



Green Tea: A Natural Source of Drug for Liver Cancer

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ABSTRACT : Green tea (*Camellia sinensis*) has anticancer activity in animal. The HBx protein of Hepatitis B virus activate AP-1 protein and it causes the down regulation of PTEN and p53 which are tumor suppressor genes, leads to the tumor formation in liver. The interactions between HBx and AP-1 have been targeted by docking of natural compounds. EGCG obtained from green tea shows highest affinity for AP-1 protein among all natural compounds. Affinity of EGCG with AP-1 protein may be used to design drug for liver cancer.

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Key Words :

Liver cancer, HBV,
Activator protein-1,
Natural compounds,
EGCG, Docking

Anticancer activity of several natural compounds has been experimental proved in research study. Green Tea (*Camellia sinensis*) is one of them and most popular beverage consumed in the world; it has anticancer activity. Hepatitis B virus infection is a leading source of liver cancer. The HBx protein of Hepatitis B virus activate AP-1 protein (Jacqueline Benn *et al.*, 1996), it causes the down regulation of PTEN and p53 gene (Wang *et al.*, 1994; Haviv *et al.*, 1998). Therefore, tumor suppressors protein are not formed, it causes the tumor formation in liver. HBx protein plays a regulatory role in HBV replication and is necessary for in viral infection. It functions by protein-protein interaction and activation of transcription factor AP-1 (Michael, 2010). When AP-1 protein is activated by HBx protein, the production of AP-1 protein is very high and cell division is uncontrolled which ultimately results in the formation of tumors in liver. Activation of AP-1 protein is repressed by several natural compounds, which modulate its target gene (Rahul Amin *et al.*, 2009). In this study docking interactions of natural compounds EGCG,

curcumin, luteoline, genistein, ellagic acid, resveratrol, lupeol, betulinic acid, lycopene have been studied by analysing protein-protein interactions between HBx and AP-1. The interactions have been targeted by docking of natural compounds. Highest affinity of EGCG (-7.97) was predicted towards AP-1 protein and which may be involved in suppression of tumor formation process.

Material :

Databases :

NCBI (<http://www.ncbi.nlm.nih.gov/>), RCSB PDB (www.rcsb.org), PUBCHEM (<http://www.ncbi.nlm.nih.gov/pccompound>). Software/Tools: LOOPP Server (<http://clsb.ices.utexas.edu/loopp/web/>), GRAMM X server (<http://vakser.bioinformatics.ku.edu/resources/gramm/grammx>), PyMOL (<http://www.pymol.org/>). Schrodinger- Maestro (<http://www.schrodinger.com>).

Methods :

Structures of natural compounds have been downloaded from Pubchem databases.

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