

Effect of protein interactions on DNA; understanding tumourigenic mutations with applications to p53 tumour suppressor

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P53 protein is a tumour suppressor encoded by a gene whose largely responsible for protecting cells from cancer causing DNA damaging agents. Functional p53 is thought to provide a protective effect against tumorigenesis, and indeed, mutations of p53 have been found in nearly all tumour types and are estimated to contribute to around 50% of all cancers. Focus of this investigation is to study the effect of protein interactions on DNA in understanding tumorigenic mutations. In this work, self-consistence field molecular orbital calculations were used to evaluate gas phase ionization potential of the isolated mononucleotides of the sequence (5'₁A-T-A-A-T₅-T-G-G-G-C₁₀-A-A-G-T-C₁₅-T-A-G-G-A₂₀-A₃) 5'-dGMP^(7,8,9,13,10'), 5'-dCMP^(9'), 5'-dTMP^(11') and dinucleotides 5'-pGpCp(10' and 9') and 5'-pTpGp(11' and 10'). To explore the impact of p53 loss on tumorigenesis in a mammalian model, most frequently mutated residues Arg²⁴⁸, Arg²⁷³, and their contacts with DNA were studied. Arg²⁷³ hot spot site was studied by considering 5'-dTMP^(11')-Arg²⁷³, 5'-dGMP^(10')-Arg²⁸⁰, and the large bridge 5'-dTMP^(11')-Arg²⁷³-Asp²⁸¹-Arg²⁸⁰-5'-dGMP^(10'). The Arg²⁴⁸ hot spot site was studied by considering 5'-dGMP⁽¹³⁾, 5'-dTMP⁽¹⁴⁾, 5'-pGpTp (13 and 14), and the small bridge 5'-dGMP⁽¹³⁾-H₂O-Arg²⁴⁸. Ionization potentials(IP) were obtained by employing Koopmans' theorem to results obtained from *ab-initio* Hartree-Fock level of calculations with a 3-21G basis set. The results guide that in the GGG sequence, the lowest ionization potential is obtained by middle Guanine among other G containing XGY sequences, indicating that the GGG triplet is most easily oxidized in DNA. The results obtained for the large bridge 5'-dTMP^(11')-Arg²⁷³-Asp²⁸¹-Arg²⁸⁰-5'-dGMP^(10') indicate that of the two base components, lowest IP is on Thymine base of 5'-dTMP^(11'). Thymine (11') bridges to Arg²⁷³, one of the frequently mutated residues of p53 in human cancer and results to obtain the lowest ionization potential. In the small bridge interaction (5'-dGMP⁽¹³⁾-H₂O-Arg²⁴⁸), Guanine (13) nucleotide was found to have the lowest ionization potential. Hence substitutions at this position cause changes in the stability of the bridge thus making Arg²⁴⁸ more prone to mutations.