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What does acetylation do to dna

What is acetylation of dna. How does acetylation affect gene expression.

Understanding: â € â € â € "Nucleosomes help regulate the transcription in eukariotic proteins is wrapped around histone proteins to form the compact nucleosomes' histone proteins Tails that determine how strongly the DNA is bundled of the histone tailstipally the histone tails have a positive charge and therefore associate firmly with the negatively loaded dnaAdding an acetyl group to the tail (acetylation) neutralizes the charge, making the DNA less curled and increasing the transcription of a methyl group to the tail (methylation) maintained positive charge, making the DNA more curled and reducing the DNA to transcriptionionation of nucleosomes types of chromatin DNA is overcome and not accessible to transcription, there is condensed heterochromatin when the DNA is loosely packed and therefore accessible to the transcription machinery, exists as the types of euchromatinidified cells will have variad segments DNA as heterochromatin and euch Romantic DNA segments can be permanently supercoiled, while other segments can change over the life cycle of CÂ © Lulase Celancing in euchromatin and heterochromatin ability: â € â € , - Â € Changes in DNA methylation patterns - DNA direct methylation (in opposition for histone tails) can also affect methylation Of genetic expression The DNA methylation decreases the constitution (preventing the connection of transcription factors), consequently, the genes that are not Transcripts tend to show more DNA methylation than genes that are actively transcriptor contributors that contribute to epigeneticsepigenetics methylation patterns is the study of changes in phenotype as a result of variations in the epigenic analysis of the genetic expression shows that the DNA methylation patterns can change along a vital authority is influence From the heritability, but it is not genetically -determinable (idless hundreds may have different DNA methylation patterns) different cell types in SAM and organism may mark differently different methylation patterns DNA methylation (for example, diet, pathogenic exposition, etc.) can influence the level of DNA methylation within the methylation patterns of DNA from CellComparatives on different ages of comparative methylation patterns in different health status (healthless â €

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