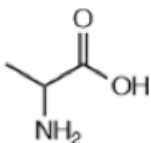
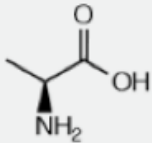
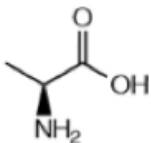
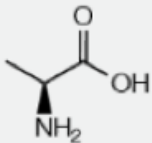
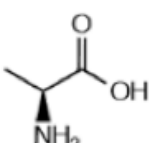
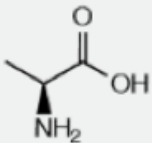
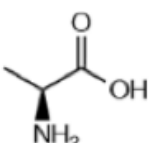
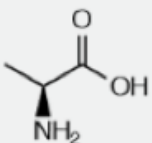
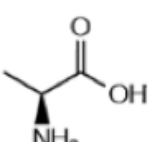
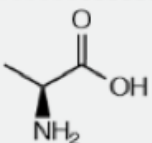
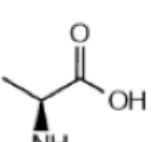
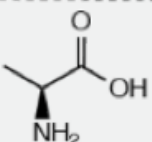
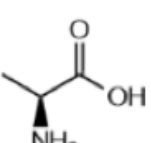
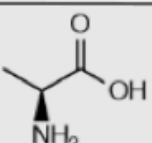
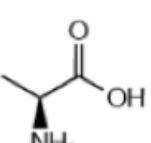
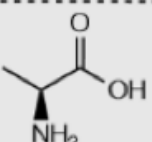
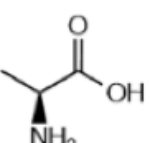
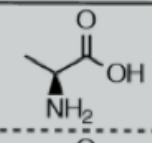
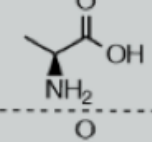
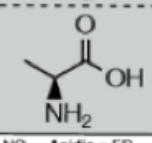


## CONCEPT: THE CLUTCH PREP AMINO ACID BREAKDOWN

Nonpolar Side Chains:	Notes:	Polar Side Chains:	Notes:
	Glycine Gly (G) COOH = 2.34 NH <sub>3</sub> <sup>+</sup> = 9.60		Cysteine Cys (C) COOH = 1.96 NH <sub>3</sub> <sup>+</sup> = 10.28
	Alanine Ala (A) COOH = 2.34 NH <sub>3</sub> <sup>+</sup> = 9.69		Serine Ser (S) COOH = 2.21 NH <sub>3</sub> <sup>+</sup> = 9.15
	Valine Val (V) COOH = 2.32 NH <sub>3</sub> <sup>+</sup> = 9.62		Threonine Thr (T) COOH = 2.09 NH <sub>3</sub> <sup>+</sup> = 9.10
	Leucine Leu (L) COOH = 2.36 NH <sub>3</sub> <sup>+</sup> = 9.60		Tyrosine Tyr (Y) COOH = 2.20 NH <sub>3</sub> <sup>+</sup> = 9.11
	Isoleucine Ile (I) COOH = 2.36 NH <sub>3</sub> <sup>+</sup> = 9.60		Asparagine Asn (N) COOH = 2.02 NH <sub>3</sub> <sup>+</sup> = 8.80
	Proline Pro (P) COOH = 1.99 NH <sub>3</sub> <sup>+</sup> = 10.60		Glutamine Gln (Q) COOH = 2.17 NH <sub>3</sub> <sup>+</sup> = 9.13
	Methionine Met (M) COOH = 2.28 NH <sub>3</sub> <sup>+</sup> = 9.21		Aspartic Acid Asp (D) COOH = 1.88 NH <sub>3</sub> <sup>+</sup> = 9.60
	Phenylalanine Phe (F) COOH = 1.83 NH <sub>3</sub> <sup>+</sup> = 9.13		Glutamic Acid Glu (E) COOH = 2.19 NH <sub>3</sub> <sup>+</sup> = 9.67
	Tryptophan Trp (W) COOH = 2.83 NH <sub>3</sub> <sup>+</sup> = 9.39		Lysine Lys (K) COOH = 2.18 NH <sub>3</sub> <sup>+</sup> = 8.95
			Histidine His (H) COOH = 1.82 NH <sub>3</sub> <sup>+</sup> = 9.17
			Arginine Arg (R) COOH = 2.17 NH <sub>3</sub> <sup>+</sup> = 9.04

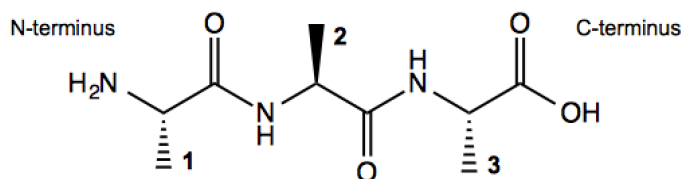
 = universally accepted as essential amino acids (FVT H\_L\_K\_M\_W) Aliphatic = GAVLP Other Nonpolar = MWF Polars = C-STY-NQ Acidic = ED Basic = KHR

## CONCEPT: THE 20 AMINO ACIDS: BLANK WORKSHEET

Nonpolar Side Chains:	Notes:	Polar Side Chains:	Notes:
	COOH = 2.34 NH <sub>3</sub> <sup>+</sup> = 9.60		COOH = 1.96 NH <sub>3</sub> <sup>+</sup> = 10.28
	COOH = 2.34 NH <sub>3</sub> <sup>+</sup> = 9.69		COOH = 2.21 NH <sub>3</sub> <sup>+</sup> = 9.15
	COOH = 2.32 NH <sub>3</sub> <sup>+</sup> = 9.62		COOH = 2.09 NH <sub>3</sub> <sup>+</sup> = 9.10
	COOH = 2.36 NH <sub>3</sub> <sup>+</sup> = 9.60		COOH = 2.20 NH <sub>3</sub> <sup>+</sup> = 9.11
	COOH = 2.36 NH <sub>3</sub> <sup>+</sup> = 9.60		COOH = 2.02 NH <sub>3</sub> <sup>+</sup> = 8.80
	COOH = 2.36 NH <sub>3</sub> <sup>+</sup> = 9.60		COOH = 2.17 NH <sub>3</sub> <sup>+</sup> = 9.13
	COOH = 1.99 NH <sub>3</sub> <sup>+</sup> = 10.60	Acidic Side Chains:	Notes:
	COOH = 2.28 NH <sub>3</sub> <sup>+</sup> = 9.21		COOH = 1.88 NH <sub>3</sub> <sup>+</sup> = 9.60
			COOH = 2.19 NH <sub>3</sub> <sup>+</sup> = 9.67
		Basic Side Chains:	Notes:
	COOH = 1.83 NH <sub>3</sub> <sup>+</sup> = 9.13		COOH = 2.18 NH <sub>3</sub> <sup>+</sup> = 8.95
	COOH = 2.83 NH <sub>3</sub> <sup>+</sup> = 9.39		COOH = 1.82 NH <sub>3</sub> <sup>+</sup> = 9.17
			COOH = 2.17 NH <sub>3</sub> <sup>+</sup> = 9.04

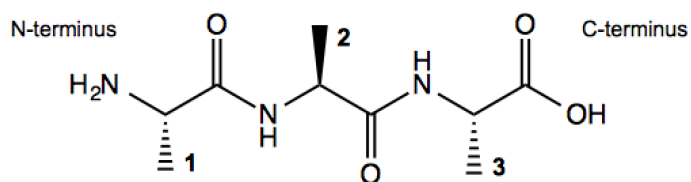
**CONCEPT: THE 20 AMINO ACIDS: NON-POLAR SIDECHAINS**

**PRACTICE 1:** Non-Polar Sidechains - Fill in the missing sidechains on the following target tripeptide.



Target Peptide: **L-A-M**

**PRACTICE 2:** Non-Polar Sidechains - Fill in the missing sidechains on the following target tripeptide.



Target Peptide: **W-I-G**

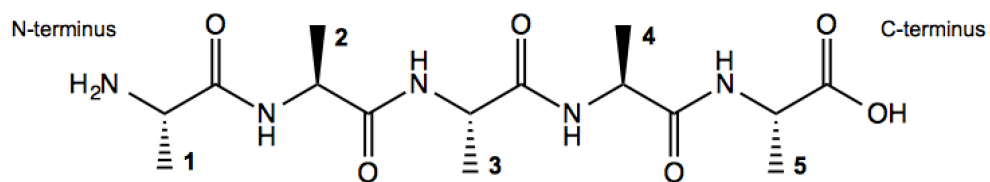
**PRACTICE 3:** Non-Polar Sidechains - Provide the complete structure of the tripeptide P-F-W

**CONCEPT: THE 20 AMINO ACIDS: POLAR SIDECHAINS**

**PRACTICE 1:** Polar Sidechains - Provide the Fischer Projection of Glutamine (Q)

**PRACTICE 2:** Polar Sidechains - Provide the complete structure of the dipeptide Asn-Cys

**PRACTICE 3:** Polar Sidechains - Fill in the missing sidechains on the following target oligopeptide.

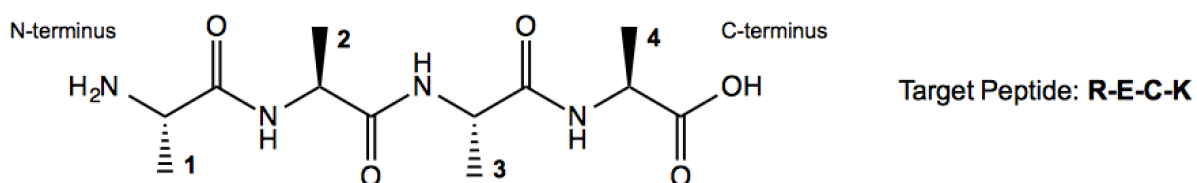


Target Peptide: **F-A-S-T-N**

**CONCEPT: THE 20 AMINO ACIDS: ACIDIC/BASIC SIDECHAINS**

**PRACTICE 1:** Acidic/Basic Sidechains - Provide the Fischer Projection of Amino Acid (H)

**PRACTICE 2:** Acidic/Basic Sidechains - Fill in the missing sidechains on the following target oligopeptide.



**PRACTICE 3:** Acidic/Basic Sidechains - Provide the complete structure of the oligopeptide V-I-D-Y. Based on your knowledge of hydrophobicity, which side of the peptide is more likely bury itself within the protein?