

第一次作业

(1)

第四列为所示特征序列在目标序列/染色体中的起始位点，第五列为该特征序列的终止位点。如果该序列是一个exon，则其长度应该为第五列减去第四列加上1。

(2)

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XI      ensembl CDS      82947  82998  .      +      0      gene_id "YKL190W";
gene_version "1"; transcript_id "YKL190W"; transcript_version "1"; exon_number
"1"; gene_name "CNB1"; gene_source "ensembl"; gene_biotype "protein_coding";
transcript_name "CNB1"; transcript_source "ensembl"; transcript_biotype
"protein_coding"; protein_id "YKL190W"; protein_version "1";
XI      ensembl CDS      83075  83547  .      +      1      gene_id "YKL190W";
gene_version "1"; transcript_id "YKL190W"; transcript_version "1"; exon_number
"2"; gene_name "CNB1"; gene_source "ensembl"; gene_biotype "protein_coding";
transcript_name "CNB1"; transcript_source "ensembl"; transcript_biotype
"protein_coding"; protein_id "YKL190W"; protein_version "1";
XI      ensembl CDS      84704  85900  .      +      0      gene_id "YKL189W";
gene_version "1"; transcript_id "YKL189W"; transcript_version "1"; exon_number
"1"; gene_name "HYM1"; gene_source "ensembl"; gene_biotype "protein_coding";
transcript_name "HYM1"; transcript_source "ensembl"; transcript_biotype
"protein_coding"; protein_id "YKL189W"; protein_version "1";
XI      ensembl CDS      86228  88786  .      -      0      gene_id "YKL188C";
gene_version "1"; transcript_id "YKL188C"; transcript_version "1"; exon_number
"1"; gene_name "PXA2"; gene_source "ensembl"; gene_biotype "protein_coding";
transcript_name "PXA2"; transcript_source "ensembl"; transcript_biotype
"protein_coding"; protein_id "YKL188C"; protein_version "1";
XI      ensembl CDS      89287  91536  .      -      0      gene_id "YKL187C";
gene_version "1"; transcript_id "YKL187C"; transcript_version "1"; exon_number
"1"; gene_source "ensembl"; gene_biotype "protein_coding"; transcript_name
"YKL187C"; transcript_source "ensembl"; transcript_biotype "protein_coding";
protein_id "YKL187C"; protein_version "1";
XI      ensembl CDS      92747  93298  .      -      0      gene_id "YKL186C";
gene_version "1"; transcript_id "YKL186C"; transcript_version "1"; exon_number
"1"; gene_name "MTR2"; gene_source "ensembl"; gene_biotype "protein_coding";
transcript_name "MTR2"; transcript_source "ensembl"; transcript_biotype
"protein_coding"; protein_id "YKL186C"; protein_version "1";
XI      ensembl CDS      94499  96262  .      +      0      gene_id "YKL185W";
gene_version "1"; transcript_id "YKL185W"; transcript_version "1"; exon_number
"1"; gene_name "ASH1"; gene_source "ensembl"; gene_biotype "protein_coding";
transcript_name "ASH1"; transcript_source "ensembl"; transcript_biotype
"protein_coding"; protein_id "YKL185W"; protein_version "1";
XI      ensembl CDS      96757  98154  .      +      0      gene_id "YKL184W";
gene_version "1"; transcript_id "YKL184W"; transcript_version "1"; exon_number
"1"; gene_name "SPE1"; gene_source "ensembl"; gene_biotype "protein_coding";
transcript_name "SPE1"; transcript_source "ensembl"; transcript_biotype
"protein_coding"; protein_id "YKL184W"; protein_version "1";
XI      ensembl CDS      98398  98607  .      -      0      gene_id "YKL183C-
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A"; gene_version "1"; transcript_id "YKL183C-A"; transcript_version "1";
exon_number "1"; gene_source "ensembl"; gene_biotype "protein_coding";
transcript_name "YKL183C-A"; transcript_source "ensembl"; transcript_biotype
"protein_coding"; protein_id "YKL183C-A"; protein_version "1";
XI      ensembl CDS      98721  99638  .      +      0      gene_id "YKL183W";
gene_version "1"; transcript_id "YKL183W"; transcript_version "1"; exon_number
"1"; gene_name "LOT5"; gene_source "ensembl"; gene_biotype "protein_coding";
transcript_name "LOT5"; transcript_source "ensembl"; transcript_biotype
"protein_coding"; protein_id "YKL183W"; protein_version "1";
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(3)

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853 start_codon
853 stop_codon
886 gene
886 transcript
895 CDS
933 exon
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