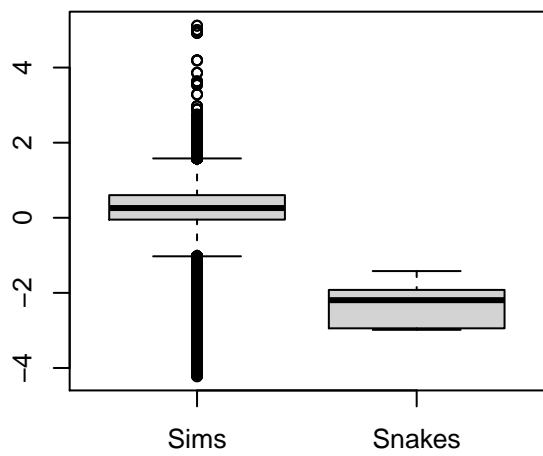
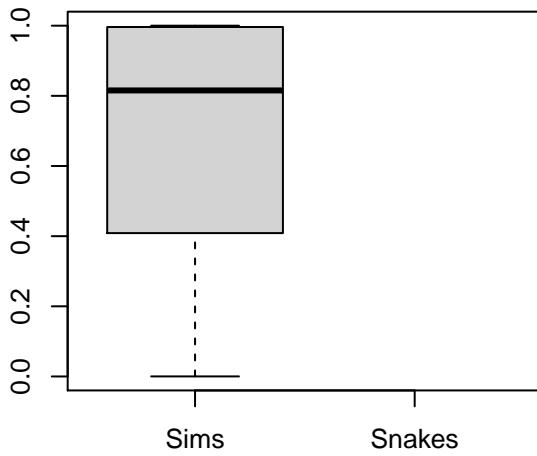


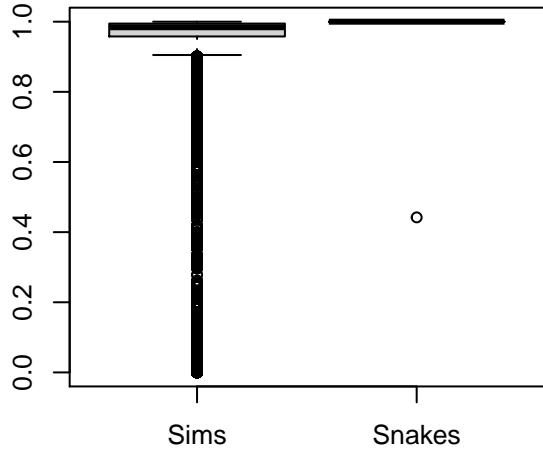
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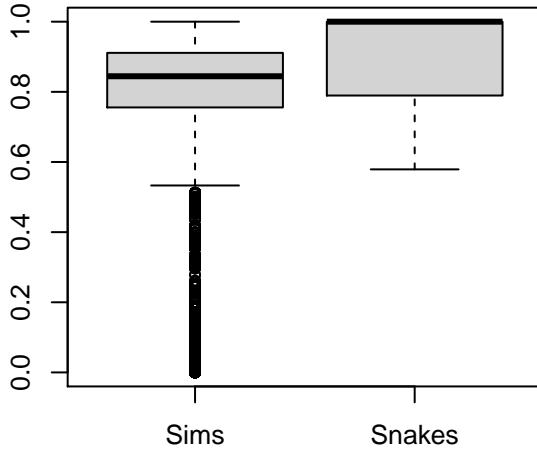
GENOME.HAPLOTYPE.DIVERSITY



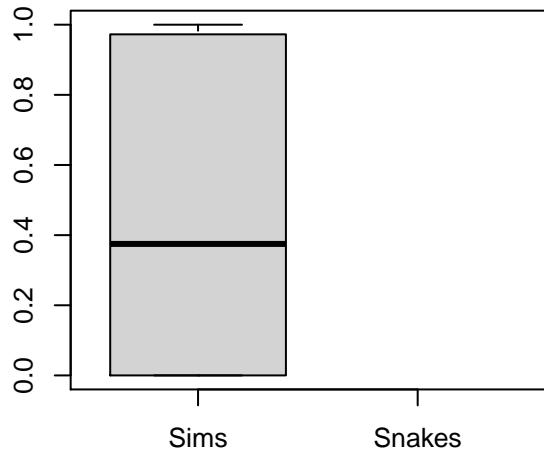
HAP.DIVERSITY.WITHIN



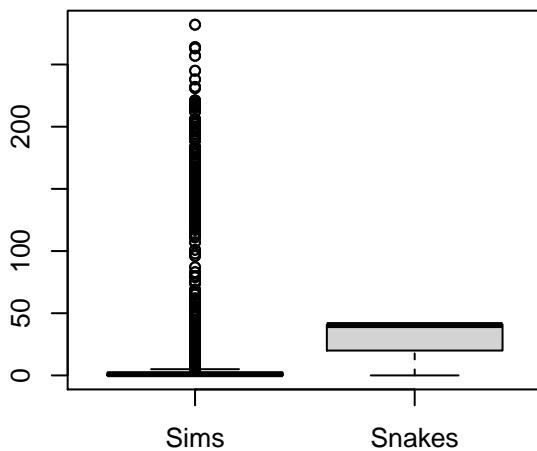
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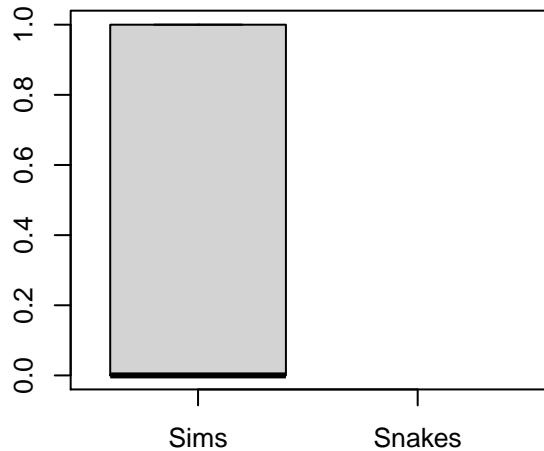
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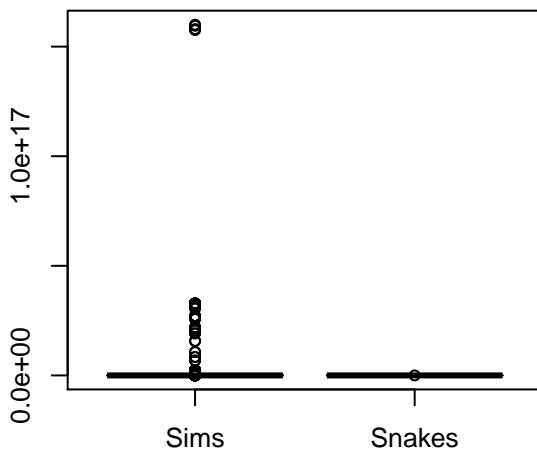
HUDSON.KAPLAN.RM



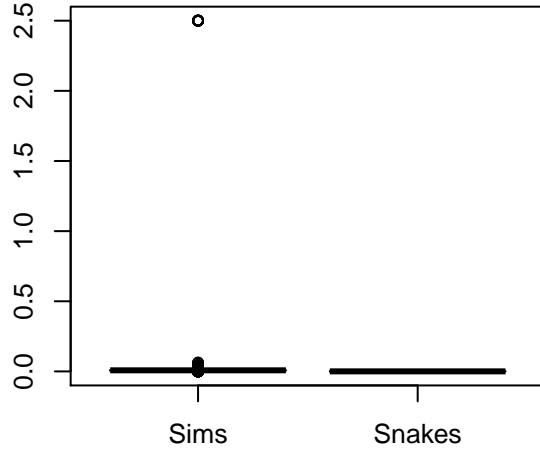
IBD



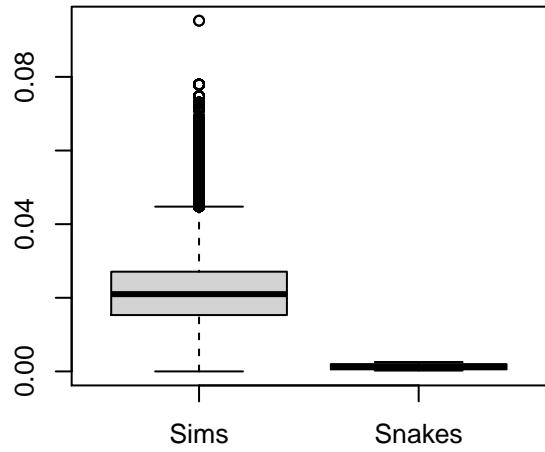
KURT_HAPLOTYPE.COUNTS



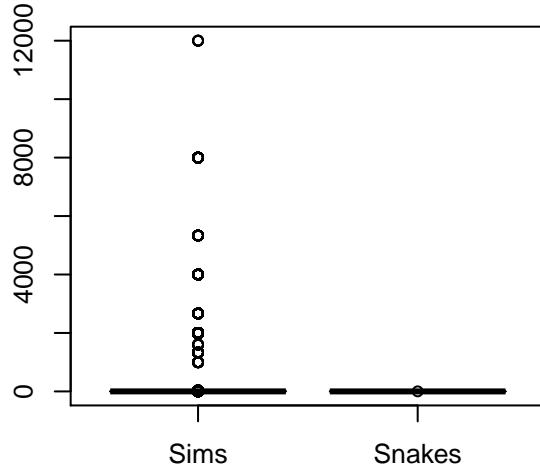
KURT_MINOR_ALLELE.FREQS



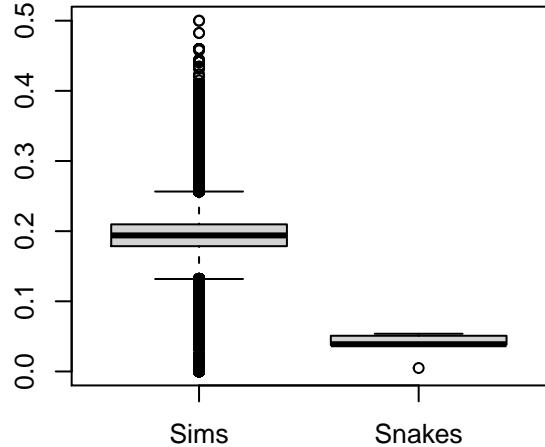
KURT_NUC.DIVERSITY.WITHIN



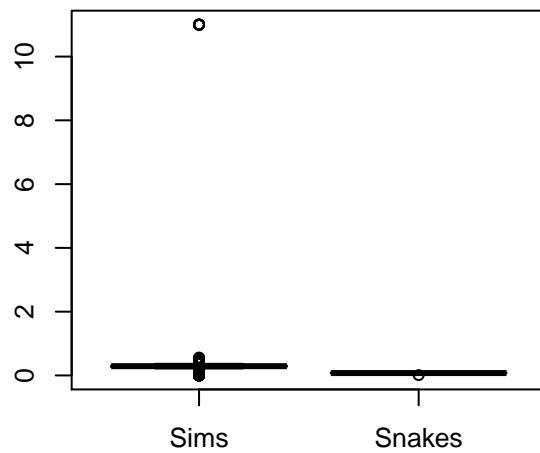
MEAN_HAPLOTYPE.COUNTS



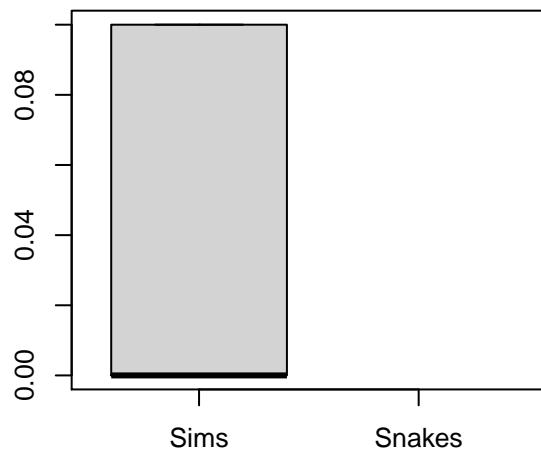
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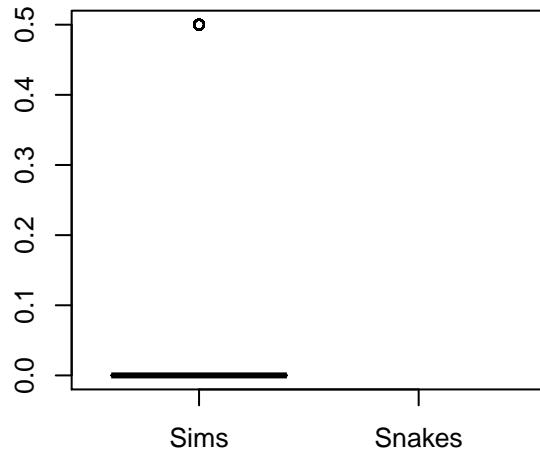
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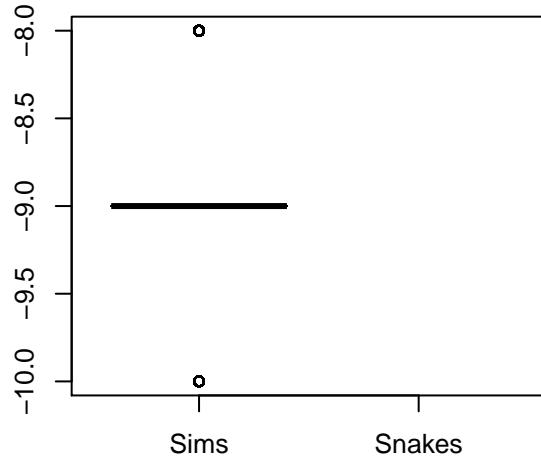
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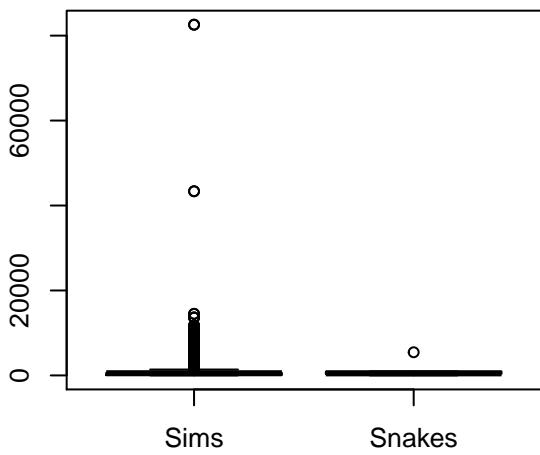
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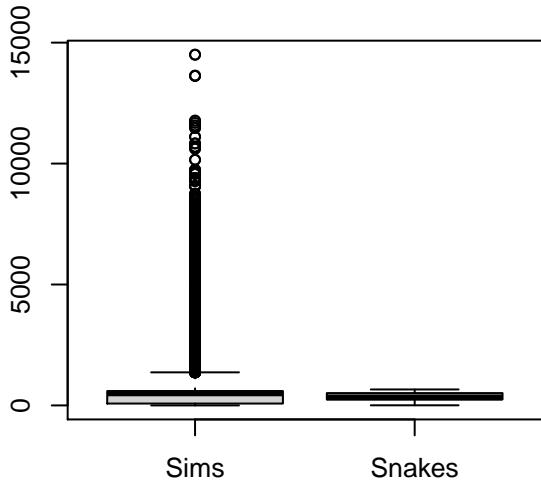
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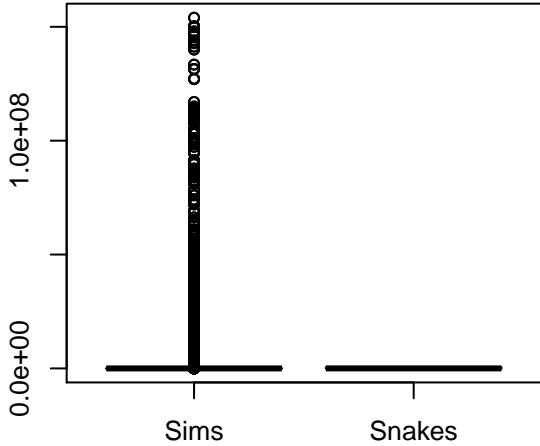
N.BIALLELIC.SITES



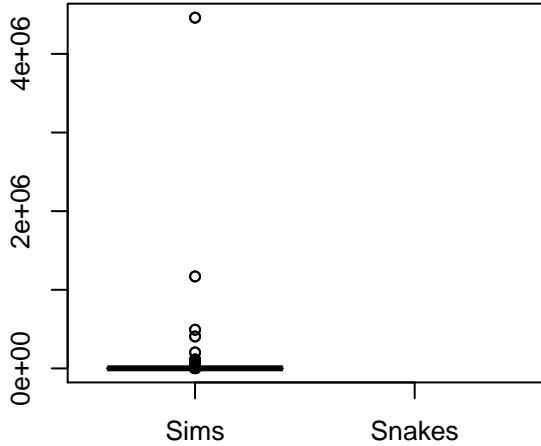
N.SEGREGATING.SITES



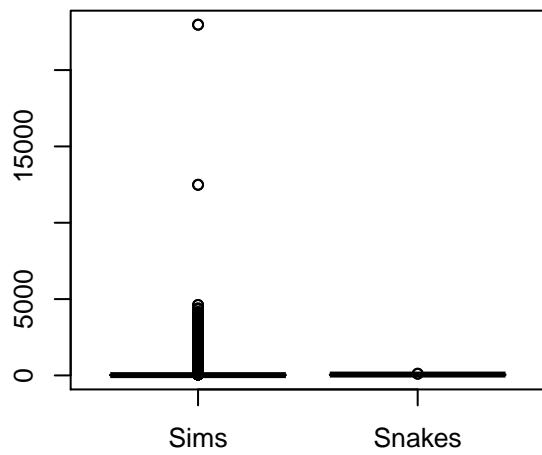
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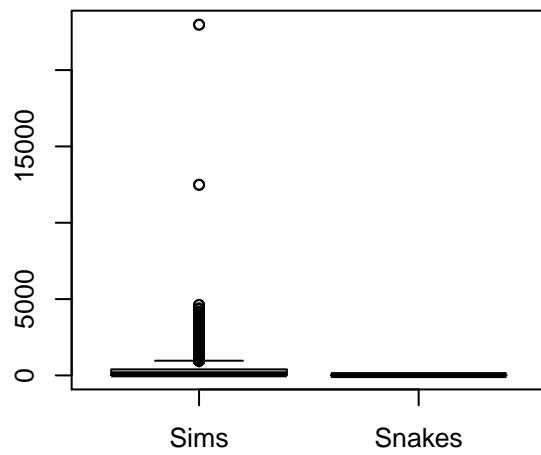
N.SITES.CORRECTED



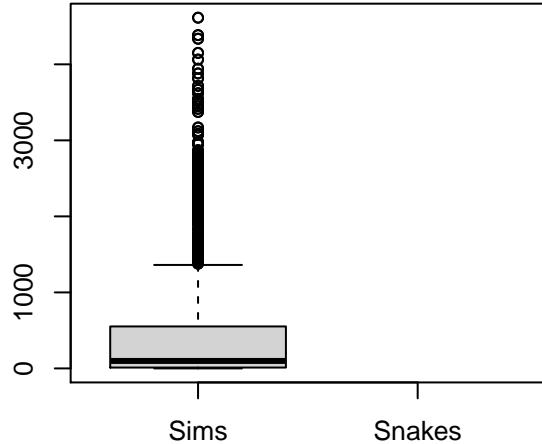
NUC.DIVERSITY.WITHIN



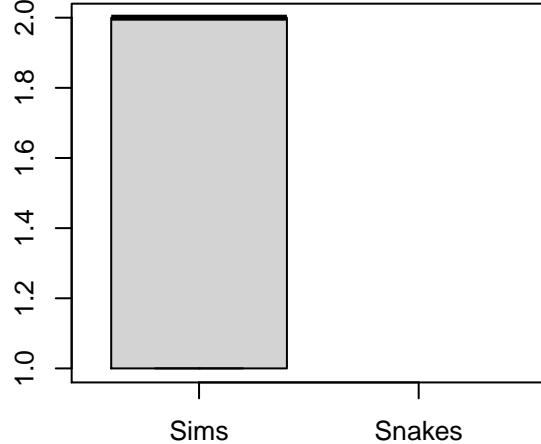
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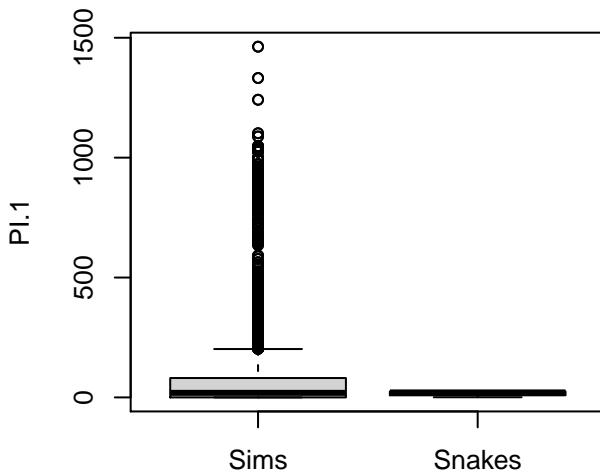
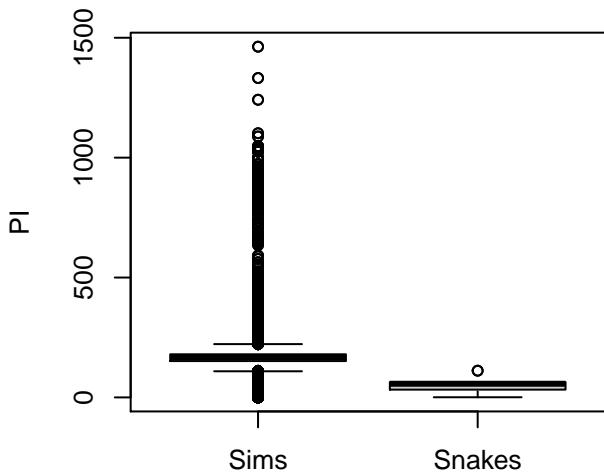
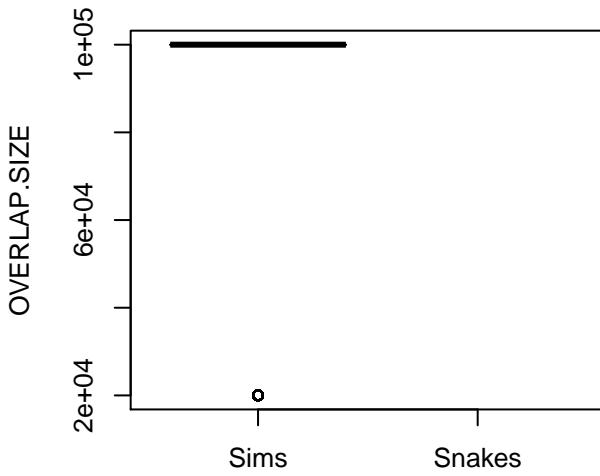


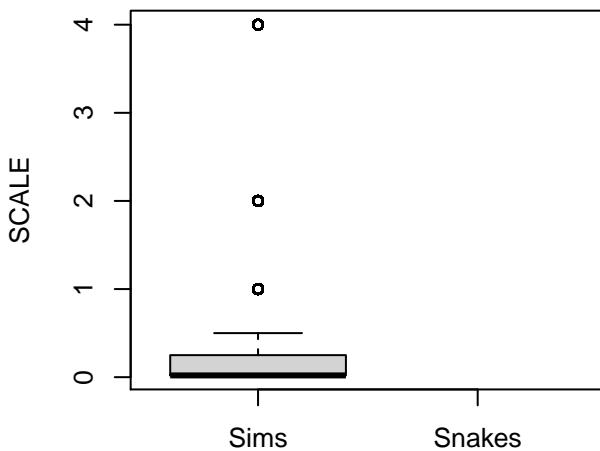
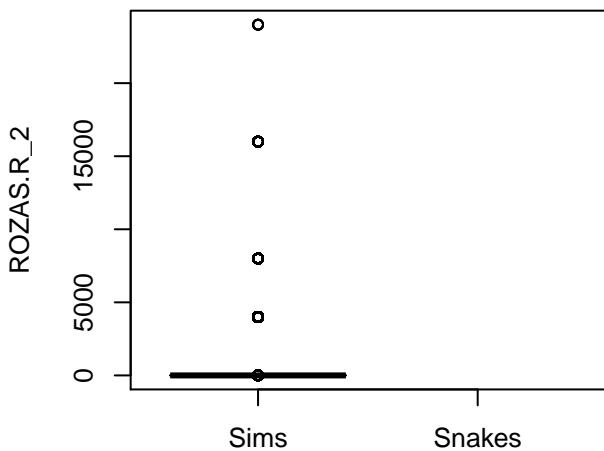
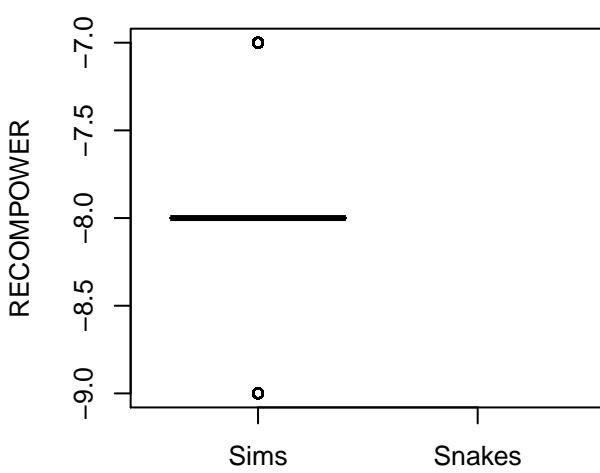
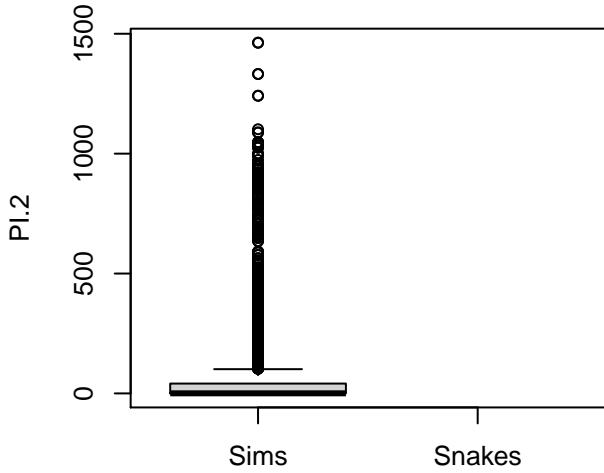
NUC.DIVERSITY.WITHIN.2



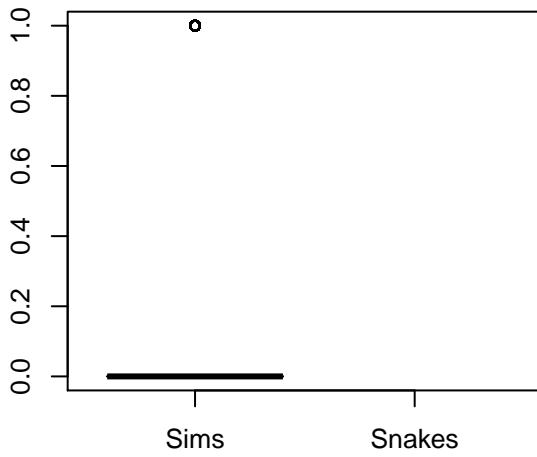
NUM.POP



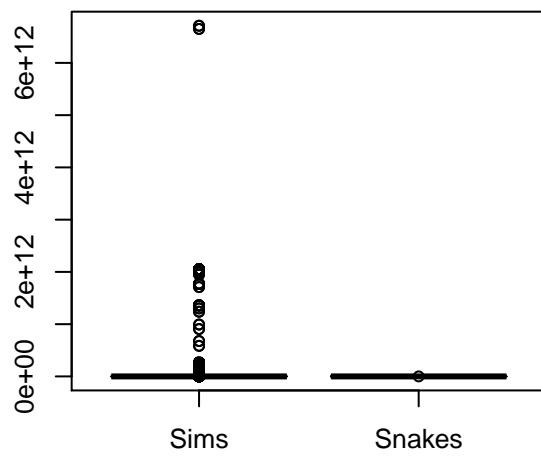




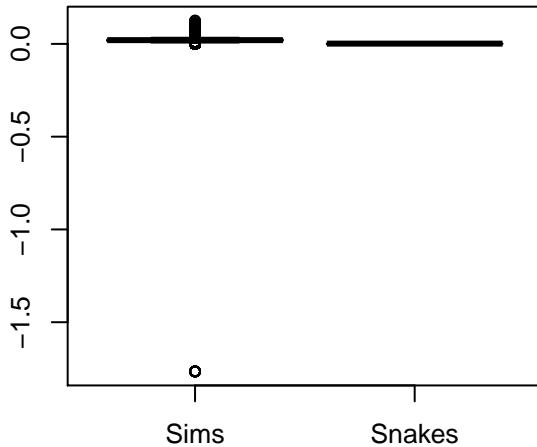
SECCON



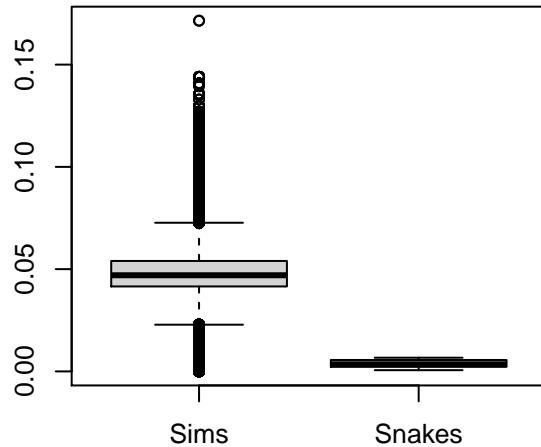
SKEW_HAPLOTYPE.COUNTS

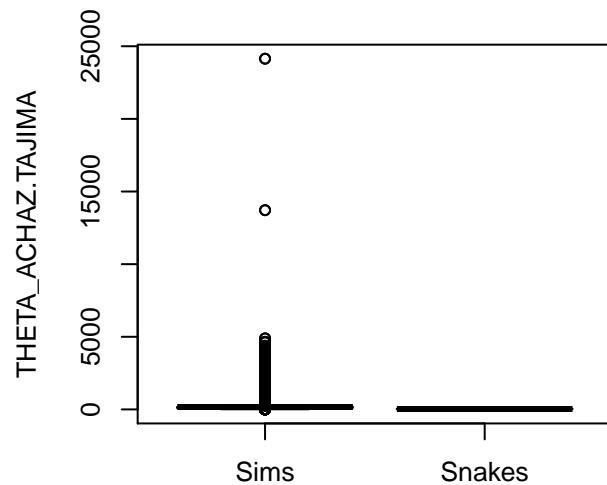
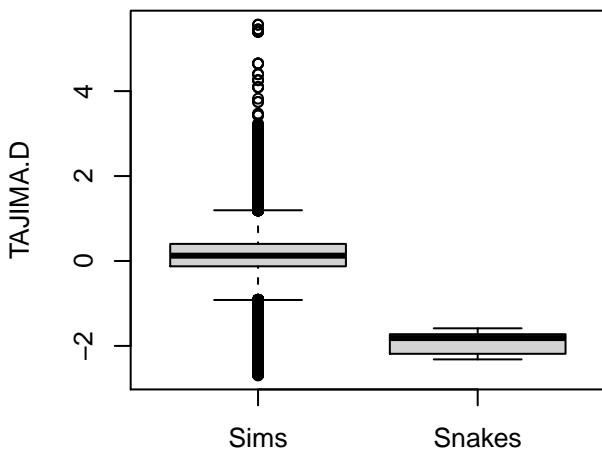
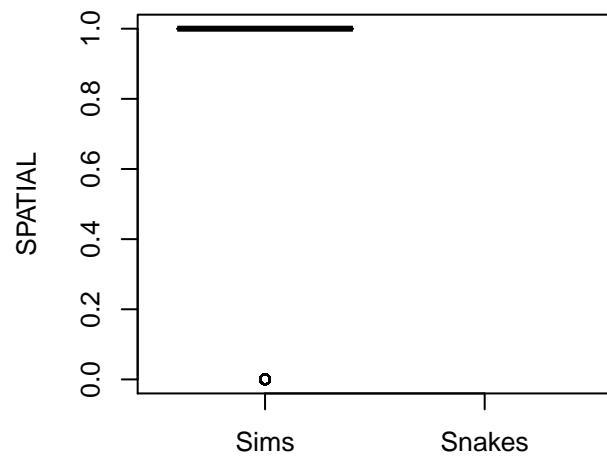


SKEW_MINOR.ALLELE.FREQS

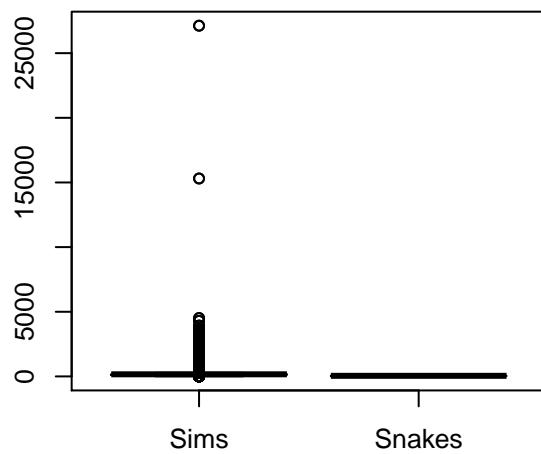


SKEW_NUC.DIVERSITY.WITHIN

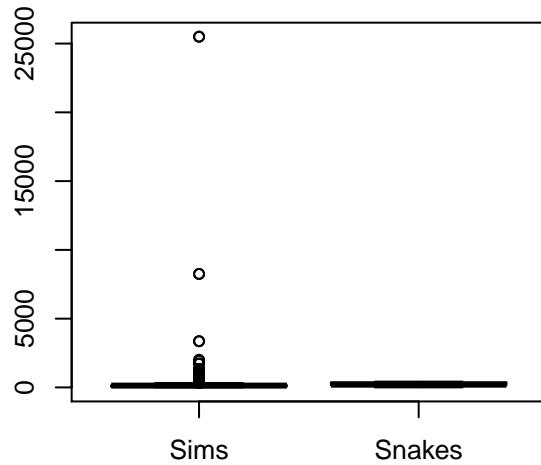




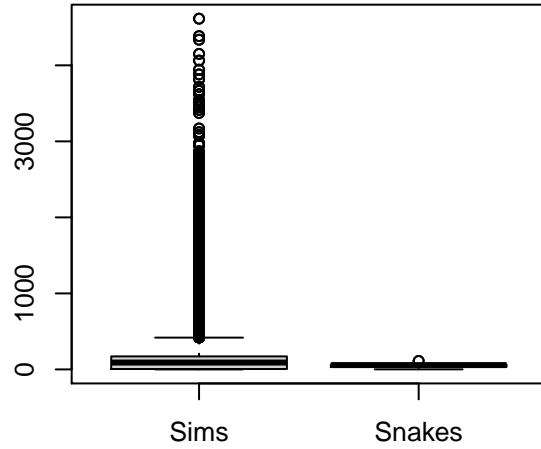
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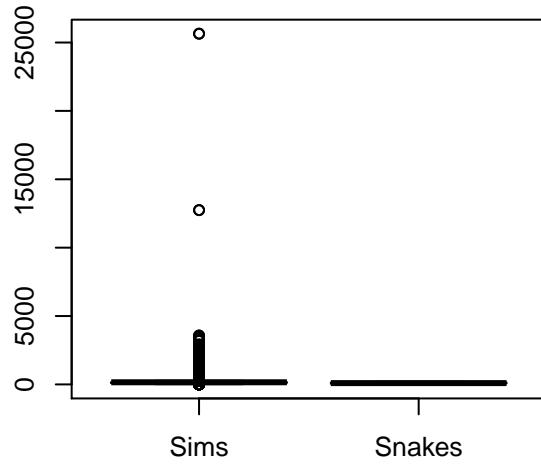
THETA_FU.LI



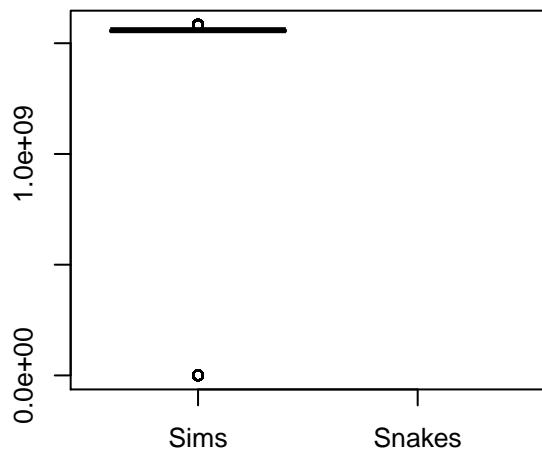
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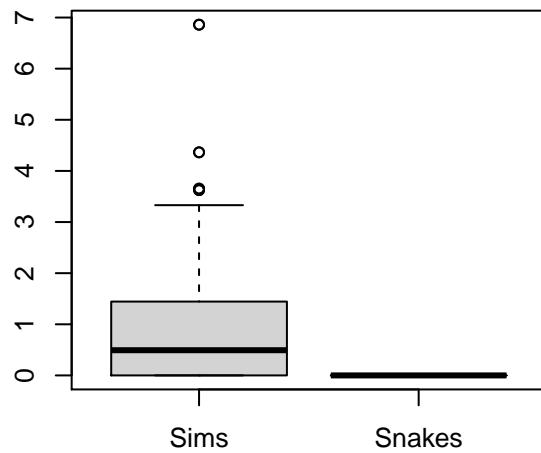
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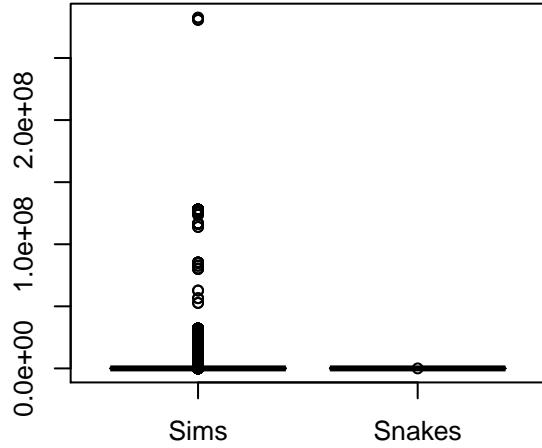
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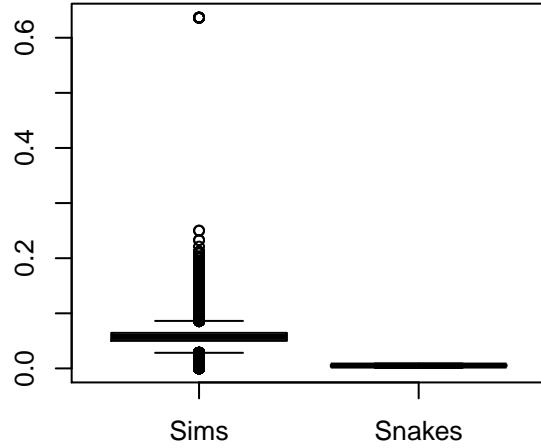
TRANS.TRANSV.RATIO



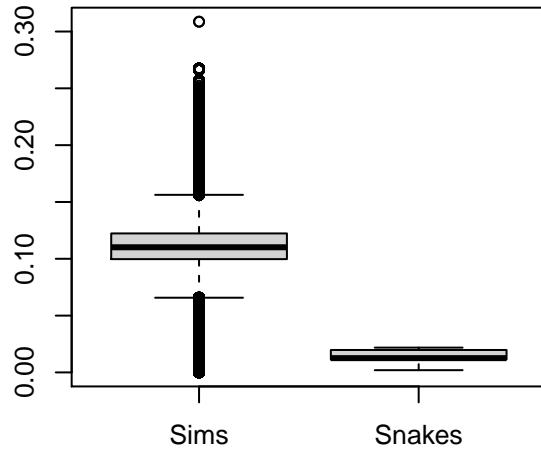
VAR_HAPLOTYPE.COUNTS



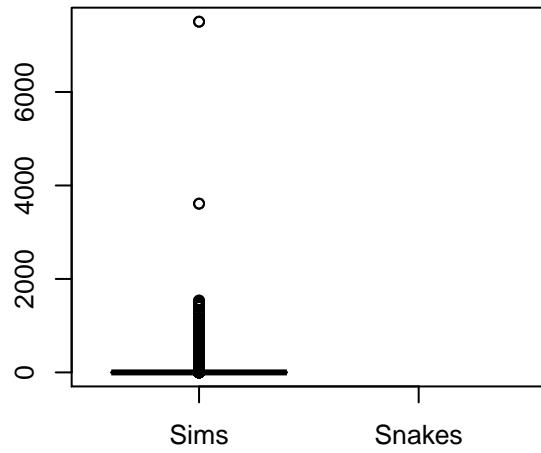
VAR_MINOR.ALLELE.FREQS



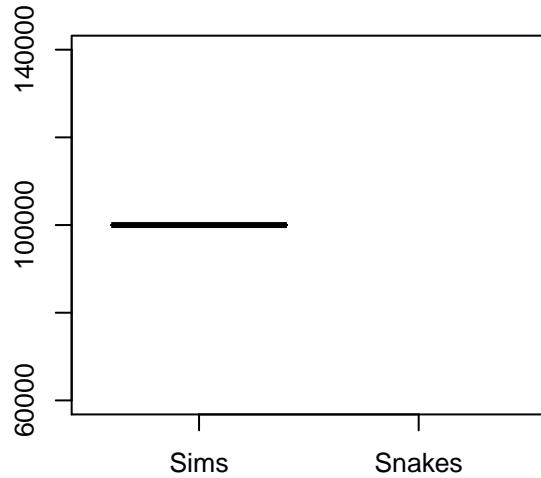
VAR_NUC.DIVERSITY.WITHIN



WINDOW



WINDOW.SIZE



YEAR

