

# Lecture 26:

# **Point mutations**

Course 281



### Lessons for life



## Jim Rohn Official

@OfficialJimRohn

"Lead the way by personal example and by personal philosophy." -- Jim Rohn



### AIMS

• Understand the various point mutations that take place in DNA and the consequences of such mutations.

• Understand the terminology given for each type of point mutation.

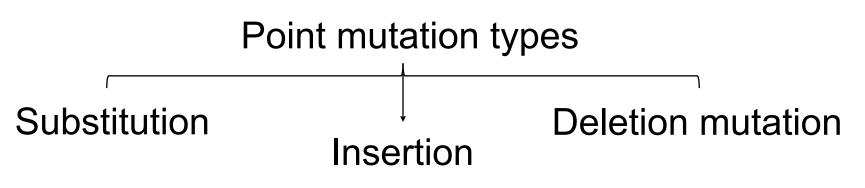
 Understand which mutation type has the least impact on protein function and structure and which has the most.

#### Mutation types

Chromosomal

Point mutation

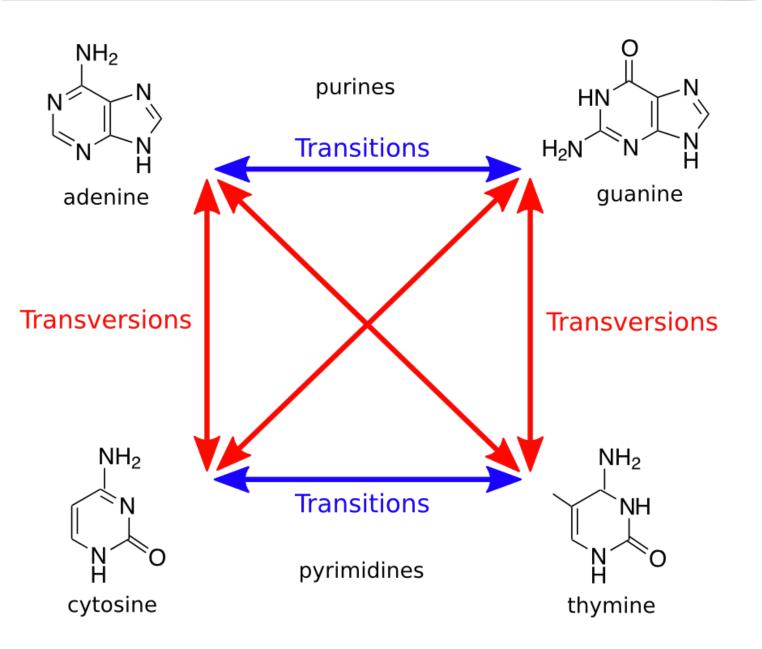
- Chromosomal mutations: are mutation that involve changes to the entire chromosome or sections of it.
- Chromosomal mutations are also called segmental mutations.
- **Point mutations:** are changes to one or few base-pairs in the DNA in the form of substitution, deletion, or insertion.



- Substitution point mutation: change of a nucleotide or few nucleotides from one type to another.
- Insertion point mutation: an insertion of one or few nucleotides in the DNA sequence.
- **Deletion point mutation:** a deletion of one or few nucleotides in the DNA sequence.

- Substituting (changing) a pyrimidine to a pyrimidine is called Transition.
- Substituting (changing) a purine to a purine is called Transition.
- Substituting (changing) a purine to a pyrimidine is called **transversion**.
- Substituting (changing) a pyrimidine to a purine is called transversion.

What does that mean?



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# What is the change from adenine to guanine called?

# What is the change from adenine to cytosine called?

# What is the change from adenine to thymine called?

# What is the change from adenine to adenine called?

#### While the change from A to T is called ....., the change from T to A is called.

### Where do mutations take place?



Point (substitution) mutations

In non-coding region

In coding region

Just a mutation No change in protein No functional consequences

Point mutations can take place at different localities in the genome (coding and non-coding regions).

 Point mutation in the non-coding region has no affects on the function or sequence of proteins.

# What are some examples of non-coding regions?

 Point mutations in the non-coding region have no biological impacts on the organism and thus can be used to study genetic variation between individuals of one species or between species.

We will be focusing one point mutations in the coding regions of the genome

Point mutations

In coding region

Synonymous mutations Non-synonymous mutations

No change in amino acid

Change in amino acid

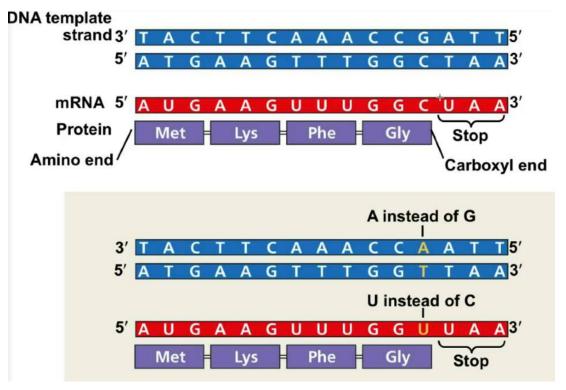
- Point mutations in coding region can be synonymous or non-synonymous.
- Synonymous mutations are also called point mutations.
- Synonymous mutations cause no change in the amino acid.

# Synonymous (silent) mutations do not change amino acid

How is that?

Synonymous point (substitution) mutations (silent mutations)

#### A change in the third nucleotide of codon GGC to GGT does not change the amino acid Glycine to another one.



# Why do synonymous (silent) mutations do not change amino acid in the coding region?

# Why most silent mutations are associated with the third base in the codon?

# Point mutations – in coding region

#### Synonymous point mutations (silent mutations) The genetic code is degenerate

| Second letter |   |                                 |                          |  |  |                  |              |  |  |  |  |  |
|---------------|---|---------------------------------|--------------------------|--|--|------------------|--------------|--|--|--|--|--|
|               |   | U                               | С                        | Α                                      | G                                      |                  |              |  |  |  |  |  |
| First letter  | U | UUU<br>UUC<br>UUA<br>UUA<br>UUG | UCU<br>UCC<br>UCA<br>UCG |  | UGU<br>UGC<br>UGA Stop<br>UGG Trp      | U<br>C<br>A<br>G | Third letter |  |  |  |  |  |
|               | c | CUU<br>CUC<br>CUA<br>CUG        | CCU<br>CCC<br>CCA<br>CCG | CAU<br>CAC<br>CAA<br>CAA<br>CAG<br>Gln | CGU<br>CGC<br>CGA<br>CGG               | U<br>C<br>A<br>G |              |  |  |  |  |  |
|               | A | AUU<br>AUC<br>AUA<br>AUG Met    | ACU<br>ACC<br>ACA<br>ACG | AAU<br>AAC<br>AAA<br>AAA<br>AAG        | AGU<br>AGC<br>AGA<br>AGA<br>AGG<br>Arg | U<br>C<br>A<br>G |              |  |  |  |  |  |
|               | G | GUU<br>GUC<br>GUA<br>GUG        | GCU<br>GCC<br>GCA<br>GCG | GAU<br>GAC<br>GAA<br>GAA<br>GAG<br>Glu | GGU<br>GGC<br>GGA<br>GGG               | U<br>C<br>A<br>G |              |  |  |  |  |  |

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# Point mutations – in coding region

Synonymous point mutations (silent mutations)

# What is the effect of synonymous (silent) mutations on the function and structure of the protein?

Point mutations In coding region Non-synonymous mutations

Missense mutation

Change an amino acid

into another amino

acid

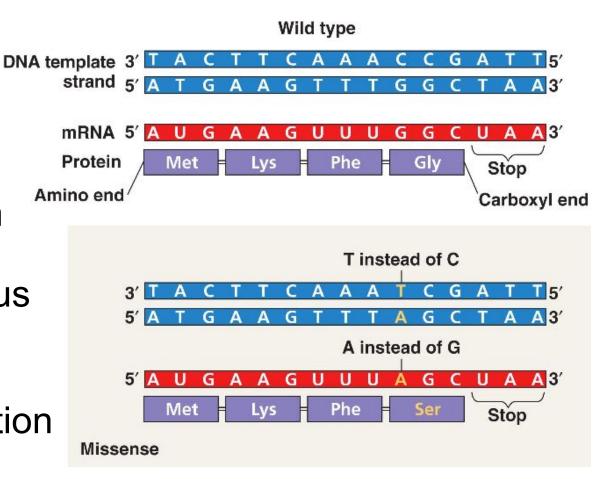
Nonsense mutation ↓ Change an amino acid into a stop codon



 Non-synonymous point (substitution) mutations in the coding region can change the amino acid or introduce a stop codon.

#### Lets focus on missense mutations first

Point mutations In coding region Non-synonymous mutations **Missense** mutation



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# Missense substitutions often occurs in the 1<sup>st</sup> or 2<sup>nd</sup> base of the codon

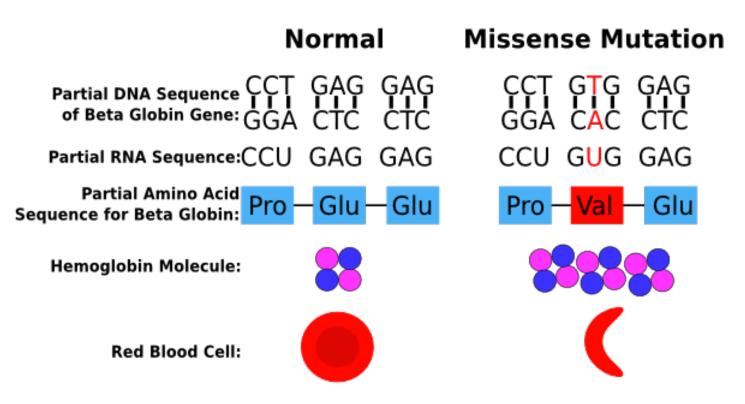
Why?

The degeneracy of the genetic code is associated with the 3<sup>rd</sup> base of the codon

#### **Second letter**

|              |   | U                            | С                        | Α                                       | G                                      |                  |               |  |  |  |  |  |
|--------------|---|------------------------------|--------------------------|---|--|------------------|---------------|--|--|--|--|--|
| First letter | U | UUU<br>UUC<br>UUA<br>UUG     | UCU<br>UCC<br>UCA<br>UCG | UAU<br>UAC<br>UAA Stop<br>UAG Stop      | <b>UGA Stop</b>                        | U<br>C<br>A<br>G |               |  |  |  |  |  |
|              | с | CUU<br>CUC<br>CUA<br>CUG     | CCU<br>CCC<br>CCA<br>CCG | CAU<br>CAC His<br>CAA<br>CAA<br>CAG GIn | CGU<br>CGC<br>CGA<br>CGG               | U<br>C<br>A<br>G | Inira         |  |  |  |  |  |
|              | A | AUU<br>AUC<br>AUA<br>AUG Met | ACU<br>ACC<br>ACA<br>ACG | AAU<br>AAC<br>AAA<br>AAA<br>AAG         | AGU<br>AGC<br>AGA<br>AGA<br>AGG<br>Arg | U<br>C<br>A<br>G | I hird letter |  |  |  |  |  |
|              | G | GUU<br>GUC<br>GUA<br>GUG     | GCU<br>GCC<br>GCA<br>GCG | GAU<br>GAC<br>GAA<br>GAA<br>GAG<br>Glu  | GGU<br>GGC<br>GGA<br>GGG               | U<br>C<br>A<br>G |               |  |  |  |  |  |

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- Missense mutations may alter the function of the protein by substituting an amino acid with an unfavorable one.
- Example: hemoglobin and sickle cell anemia.

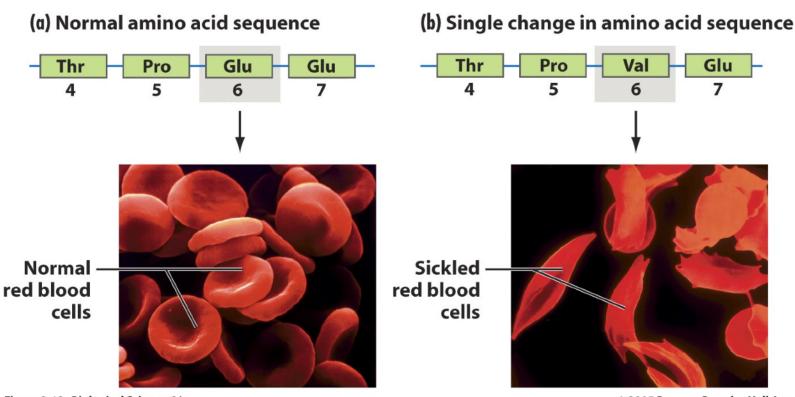


Figure 3-13 Biological Science, 2/e

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A change from glutamic acid (negatively charged) to a valine (non-polar) causes a severe change in the protein function and structure.

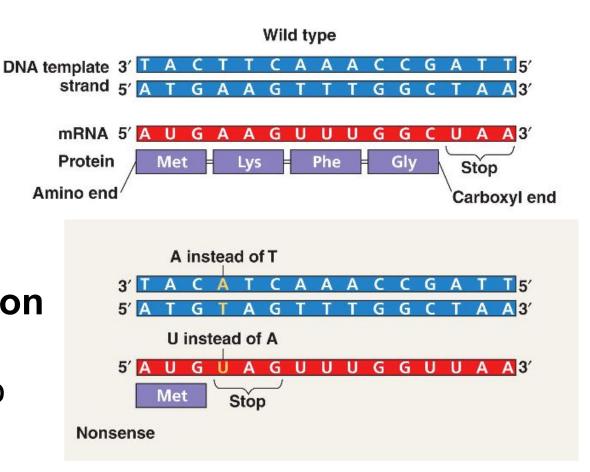
#### Point mutations

In coding region

Non-synonymous mutations

Nonsense mutation

Change an amino acid into a stop codon

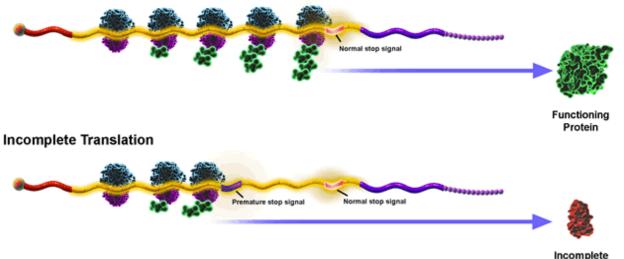


# Nonsense mutations introduce premature stop codon

### What happens to the protein?



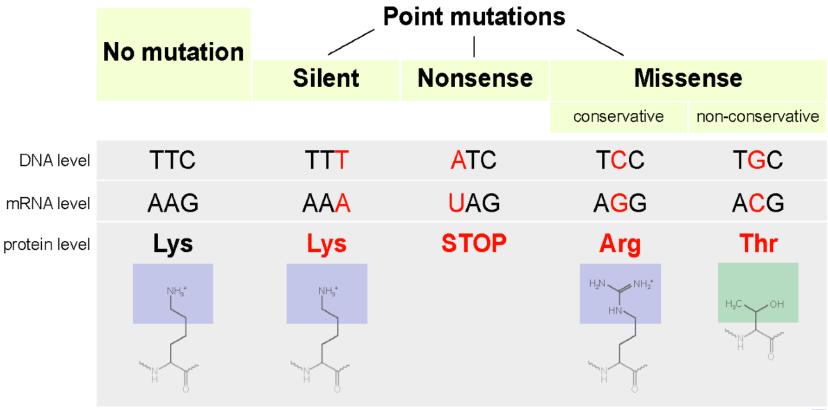




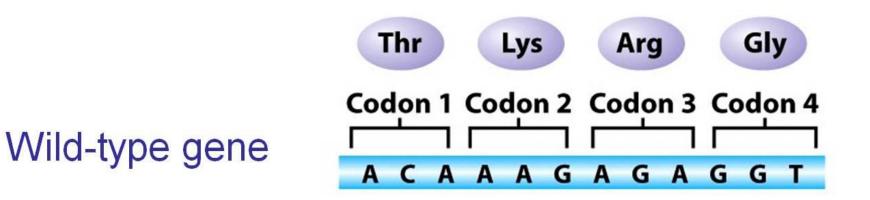
Incomplete Protein

- Nonsense mutation occurs early the gene results in the truncation (shorter) of protein and loss of function (**truncated protein**).
- Nonsense mutation occurs later in the sequence of the gene results in the protein loss of function or reduction in function.

#### Summary of substation mutations in coding region

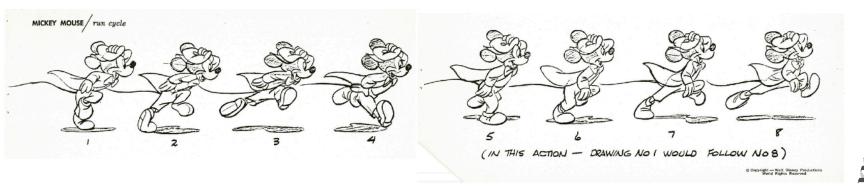


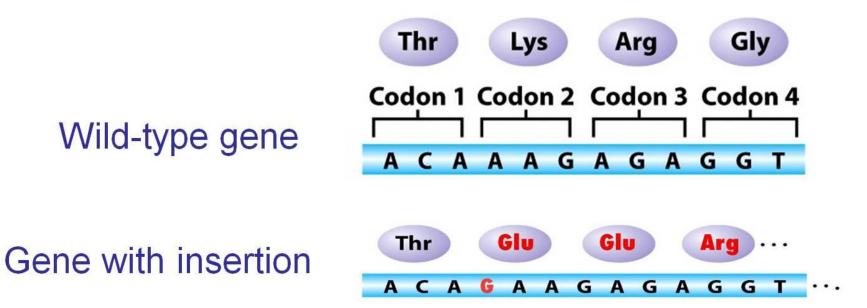
basic polar



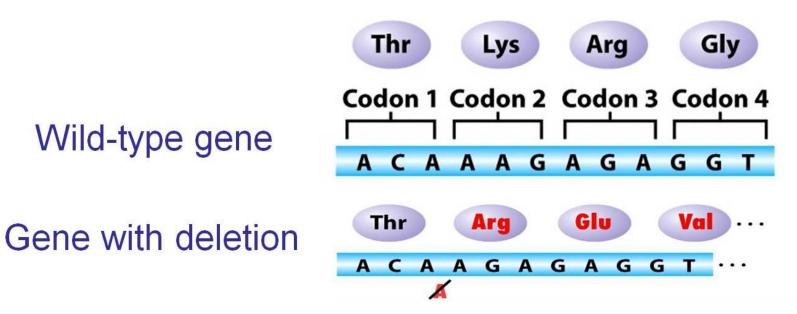
- Normal coding mRNA is called in frame when codons are arranged in specific sequence from start codon to stop codon.
- Insertion and deletion mutations introduce a disruption to the codon sequence called frame-shift.







- Insertion point mutations introduce a base-pair or more to the sequence of the gene which causes a frame-shift downstream.
- The frame-shift may result in a premature stop codon down stream or changes in amino acid sequence.



- Deletion point mutations remove a base-pair or more to the sequence of the gene which causes a frame-shift upstream.
- The frame-shift may result in a premature stop codon down stream or changes in amino acid sequence.

# Is there a difference when 1 bp is inserted vs. 2 or three?

# Is there a difference when 1 bp is deleted vs. 2 or three?

When an insertion or deletion happens and no frame-shift occurs (hint: numbers nucleotides in codon)?

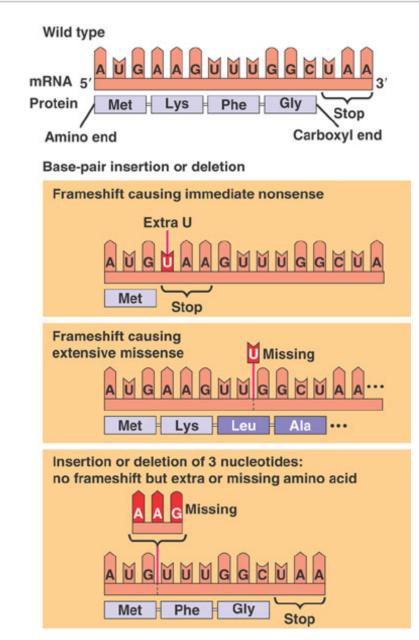
### Frame-shift by deletion or insertion causes:

1. An immediate nonsense mutation and presence of a stop codon.

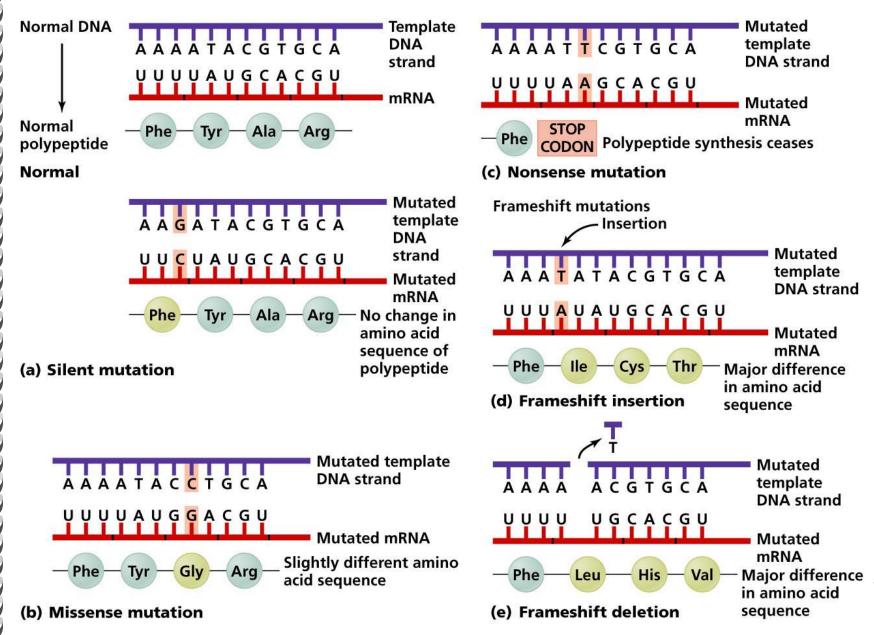
2. Causing missense mutation to downstream codons and change in all amino acids.

3. No frame-shift only a missing or added amino acid to the sequence.

When insertion or deletion happens with three base-pairs or multiple of three, only amino acids are added or removed

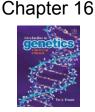


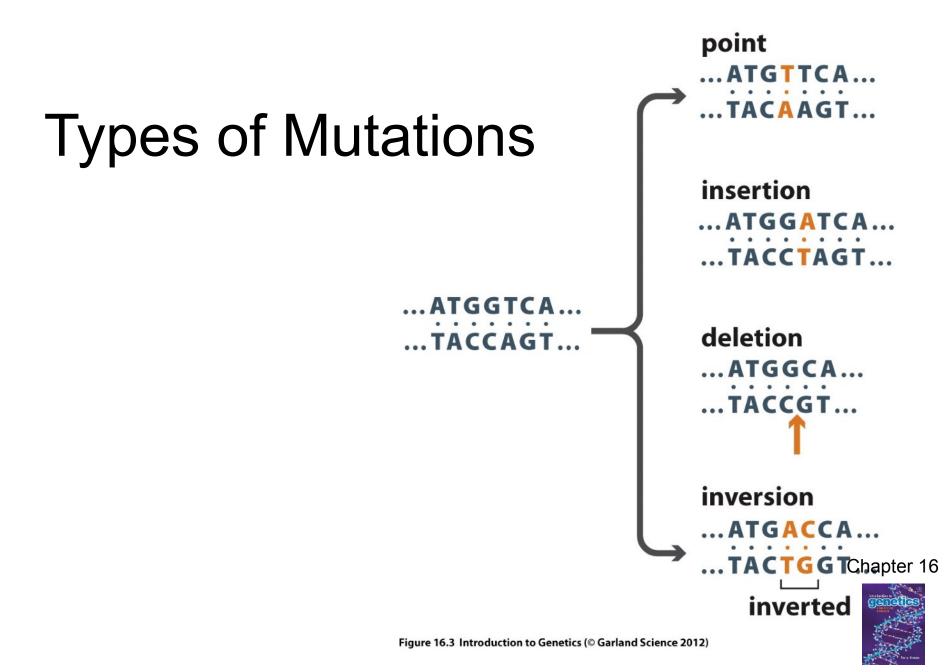
### Summary



# Types of Mutations

- **Point mutation:** one nucleotide is replaced by another
- **Insertions:** addition of base(s)
- **Deletions:** removal of base(s)
- Inversions: excision and reinsertion of a portion of DNA at the same position in the reverse orientation





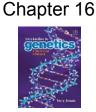
# **Point Mutation**

- Also called simple mutations, or single-site mutations
- Two types of point mutation;
  - Transition: purine to purine (A -> G) or pyrimidine to pyrimidine (T -> C) change
  - Transversion: purine to pyrimidine or vice versa (A or G -> T or C)



# Insertion and Deletion Mutations

- Insertion and deletion mutations result from aberrant DNA replication
- Insertion or deletion within the coding region may change the reading frame (frameshift mutation)
- Remember, inserting or deleting three nucleotides, or multiples of three, add or remove codons, or parts of adjacent codons, without affecting the reading frame
- Insertions and deletions also occur within <u>intergenic</u> regions





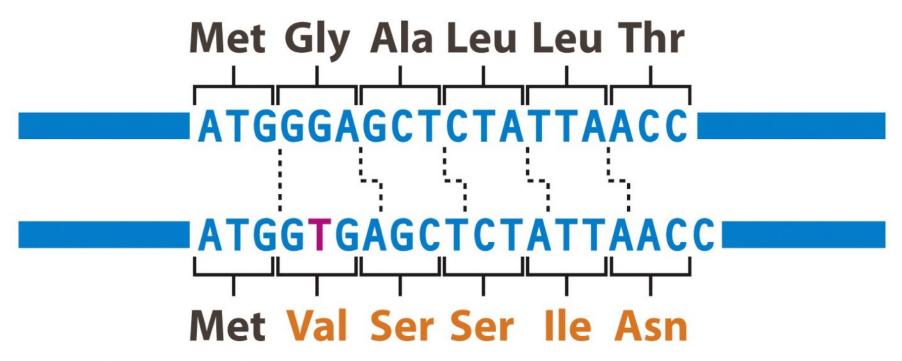


Figure 16.8 Introduction to Genetics (© Garland Science 2012)

Chapter 16



### To know

Nonsense mutation -

Transition

Point mutation

insertion

Non-coding region

Premature stop codon

substitution

Non-synonymous mutation

Missense mutation

deletion

Silent mutation

transversion

Segmental mutation

Chromosomal mutation

Frame-shift mutation

Truncated protein

Coding region

Synonymous mutation

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• You know the difference between point and segmental mutations.

• You know the various types of point mutations, the location of their occurrence, and their names.

• You know the consequences of frame-shift mutations and when they are least likely to affect protein function.

### For a smile



July .