



AP Bio

FRQ Fridays

2017 #3
Enzyme Structure, Mutations,
& Dominance



FRQ Friday #15

2017 #3

Gibberellin is the primary plant hormone that promotes stem elongation. GA 3-beta-hydroxylase (GA3H) is the enzyme that catalyzes the reaction that converts a precursor of gibberellin to the active form of gibberellin. A mutation in the *GA3H* gene results in a short plant phenotype. When a pure-breeding tall plant is crossed with a pure-breeding short plant, all offspring in the F_1 generation are tall. When the F_1 plants are crossed with each other, 75 percent of the plants in the F_2 generation are tall and 25 percent of the plants are short.

		Second Base in Codon				
		U	C	A	G	
U	UUU	UCU } Phe	UUA } Tyr	UGU } Cys	U	
	UUC				U	
	UUA	UCA } Ser	UAA Stop	UGA Stop	C	
	UUG				UAG Stop	UGG Trp
C	CUU	CCU } Leu	CAU } His	CGU } Arg	C	
	CUC				C	
	CUA	CCA } Pro	CAA } Glu	CGA } Arg	A	
	CUG				CCG } Arg	CGG } Arg
A	AUU	ACU } Ile	AAU } Asn	AGU } Ser	U	
	AUC				U	
	AUA	ACA } Thr	AAA } Lys	AGA } Arg	C	
	AUG Met or Start				ACG } Thr	AAG } Arg
G	GUU	GCU } Val	GAU } Asp	GGU } Gly	G	
	GUC				G	
	GUA	GCC } Ala	GAA } Glu	GGA } Gly	C	
	GUG				GCA } Ala	GAG } Glu
	GCG } Ala		GGG } Gly	G		

Figure 1. The universal genetic code



FRQ Friday #15

2017 #3

- (a) The wild-type allele encodes a GA3H enzyme with alanine (Ala), a nonpolar amino acid, at position 229. The mutant allele encodes a GA3H enzyme with threonine (Thr), a polar amino acid, at position 229. **Describe** the effect of the mutation on the enzyme and **provide reasoning** to support how this mutation results in a short plant phenotype in homozygous recessive plants.

Recall, GA3H is the enzyme that catalyzes the conversion to active gibberellin which promotes stem elongation

Description (1 point)	Reasoning (1 point)
The amino acid substitution changes the shape/structure/function of the protein.	The mutation decreases/eliminates gibberellin production.



FRQ Friday #15

2017 #3

a) The change from a nonpolar amino acid ^(Ala) to a polar amino acid (thr) disrupts the structure of the enzyme (tertiary structure). Because the molecule is non charged, weak interactions (such as van der Waals ~~ester~~ interactions) will occur between the amino acid and other polar amino acids (will be attracted to each other, thus altering the folding of the protein). If the folding and twisting of the enzyme is changed, the active site will change and the precursor to gibberellin won't be able to bind and convert to the active form of gibberellin. This means that stem elongation will not be promoted due to decreased amounts of the active form of gibberellin.



FRQ Friday #15

2017 #3

(b) Using the codon chart provided, **predict** the change in the codon sequence that resulted in the substitution of alanine for threonine at amino acid position 229.

		Second Base in Codon				
		U	C	A	G	
U	UUU	UCU } Phe	UUA } Tyr	UGU } Cys	U	
	UUC					
	UUA	UCA } Ser	UAA Stop	UGA Stop		
	UUG					
C	CUU	CCU } Leu	CAU } His	CGU } Arg	C	
	CUC					
	CUA	CCA } Pro	CAA } Glu	CGA } Arg		
	CUG					
A	AUU	ACU } Thr	AAU } Asn	AGU } Ser	A	
	AUC					
	AUA		AAA } Lys	AGA } Arg		
	AUG Met or Start					
G	GUU	GCU } Ala	GAU } Asp	GGU } Gly	G	
	GUC					
	GUA		GAA } Glu	GGA } Gly		
	GUG					

Figure 1. The universal genetic code

Prediction (1 point maximum)

- G ↔ A in the first position (of the codon)
- 5'-GCN-3' ↔ 5'-ACN-3'
- 5'-NGC-3' ↔ 5'-NGT-3' in the template strand of DNA



FRQ Friday #15

2017 #3

(b) Using the codon chart provided, predict the change in the codon sequence that resulted in the substitution of alanine for threonine at amino acid position 229.

Prediction (1 point maximum)

- $G \leftrightarrow A$ in the first position (of the codon)
- $5\text{'-GCN-}3' \leftrightarrow 5\text{'-ACN-}3'$
- $5\text{'-NGC-}3' \leftrightarrow 5\text{'-NGT-}3'$ in the template strand of DNA

b) GCA to ACA (replace first ^{base} ~~base~~ G with a A base).



FRQ Friday #15

2017 #3

(c) Describe how individuals with one (heterozygous) or two (homozygous) copies of the wild-type *GA3H* allele can have the same phenotype.

Description (1 point)

- Enough active enzyme is produced from one wild-type/dominant allele.
- Enough gibberellin is produced in the presence of one wild-type/dominant allele.

c) The ~~GA3H~~ wild-type *GA3H* could be dominant to the mutant allele, so the wild-type gene in a heterozygous plant could be expressed rather than the mutant gene. In a homozygous wild-type, the wild-type gene would be expressed as well. Both heterozygous and homozygous would be able to ^{convert} ~~produce~~ gibberellins to promote stem elongation, resulting in the same phenotype.

