Introduction

The human body is made up of millions of cells. The growth of body cells in a normal mechanism takes place in a systematic and coordinated system. These organized cells in the early age of an individual multiply more swiftly so that the growth of the individual may be increased. Mostly cell division after the maturity proceeds the repairing of the dead tissues in the body to accomplish the repair mechanism. When the number of these dead or injured cells in the body increased considerably then these cells ultimately change into cancerous conditions. There are different types of cancers, but the principal occurrence of all kinds of cancers is abnormal and unusual cell growth. One of the emerging issues is prostate cancer occur in men because the prostate is a ductal small walnut shaped gland situated in men below the urinary bladder that produces the seminal fluid for sperms provision and transportation. The risk of emerging prostate cancer during the man’s lifetime is one out of seven. Naturally prostate enlarges with age causing a condition known as benign prostatic hyperplasia (BPH), in one-third of men over 60 and about half over 80 and symptoms with frequent urination. Due to common histopathology and molecular techniques, BPH has been specified by cancer tumor genesis, but their accurate connection remains vigorously contested.
Environmental Factors

Environmental factors associated with prostate cancer are studied in different countries among the different communities and it was concluded that different environmental factors may cause risk of prostate cancer such as smoking, a diet with an enhanced quantity of fat, workplace exposure such as stokers who have direct contact with different toxic ignition products, obesity, and diabetes, inflammation of the prostate, STDs and vasectomy, etc..

Genetic Basis of Prostate Cancer

Recently approximately 5000 somatic mutations in prostate growth have been reported, including PTEN-interacting protein MAGI2 or repeated translocations assuming the cell-adhesion molecule CADM2, besides the periodic mutations of the Speckle-type POZ Protein (SPOP). The 12 most seriously mutated genes are C14orf49, THSD7B, CDKN1B, TP53, NIP2, SPOP, PTEN, ZNF595, SCN11A, PIK3CA, MED12 and FOXA1. SPOP with a ratio of 13% was the most regularly mutated gene in these tumors. The individuals having BRCA mutations are at a 15-25% increased risk of prostate cancer. A study suggested that BRCA1, BRCA2 and HOXB13 might be possible of life alarming prostate cancer. Many different gene integrations have been observed in prostate cancer until now. Following ESRP1-RAF1, SLC45A3-BRAF and TMRPSS2-ERG, gene integrations are the leading molecular subtype of PCa. Due to hereditary factors, almost 10% of men get prostate cancer which causes the initial emergence of disease. To the X chromosome on the position of chromosome 1q, two familial susceptibility loci have been mapped. Different studies revealed the correlation of prostate cancer with breast cancer. The allelic disappearance exhibit the failure of function or depletion of tumor suppressor genes in PCa. Disappearances of heterozygosity were commonly identified at 17p, 13q, 10q and 8p, whereas some research reported deprivation of 18q, 16q, 7q and 6q. The initial investigators made it logical that hereditary prostate cancer (HPC) genes were located on chromosome 1; meanwhile, the second prostate cancer gene was located on the X chromosome (Xq27-28) suggesting the X-linked pattern of HPC inheritance. On the mapped region of chromosome 16 was observed that DNA sequence KIAA 0872 and 17-β hydroxyl steroid dehydrogenase present which showed that not any genes shift mutations in the protein-coding region, and this knowledge reported that these are limited to become the source of familial PCa. On different chromosomes, prostate cancer risks have been reported particularly on chromosome 1. The selected possible gene CAPB on chromosome 1p36, PCAP on chromosome 1q42-43, HPCX on chromosome Xq27-28, HPC2 on chromosome 17p, HPCI on chromosome 1q23-25, HPC20 on chromosome 20q13 and correlation to chromosome 8p22-23. These correlation studies expressed the mutation detection and mapping of strong possible genes such as MSRI, RNASEL and ELAC2. The repeated and limited germline mutation GB4E in HOXB13 has been linked with the higher threat of familial PCa. It has been observed that individuals holding both polymorphisms in the HPC2/ELAC2 gene encounter a significant increase in prostate cancer.

Risk Factors of Prostate Cancer

Anything that determines the chances of getting an illness is known as a risk factor, for example, disease. Different diseases have different risk factors. Some kinds of risk factors can be replaced such as smoking, but others related to an individual’s age or familial history cannot be replaced. But it was observed risk factors don’t express everything about the individual disease. Many individuals observed with one or more risk factors, but never get the disease while many individuals who have not identified with risk factor even though develop infections. It is still not identified the obvious possible risk reason for the development of prostate cancer, but professionals have found some of the risk factors associated with prostate cancer developed in men.

Age

Elder males are much more affected by prostate cancer. Elderly are being diagnosed with prostate cancer due to an increase in the prostate-specific antigen (PSA) testing and life span. The risk of the prostate has been observed in African Americans, or those individuals with familial positive history after 40 years and after 50 years among those who have no positive familial history. After the age of 65, the risk of the prostate is 6 in 10. The men who have a different percentage of prostate cancer during the different periods are shown in Table 1. However, it is greatly suggested for older individuals to perform digital rectal examination (DRE) and prostate-specific antigen (PSA) testing diagnosis.

Table 1: Percent of US. men who develop PCa over 10-, 20-, and 30-year intervals according to their current age, 2008-2010†

| Current Age | 10 Years | 20 Years | 30 Years |
|-------------|----------|----------|----------|
| 30          | 0.01     | 0.35     | 2.54     |
| 40          | 0.34     | 2.57     | 8.18     |
| 50          | 2.31     | 8.12     | 13.74    |
| 60          | 6.41     | 12.59    | 14.92    |
| 70          | 7.73     | 10.64    | N/A      |

Adapted from SEER Cancers Statistics Review 1975-2012
Ethnicity

Different recently researches identified that ethnicity is the major risk factor associated with prostate cancer. Prostate cancer is observed to have strong ethnic relations. African American male individuals have greater chances of this disease. African Americans are more probably identified with PCa in the United States and the death rate was observed 2.5 times more among them. According to the Surveillance, Epidemiology, and End Results (SEER) US population registry, the black ethnic population was related to a major risk of prostate cancer death rate. Individuals in Sub-Saharan Africa were probably died 5-fold more with PCa as compared to African Americans in the United States. Prostate cancer risks have been reported is correlated with chromosomal 8q24 variants and are much more familiar among the population of African American male individuals. A few studies have also suggested the increased rate of variations in cells programmed death-associated genes such as BCL2 and EphB2 tumor suppressor genes in the African American population.

Family history

To the occurrence of prostate cancer both genetic and environmental factors are also associated with them. The chances of prostate cancer occurrence are more in those patients with familial member’s history and diagnosis at a young age and it can be more by two to three-folds with first-rank relatives get prostate cancer. HPC1 found at chromosomal location 1q24–25 and HPCX at Xq27–28 are hereditary prostate cancer (HPC) genes associated with PCa. BRCA1 & BRCA2 mutations have been supportive in facilitating the sharing of inherited alleles in cancer-associated families. In table 2 the greater risk caused by the hereditary factors is mentioned.

Table 2: Main genes associated with the hereditary formation of familial prostate

| Chromosomal location | Genes | Characteristics |
|----------------------|-------|-----------------|
| 1q24-25              | HPC1  | Autosomal dominant inheritance, associated with brain tumors |
| 1q42-43              | PCaP  | Autosomal dominant inheritance |
| Xp11                 |       | Sex-related inheritance |
| Xp27-28              | HPCX  | Sex-related inheritance |
| 17p11                | HPC2  | Very aggressive clinical evolutionary forms |
| 20q13                | HPC20 | Associated with brain tumors |
| 7q32                 |       | Autosomal dominant inheritance |
| 19q12-13             |       | Associated with brain tumors |
| 1p36                 | CAPB  | Very aggressive clinical evolutionary forms association with colorectal, breast, ovary, and urinary bladder cancers |
| 10q25                | BRCA1 | Autosomal dominant inheritance |
| 8p22-23              | PG1/MSR1 | Autosomal dominant inheritance |
| 17q24                | ELAC2 | Very aggressive clinical evolutionary forms association with colorectal, breast, ovary, and urinary bladder cancers |
| 16q23                |       | Autosomal dominant inheritance |
| 8q24 (region 1)      |       | Autosomal dominant inheritance |
| 8q24 (region 2)      |       | Autosomal dominant inheritance |
| 8q24 (region 3)      |       | Autosomal dominant inheritance |
| 2p15; 3p12; 6q25     |       | Autosomal dominant inheritance |
| 7p21; 10q11; 10q26   |       | Autosomal dominant inheritance |
| 11q13; 17q12         |       | Autosomal dominant inheritance |
| 17q21; 13q12-13      |       | Autosomal dominant inheritance |

Adapted from Tortatiada and castell.

Genetic conditions

BRCA1 & BRCA2 mutations

Many hereditary mutations but most of the notable is the BRCA2 gene has been identified with a major risk factor of PCa. BRCA1 and BRCA2 are both homologous recombination proteins. Ashkenazi Jewish populations are much more affected with these proteins mutation and commonly related with more chances of ovarian and breast cancer. However, the men with the age over 65 are at more risk by 8.6-fold with the BRCA2 mutation prostate cancer and 2.64-fold more among all the aged individuals. But the lower risk is associated with BRCA1. A cohort study illustrated that the men with BRCA2 transformation have a greater threat up to 5 times more with prostate cancer meanwhile, the risk of prostate cancer with BRCA2 change contrasted and the overall public among the individual under 65 years of age is more than 7 times higher. Prostate tumor threat might be
greater in men with BRCA1 variation, but still not confirmed. The individuals with BRCA2 carriers at the age of 40 are suggested for prostate cancer screening. In BRCA mutant breast and ovarian cancers, Poly-ADP Ribose Polymerase (PARP) inhibitors are used due to synthetic fataly, are recently go through clinical experiments for BRCA mutant PCs proteins. Prostate malignancy risk may be increased due to some other hereditary mutations which are under examination. Lynch syndrome

A cohort study illustrated that the risk of prostate cancer in men with Lynch syndrome is 2.1-4.9 times higher.

Endogenous hormones

Insulin-like Growth Factor-1 (IGF-1)

Mitotic and anti-apoptotic effects are associated with a polypeptide known as the Insulin-like growth factor (IGF-1). It is involved in both anti-apoptotic and mitogenic processes in prostate cancer cell lines and also plays a major role in its biology. Meta and pooled analysis have indicated prostate tumor risk is 38-83% more in men with the increased abnormal level of insulin-like development element 1 (IGF-1). Meta and pooled analysis have also indicated that prostate tumor risk is not associated with insulin-like development element 2 (IGF-2) levels. Furthermore, Meta and pooled analysis have also indicated prostate disease risk is by and large not associated with insulin-like development variable typing protein (IGFBP) levels, but this may fluctuate in between IGFBPs. It was observed that increased circulating IGF-I accumulation is positively connected with the danger of PCa over the short and long period in the European population therefore circulating IGF-I accumulation is connected with a higher risk for prostate cancer.

Testosterone

International Agency for Research on Cancer (IARC) reported the appropriate explanation of the prostate tumor that androgenic steroids have significant impacts on testosterone production in the body, taking into description limited proof. As prostate tumors depend on testosterone to develop, the Prostate growth treatment can involve manipulation of solutions or surgery to reduce the testosterone production levels.

Ionizing Radiations

Thorium-232 and its decay items, X radiation, and gamma radiation are arranged by the International Agency for Research on Cancer (IARC) as likely explanations of prostate cancer, in light of restricted proof. The prostate malignancy danger is higher in nuclear bomb survivors contrasted and the all-inclusive community, an accomplished study has demonstrated.

Obesity

During the epidemiological evaluation, body mass index is used to measure the overall obesity and abdominal obesity is measured by the ratio of waist to the buttocks edge. Among some of the recognized adoptable risk factors, obesity is well characterized for prostate cancer. Obesity has been involved in the deregulation of the insulin level, oxidative stress of DNA impair and inflammatory cytokine signaling, enlarging the threat of different neoplasm, including colorectal, breast and prostate cancer. Moreover, the elevated serum concentration of insulin is also associated with the risk of prostate cancer. Therefore, a higher concentration of serum leptin which is the product of the obesity gene Ob has been connected to immense tumor volume (>5 cm). Male individuals with type I obesity was identified with a 20% increased risk of PCa death ratio, while type II obese individuals with 34% of higher risk. One of the important particular metabolic results of obesity is the combination of physical lethargy, which ultimately causing lower tissue feedback to insulin, particularly in the term of lower absorption of glucose. This insulin resistance gives on to a chronically higher concentration of blood insulin, which is a growth-stimulating hormone and therefore it’s a possible risk factor to developing progression and cancers.

Smoking

Different types of mutagens present in cigarettes increased the production of tumorigenesis of prostatic epithelial cells, which is the possible risk factor to causing tobacco smoking injurious for health by developing prostate cancer disease. According to the International Agency for Research on Cancer (IARC), tobacco and cigarette smoke consist of more than 4,000 chemicals in which more than 60 are recognized as class 1 or class 2 carcinogens. Therefore, different ingredients of cigarette smoke such as polycyclic aromatic hydrocarbons (PAH), requisite metabolic stimulation, evasion of detoxification mechanism and successive attachment with DNA to apply their carcinogenic action. However, functional polymorphisms in genes that take part in PAH metabolism and detoxification may alter the consequence of smoking on prostate cancer. The correlation with smoking may also have a hormonal background, it was diagnosed male smokers have a higher level of testosterone, roaster one and circulating, which may cause an increased risk of prostate cancer or cancer advancement. The recent studies described that young age men smokers diagnosed with prostate cancer have an OR of 1.4, while an OR of 1.6 was observed in smokers with the age of above 40 years. One more study suggested that the risk of mortality among smokers was 1.6× more associated with prostate cancer. Smoking termination was observed to have an appropriate impact on prostate cancer extent and mortality, with a higher impact as time since termination increased.

Diet

Ecologic observations have expressed a strong association between the occurrence of prostate cancer and dietary fat uptake. A western diet, particularly excessive in fat, has been connected to increase the risk of prostate cancer by enhancing the production and accessibility of both estrogen and androgens, while Asian and vegetarian diets have low-fat content which is connected with less circulating concentration of hormones. The most intentional nutritional risk factor for prostate cancer is the excessive uptake of fat content. The role of the total, saturated, and animal fat has been observed in most epidemiological studies. The results of these studies suggested that monounsaturated, saturated and animal fat have a positive correlation and omega-3 fat has a negative correlation with prostate cancer. The findings for polyunsaturated fat are less reconcileable. The dietary products high in dairy items and calcium with low in selenium and alpha-tocopherol have been observed to raise the chances of PCa. One of the meta-analyses about the Prostate Cancer SIR, in which 4 of 15 observations were revealed a significant positive correlation between dairy and PCa risk, while 5 of 15 observations were, reported a 7% higher risk per 400 g of dairy products per day. They also reported a significantly higher risk with calcium intake about 13 of 15 studies, with 5% higher risk per 400 mg per day about 15 of 16 studies. Calcium is identified to deregulate the effective form of vitamin D3.
which ultimately causing the excessive growth of prostatic cells. Meanwhile, in the same studies, it was also observed low uptake of selenium and alpha-tocopherol was significantly associated with increased risk of PCa among 2/10 and 2/11 observations, respectively. According to the meta-analysis, Beta-carotene a precursor of vitamin A was observed to have no reasonable impact on prostate cancer risk 63.

Sexually transmitted diseases

Prostate cancer also has an association with sexually transmitted diseases (STDs). Prostate cancer risk about 2-3 fold correlated with STDs specifically with gonorrhoea and syphilis infections recently suggested one large population-based study 64. Human papillomavirus-16, -18, and -33 diagnoses have been also shown a higher risk of PCa suggested some other studies 65. Meanwhile, the duration of HIV infection was connected with a higher risk of prostate cancer, reported a study based on human immunodeficiency virus (HIV) infected population 66. A current meta-analysis of 17 observations expressed that an increased ratio of sexual partners is correlated with higher PCa risk, especially through the more activities of sexually transmitted infections 67. However, the contraposition is not clear; bacterial or viral pathogens through sexual activities have been involved in the initiation of chronic inflammation of the prostate which possibly becomes the source of prostate cancer.

Vasectomy

In the United States, vasectomy is the most frequently adapted procedure with approximately 500,000 achieved 68. But there is not an exact biological procedure that might illustrate the correlation between vasectomy and prostate cancer has been recognized. Different researches indicating low relative danger; data were negligible by methodological limitations and potential intolerance, including diagnosis and misclassification bias annually. In some studies, it has been related to a higher risk of PCa 69.

Other factors

Many other kinds of risk factors are associated with prostate cancer such as diabetes, physical activity, alcohol consumption, profession and hepatic cirrhosis have been observed, but their role is less or indefinite in prostate cancer which depends on the scientific knowledge in the present literature 70.

Conclusion

Globally prostate cancer is the second most familiar and fifth-most hostile neoplasm among male individuals. It is the disease of the male genital system and particularly the abnormal growth of the prostate gland. According to the epidemiological studies different environmental and genetic factors are associated with the progression of abnormal prostate cell growth which ultimately causes the development of cancerous cells. Prostate cancer is specified by vast fluctuations in prevalence and mortality in the world. Epidemiological surveillance provides crucial indications to the etiology of PCa. In the advancement of molecular techniques, identification of the individual as well as the combined consequences of these factors has been initiated by a new generation of a wide range of population-based studies. Such studies may provide significant outcomes for risk factors which may be effective in recognizing the more adaptable subcategory of the population to observing prostate cancer.

Authors Contribution

All the authors contributed equally to the collection of data and to help in the formatting of the manuscript. All the authors support technically and responsible for manuscript reviewing, data analysis.

Conflict of Interest

All the authors declare no conflict of interest.

References

1. Skolarus TA, Wolf AM, Erb NL, et al. American Cancer Society prostate cancer survivorship care guidelines. Cancer Journal for Clinicians. 2014; 64(4):225-249.
2. Seisen T, Roupert M, Faix A, Droupy S. The prostate gland: a crossroad between the urinary and the seminal tracts. Progr en urologie: journal de l'Association francaise d'uropolegie et de la Societe francaise d'urologie. 2012; 22:S2-6.
3. Gunha GR. Role of mesenchymal-epithelial interactions in normal and abnormal development of the mammary gland and prostate. Cancer. 1994; 74(5):1050-1044.
4. McNeil JE. Origin and evolution of benign prostate enlargement. Investigative Urology. 1978; 15(4):340-345.
5. Hayward SW, Gunha GR, Dahia R. Normal Development and Carcinogenesis of the Prostate: A Unifying Hypothesis. Annals of the New York Academy of Sciences. 1996; 784(1):50-62.
6. Ferlay J, Soerjomataram I, Dikshit R et al. Cancer incidence and mortality worldwide: sources, methods and major patterns in GLOBOCAN. International Journal of Cancer. 2015; 136(5):E359-E386.
7. Jain D, Chaudhary P, Varshney N, Jameda P. Carcinogenic effects of Nitrosro compounds in the environment. Environment Conservation Journal. 2020; 21(3):25-41.
8. Favoriti P, Carbone G, Greco M, Pirozzi F, Pirozzi REM, Corcione F. Worldwide burden of colorectal cancer: a review. Updates in Surgery. 2016; 68(1):7-11.
9. Langan RC. Benign prostatic hyperplasia. Primary Care: Clinics in Office Practice. 2019; 46(2):223-232.
10. McVary KT, Roehrborn CG, Avins AL, et al. Update on AUA guideline on the management of benign prostatic hyperplasia. The Journal of Urology. 2011; 185(5):1793-1803.
11. Habib A Anjum KM, Ashraf Z, et al. Global Epidemiology of COVID-19 and the Risk of Second Wave. Journal of Drug Delivery Therapeutics. 2021; 11(2):188-193.
12. Mahmood S, Qasmi G, Ahmed A, et al. Lifestyle factors associated with the risk of prostate cancer among Pakistani men. Journal of Ayub Medical College Abbottabad. 2012; 24(2):122-126.
13. Grasso CS, Wu V-M, Robinson DR, et al. The mutational landscape of lethal castration-resistant prostate cancer. Nature. 2012; 487(7406):239-243.
14. Helfand BT, Catalonia WJ. The epidemiology and clinical implications of genetic variation in prostate cancer. The Urologic Clinics of North America. 2014; 41(2):277-297.
15. Ewing CM, Ray AM, Lange EM, et al. Germline mutations in HOXB13 and prostate-cancer risk. New England Journal of Medicine. 2012; 366(2):141-149.
16. Tolumis SA, Laxman B, Varambally S, et al. Role of the TMPRSS2-ERG gene fusion in prostate cancer. Neoplasia. 2008; 10(2):177-180.
17. Carter BS, Ewing CM, Ward WS, et al. Allelic loss of IN179. Neoplasia. 2008; 10(2):177-180.
18. Xue J, Meyers D, Freije D, et al. Evidence for a prostate cancer susceptibility locus on the X chromosome. Nature Genetics. 1998; 20(2):175-179.
19. Anderson DE, Badzioch MD. Breast cancer risks in relatives of male breast cancer patients. Journal of the National Cancer Institute 1992; 84(14):1114-1117.
20. Schaid DJ. The complex genetic epidemiology of prostate cancer. Human Molecular Genetics. 2004; 13(suppl_1):R103-R121.
21. Huang H, Cai B. GBAC4 mutation in HOXB13 is firmly associated with prostate cancer risk: a meta-analysis. Tumor Biology. 2014; 35(2):1177-1182.
22. Jain D, Chaudhary P, Kotnala A, Hossain R, Biecht K, Hossain MN. Hepatoprotective activity of medicinal plants: A mini review. Journal of Medicinal Plants Research. 2020; 14(2):185-188.

23. Dasgupta P, Baede PD, Attkin JF, Ralph N, Chambers SK, Dunn J. Geographical variations in prostate cancer outcomes: a systematic review of international evidence. Frontiers in Oncology. 2019; 9:238.

24. Rawla P. Epidemiology of prostate cancer. World journal of oncology. 2019; 10(2):63.

25. Grossman DC, Curry SJ, Owens DK, et al. Screening for prostate cancer: US Preventive Services Task Force recommendation statement. The journal of the American Medical Association. 2018; 321(18):1901-1913.

26. Howlader N, Noone A, Krapcho M, et al. Cronin KÄe. SEER Cancer Statistics Review. 1975:2010.

27. Habib A, Jaffer G, Khalid M. Letter to Editor: Challenges in the Development of Hepatitis C Vaccine. International Journal of Biomedical Investigation 2020; 3:127.

28. DeSantis CE, Siegel RL, Sauer AG, et al. Cancer statistics for African Americans, 2016: Progress and opportunities in reducing racial disparities. Cancer Journal for Clinicians. 2016; 66(4):290-308.

29. Barouk A, Padala SA, Vakiti A, et al. Epidemiology, staging and management of prostate cancer. Medical Sciences. 2020; 8(3):28.

30. Robbeek TR, Devesa SS, Chang B-L, et al. Global patterns of prostate cancer incidence, aggressiveness, and mortality in men of african descent. Prostate Cancer. 2013; 2013.

31. Wu L, Modlin LS. Disparities in prostate cancer in African American men: what primary care physicians can do. Cleveland Clinic Journal of Medicine. 2012; 79(5):313-320.

32. Zeegers MP, Jelena A, Ostrea H. Empiric risk of prostate carcinoma for relatives of patients with prostate carcinoma: A meta-analysis. Cancer: Interdisciplinary International Journal of Cancer. 2009; 1177:2014-2015.

33. Xu J, Meyers D, Freije D, et al. Evidence for a Prostate Cancer Susceptibility Locus on the X Chromosome. The Journal of Urology. 1999; 161(4):1386-1386.

34. Carpten J, Nuppenon N, Isaacs S, et al. Germline mutations in the ribonuclease L gene in families showing linkage with HPC1. Nature Genetics. 2002; 30(2):181-184.

35. Zhou A, Paranjape J, Brown TL, et al. Interferon action and apoptosis are defective in mice devoid of 2′, 5′‐oligoadenylate‐dependent RNase L. The EMBO journal. 1997; 16(21):6355-6363.

36. Erkko H, Xia B, Nikkilä J, et al. A recurrent mutation in PALB2 in Finnish cancer families. Nature. 2007; 446(7133):316-319.

37. Ferris‐Tortajada J, García‐Icastizá J, Berbel‐Torner O, Ortega‐García J. Constitutional risk factors in prostate cancer. Actas Urológicas Españolas. 2011; 35(5):235-238.

38. Oh M, Alkhushayn M, Fallahat S, et al. The association of BRCA1 and BRCA2 mutations with prostate cancer risk, frequency, and mortality: A meta-analysis. The Prostate. 2019; 79(8):880-895.

39. Kibel AS, Schutte M, Kern SE, Isaacs WB, Bova GS. Identification of 12p as a region of frequent deletion in advanced prostate cancer. Cancer Research. 1999; 59(24):5652-5655.

40. Habib A, Razzaq KSB, Imran M, Khalid B. Elimination of Hepatitis C Virus: A Goal of WHO. Vigan Varta. 2020; 1:56-59.

41. Cuzzick J, Thorat MA, Andriole G, et al. Prevention and early detection of prostate cancer. The Lancet Oncology. 2014; 15(11):484-492.

42. Ashworth A, Lord CJ. Synthetic lethal therapies for cancer: what’s next after PARP inhibitors? Nature Reviews Clinical Oncology. 2018; 15(9):564-576.

43. Kicinski M, Vangronsveld J, Nawrot TS. An epidemiological reappraisal of the familial aggregation of prostate cancer: a meta-analysis. PloS One. 2011; 6(10):e27130.

44. Kleihues P, Schaufel B, zur Hausen A, Esteve J, Ohgaki H. Tumors associated with p53 germ line mutations: a synopsis of 91 families. The American Journal of Pathology. 1997; 150(1):3.

45. Gennings C, Menetrier-Caux C, Droz J. Insulin-like Growth Factor (IGF) family and prostate cancer. Critical reviews in Oncology/Hematology. 2006; 58(2):124-145.

46. Liu X, Wang P, Fu J, et al. Two-photon fluorescence real-time imaging on the development of early mouse embryo by stages. Journal of Microscopy. 2011; 241(2):212-218.