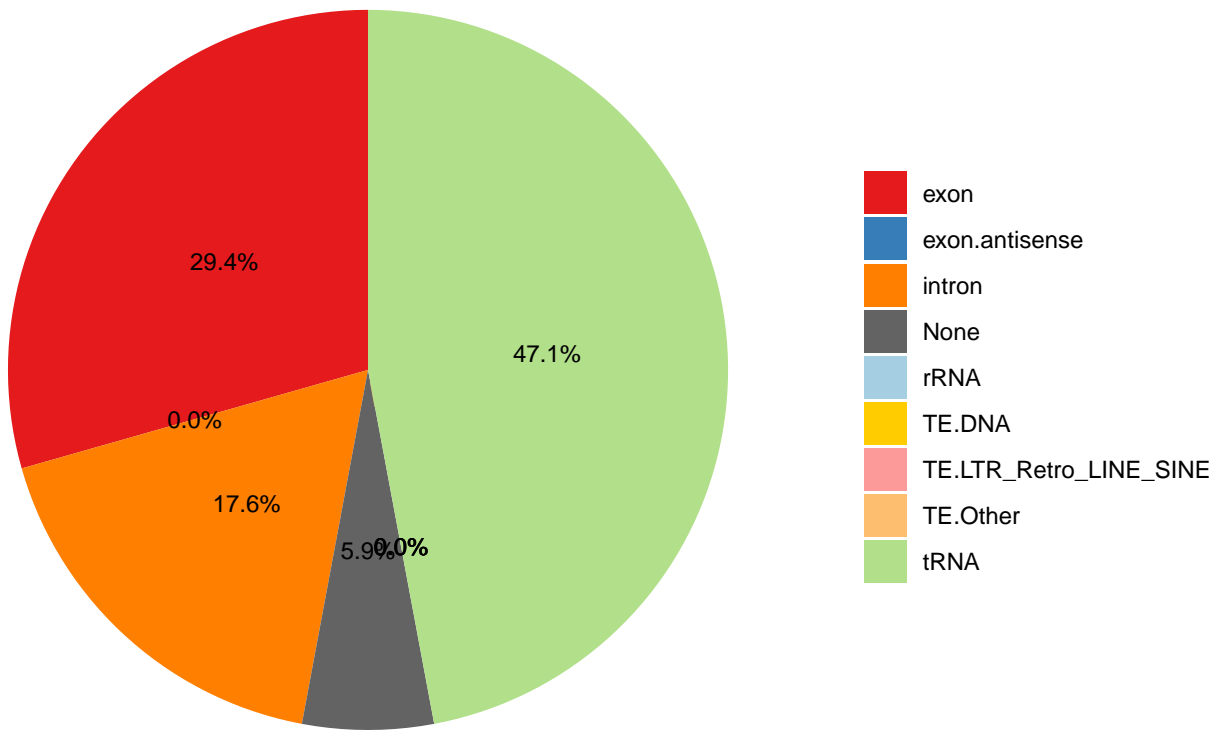
Botrytis Mycelium sRNA profile for Unique Mapping

Legend:
- exon
- exon.antisense
- intron
- None
- rRNA
- TE.DNA
- TE.LTR_Retro_LINE_SINE
- TE.Other
- tRNA

29.4%
0.0%
17.6%
5.9% 0.0%
47.1%

Tomato Infection sRNA profile NonCoding Feature for Unique Map
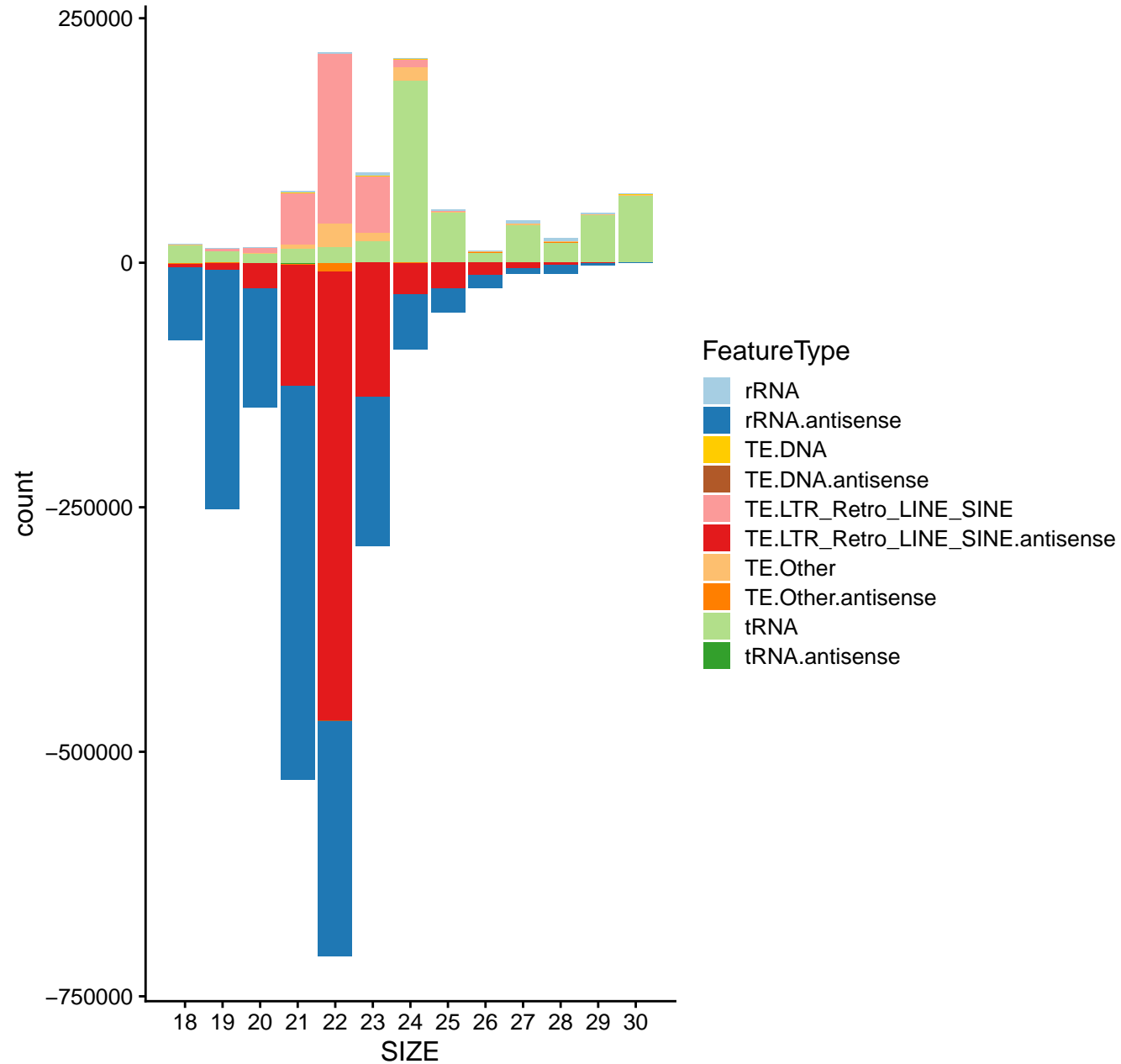
Tomato Infection Libraries sRNA profile Exon,Intron Features for Un
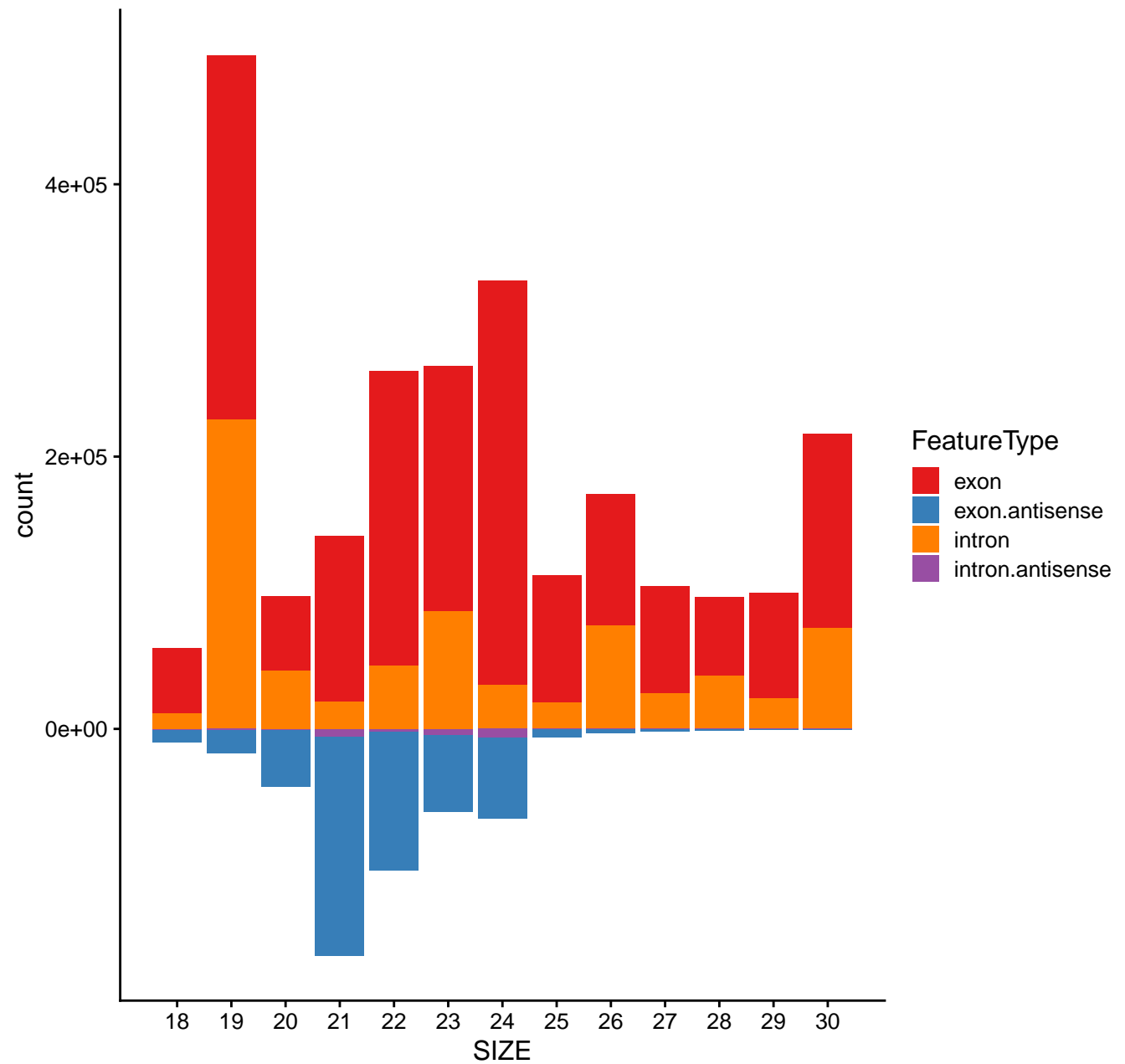
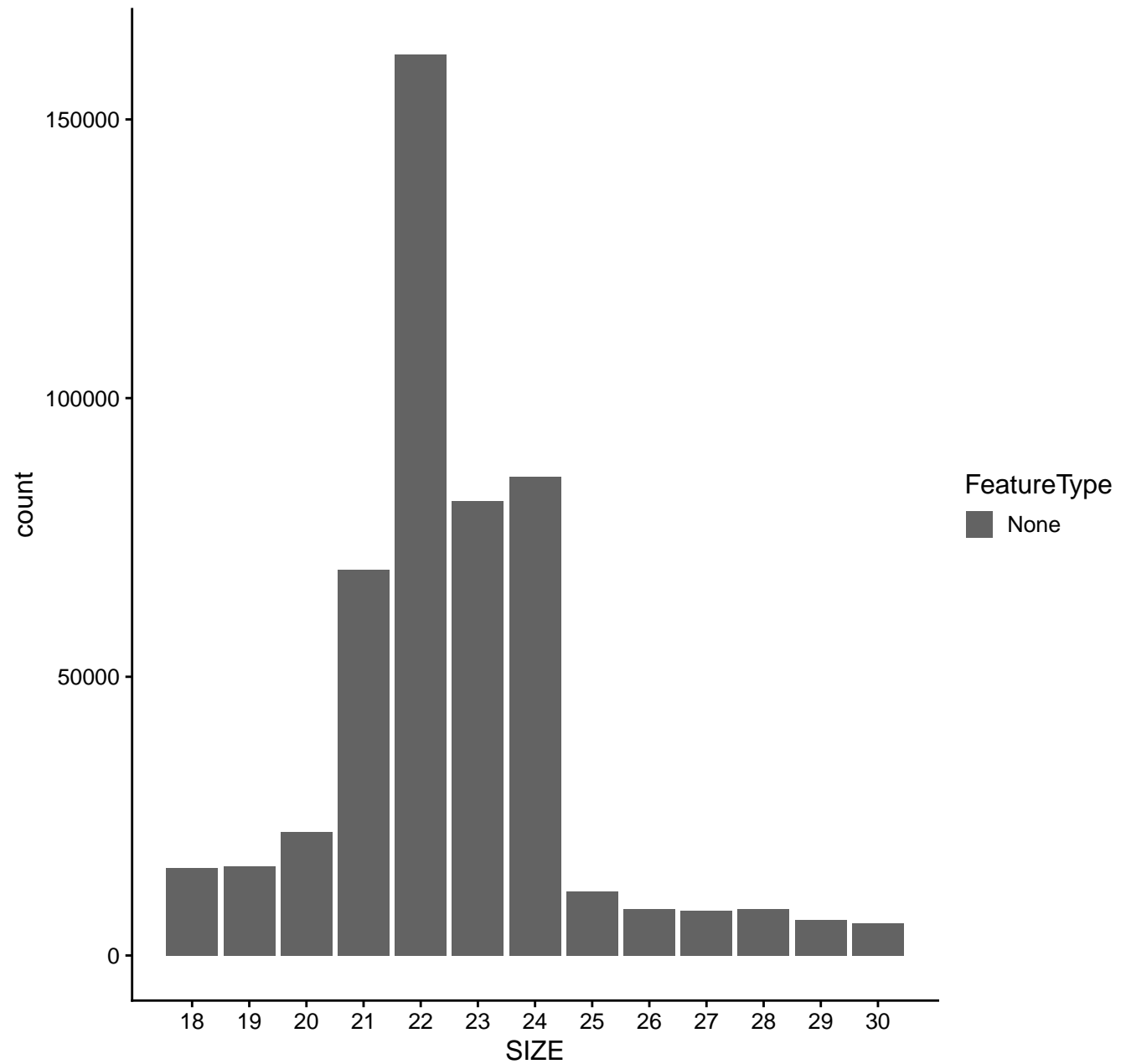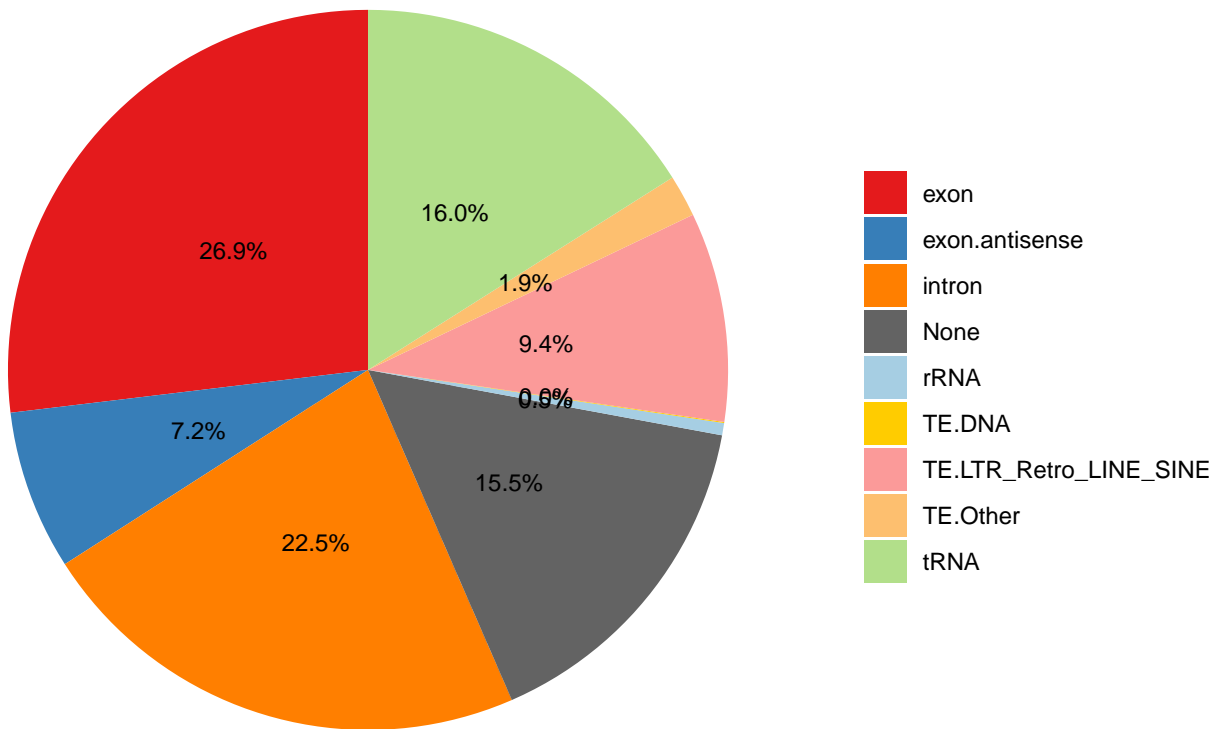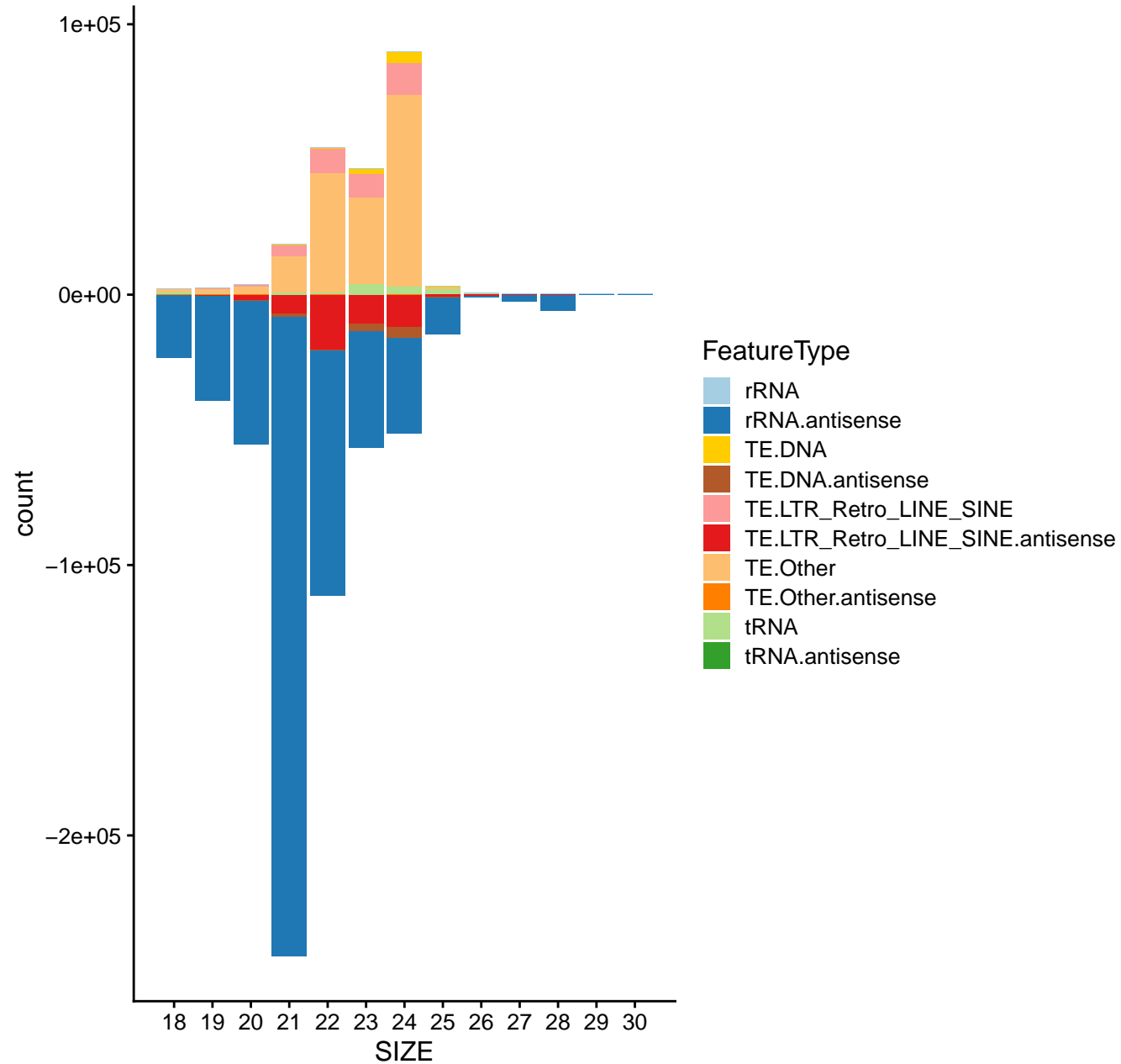**Tomato Infection Libraries sRNA profile for Nongenic location for U**

Botrytis Tomato Infection sRNA profile for Unique Mapping

Legend:
- exon
- exon.antisense
- intron
- None
- rRNA
- TE.DNA
- TE.LTR_Retro_LINE_SINE
- TE.Other
- tRNA

Percentages shown in pie chart: 26.9%, 7.2%, 22.5%, 15.5%, 0.0%, 0.6%, 9.4%, 1.9%, 16.0%

Tomato Mock Infection sRNA profile NonCoding Feature for Unique

FeatureType
- rRNA
- rRNA.antisense
- TE.DNA
- TE.DNA.antisense
- TE.LTR_Retro_LINE_SINE
- TE.LTR_Retro_LINE_SINE.antisense
- TE.Other
- TE.Other.antisense
- tRNA
- tRNA.antisense

Tomato Mock Infection Libraries sRNA profile Exon,Intron Features

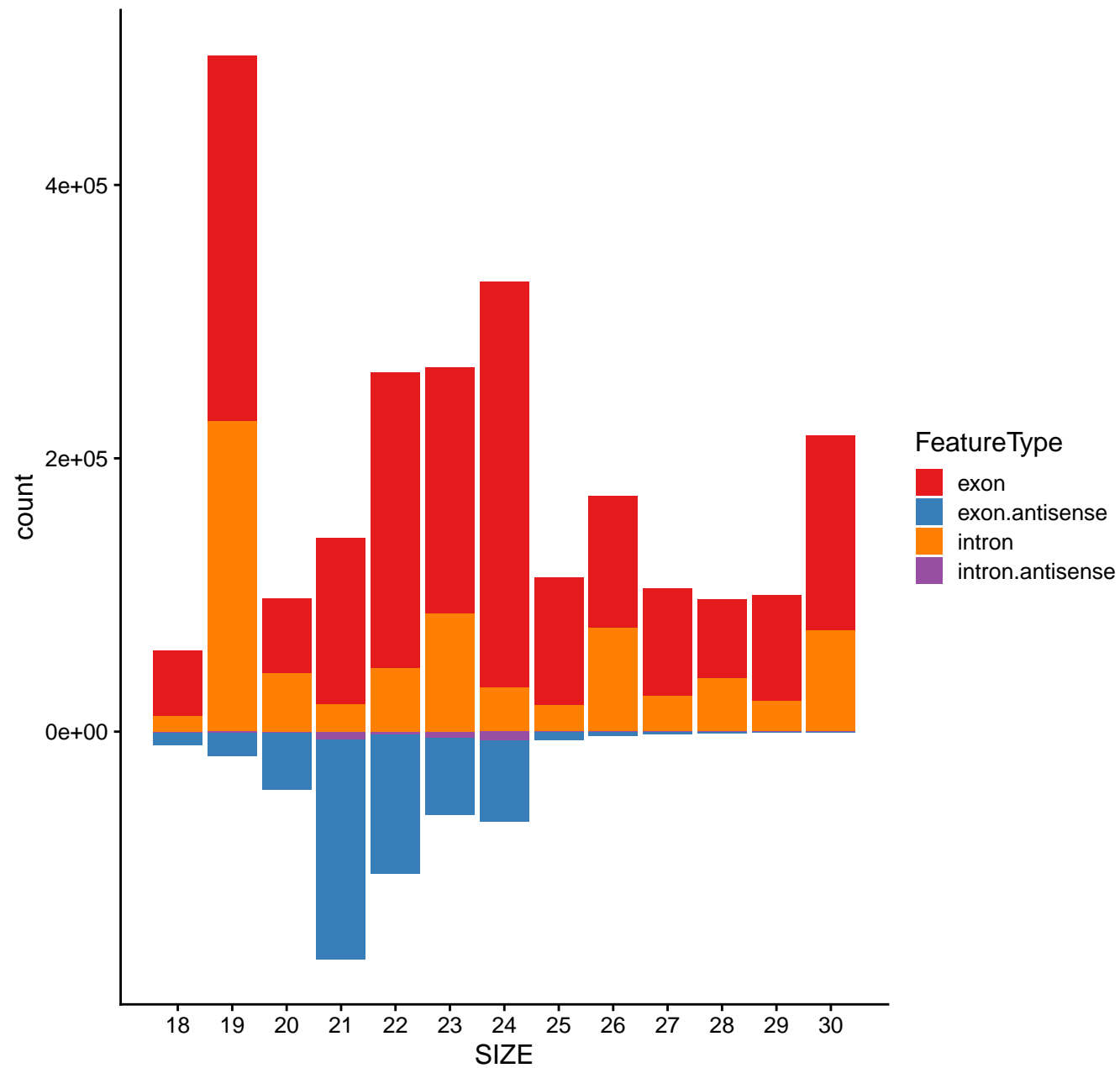Tomato Mock Infection Libraries sRNA profile for Nongenic location
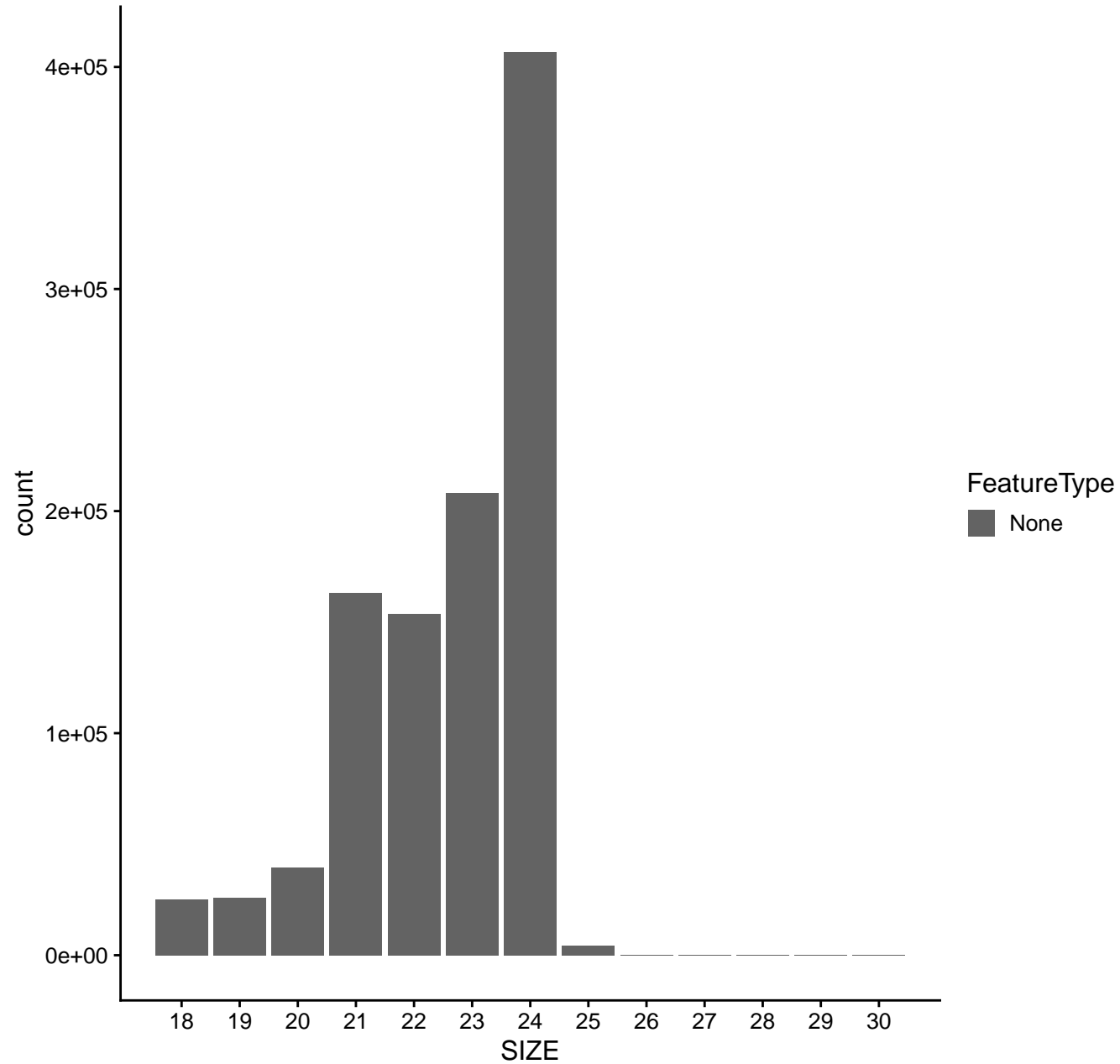
Mock Tomato Infection sRNA profile for Unique Mapping

■	exon
■	exon.antisense
■	intron
■	None
■	rRNA
■	TE.DNA
■	TE.LTR_Retro_LINE_SINE
■	TE.Other
■	tRNA

Percentages shown: 16.2%, 18.5%, 15.7%, 28.6%, 0.3%, 6.3%, 4.2%, 9.9%