

Solutions to 7.012 Central Dogma Problems

Prokaryotic Gene (β -galactosidase)

- a) 5' AAUUGUGAGC.....3'
- b) H_3N^+ -met-ser-his-arg-lys-... COO^-
- c) No, because it is read out of frame; it is read as: CUA, AGA
- d) H_3N^+ -met-ser-**thr-gly-asn-.....** COO^-
- e) H_3N^+ -met-ser-his-**lys-glu-...** COO^-
- f) No change, silent mutation
- g) Delete 1 or add 2 after AUG and before the TAA (and do not make another in-frame stop).
- h) UGG (trp) ---> UGA (stop)
- i) AGG (arg) ---> GGG (gly)
- j) AGG (arg) ---> AGA (arg)

Eukaryotic Gene (con-6)

The structure of con-6 is shown below (numbers as in the genomic sequence).

