Pathway Leading to GST Inactivation

Eukaryotic Gene Structure

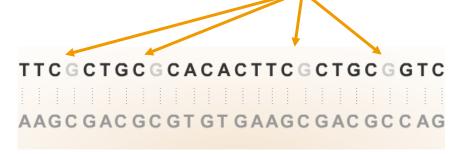


http://dynamicgene.org/structure/structure.html

Key components:

- Promoter site to which RNA Polymerase binds for transcription
- Exons those sequences of nucleotides expressed in the final RNA molecule
- Introns those sequences of nucleotides removed during RNA processing

- The promoter regions of some genes have been found to be rich in C and G nucleotides
 - When C is found next to G (in the 5' to 3' direction) the dinucleotide is referred to as a CpG dinucleotide.



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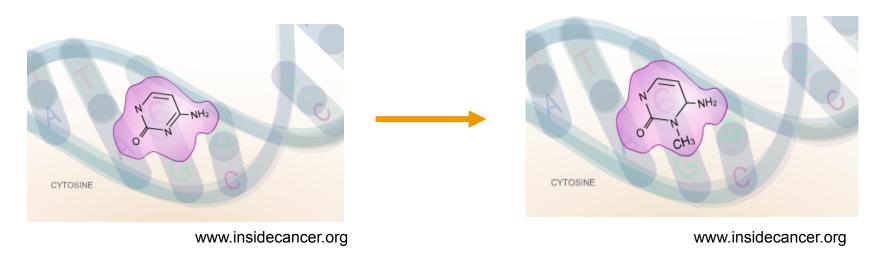
- The "p" indicates the two nucleotides are separated by a phosphate and are therefore on the same strand.
- When this dinucleotide is present numerous times the region is referred to as CpG island

Regulating Gene Expression

- Methylation: a key mechanism cells use to regulate gene expression.
 - Click <u>here</u> to see the role of methylation in gene expression

Conclusion:

■ Methyl groups, -CH₃, are added to cytosine nucleotides in the CpG islands found in the promoter regions of genes



Unmethylated Cytosine

Methylated Cytosine

■ This results in a suppression of the expression of the gene



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Methylation and Prostate Cancer

- The suppression GST expression is due to excessive methylation of the CpG islands in the promoter region of the gene
 - As a result molecules such as oxidants and PAH's remain in the cell