

RESEARCH ARTICLE

EVALUATION OF MUTATION OF TP53 GENE WITH RISK OF SQUAMOUS CELL CARCINOMA HEAD AND NECK (HNSCC) IN KASHMIRI POPULATION.

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Manuscript Info

Abstract

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Manuscript History	Objective:- To find relationship between TP53 mutations and
Received: 29 September 2016 Final Accepted: 30 October 2016 Published: November 2016	squamous cell carcinoma head and neck (SCCHN). Study design: Case control study. Setting: Tertiary care hospital (SMHS associated Medical College, Srinagar, Kashmir, India)
<i>Key words:-</i> Squamous cell carcinoma Head and neck, Tp53 mutation, Immunohistochemistry.	 Participants: 50 cases and 50 controls of squamous cell carcinoma head and neck reported our hospital from 2013-2016. Conclusion: In the present study, HNSCC patients showed 15(30%) GG (Arginine/Arginine), 26(52%) GC (Arginine/Proline) and 9 (18%) CC (Proline/Proline) genotypes and controls shows 31(62%) GG, 11(22%) GC and 8(16%) CC. It was observed that TP53 codon 72 polymorphism was significantly associated with HNSCC. The frequency of homozygous CC (Proline/Proline) genotype was elevated in HNSCC cases compared to controls (P<0.0003,RR 1.845,95% CI 1.264-2.693,OR 3.845), where as frequency of Arg/Arg (P Value = 0.0024,RR 0.563,95% CI 0.3177-0.7967,OR. 0.2627) were elevated in controls compared to cases.
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Background:-

P53 was identified in 1979 by Lionel Crawford, David P. Lane, Arnold Levine, and Lloyd Old, working at Imperial Cancer Research Fund (UK) Princeton University/UMDNJ (Cancer Institute of New Jersey), and Memorial Sloan-Kettering Cancer Center, respectively. It had been hypothesized to exist before as the target of the SV40 virus, a strain that induced development of tumors. The TP53 gene from the mouse was first cloned by Peter Chumakov of the Russian Academy of Sciences in 1982¹ and independently in 1983 by Moshe Oren in collaboration with David Givol (Weizmann Institute of Science).^{2,3}The human TP53 gene was cloned in 1984 and the full length clone in 1985^{4}

In humans, the TP53 gene is located on the short arm of chromosome 17 (17p13.1)^{5,6,7,8} the gene spans 20 kb, with a non-coding exon 1 and a very long first intron of 10 kb. The coding sequence contains five regions showing a high degree of conservation in vertebrates, predominantly in exons 2, 5, 6, 7 and 8, but the sequences found in

Corresponding Author:- Dr Shakeel Mohmad Wani. Address:- Resident ENT HNS, SMHS Hospital, Srinagar, Kashmir, India. invertebrates show only distant resemblance to mammalian TP53TP53orthologs have been identified in most mammals for which complete genome data are available.

In humans, a common polymorphism involves the substitution of an arginine for a proline at codon position 72. Many studies have investigated a genetic link between this variation and cancer susceptibility; however, the results have been controversial. For instance, a meta-analysis from 2009 failed to show a link for cervical cancer⁹. A 2011 study found that the TP53 proline mutation did have a profound effect on pancreatic cancer risk among males.¹⁰ A study of Arab women found that proline homozygosity at TP53 codon 72 is associated with a decreased risk for breast cancer.¹¹

Meta-analyses from 2011 found no significant associations between TP53 codon 72 polymorphisms and both colorectal cancer risk¹² and endometrial cancer risk.¹³ A 2011 study of a Brazilian birth cohort found an association between the non mutant arginine TP53 and individuals without a family history of cancer.¹⁴ Another 2011 study found that the p53 homozygous (Pro/Pro) genotype was associated with a significantly increased risk for renal cell carcinoma.¹⁵

The TP53 gene can also be modified by mutagens (chemicals, radiation, or viruses), increasing the likelihood for uncontrolled cell division. More than 50 percent of human tumors contain a mutation or deletion of the TP53 gene.¹⁶ Loss of p53 creates genomic instability that most often results in an aneuploidy phenotype.¹⁷

Increasing the amount of p53 may seem a solution for treatment of tumors or prevention of their spreading. This, however, is not a usable method of treatment, since it can cause premature aging.¹⁸ Restoring endogenous normal p53 function holds some promise. Research has shown that this restoration can lead to regression of certain cancer cells without damaging other cells in the process. The ways by which tumor regression occurs depends mainly on the tumor type. For example, restoration of endogenous p53 function in lymphomas may induce apoptosis, while cell growth may be reduced to normal levels. Thus, pharmacological reactivation of p53 presents itself as a viable cancer treatment option.^{19,20}The first commercial gene therapy, Gendicine, was approved in China in 2003 for the treatment of head and neck squamous cell carcinoma. It delivers a functional copy of the p53 gene using an engineered adenovirus.²¹

Aim of the study:-

✤ To find relationship between TP53 mutations and squamous cell carcinoma head and neck (SCCHN.

Materials and methods:-

Study subjects:-

- This study was done in tertiary health centre of kashmir in Department of ENT and Head & Neck surgery SMHS and included patients with histologically confirmed squamous cell carcinoma Head and neck (SCCHN) over a period of 18 months. Patients with squamous cell carcinoma of oral cavity, oropharynx, nasal cavity, nasopharynx, hypopharynx and larynx, identified at the Department of ENT and head&neck surgery SMHS Hospital were included. The patients with secondary SCCHN, Thyriod malignancies, primaries outside the upper aero digestive tract, cervical metastases of unknown origin or histopathologic diagnoses other than squamous cell carcinoma were excluded. All cases were from Kashmir and had not received any treatment at the time of recruitment. Controls were also taken from Kahmiri population who were admitted to our hospital for some other non-neoplastic disease. After verbal and written consent 2ml of blood sample was taken from controls and analysed for genetic mutation.
- From Histologically proven patients of squamous cell carcinoma. 2 ml of blood was taken from each patient after proper consent and was analysed at the Department of Biochemistry (DNA extraction and genotyping). Samples were stored at temperature -80 °c.

Genotyping:-

From each blood sample of case and control, a leukocyte cell pellet was obtained from the buffy coat by centrifugation of 2 ml of whole blood for DNA extraction. Genomic DNA was isolated with the help of DNA extraction kit (Biotools Spain).

After the isolation of DNA from the samples, Polymerase Chain Reaction (PCR) using Thermal cycler (Eppendorf) was performed to amplify TP53 genes to see whether there is any mutation in this gene using specificprimers Listed in table1.

Table 1:- Shows Forward and Reverse Primers for TP53 Gene Used For Polymerase Chain Reaction of the above three different genes in Head & Neck Carcinoma cases and controls

Gene	Forward primer	Reverse primer
	5'- ATCTACAGTCCCCCTTGCCG-3'	5'-GCAACTGACCGTGCAAGTCA-3'

The reaction volume was 25 μ L containing the reagents listed in table 2 and the PCR conditions for the above three genes are listed in table 3 below

Table 2:- Shows Reagents Used For Polymerase Chain Reaction of the above three different genes in Head & Neck

 Carcinoma cases and controls

Reagents	TP53 gene
Modified DNA	2 μl
dNTPs	1 μl
Forward & Reverse Primer	1.5µl each
Taq Buffer	2 μl
DMSO	3 µl
Taq DNA Polymerase	0.2 μl (5U/ μl)
Deionised Water	15.3 μl
Total Volume	25µl

Table 3:-

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Condition	TP53 gene	
Initial Denaturation	96°C, 5min	
Denaturation	96°C,45sec	
Annealing	56°C,45sec	35 cycles
Extension	72°C,30sec	
Final elongation	72°C,5min	

After the PCR was over, 10 μ L of the PCR products were run on 2.5% agarose gel using electrophoresis apparatus and PCR products were verified 296bp for Tp53 gene .

For analysing TP53 (Arg72Pro) polymorphism the PCR product was digested by MspI (Biotools Spain) overnight at 37°C. The digested product was separated on 2.5% agarose gel with ethidium bromide and was photographed with the help of gel documentation system present in Biochemistry lab. The wild-type allele (GG) produced two bands (169 and 127 bp); wild-type/variant allele (GC) produced 169, 127 and 296 bp and the variant allele (CC) which lacks the MspI restrictionsite produced a single 296 bp band.

Results and observations:-

This study was taken to understand the affect of TP53 gene polymorphism in Head & Neck squamous cell Carcinoma patients. Genomic DNA was isolated from all Samples by using kit provided by Zymo-Research.



Figure 1:- Representative gel picture run on 0.8% agarose showing isolated Genomic DNA L1-L5= Cases of Head & Neck carcinoma,L6-L8= Controls.

. Negative control with ddH2O instead of DNA template was included in each PCR run. Statistical analysis was performed using fishers exact test.



Figure 4:- PCR amplification of TP53 (Arg 72 pro) gene showing band size of 296bp LM=50bp Molecular Marker DNA ladder .



Figure 7:- BstUI Restriction digestion of PCR product with TP53 (Arg72 Pro) polymorphic sites L1,L2= Heterozygous GA genotype (169bp, 127bp & 296bp) L3, L4=Homozygous AA genotype (296bp)

L1-L7=Cases of Head & Neck Carcinoma

L5, L6=Homozygous AA genotype (169bp & 127bp) L7=Negative Control (water) and LM= Molecular Marker 50bp DNA Ladder

Results:-

In the present study, HNSCC patients showed 15(30%) GG (Arginine/Arginine), 26(52%) GC (Arginine/Proline) and 9 (18%) CC (Proline/Proline) genotypes and controls shows 31(62%) GG, 11(22%) GC and 8(16%) CC. It was observed that codon 72 polymorphism was significantly associated with HNSCC. The frequency of homozygous CC (Proline/Proline) genotype was elevated in HNSCC cases compared to controls (P<0.0003,RR 1.845,95% CI 1.264-2.693,OR 3.845), where as frequency of Arg/Arg (P Value = 0.0024,RR 0.563,95% CI 0.3177-0.7967,OR. 0.2627) were elevated in controls compared to cases (Table 5).

Table 5:- Frequency of TP53 gene genotypes.

Results							
	Wild type GG	Homozygous CC	Heterozygous GC	Row Totals			
	(Arg/Arg	(Pro/Pro)	(Arg/Pro)				
CASES	15	26	9	50			
CONTROLS	31	11	8	50			
Column Totals	46	37	17	100 (Grand			
				Total)			

Data was analysed using Graph-pad prism softwear version-6 and the association was found statistically significant inrelation to TP53 gene as P Value < 0.05 (0.002), RR = 1.988, Odds ratio= 3.807 and 95% CI= 1.255 to 3.148

Conclusion:-

TP53 mutation has got definite association with head and neck squamous cell carcinoma in Kashmiri population with special reference to Laryngeal squamous cell carcinoma.

Most of the mutations of Tp53 has been found in aggressive tumours, advanced stage of disease so there is need to screen the population at risk early to prevent dreadful morbidity and mortality associated with disease.

Non-invasive simple blood based detection of TP53 mutation may serve as good screening tool The presence of TP53 mutation in these tumours represents a potential Tumour specific targeting opportunity for the development of new therapeutics

Ethical Clearance:- Sought from ethical committee GMC Srinagar. Conflict of interest:- Nil Source of funding:- Hospital

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