Knowledge of Genomics

1. For each of the following statements, please select one response per row.

	True	False	l don't know
A gene encodes a protein.	0	0	0
Each gene has its own unique promoter.	0	0	0
Each gene encodes an RNA.	0	0	0
If translated, the DNA sequence of the gene controls protein structure and function.	0	0	0
A gene is transcribed.	0	0	0
If translated, the DNA sequence of a gene has no impact on the cellular localization of the encoded protein.	0	0	0
Each gene has its own Shine-Dalgarno sequence.	0	0	0
If translated, any change in the DNA sequence of the gene leads to a change in the amino acid sequence of the encoded protein.	0	0	0
A gene's DNA sequence is usually no more closely related to DNA sequences from its closest relatives than to DNA sequences of unrelated organisms.	0	0	0
A gene has homologs in other organisms.	0	0	0

2. For each of the following statements, please select one response per row.

	True	False	l don't know
There are 6 possible reading frames for protein translation for any double stranded DNA molecule.	0	0	0
The line below is the correct format for a FASTA header that is added to the beginning of a DNA or amino acid sequence prior to computer analysis.	C	0	С
>Ksed_00010 nucleotide sequence			
T-Coffee analysis predicts the cellular location of a gene product.	0	0	0
A Shine-Dalgarno sequence in bacterial DNA indicates the site where DNA polymerase binds to start DNA replication.	0	0	С
A signal peptide directs proteins to be secreted from the cell in bacteria.	0	0	0
An EC number is used to describe the function of an enzyme.	0	C	С
All genes in a bacterial genome are functional.	0	0	0
Some genes in a species of bacterium may have been obtained from an entirely different species of bacterium without being inherited.	0	0	0
Gene annotation is the process of assigning function to a particular DNA sequence.	0	0	0
Refer to the TMHMM output below. The results indicate that the protein being analyzed is located outside of the cell. TMHMM posterior probabilities for 644990854 12 13 0.8 0.4 0.2 0.5 100 150 200 250 300 transmembrane outside outside	C	C	C