



EMERGING CANCER MODELS FOR DRUGS AND NOVEL DOSAGES DEVELOPMENT

P. DAS, M. K. DAS

Department of Pharmaceutical Sciences, Dibrugarh University, Dibrugarh, Assam, India

Abstract – Objective: Cancer is the primary cause of death all over the world, despite of its early detection and novel therapies available. The process of cancer drug development is growing tremendously. With respect to many limitations in human studies, there is the need to develop experimental models for screening of efficacy and toxicity of developing drugs. The purpose of this study is to investigate the cancer model for drug development.

Materials and Methods: In this review, articles are extracted with selected keywords from the PubMed, SID, Springer, Medlib, Web of Science (Clarivate) and ScienceDirect databases without any language restrictions.

Results: The efforts to reduce global cancer burden are mainly focused on developing innovative diagnostic and therapeutic tools. A plenty of in vitro and in vivo models are utilized in cancer drug discovery process. Successful bench-to-bedside translation of basic scientific findings about cancer into therapeutic interventions for patients depends on the selection of appropriate experimental models for testing.

Conclusions: The presented models have both advantages and disadvantages. None of them are absolutely ideal while they are chosen according to the purpose of the investigator and the advantages of each method for cancer research.

KEYWORDS: Animal models, Cancer, In vitro and in vivo models, Clinical translation.

INTRODUCTION

Cancer is an utmost serious malignant disease in today's world which represents uncontrolled growth and spread of abnormal cells¹. Due to the advancement in medical science the mortality rates related to many different illnesses have declined in current decades, but cancer-associated deaths have remained surprisingly constant². Recent worldwide data from the World Health Organization (WHO) Global Cancer Observatory (GLOBOCAN) estimate that till 2018 there is an approximately 18.1 million of new case and deaths of 9.6 million occur because of cancer^{3,4}. The major problem of cancer is the lack of its early detection testing which corresponds to diagno-

sis of disease in more complex condition that may enhance death rate. Almost 50% cases of all cancer types are diagnosed at distant stages which results meagre survival of patients. The pathway of developing advanced cancer diagnostics and therapeutics is more adventurous, it is both lengthy as well as highly priced⁵. A variety of therapies and strategies has been flourished to fight against this deadly disease such as systemic chemotherapy, radiation therapy and also more targeted approaches including immunotherapy and antibody-based therapies. But still in a petty number of new drugs have received FDA approval which shows that there may be something missing in the way of discovering new agents and testing protocol of these agents prior to clinical trials⁶.



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Early detection of tumours and accurate monitoring of responses during treatment are pivotal to patient survival. For upgrading the detection and productive treatment of early cancer there is a need to discover specific molecular targets for tumours. Thus, the appropriate use and development of *in vitro* and *in vivo* cancer models is extremely desirable⁷. Drug discovery is a monotonous process and encompasses a high failure rate. The new compounds shown to be effective in laboratory *in vitro* cell culture system might not be able to reproduce their activities in animal tumor models. Albeit they are acting in animal tumor systems most of these drugs are not very effective in humans⁸. The animal models used in the preclinical research must truly reflect the microenvironment of tumour in human. Additionally, the therapeutic effectiveness and toxicity profile of the agent at the specific doses that may be rationalized to human ought to be thoroughly investigated. Often, many new small molecules are tested at doses that do not show any significant pharmacological affect. There is a need for better design of preclinical animal model studies in order to enhance the success rate of new drugs introduced into clinical trials⁹.

The discovery of anti-cancer drugs continuous to develop at an alarming rate, and a large amount of recourses are affianced for drug discovery and design. The evaluation of designed drug molecule is a key step in drug discovery program. Inappropriate selection of evaluation method may result in potential drugs being excluded from further development¹⁰. The use of cell lines with high throughput screening is the primary screening method, but due to limitations such as low relevance to clinical condition there is an additional

need to use appropriate *in vivo* model for screening. The selected animal model should reflect the human tumour microenvironment as closely as possible. At the same time, their feasibility and cost efficiency for large scale drug screening programs are also important factors¹¹. The most common cancers leading to cause of death globally are depicted in Figure 1.

Animal model for cancer

Animal models are indispensable tools for investigating the cancer pathogenesis, tumor invasion and metastasis mechanisms and also for new therapeutic approaches. Various types of *in vitro* and *in vivo* tumor models have been developed to understand the biological changes associated with tumorigenesis. These models are also integral in drug development¹². The development of *in vivo* animal models that mimic the natural history of human cancers and its clinical response to treatment is an important prerequisite for rapid translation of anticancer therapies¹³.

In-vitro models for cancer

In vitro cancer research utilized animal and human cancer cell lines to reveal biochemical pathways associated with cancerous cells¹⁴. These cancer cell lines are derived from high-grade, high-stage cancers. At present, clinical trials are the eventual determinant of drug efficacy but due to limited ethical and safety considerations, pre-clinical studies using *in vitro* tumor models are

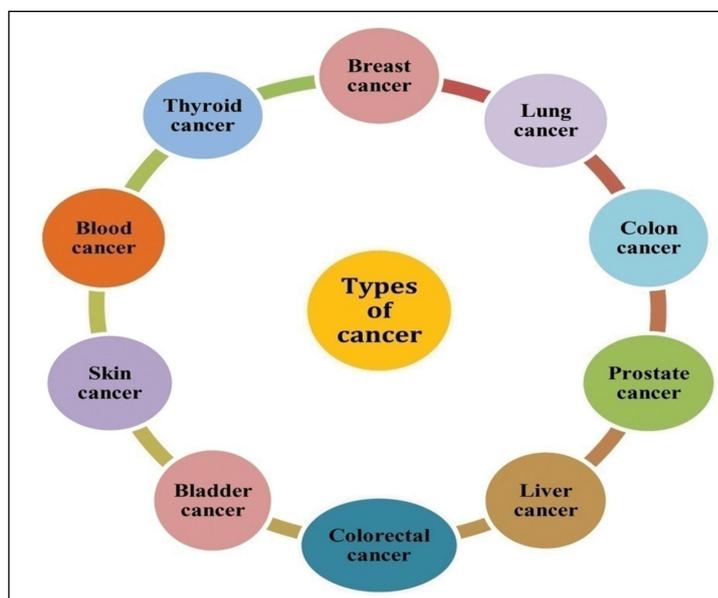


Figure 1. Types of most common cancer which lead to increasing death rate globally.

particularly important. The advantages of *in vitro* cancer models are its highly controlled conditions, homogeneity, revealing underlying molecular mechanisms, and reproducibility¹⁵. *In vitro* human cancer models have varying degree of complexity, ranging from 2D single-layer models to 3D tumor models. *In vitro* tumor models can be developed from patient cells, cell lines, stem cells, stromal cells, immune cells, and expose to various physical and chemical conditions according to experimental requirements¹⁶. With the help of *in vitro* tumor models one can explore the important cellular and molecular mechanism associated with carcinogenesis including survival, proliferation, migration, invasion, matrix remodelling, latency, and angiogenesis¹⁷. The limitations of two-dimensional *in vitro* model are selection of phenotypic and genotypic cells in the *in vitro* adaptation process, a homogeneous cell population, accumulation of cell mutations over time during the culture process, and separation of cells in the tumor microenvironment¹⁸. Therefore, in order to overcome the limitations of 2D cells, a 3D *in vitro* cancer model is proposed as an alternative method, which is able to accurately simulate some characteristics of solid tumors, such as their spatial structure, physiological responses, soluble mediators secretion, pattern of gene expression and mechanism of drug resistance^{19,20}. The 3D model over 2D is depicted in Figure 2.

Here we discuss some of the models that are mostly used such as spheroids, scaffolds, organoids, and microfluidic devices. They are also used in combination with another such as scaffolds are often placed inside microfluidic devices or spheroids can be placed inside scaffolds.

a) Spheroids: Three-dimensional (3D) spheroid models contain complex structures composed by one or more cell type. When these cell types are grown and large enough, they form a necrotic cell nucleus surrounded by a layer of actively

proliferating cells²¹. The complex formed also includes an extracellular matrix (ECM) composed of the produced protein²². These models are very similar to tumors *in vivo* because they have similar gene expression, growth kinetics, and cell heterogeneity; in addition, the presence of natural ECM allows the simulation of natural barriers for drug penetration studies, and the internal structure of the spheroids has also great *in vivo* environment mimicking ability, where the internal region appears necrotic due to lack of nutrients and oxygen²³.

Several methodologies proposed to produce 3D spheroids are as follows^{24,25}:

- I) Spinner flask technique: stir the cell suspension continuously to promote the formation of cell aggregates.
- II) Liquid overlay technique: cells grow on an on-stick surface and promote aggregation instead of adhering to the surface of the flask.
- III) Hanging drop technique: the cells grow in the form of droplets of culture medium on the lid of the petri dish, the lid is turned upside down and placed on the petri dish below, and then the cells are cultured until the sphere reaches the desired size.

b) Scaffold based model: The three-dimensional scaffold-based model is based on the growth of cells in a three-dimensional structure, and in some respects mimics the extracellular matrix²⁶. The structures used in the model can be made of natural (e.g., collagen), semi-synthetic (e.g., chitosan), or synthetic (e.g., polycaprolactone) biomaterials. In addition, cells can be cultured on prefabricated/synthesized structures, or 3D bio-printing²⁷. The use of structural elements makes it possible to simulate tiny and complex shapes, and at the same time, the simulation of ECM produces a certain resistance to drug penetration, similar to tumors in the body²⁸. In addition, these models usually have the following

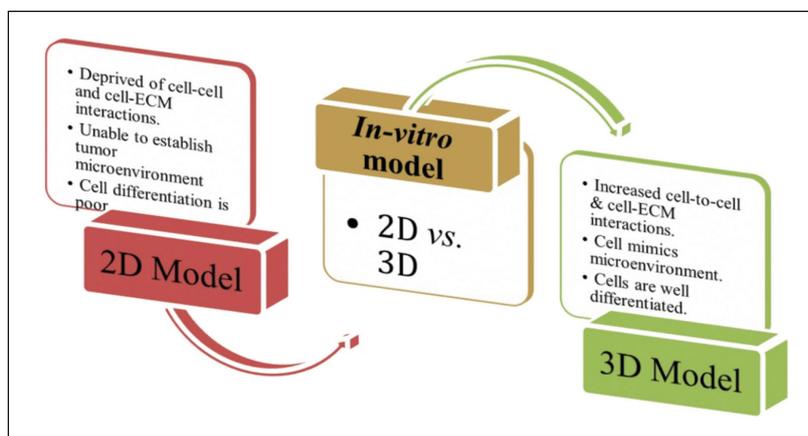


Figure 2. *In-vitro* 2D vs. 3D cell culture. 3D models have several advantages over 2D models. Thus, 3D models are preferable for cancer drug screening.



characteristics: spontaneous cell organization, possible heterogeneity, gene expression, and tumor-like cell phenotype *in vivo*. The main obstacle of this model is the use of artificial ECM structure. In addition, it also requires the use of highly biocompatible cell-containing building materials. Because of the bioprinting structure, it must be printed efficiently²⁹.

- c) Organoids based model:** Organoid tumor model is a new method of tumor research that has emerged in recent years. Organoids are derived from stem cells and can mimic all aspects of organ structure and function, including differentiation into different cell types³⁰. Organelles can be cultured for a long time (especially for patient-specific tumor cells and establishment of *in vitro* cell banks) and transferred under three-dimensional conditions. In addition, the organelles can retain the histology, immunohistochemistry, and genetic heterogeneity of the original tumor tissue, which makes it suitable for high-throughput drug screening³¹. Although this is a relatively new approach, these two organelles derived from normal tissue stem cells, and those derived from cancer stem cells, have made significant contributions to cancer biology and personalized medicine³². Organoid models have been developed for many different normal tissue types (for example, gut, salivary glands, mammary glands and liver), and an increasing number of cancer types (including for example breast, prostate, pancreatic, colon, bladder and gastrointestinal cancers)³³. The only disadvantage of this model is the lack of interstitial cell components and the absence of interactions between cells³⁴.
- d) Microfluidic models:** Microfluidics has become a powerful platform for cancer metastasis research and drug discovery³⁵. Microfluidics technology has completely changed our ability to simulate the natural biophysical/chemical conditions of cells in *in vitro* models. The goal of microfluidics is to create a platform that can simulate the pathophysiological functions of tissues and organs, so-called “organ-in-chips.” These platforms are composed of a network of microfluidic channels that can cultivate contin-

uously perfused cells³⁶. The main advantage of microfluidics is the ability to create complex three-dimensional culture systems in which various parameters can be changed and controlled. A variety of cell types can be grown on microfluidic chips to analyze specific interactions, which is particularly interesting when studying the relationship between cancer and stromal cells³⁷. Microfluidic devices are usually made of polydimethylsiloxane (PDMS), which has many outstanding properties such as, due to its transparent nature *in vivo*, real time imaging, but requires new materials to fabricate the chips because PDMS can absorb small molecules in a non-specific manner^{38,39}. There are several challenges in using this method to translate results into the clinic, including validating *in vitro* function in *in vivo* tumor models and correlating the results with patient samples⁴⁰.

Some advantages and disadvantages of *in-vitro* model are discussed in Table 1.

In-vitro evaluation analysis

A great number of *in vitro* methods has been developed to measure the efficiency of anticancer compounds. The different parameters analyzed by different assay/detection methods are described below.

I. Cell viability/cytotoxicity determination: Different methods used for detection of cell viability are:

- a) MTT assay:** MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] assay is used to measure cellular metabolic activity as an indicator of cell viability, proliferation, and cytotoxicity. MTT assay is based on the conversion of the yellow tetrazolium salt-MTT to purple-formazan crystals by metabolically active cells and quantitative determinations of viable cells. Cells are seeded in 96 well plates at 5000 cells/well and allowed to grow in incubator at 37°C with 5% CO₂ for 24 h. The medium is then removed and replaced by fresh medium containing different concentrations of sample and then they are incubated at same

TABLE 1. Advantages and disadvantages of *in vitro* model in cancer research.

Model	Advantages	Disadvantages
<i>In vitro</i> model	<ul style="list-style-type: none"> – Cheap, relatively easy, and fast maintenance – Wide range of cell lines available – Highly controlled experiment – High degree of similarity with initial tumor – Less ethical containments 	<ul style="list-style-type: none"> – Less mimic tumor environment – Different expression of genes and markers – Less sensitivity to same environmental features – Inevitable mutations of certain cellular features – Genomic instability

condition for 24-48 h. Then, 20 μ L MTT dye stock solution (5 mg/ml in PBS) is added to each well and incubated for 2-3 h. The medium is removed and 200 μ L dimethyl sulfoxide (DMSO) is added to each well to dissolve the MTT metabolic product. Then the plate is shaken at 150 rpm for 5 min and the optical density is measured at 560 nm. Untreated cells (basal) are used as a control of viability (100%) and the results are expressed as % viability (log) relative to the control^{41,42}.

b) LDH (Lactate dehydrogenase) Assay: Another method for determining cytotoxicity is based on measuring the activity of cytoplasmic enzymes released by damaged cells. LDH is a stable cytoplasmic enzyme found in all cells and it is rapidly released when the plasma membrane is damaged, a key feature of cells undergoing apoptosis, necrosis, and other forms of cellular damage. LDH activity can be easily quantified by using the NADH (nicotinamide adenine dinucleotide + hydrogen) produced during the conversion of lactate to pyruvate. The protocol includes: the cell cultures suspension is centrifuged for 4 min, and then transfer 50 μ L of the supernatant into a 96-well plate. Add 50 μ L of LDH assay substrate to the medium. Cover the plate with foil or a small opaque box to protect it from light and incubate at 37°C for 15-30 min. Then add 100 μ L of Stop solution (50% dimethylformamide and 20% Sodium dodecyl sulphate at pH 4.7). Measure the absorbance at 490 nm within 1 h of adding the Stop solution. Set the background absorbance at 690 nm and subtract this value from the primary wavelength measurement (490 nm). The percentage of LDH released is calculated as percentage of the total amount, considered as the sum of the enzymatic activity present in the cellular lysate and that in the culture medium^{43,44}.

c) XTT assay: In order to measure the proliferation response, the (2,3-bis[2-Methoxy-4-nitro-5-sulphophenyl]-2H-tetrazolium-5-carboxyanilide inner salt (XTT) assay is used. The tetrazolium salt, XTT is especially useful in quantifying viable cells. This assay is based on the cleavage of yellow tetrazolium salt, XTT, to orange formazan dye by metabolically active cells. XTT cleavages into an orange formazan dye by the mitochondrial enzyme, dehydrogenase, occurs exclusively in living cells. Cells are grown in growth medium containing 10% fetal bovine serum (FBS) in 96-well plates until 70-80% confluence and then treated with drug sample. 50 μ L of XTT labelling mixture solution is added to each well

and the cells are incubated at 37°C for 4 h. The formazan dye formed is soluble in aqueous solutions and quantified optical density at 450 nm, compared with that of control wells with a screening multiwell spectrophotometer enzyme-linked immunosorbent assay (ELISA) reader. The reference wavelength should be more than 650 nm^{45,46}.

d) Sulforhodamine B (SRB) assay: Sulforhodamine B assay is a bright pink aminoxanthine dye that binds to basic amino acids in mild acidic conditions and dissociates under basic conditions. Cells are plated in 96-well-flat bottom plates at 5000-10000 cell/well. The difference in cell numbers plated adjusts for differences in the growth rates of the various cell lines. Cells are allowed to adhere to the wells overnight, then the samples are added to triplicate wells in serial 3-fold dilutions. Water is added to the control wells at a 1:10 dilution in medium. Plates are incubated in humidified atmosphere (37°C, 5% CO₂) for 3 days, and then assayed for growth inhibition using a Sulforhodamine B (SRB) assay. The cells are fixed with 10% (w/v) trichloroacetic acid and then stained with 0.4% SRB dissolved in 1% acetic acid for 15-30 min and subsequently washed with 1% acetic acid to remove excess stain. After the plates are air dried at room temperature, the bound dye is solubilized with 10 mM Tris base and the plates are analysed for optical density at 595 nm using a microplate reader^{47,48}.

e) WST-1 Assay: This assay is designed for evaluation of intracellular metabolic activity of cells. The reduction of the WST-1 into a water soluble formazan occurs by the NADH dehydrogenase and plasma membrane electron transport. Here, 10 μ L of WST-1 reagent is added to 100 μ L of media in a 96-well plate. Seeded cells are incubated overnight at 37°C with 5% CO₂ in a humidified atmosphere for 24 h. Then cells are treated with apoptotic inducers, cell proliferation triggers, or cytotoxic reagents and then 10 μ L/well of WST-1 is added and incubation is continued for an additional 1-2 h at same humidified atmospheric condition. The absorbance is measured at 450 nm vs. a 650 nm reference by using a plate reader^{49,50}.

f) Acid Phosphatase assay: It is the quantification of cytosolic acid phosphatase activity on the cells. The assay is based on hydrolysis of p-nitrophenyl phosphate to p-nitrophenol by the intracellular acid phosphatase enzyme. The absorbance is measured at 405 nm and absorbance at this wavelength is monitored as a measure of cell number^{51,52}.



II. Cell death/ Apoptosis: Apoptosis or programmed cell death is the process of cell death that occurs as a normal and controlled part of an organism's growth or development. Apoptosis takes place through a complex signalling mechanism and is dedicatedly balanced or regulated in a physiological context⁵³. Apoptosis can be detected by the following assay:

- a) **Trypan blue dry exclusion assay:** In this assay, cell suspension is mixed with trypan blue and visually examined to determine whether cells take up or exclude the dye. Viable cells have clear cytoplasm and non-viable cell have blue cytoplasm. The cell suspension is washed with buffered Salt Solution and centrifuged for 5 min at 10,000 rpm. Then resuspend the cell pellets in 1 mL PBS. The cells are exposed to drug dilutions and incubated at 37°C for 3 h. After 3 h, equal quality of the drug treated cells are mixed with trypan blue (0.4%) and left for 3 min at room temperature. Then a drop of trypan blue/cell mixture is loaded in a haemocytometer and viable and non-viable count are recorded within 2 min. Viable cells do not take up colour, whereas dead cells take up colour⁵³. The percentage of growth inhibition is calculated using the following formula:

$$\text{Growth inhibition (\%)} = 100 - \frac{(\text{Totalcells} - \text{Deadcells})}{\text{Totalcells}} \times 100$$

- b) **Live/Death staining:** It is done for identification and quantification of live and dead cells. The cells are stained with fluorescent dyes such as calcein-acetoxymethyl (calcein-AM), intercalating agent- propidium iodide (PI) and ethidiumhomodimer (EthD-1)). Live cells are stained in green following intracellular cleavage of the acetomethoxy group of calcein-AM. Dead cells are stained in red following penetration of the intercalating agents through their permeable membrane⁵⁴.
- c) **Annexin V-FITC staining:** It is used to detect apoptosis marker phosphatidylserine molecules which have translocated to the outside of the cell membrane. In normal cells, phosphatidylserine (PS, membrane phospholipids) is held on the inner layer of the cell membrane, so Annexin V does not attach to the cells. During early apoptosis, the PS is exposed on the outer layer, where they attach to the FITC (Fluorescein isothiocyanate)-labelled Annexin V and stain the cell surface green. During late apoptosis, propidium iodide (PI) enters the cell and stains the contents red. Analysis is done by flow cytometry or fluorescence microscopy⁵⁵.

In-vivo model of cancer

The development of an *in vivo* animal model that simulates the natural history of human cancers and its clinical response to treatment is an important prerequisite for laboratory to clinical transformation in which investigational cancer treatment and imaging technologies have shown promise in *in-vitro* models. There are different animal species which are used as an *in vivo* model for cancer screening. It is important to select the appropriate animal model to evaluate a specific cancer. Animal model should reflect the human disease as much as possible. At the same time its feasibility and expenses to be used in large drug screening programs is also important factors⁵⁶. Commonly animal models are divided in to two types (I) Spontaneous tumor model and (II) Induced tumor model. The models are depicted in Figure 3 and briefly discussed below:

I) Spontaneous tumor model

It includes the selection and use of animals that naturally occurs cancer. For example, some innate mouse and rat strains are particularly susceptible to various cancers, especially leukemia, breast cancer, adenoma and lung hepatoma⁵⁷. In DA/Han rats more than 60% of female animals die from endometrial adenocarcinoma. In BDII/Han rats, 87 to 90% of animals die of endometrial adenocarcinoma⁵⁸. In DA/Han inbred rats, 53.9% of males and 14.4% of females developed spontaneous urinary bladder tumors. These models mimic the clinical situation most closely. They are similar to human cancers in kinetics and antigenicity, but these systems have many limitations. It is impossible to obtain a sufficient number of tumors of the same size for screening at a time. Tumors can usually only be measured at the end of their development, but the mode of metastasis is not, and it is difficult to accurately determine homogeneity at this stage. These models are usually not replicable, and most of them are viral origin⁵⁹.

II) Induced tumor model: Cancer can be induced by various agents such as some chemicals and viruses can act as a cancer inducing agent. Different induced cancer models are discussed below in this section.

a) Chemically induced tumor model

Tumors caused by chemical carcinogens originate from the host's own cells and therefore resemble human clinical cancer more closely than transplantable tumors. The limitation of chemically induced tumors is that carcinogens may have an impact on tumor behaviour, and the excretion of carcinogens and their metabolites in animal feces and urine may cause danger to other animals and personels^{60,61}.

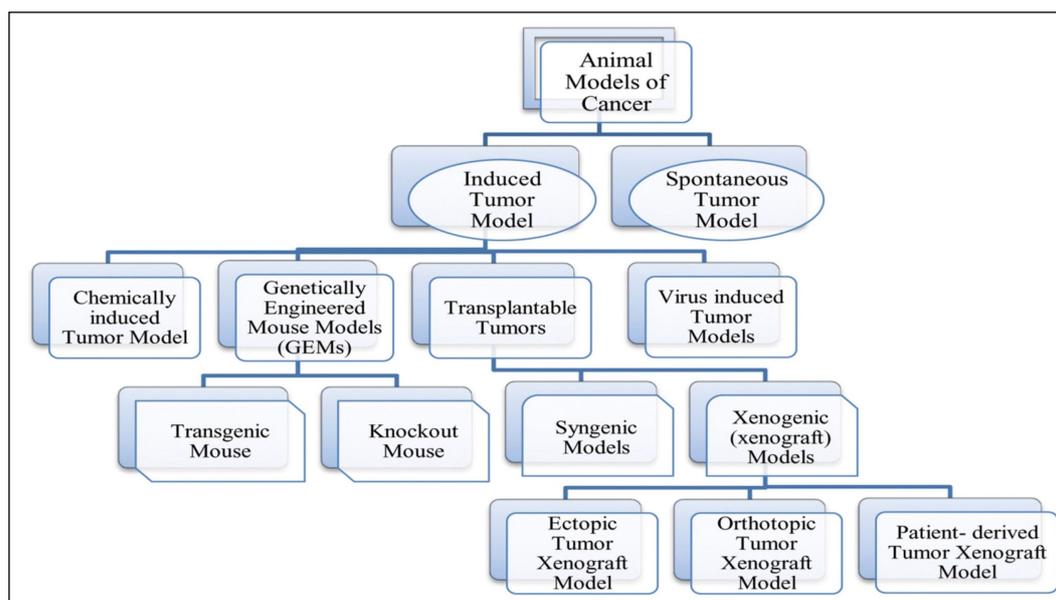


Figure 3. Commonly used animal models in cancer. The models portrayed here are mostly used to determine safety and efficacy in drug and dosage development process.

Chemical carcinogens can be divided into two categories:

- i. Direct acting agents: require no chemical transformation to induce carcinogenicity.
- ii. Indirect acting agents: become active only after metabolic conversion. Also known as procarcinogens and their active end products are called ultimate carcinogens.

Although any gene may be the target of chemical carcinogens, RAS gene mutations and TP53 genes are the important targets. Few examples of chemical carcinogenesis are mentioned below:

Urinary bladder cancer induced using *N*-butyl-*N*-(4-hydroxybutyl)nitrosamine (BBN)

Mice are treated with BBN at 500 ppm in drinking water, after 2-6 weeks mutagenesis in urothelial cells are greater than the spontaneous mutation background and that in the smooth muscle cells of the urinary bladder; after 20 weeks of BBN treatment at the same concentration, bladder cancer developed in all treated mice. The detailed histopathological and immunohistochemistry examination of tumour is valuable in determining the drug action⁶².

Colon cancers induced using DMAB (3, 2-dimethyl-4-aminobiphenyl)

These tumours occurred after subcutaneous injection of the carcinogens for 6 consecutive weeks. It

can cause multiple colon tumours in approximately 26-30% of animals fed on modified fatty diet. Induced tumours are adenoma (benign tumours) and adenocarcinoma (malignant tumours) in the large bowel. This model provides good information about tumour induction and the shape, size and histological properties of tumour tissue. The only drawbacks of this model is that it requires multiple injections of carcinogens and form secondary tumours in other areas such as skin, mammary glands, stomach, salivary glands, lymph nodes, ear canal and urinary bladder, which will affect the effectiveness of the drug and survival rate and make it difficult to compare⁶³.

Fibrosarcoma Model in Mouse Using 3,4,9,10-dibenzopyrene

Single subcutaneous injection of carcinogen is sufficient to form uniform subcutaneous tumours at the site of injection within 30-40 days post injection. The most important advantage of this model is induction of tumour with single dose of carcinogen and no excretion of carcinogen through faeces or urine.

b) Genetically Engineered mouse model (GEM)

Cancer originates from genetic changes, and hence the animals genetically engineered depict cancer scenario in humans better than other animal models⁶⁴. Compared with tumor models with subcutaneous xenograft, these animals showed spontaneous development of cancer in their natural anatomical site. In GEM, tumours



show natural progression and metastatic biology similar to human counterpart. Importantly these tumours are originated within self; hence, there is no need for immunosuppressed animal to grow them⁶⁵. GEM animals include (i) transgenic and (ii) knockout mice.

(i) **Transgenic mouse model:** These animals are the result of introducing foreign genes into the pronucleus after the eggs are fertilized. Later offspring developed from this modified egg vector and expresses the foreign gene which was inserted and passes to further offspring. The gene to be studied can be introduced into the pronuclei by microinjection, retroviral vector or embryonic stem cell (ESC) transfer, which has proven to be an excellent animal model system for studying oncogenic phenotype, resulted from the dysregulation of a known gene⁶⁶. One can study the characteristics of oncogenes such as TP53, C-myc, E2F1, retinoblastoma (RB), neurofibromin 1 (NF1) and others in these animals. Due to known pathway defects, these animals expressing oncogene will develop spontaneous tumours, so it is most suitable to evaluate drugs/molecules specifically targeting these molecular signalling pathways in transgenic mouse models⁶⁷. Some examples of transgenic mice are as follows:

1. **The TRAMP (Transgenic Adenocarcinoma of the Mouse Prostate) Mice:** This consists of a minimal rat probasin promoter that drives expression of simian virus40 (SV40) tumor antigens. These mice develop prostate cancer within 12 weeks of age and ultimately develop metastasis by 30 weeks. The TRAMP mice recapitulate many salient aspects of human prostate cancer⁶⁸.
2. **p53^{+/-} Wnt-1 transgenic mice:** p53^{+/-} mice have been crossed with mWnt-1 transgenic mice, which express Wnt-1 transgene in the mammary epithelial cells (MECs) under the control of the mouse mammary tumor virus (MMTV). These combined transgenic mice develop a model of mammary tumorigenesis⁶⁹.
3. **Apc deficient mice:** These mice spontaneously develop preneoplastic intestinal polyps due to a dominant mutation of an *Apc* (adenomatous polyposis coli) gene. Mutation of this gene is common to most human colon cancers⁷⁰.

(ii) **Knockout mouse model:** In these animals, the gene is deleted or inactivated artificially using synthetic DNA segment.

These knockout mice are the other type of genetically engineered animals generated by deleting both the alleles of a gene of interest⁷¹. Some examples of knockout mice are as follows:

1. **The *Nkx 3.1* knockout mice:** *Nkx 3.1* is a prostate specific tumor suppressor gene. It is important for differentiation and function of prostate. Loss of function of this gene leads to histopathological defects that resemble human prostate cancer. This model also provides a model for studying mechanism of prostate cancer initiation as well as to explore the tissue specific features of the disease⁷².
2. **Homozygous p53 knockout mice:** Mutation of *p53* tumor suppressor gene is the most frequently observed genetic lesion in human cancer. Over 50% of all human tumors have identifiable *p53* gene point mutation or deletions. These mice are highly susceptible to spontaneous tumorigenesis particularly lymphomas⁷³.
3. **Brca1 conditional knockout model:** Breast cancer suppressor gene 1 (*Brca1*) mutation is the main reason for 45% of breast cancer cases. Animal develop mammary tumor by the age of 10 to 15 months⁷⁴.

c) **Transplantable tumors:** These models are the most used model systems utilizing cancer cell lines or tissues of human or mouse origin. These models are divided into two types depending on methods of transplantation⁷⁵.

i) **Heterotopic tumor transplantation**

In this case, implantation of tumour cells or tissues is done at a site other than its anatomical site. For example, breast cancer cells are transplanted subcutaneously or intraperitoneally. This method generally involves subcutaneous or intraperitoneal transplantation of tumour cells; the tumour proliferates in that site to form solid nodules or as cists, respectively. The procedure of inoculation is simple, quick, and precise and allows inoculation in a large number of animals for screening of anticancer agents⁷⁶.

ii) **Orthotopic tumor transplantation**

It refers to the transplantation of cancer cells into the anatomical site or tissue where the tumor appears. For example, transplanting lung tumors into the lungs. Utilization of this method leads to the development of tumor models which more closely resemble tumor histology, vas-

cularity, gene expression, sensitivity to chemotherapy and metastatic biology of human cancers. In the orthotopic tumour model, the interaction of the host microenvironment in tumorigenesis, progression, and metastasis behaviour is more strongly perceived, so it can be considered superior over conventional flank mouse models⁷⁷.

These models are developed by inoculation of cancer cells by direct injection or by surgical procedure in anatomical site, i.e. by implanting cancer cells or intact tumor fragments orthotopically by invasive procedures. Surgical implantation is invasive as it improves the reproducibility and metastatic behaviour of the model⁷⁶.

Transplantable tumour models are further categorized into two categories depending upon the origin of the tumour and the host used.

I) **Syngeneic model:**

In these animal models, the cancer cell lines or tissues of murine (mouse or rat) origin are used for transplantation. Syngeneic models are obtained by injecting a recipient of a specific genetic background with cell lines previously established through isolation of tumor cells from a mouse of the same genetic background. The advantage of this model is that the transplanted tissues, the tumor microenvironment, and the host are from the same species⁷⁸. Some of the examples of syngeneic tumor models are given below^{79,80}:

- B16-F10 melanoma cell line obtained from C57BL/6J mouse is used to develop cancer models in animals of same species.
- L1210 leukemic cell line obtained from DBA/2 mouse grown in animals of same species.
- Lewis lung carcinoma model was developed by isolation from a spontaneous epidermoid carcinoma of the lung in C57BL/6 mouse.

II) **Xenogenic (xenograft) models:**

This tumor model closely resembles the clinical scenario because they are designed with transplanted tumor of human origin. Since the transplanted cancer cells are from different host origin such as human tumor in mice, which may result in severe immune rejection. To overcome this problem, mice immunosuppressed by thymectomy or radiation are used. But there are several limitations associated with this method. Therefore, athymic (nude) mice or mice with severe com-

bined immunodeficiency (SCID) are used. These animals lack an immune response to the transplanted foreign material⁸¹.

Transplantation of tumor cell lines into nude mice can be accomplished via multiple routes like subcutaneous, intraperitoneal, intravenous, intracranial, intrasplenic, and renal subcapsular, or through a new orthotopic model by site-specific organ inoculation. Each site has specific advantages and limitations⁸².

When human tumor cells are transplanted in nude mice, they undergo kinetic changes. The doubling time is usually shorter than that of the original tumor and decreases with subsequent passages. Despite this, many xenografted human tumors maintain original morphologic and biochemical characteristics. Therefore, human tumor xenografts are mainstay of cancer drug discovery programs⁸³.

Xenograft tumor models are further classified based on the transplant site, which are as follows:

1. *Ectopic tumor xenograft model*

Generally, human cancer cells are subcutaneously injected into the hind leg or back of mice. In an ectopic tumor xenograft model (ectopic model), the transplanted site is different from the origin of the cultured cells. The ectopic model is considered as the standard model for evaluation of anticancer efficacy because it can be used to monitor tumorigenicity and tumor growth easily. Furthermore, the ectopic model is very reproducible, homogeneous, and amenable to use. But they have some limitations such as limited histological and phenotypic similarities to primary cancer, low metastatic rates and lack of native tumor microenvironment^{84,85}.

2. *Orthotopic tumor xenograft model:*

In orthotopic tumor xenograft model (orthotopic model), variety of human tumor cells are transplanted into the same origin site of the tumor, so they develop the same anatomical microenvironment. For instance, lung cancer cells are directly injected into the mouse lung for the orthotopic model. Orthotopic tumors show faster early-stage tumor growth, angiogenesis and hyper-permeability of blood vessels compared to ectopic tumors. But the limitation associated with this model is determination of drug efficacy therapies endpoints is more complex unlike subcutaneous ectopic models^{86,87}.



3. *Patient derived tumor xenograft model:* Although the xenograft model has advantages, it has limited ability to show the response of cancer patients to specific treatments. Clinical trials require reliable prognosis of drug response, and existing models are insufficient⁸⁸. To overcome the shortcomings of these models, patient-derived tumor xenografts (PDTX) were developed and utilized⁸⁹. Because PDTX involves directly transplanting cancer patients' tissue into immunosuppressed mice; genetic information and immunohistological markers are relevant to patient and can be used to evaluate new anticancer drugs and personalized cancer therapies⁹⁰. The several advantages of PDTX can be summarized as follows: 1) it maintains and stabilizes genetic, histological and phenotypic characteristics of tumors, 2) maintains stromal and stem cell components of the tumor, 3) facilitates the evaluation of biomarker, and 4) can be used to predict the response to an anticancer drug⁹¹.

However, the PDTX model has technical constraints, and is expensive and time-consuming. In this scenario the freshly excised human primary tumors must be transfer from operating room to the laboratory within a few hours. At the same time, samples of the primary human tumors should be examined through immunohistological analysis⁹². Therefore, cooperation between surgeons, histologist, scientist and laboratory staff is required to collect specimens correctly and subsequently obtain approval. In addition, approval from institutional review board (IRB) is required because utilization of patient-derived tumor tissue entails clinical and ethical considerations.

Notwithstanding these efforts, the take rate of PDTX is about 25%, and establishment of PDTX takes approximately three months before the first passage⁹³. Like xenotransplantation, it is necessary to transplant to SCID mice for the first time to avoid acute immunorejection, which is expensive. Additionally, the volume of tumor tissue obtained from patients is very limited so, the size of the PDTX population must increase with the passage of tumor tissue. Simultaneously, each passage of tumor tissues should be histopathologically analysed and compared with the original tissue. From the second passage, nude mice can be utilized. The tumor tissue fragments can be frozen and preserved in liquid nitrogen. PDX has been explored as a promising model for personalized cancer therapy⁹⁴. In Table 2 we report different tumor types and anatomical sites utilizing various mouse strain and engraftment rate in PDX model.

d) Virus induced tumor model

The carcinogenic effects of viruses are related to miscarriage and non-productive infections. The infectious nature of oncogenic viruses sets them apart from other carcinogenic agents. Therefore, a comprehensive study of the pathogenesis and host response of viral infection is crucial to a full understanding of the resulting cancers¹⁰⁵. This knowledge in turn expands our understanding of cellular pathways involved in growth and disease, differentiation and neoplasia. Human oncogenic viruses belong to different virus families and utilize diverse strategies to promote cancer development, but they have many common characteristics. A key feature is that it can infect but not kill their host cell¹⁰⁶. Unlike many other

TABLE 2. Different tumor types and anatomical sites utilizing various mouse strain and engraftment rate in PDX model.

Tumor type	Mouse strain	Implantation site	Engraftment rate (%)	References
Colorectal cancer	SCID	s.c	87	[95]
Colorectal cancer	Nude	s.c	64	[96]
Colorectal cancer	Nude	orthotropic	89	[97]
Breast cancer	Nude	s.c	13	[98]
Prostate cancer	NOD/SCID	Subrenal capsule	95	[99]
Pancreatic ductal carcinoma	Nude	orthotopic	62	[100]
Pancreatic ductal carcinoma	Nude	orthotopic	62	[101]
Gastric cancer	Nude and NOG	i.p	31	[102]
SCCHN/SCC	Nude	s.c	54	[103]
Renal cell carcinoma	Nude	s.c	8.9	[104]

SCCHN = Squamous cell carcinoma of head and neck; SCC = Squamous cell carcinoma; NOD = Non-obese diabetic; SCID = Severe combined immunodeficiency; s.c = Subcutaneous; i.p = Intraperitoneal

TABLE 3. Viruses associated with human malignancies.

Name of Virus	Type of cancer
Human T-cell leukemia virus (HTLV-1)	Adult T-cell leukemia (ATL)
Human Papillomavirus (HPV)	Cervical cancer, skin cancer in patients with epidermodysplasia verruciformis (EV), head and neck cancers, and other anogenital cancers
Human herpes virus 8 (HHV8)	Kaposi's sarcoma (KS), primary effusion lymphoma
Epstein-Barr Virus (EBV)	Burkitt's Lymphoma (BL), nasopharyngeal carcinoma (NPC), post-transplant lymphomas, and Hodgkin's disease
Hepatitis B virus (HBV) and Hepatitis B virus (HBV)	Hepatocellular carcinoma
Simian vacuolating virus 40 (SV40)	Brain cancer, Bone cancer, and mesothelioma
BK virus (BKV)	Prostate cancer
JC virus (JCV)	Brain cancer
Human endogenous retroviruses (HERVs)	Breast cancer, Ovarian cancer, melanoma
Human mammary tumor virus (HMTV)	Breast cancer
Torque teno virus (TTV)	Gastrointestinal cancer, lung cancer, breast cancer, and myeloma

disease causing viruses, cancer causing viruses tend to establish long-term persistent infections. The long-term interactions between virus and host are key features of the oncogenic viruses, as they set the stage for a variety of molecular events that may contribute to eventual virus-mediated tumorigenesis¹⁰⁷. Viruses that are compellingly associated with human malignancies are depicted in Table 3.

There is no more perfect animal model than a human virus. All animal models, in terms of their nature and design represent an aspect of the disease rather than the entire life cycle of human tumor virus. Different rodent models are used to stimulate virus induced human cancer. Rabbit serves as a useful animal model to study HTLV-1 induced cancer. Immunodeficient and genetically engineered mouse models are used to study Epstein - Barr virus (EBV) induced cancer¹⁰⁸.

The methods for development of different *in vivo* anticancer models are pictorially represented in Figure 4.

Zebrafish as a novel *in vivo* model in cancer research

Zebrafish (*Danio rerio*) is a small vertebrate tropical fish that has emerged as a powerful model to study cancer biology, process for cancer drug discovery and also for toxicity screening¹⁰⁹. The growing interest of this model leads to the existence of a bunch of advantages that make it a good model for cancer drug development and its safety assessment¹¹⁰. Zebrafish (ZF) can be used at any stage of development to conduct experiments whether it's in a larval stage (3 to 29 day after fertiliza-

tion, dpf) or the adult stage (90 dpf-2 years)¹¹¹. The husbandry cost of zebrafish is less expensive and due to smaller size of the adult fish, its breeding can be done in a small freshwater tank hence less space is required. It can absorb compounds, small molecules, or drugs directly from the water environment which make drug administration easy and feasible¹¹². Zebrafish contribute a high level of physiological and genetic resemblance with human, almost 70% of human genes are functional analogous to zebrafish¹¹³. Furthermore, zebrafish gestation period is so fast, it is capable of fertilizing 200-300 eggs in a week thus enough number of animals are available to utilize in the experiment although the zebrafish are fully developed by 24 hours post fertilization (hpf), and they are ready for use in larvae experiments by 3 days post fertilization¹¹⁴. Their extrauterine development is rapid and require less time for development of zebrafish organ system¹¹⁵. Moreover, other benefit of utilizing zebrafish model is transparency of its embryos and larvae, which allows visualization of organ and tumor growth inside the fish body and dynamic evaluation at the stages of drug development¹¹⁶. Embryos of ZF stay transparent to 60 hpf, their transparency period can be extended by treating embryos with 1-phenyl 2-thiourea (PTU), which inhibits the pigmentation process¹¹⁷. The transparency of ZF embryo also allows investigating anticancer nanomedicines biological effects, therapeutic efficacy, and safety by simply utilizing *in vivo* imaging¹¹⁸. A diversiform of cancer models have been developed in zebrafish which pathophysiologically or in molecular level also resemble human counterparts. Scrutinizing these features of zebrafish it is considered as a devoted model for cancer drug discovery and toxicity screening.

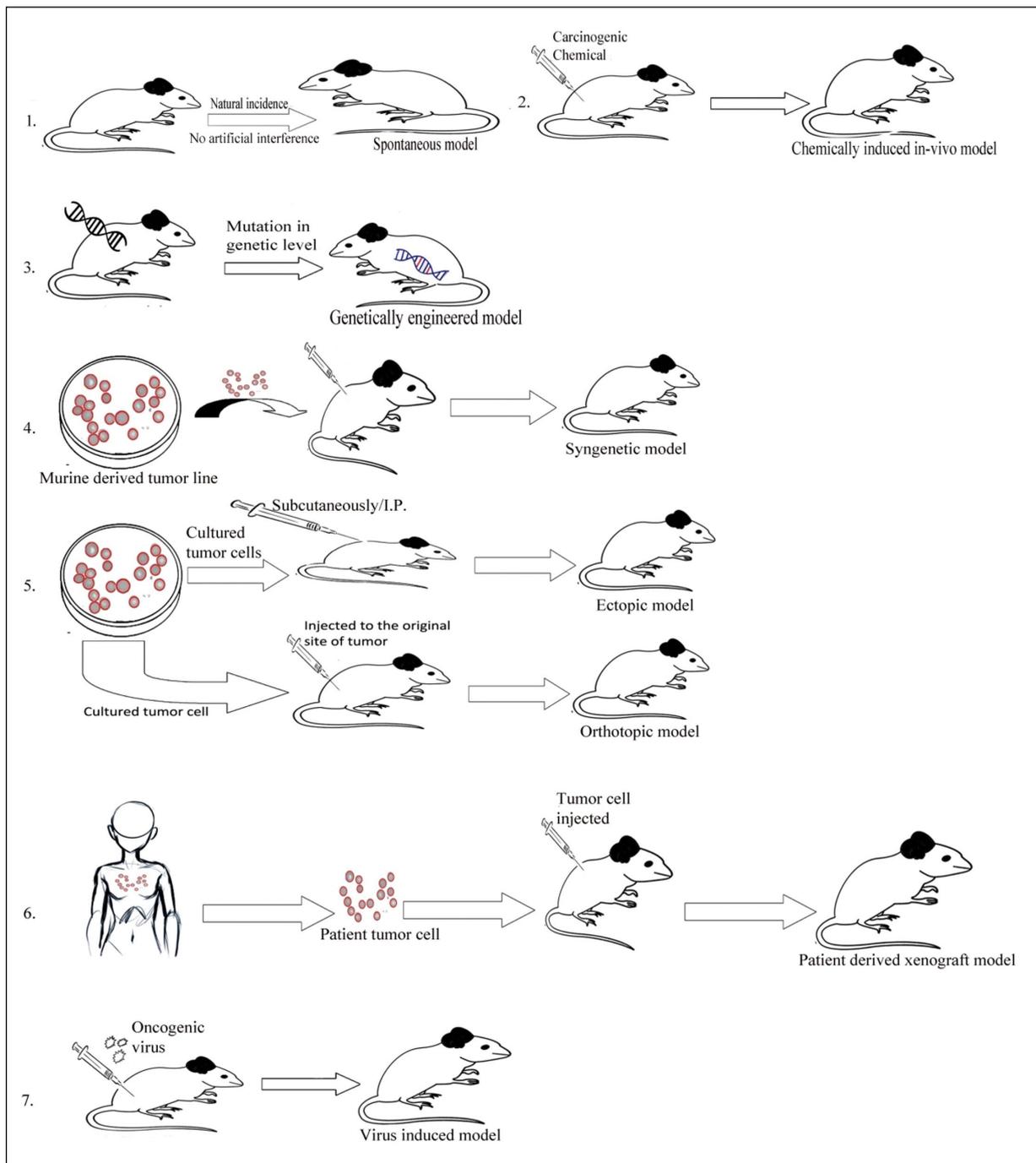


Figure 4. Pictorial presentation of different procedures for the development of *in-vivo* anticancer models. (1) *spontaneous model* are naturally cancer developing model i.e no artificial treatment required for cancer induction (2) *chemically induced model* is developed by introducing chemical carcinogens to the animal (3) *genetically engineered model* is developed by altering the targeted genetic makeup of the animal which induces cancer susceptibility (4) In *syngeneic model* tumour cell line of a murine species is injected to another animal of same species to produce cancer (5) *ectopic model*- human cancer cell lines subjected to murine model s.c/i.p and in *orthotopic model*- cancer cells of human specific cancer is injected to original site of tumour (6) *patient derived xenograft model* developed by collecting tumor tissue of patient and injecting the same to the animal (7) In *virus induced model* oncogenic viruses are introduced to animal which interfere with the DNA synthesis process and produce cancer.

Zebrafish is a convenient model for induction of tumor by the utilization of different methods such as (a) chemical exposure (b) genetically engineered and (c) xenotransplantation. Different approaches are used to generate these zebrafish models which

investigate cancer progression, molecular mechanism involved and study novel anticancer drugs efficacy¹¹⁹. Methods for induction of zebrafish tumor model are briefly explained in the following and schematically represented in Figure 5.

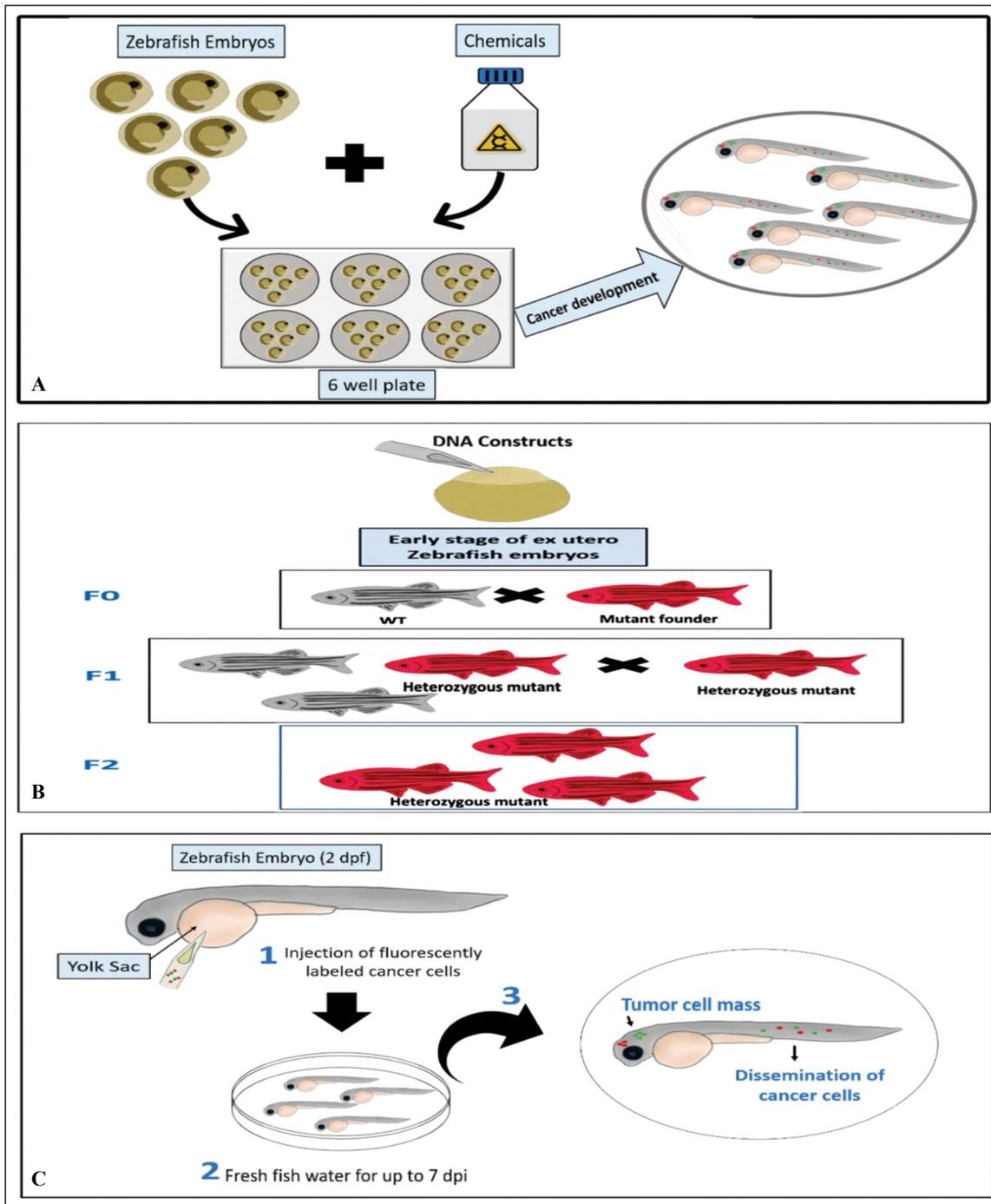


Figure 5. Different zebrafish model for cancer screening. *A*, *Zebrafish* chemical carcinogenesis model; Zebrafish embryos are collected, and healthy embryos are arrayed into a multi-well plate. Chemical carcinogens are then added to the multi-well plate following dissolving or suspending them in the fish water. After appropriate incubation period, embryos are screened for cancer growth. *B*, *Transgenic cancer zebrafish* lines; DNA constructs of cancer gene are microinjected into the cell of fertilized egg which is an early stage of zebrafish embryos. The adult mutant founder fish (Red) that have the germ cells, in which the construct was integrated into the genome, is mated with a wild-type (WT) zebrafish (Gray) at F0 to produce the F1 generation of fish, in which some of the progenies are heterozygous (Red) for the constructs and these heterozygous fish are then mated with each other to produce the F2 generation of fish that are homozygous to the construct (Red). *C*, *Xenotransplantation assay in zebrafish embryos*; Human cancer cells stained with red- or green- fluorescent dye and injected alone or in combination using a glass micropipette or a glass capillary needle in the yolk sac of zebrafish embryos at 2 dpf. The xenografted embryos are then maintained at a specific temperature between 28°C to 37°C. After 2 to 7 days post-injection (dpi), the number of tumor cells increases, and cancer cells disseminate at distance sites such as head and tail¹⁸.



(a) Chemically induced zebrafish cancer model

Zebrafish animal model has shown terrific diversity in mutation of neoplasm as compared to other fish species¹²⁰. Induction of tumor by chemical exposure was the initial approach to develop zebrafish as a cancer model. Zebrafish treated with carcinogen can provide a robust cancer induced model of any type and are substantially less difficult to carry out as compared to murine mouse model¹²¹. The spontaneous development of almost any type of cancer by these carcinogens resembles similar morphology and mechanical pathway to human. The most prevalent process for chemical induction of cancer is dissolving the carcinogen in the water and then soaked the fish for varied period of time. Carcinogen with different concentration for a changeable duration is utilized such as smaller doses soaked up to 24 h and large doses for 8 h or less. Additional method for induction is injection of carcinogen directly to embryo and topical application¹²². Allied numbers of carcinogens are used to develop cancer in mammals and zebrafish. For example, N-methyl-N-nitro-N-nitrosoguanidine (MNNG) dimethylbenzanthracene (DMBA), N-nitrosodiethyl-amine (DEN), N-nitrosodimethylamine (NDMA) when exposed to zebrafish, can induce a variety of cancers such as rhabdomyosarcomas, leiomyosarcoma, chondromas, seminomas, hemangiosarcomas, papillomas, and hepatic cancer. These chemically induced zebrafish exhibit molecular and histologic similarities with human cancer¹²³.

N-nitrosodiethyl-amine (DEN) induced pancreatic cancer: zebrafish was immersed in DEN solution (100 ppm) in an acrylic tank for a period of 8 weeks. The water temperature maintained at $26 \pm 0.5^\circ\text{C}$ and the water containing the carcinogen was exchanged every 2 weeks. After exposure to 8 weeks, the fish was transferred to fresh water tank and the formation of tumor was observed¹²⁴.

N-nitrosodimethylamine (NDMA) induced hepatocellular carcinoma: 50 ppm NDMA was dissolved in dechlorinated tap water and then transferred to 20 L acrylic tank and immersed 50-60 fish in a tank. The tank was thermo-regulated and attached with mechanical filtration and water exchanged carried out every week. Then after exposure of fish to NDMA for 8 weeks, it was transferred to a fresh carcinogen water until observe the carcinoma produced. During the experiments diet, temperature, and light/dark cycle remained at the standard conditions¹²⁵.

7,12-dimethylbenzanthracene (DMBA) induced intestinal carcinoma: DMBA was initially dissolved in DMSO and then mixed into the oil

component and then finally mixed to a purified casein diet. Gelatin (2%) added to this modified diet (MPC). Diet was prepared in moist form by mixing 65% hot (55°C) distilled water with 35% dry mix. Then juvenile fish (2 months of age) were fed MPC diet containing 0, 100, 500, or 1,000 ppm DMBA for 4 months. Fish were then fed basal MPC diet for an additional 3 months, for a total of 7 month. After initiation of DMBA exposure at different interval of 4, 8, 12, and 16 weeks, histopathological study was carried out to evaluate intestinal tumour¹²⁶.

(b) Genetically engineered zebrafish cancer model:

Genetic manipulation is another significant method for developing zebrafish as a cancer model. Different mutant and transgenic zebrafish models are utilized as potent model for cancer research which are generated by using different techniques such as forward or reverse genetic approaches which directly assess the role of various genes in cancer related phenotypes¹²⁷.

The reverse genetic approach in zebrafish through the specific knockdown or knockout genes of interest has previously been limited but with the advancement of new techniques this area is progressed¹²⁸. Furthermore, another reverse genetic approach to induce gene knockout is antisense morpholino oligonucleotides technique, the microinjection of the antisense morpholino oligonucleotides (MO) into the zebrafish embryos' blastomeres or yolks in early stages binds to endogenous mRNAs and would prevent the targeted gene translation¹²⁹.

In forward genetic screening mutation in the zebrafish genome has been done by introducing chemical (ethylnitrosourea) and viral mutagenesis strategies¹³⁰. Mutagenized zebrafish are then screened for identifying mutation in respective gene by gene mapping and high-throughput sequence analysis¹³¹. Targeting induced local lesions in genomes (TILLING) by using chemical mutagens such as N-ethyl-N-nitrosourea (ENU) techniques have been used for the target-selected mutagenesis in the zebrafish embryos' genome¹³². Another targeted genome modification technology includes ZFNs (zinc finger nucleases), CRISPR (clustered regularly interspaced short palindromic repeats) system, and TALENs (transcription activator-like effector nucleases)¹³³⁻¹³⁵. As a result of utilizing this mutation techniques, a numerous number of stable mutant zebrafish line established that harbour mutation in a ribosomal protein (RP) or specific tumor suppressor genes such as p53, adenomatous polyposis coli (APC), PTEN, Brca2, and neurofibromatosis type 2 (NF2) genes¹³⁶. This genetically mutated

model can distinguish from chemical induced mutations because higher rates of cancer incidence observed in genetic model such as 28% for p53 (17% for Apc, up to 100% for ribosomal proteins mutants) and that each of these mutant lines shows predisposition for a defined set of cancer types¹³⁷.

Transgenic zebrafish model: Zebrafish can also be used as convenient model for human cancer by targeting misexpression of a known oncogene under a zebrafish tissue-specific promoter. Microinjection of foreign DNA constructs in early zebrafish embryos' driven by zebrafish tissue-specific promoter accomplish the spatial and temporal control of the expression of some transgenes such as the GAL4-UAS and Cre-LoxP and Tol2 transposon and LexPR binary systems had led to the development of several transgenic zebrafish cancer models. Modeling human cancers in zebrafish through transgenesis becomes much easier¹³⁸.

A few examples of genetically engineered zebrafish cancer model are given below:

Myc and TEL-AML1 transgenic zebrafish leukemia models: T cell acute lymphoblastic leukemia in transgenic zebrafish expressing mouse *c-myc* and chimeric *EGFP-mMyc* transgenes under control of the zebrafish *Rag2* promoter are established. The resulting transgenic *zRag2 EGFP-cMyc* fish line showed rapid (21-42 days) onset of T-cell acute myeloid leukemia (AML)¹³⁹.

Apc mutated zebrafish colon cancer model: The *APC* gene mutant in zebrafish leads to colon adenoma initiation and progression, suggesting an association with the activation of the *Wnt* signaling pathway¹⁴⁰.

BRAF^{V600E} transgenic zebrafish melanoma model: *BRAF^{V600E}* oncogene under the control of melanocyte-specific *mitf* promoter injected into wt fish, this transgene led to an increased formation of nevi. However, when injected into *p53*-homozygous mutants, 7% of fish developed melanoma by 4 months of age, indicating cooperation between *BRAF* and *p53* pathways in the pathogenesis of these neoplasms¹⁴¹.

HBx (hepatitis B virus X antigen) transgenic Hepatocellular carcinoma model: Overexpression of *HBx-mCherry* under the control of the liver-specific *fabp10* promoter takes place. This transgenic zebrafish develops hepatocellular carcinoma in *TP53*^{-/-} background at 11 month post fertilization via Src (non-receptor tyrosine kinase) activation¹⁴².

P53 knockdown malignant peripheral nerve sheath tumors (MPNSTs) model: Mutation in the wild-type p53 DNA-binding domain of the

tp53^{M214K} mutant zebrafish model line could spontaneously develop the malignant peripheral nerve sheath tumors at eight and half months and 16.5 months with an incident of 28%¹⁴³.

(c) Xenograft zebrafish model:

A foremost method for establishment of cancer model involves transplantation of human cell lines into zebrafish embryos. The primary advantage of using zebrafish for xenotransplantation is its lack of an innate immune system until 72 hpf and a mature adaptive immune response until 4 weeks of life which overcome the limitation of immune rejection and also the transparency of embryo allow detailed *in vivo* environment imaging for real-time visualization of cell-cell interactions¹⁴⁴. A numerous types of human cancer cell lines are transplanted into different zebrafish embryonal stages mainly 48 hpf stage and blastula stage of the embryo are considered¹⁴⁵. Xenotransplantation of patient-derived tumor cell into Zebrafish can be utilized as a pre-clinical screening model for personalized cancer therapy. Different sites such as yolk sac, vein, duct of Cuvier and hindbrain are utilized for transplantation and established a good model for studying cancer pathogenesis as well as screening of novel drug and therapeutics¹⁴⁶.

Zebrafish melanoma xenograft model: Melanoma cell line WM-266-4 labeled with CM-DiI, a lipophilic fluorescent tracking dye inject on 2 dpf, albino zebrafish maintaining at 28°C into the yolk sac, hindbrain ventricle or circulation and then transferred to fresh fish water and maintained at 35°C for up to 7 dpi and activated xenograft zebrafish formed melanoma¹⁴⁷.

Zebrafish glioblastoma xenograft model: Stelletin B, a naturally occurring marine triterpenoid, was tested in a zebrafish xenograft model of glioblastoma and it shown significant inhibition of angiogenesis *in vitro* as well as *in vivo* in zebrafish¹⁴⁸.

Zebrafish breast cancer xenograft model: The zebrafish breast cancer xenograft model is pertinent for revealing underlying mechanism, screening and development of anti-transforming growth factor- β drugs for the treatment of metastatic breast cancer¹⁴⁹.

Zebrafish prostate cancer (PCa) xenograft model: PC3-CTR cell line injected subcutaneously to zebrafish larvae at 48 hpf and then migration and proliferation of cells evaluated. This model serves as a best model for drug screening¹⁵⁰.

Some examples of the *In vitro* and *In vivo* models already used for evaluation of several cancers are depicted in the Table 4.



TABLE 4. Application of *in vitro-in vivo* model in anticancer drug development.

Name of model	Type of cancer	Key findings	References
In vitro model used for cancer research			
Spheroids model	Non-small cell lung cancer (NSCLC)	Establishment of an expandable model for drug screening and drug resistance	[151]
	Colorectal cancer	Mimicked different features of the tumor, such as ECM production, spatial organization, formation of a necrotic core, and ability to differentiate and polarize monocytes to M2-like macrophages	[152]
	Colorectal adenocarcinoma	Represents patient to patient differences in response to chemotherapy	[153]
	Breast cancer	Developed advanced model of breast cancer provides a possibility to study fibrosis and optimize antifibrotic therapies in cancer treatment.	[154]
	Prostate cancer	Spheroid formation from human prostate tumor cells exhibits tissue-like features.	[155]
	Hepatocellular carcinoma	Tumor spheroids not only mirror the 3D cellular context of the tumors but also exhibit therapeutically relevant pathophysiological gradients and heterogeneity of <i>in vivo</i> tumors	[156]
	Squamous cell carcinoma	Present a unique opportunity to evaluate growth properties that may not be realized when studying tumor cell growth as a monolayer.	[157]
Organoids model	Breast cancer	Predict the therapeutic response of anti-tumor drug in individual patients	[158]
	Liver cancer	Establishment of hepatocellular carcinoma organoids from needle biopsies, and cancer organoids maintain the genomic features of the original tumors for up to 32 weeks	[159]
	Gastric cancer	First reveal the potential metastatic role of TGFBR2 loss-of-function in diffuse gastric cancer	[160]
	Colorectal cancer	Verify the deficient of key DNA repair gene MLH1 role in drives tumorigenesis	[161]
	Prostate cancer	Show the role of nucleoporins in the progression of pancreatic cancer	[161]
	Pancreatic cancer	Evaluate cancer-stroma cell interactions	[163]
	Glioblastoma	Patient-derived organoids display histological features and recapitulate the hypoxic gradients	[164]
Scaffold model	Breast cancer	Cancer cell-laden polymeric scaffolds support consistent tumor formation in animals and biomarker expression as seen in human native tumors	[165]
	Osteosarcoma	Improved the osteosarcoma stem cell niche micro-environment simulation, providing precise inputs supporting cell-cell and cell-ECM interactions and tumour signalling pathways <i>in vitro</i> .	[166]
	Pancreatic ductal adenocarcinoma	Developed hybrid zonal model was able to support long-term growth and proliferation of cancer (PANC-1), endothelial (HMEC), and stellate (PS-1) cells for up to 35 days	[167]
	Prostate cancer (PCa)	Chitosan-alginate scaffold cultures could be used to identify PCa phenotypic characteristics, with potential applications for <i>in vitro</i> anti-cancer drug screening.	[168]
	Lung cancer	PLGA microparticles show better responses in the prepared model.	[169]
Microfluidic model	Cervical cancer	Study the efficacy and mechanism of action of tannin from <i>Spatholobi Caulis</i> as anticancer agent	[170]
	Lung cancer	Mimic the <i>in vivo</i> microenvironment of cancer metastasis and to investigate cell-cell interactions during metastasis	[171]
	Colon cancer	Developed tumor-on-chip platform used for high-content image-based screenings or gene expression analysis to study drug-dose responses	[172]
	Breast cancer	Mimicking the Ductal carcinoma <i>in situ</i> (DCIS) structure, identifying multiple cellular adaptations to endure the hypoxia and nutrient starvation generated within the mammary duct	[173]

Continued

TABLE 4 CONTINUED. Application of *in vitro-in vivo* model in anticancer drug development.

Name of model	Type of cancer	Key findings	References
In vivo models used for cancer research			
<i>Spontaneous tumor model</i>	Breast cancer	Loss of estrogen and progesterone receptors and low expression of Her2/neu and overexpression of c-Myc, cyclin D1 and VEGF were observed	[174]
Chemically induced tumor model			
<i>7,12-dimethylbenz [a]anthracene (DMBA) induced cancer</i>	Squamous cell carcinoma	Skin lesions were induced by twice-weekly topical application of DMBA (100 nmol/100 μ L acetone) for 8 weeks	[175]
<i>Diethylnitrosamine (DEN) induced</i>	Lung adenocarcinoma	A single dose (15 μ g/g body weight) of DEN injected intraperitoneally & serve as a useful model for targeting KRAS and EGFR negative tumors	[176]
<i>7,12-dimethylbenz [a]anthracene (DMBA) induced cancer</i>	Breast cancer	50 mg/kg of DMBA once a week for 4 weeks to induced cancer	[177]
<i>Benz[a]pyrene induced cancer</i>	Liver cancer	Twice a week for a period of 60-90 days (50mg/kg body weight). There is a marked elevation in lipid peroxidation, diminished level of glutathione peroxidase (GPx)	[178]
<i>2-amino-1-methyl-6-phenylimidazo [4,5-b]pyridine (PhIP)</i>	Breast cancer	PhIP dissolved in 5% DMSO (75 mg/kg body wt.) for 1-5 and 8-12 days. Mechanism of metformin in PhIP induced breast cancer was observed.	[179]
<i>N-nitroso-N-methylurea (MNU)</i>	Prostate cancer	Single i.v. dose (50 mg/kg body weight) of MNU (dissolved in saline at 10 mg/ml) and after induction observe the chemopreventive effect of quercetin.	[180]
Genetically Engineered mouse model			
<i>MT/ret transgenic mouse model</i>	Skin melanoma	Development of cancer appears to resemble that of the malignant transformation of a human giant congenital melanocytic nevus.	[181]
<i>p53val135/wt transgenic mice</i>	Lung cancer	p53perhaps, the most important cancer genes in cancer development and progression	[182]
<i>MMTV-PyMT transgenic mice</i>	Breast cancer	Increased metastatic potential depend on colony-stimulating factor-1 (CSF-1) and EGF ligands between macrophages and tumor cells	[183]
<i>BRCA1 knockout mice</i>	Breast cancer	Disruption of Brca1 causes genetic instability and triggers further alterations, including inactivation of p53, that leads to tumour formation	[184]
<i>p53 knockout mice</i>	Skin cancer	Absence of p53 does not augment the frequency of initiation or the rate of promotion but greatly enhances malignant progression.	[185]
<i>PTEN (Phosphatase and tensin homolog) knockout mice</i>	Prostate cancer	PTEN-deficient mouse model represents useful tool for the preclinical evaluation and characterization of chemopreventive agents.	[186]
<i>Syngeneic tumor model</i>	Lung carcinoma	Kras ^{G12D} .p53 ^{-/-} line derived in a C57BL/6 background that forms lung tumors in C57BL/6 mice	[187]
	Triple negative breast cancer	Mvt-1and 4T1 TNBC mouse cell lines were injected into the mammary ducts via nipples of FVB/N mice and BALB/c mice	[188]
	Ovarian cancer	ID8 murine ovarian cancer cells transduced with pFB-neo-Luciferase (ID8-luc cells) to induce intra-peritoneal tumors in mice	[189]
<i>Orthotopic mouse xenograft models (HT-29 and HCT-8)</i>	Colorectal cancer (CRC)	The human HT29 cell line and HCT8 cell line are phenotypically similar to human CRC	[190]
<i>Patient-derived xenograft model</i>	Triple negative breast cancer	Phosphatidylinositide 3-kinase (PI3K)/mammalian target of rapamycin (mTOR) pathways were observed	[191]
	Lung cancer	PDX model is sensitive with EGFR mutations to EGFR-TKI and induced acquired resistance to EGFR-TKI	[192]
	Melanoma skin cancer	PDX can help guide vemurafenib treatment for metastatic melanoma	[193]
	Cervical cancer	Ability to detect cervical dysplasia and normal cervical tissue cells is novel and provides models for the study of tumour initiation and progression	[194]

Continued



TABLE 4 CONTINUED. Application of *in vitro-in vivo* model in anticancer drug development.

Name of model	Type of cancer	Key findings	References
Zebrafish <i>in vivo</i> model			
Chemically induced zebrafish model			
<i>Ethyl nitrosourea (ENU) induced cancer</i>	Skin cancer	Immersion of zebrafish in 2.5-3 mM ENU for 8 weeks develop skin tumor	[195]
<i>N-methyl-N-nitro-N-nitrosoguanidine (MNNG) induced</i>	Liver cancer	Zebrafish embryo (80 hpf) immersed for 1 hr containing 10 ppm (mg/L) MNNG, transferred to fresh water. After 3 months produce liver carcinoma.	[196]
Genetically engineered zebrafish model			
<i>Twist1-ERT2 zebrafish</i>	Hepatocellular carcinoma	80% of Twist1-ERT2/xmrk double-transgenic zebrafish spontaneously show metastatic dissemination of hepatic cells	[197]
<i>zRag2-EGFP-mMyc transgenic zebrafish</i>	T cell acute lymphoblastic leukemia (T-ALL)	Screening efficacy of cyclophosphamide (CY), vincristine (VCR) and prednisolone (PRE) in leukemia therapy	[198]
<i>eGFP-KRAS(G12V) transgenic zebrafish</i>	Pancreatic carcinoma	Cells expressing the transgene undergo minimal differentiation during embryogenesis and subsequently instigate several types of malignant pancreatic neoplasia in adult zebrafish	[199]
<i>rag2-KRAS^{G12D} transgenic zebrafish</i>	Rhabdomyosarcoma	Molecular pathway of disease analysed, identification of tumour-initiating cell populations	[200]
<i>Neurofibromin 1 (NF1) knockout zebrafish</i>	Malignant peripheral nerve sheath tumors (MPNSTs)	Identifying the regulatory pathway of tumorigenesis	[201]
Xenograft zebrafish model			
Cell line derived Zebrafish xenotransplantation			
	Lung cancer	Inhibition of proliferation of tumor cells with EGFR mutation and T790M resistance mutation by Osimertinib was observed	[202]
	Ovarian cancer	Suggested that targeting extracellular signal-regulated kinase (ERK2) in the presence of cisplatin may reduce the burden of residual tumor	[203]
	Triple negative breast cancer	Inhibition of CXCR4 (chemokine receptor) signalling with IT1t (CXCR4 antagonist) led to a 39-60% decrease in tumor burden at 4 dpi.	[204]
Patient-derived xenograft (PDX) model	Breast cancer	PDX zebrafish gives better understanding of drug sensitivity and identify both prognostic markers and markers that are predictive of response to therapy	[205]
	Gastric cancer	PDX model showed potential proliferating, angiogenic and metastatic activity	[206]
	Pancreatic cancer	Develop same tumor environment to the original tumor, efficient model	[207]
	T-ALL	Elucidate molecular pathogenesis and targetable gene mutation	[208]

Parameter evaluated for *in vivo* anticancer efficacy analysis

Various parameters are evaluated after induction of carcinogenesis in different tumor models and evaluate the anticancer efficacy^{209, 210}. For example,

- i) Tumor volume: it is measured by the following equation:
Tumor volume = length x width²/2 where length represents the largest tumor diameter and width represents the perpendicular tumor diameter.

Relative tumor volume is calculated by using the formula:

$$\text{Relative tumor volume} = \frac{T_x \times 100}{T_0}$$

T_x = absolute tumor volume of the respective tumor on day x

T_0 absolute tumor volume of same tumor on day 0, when the treatment started.

- ii) Tumor weight: the weight of each tumor is measured at the end of the experiment.

- iii) Body weight: the weight of the animal is measured each week and percent increase in body weight is calculated by the following formula:

$$\% \text{ increase in body weight} = \frac{[(\text{animal weight on respective day} / \text{animal weight on day 0}) - 1] \times 100}{100}$$
- iv) Haematological parameters: in order to assess the influence of treatment on the haematological status of animals is observed by determining the following parameters:
- White blood cell total count
 - Red blood cell total count
 - Haemoglobin contents
- v) Biochemical estimation: different set of methods, assays, and procedures are utilized to determine the substances present the serum glutamic pyruvic transaminase (SGPT), serum glutamic-oxaloacetic transaminase (SGOT), alkaline phosphate level (ALP), blood urea, serum creatinine, total protein, albumin.

Evaluation of in vivo pharmacokinetic parameters

Pharmacokinetic (PK) understanding is an essential component of drug discovery process because it directly associated with the efficacy and safety profile of the drug. Different techniques have been developed to measure PK in both preclinical and clinical developmental stages of new drug discovery process. Pharmacokinetics can be determined by conducting two methods- either invasively, i.e., by collecting blood, tumor or tissue samples, or non-invasively, i.e., by using various imaging techniques²¹¹.

Bioanalytical method

PK profiles can also be obtained by collecting plasma, urine and tissue samples at various time points from the dosed (via different route) animal models through either serial or composite sampling and by analyzing the samples for the drug using analytical methods such as high-performance liquid chromatography coupled with mass spectrometry²¹². The drug concentration against the time curve performed using non-compartmental analysis. Pharmacokinetic parameters such as the time taken to reach the maximum plasma concentration (t_{max}), half-life time ($t_{1/2}$), peak plasma concentration (C_{max}), the area under the curve ($AUC_{0-\infty}$ h), volume of distribution (V_d) and the mean residence time (MRT) were determined. The drug concentration is usually expressed as ng/mL for plasma and ng/g for various tissues²¹³.

Molecular imaging method

Various non-invasive imaging methodologies are available for assessing pharmacokinetic. Molecular imaging can give information about biodistribution and presence of drug to the site. Different imaging techniques for PK assessment are:

Magnetic resonance spectroscopy (MRS) and Magnetic resonance imaging (MRI): MRS is the non-invasive *in vivo* method for measuring the concentrations of biochemical compounds (or drugs and their metabolites). MRI is used to define tumor location and detecting drug accumulation, but MRS can be used to measure the concentration of drugs in that tumour in real time. For PK studies, drug concentration in the tumor and also accumulation of drug in other organs can also be measured by this technique. Thus, drug retention, metabolism, and elimination can be studied in the target tissue by using MRI techniques followed by MRS²¹⁴.

Positron emission tomography (PET): The PK profiles can be obtained by radiolabeling the drug of interest with different radioisotopes. Many positron emitting isotopes of carbon, oxygen, nitrogen and fluorine are available which has half-lives of few seconds to several days. When absorption, distribution, metabolism and excretion of the analyte take place, the concentration of these radiolabeled compounds in different compartments such as blood, plasma, urine, bile, faeces and tissues correspond to the amount drug and its metabolites. PET can provide valuable information on drug pharmacokinetics in tumour and normal tissue by mathematical modelling of data. Rapid plasma radioactive profiling during the PET scan can be used for calculation of plasma input function and to provide evidence of specific metabolic processes in preclinical and clinical drug screening. PET/MRI scans will yield sensitive images of probe uptake with excellent soft-tissue contrast^{215,216}.

Computed tomography: X-ray computed tomography (CT) is used for visualizing three-dimensional tomographic imaging and diagnosis of soft-tissue and bone cancer treatment response. In PD studies, CT is used mainly to measure changes in the volume of disease with treatment²¹⁷.

Ultrasound imaging (US): US have high sensitivity for measuring tumor vasculature and blood flow. Tumor perfusion rates can be measured within a field of view by acoustically bursting the bubbles and then calculating the time spent to repopulate and regain contrast²¹⁸.

Fluorescence imaging (FRI): Fluorescence imaging is the most commonly used powerful imaging tool for pharmacokinetic assessment. A plentiful of suitable fluorescent dyes are available that are excitable in the visible to near-infrared



(NIR) wavelengths of the spectrum. FRI gives information on biodistribution of drug to the tumor environment and also accumulation of drug to the other organ parts. It is also accessible for gathering information on elimination or clearance of drug from kidney. Laser scanning confocal microscopy (LSCM) is a commonly used technique in preclinical research to obtain high-resolution optical images at controllable depths. By measuring the intensity of the fluorescent, the quantitative determination is monitored, higher intensity accomplish accumulation of more drug. Fluorescence molecular tomography (FMT) is an advanced tool for quantification of drug into the deeper tissues²¹⁹.

Translation of animal models to clinic

Due to practical and ethical issues associated with human experimentation, animal models play an essential role in cancer research. Animal models are needed to bridge the translational gap between preclinical and clinical research. However, the average rate of successful implementation of animal models to clinical cancer trials is less than 8%²²⁰. Preclinical testing in animal models is an integral component of the drug discovery and development process, which can predict the clinical efficacy and safety profile of the drug. The failure to translate from animals to humans is likely due in part to poor methodology and failure of the models to accurately mimic the human disease condition. A major obstacle to adequate prediction of clinical outcome based on preclinical animal studies may be potentially attributed to insufficient external and internal validity within the study design²²¹.

Table 5 depicts some reported clinical trials on the model used for anticancer screening.

Patents and other forms of intellectual property protection are generally thought to play essential roles in encouraging innovation in biopharmaceuticals. Patenting is the most effective mean of excluding market competitions²²². Standard essential patents and patent pooling are seldom seen in the biopharma industry. The key to success of a biopharmaceutical product is to prove its safety and therapeutic benefits in clinical trials, which are stringently regulated and the outcomes are unpredictable. Such stringency and unpredictability have consumed enormous human and financial resources, and greatly increased the risk in product development. To protect potentially massive returns, drug makers have raced to build strong patent portfolios. Patent analysis can provide a snapshot of patenting activities of the industry²²³. Some of patented *in vitro-in vivo* model used in cancer research are depicted in Table 6.

CONCLUSIONS

In vitro and *in vivo* models are important tools in cancer research because they allow the identification of genetic manipulation culpable for cancer, the development of cancer therapies, drug screening, and provide information on the molecular mechanisms of tumor growth and metastasis. Although conventional human tumor models, including two-dimensional (2D) monolayer culture, are widely used in experimental models, they come with a variety of limitations. Therefore, the three-dimensional (3D) *in vitro* tumor models are most widely used as a bridge between

TABLE 5. Clinical trial on cancer management using different *in vitro-in vivo* model.

Type of Study design	Clinical phase	Investigated application	CT Identifier
Developing mouse model	Completed	Breast cancer	NCT00897468
Constitution of ex-vivo model	Recruiting	Ovarian cancer	NCT03831230
Establishment of Patient Derived Cancer Cell Models	Recruiting	Metastatic cancer	NCT02646228
Patient-derived Organoid Model	Recruiting	Lung cancer	NCT03655015
Patient-derived xenograft model	Recruiting	Head and neck cancer	NCT02572778
Three-dimensional tumor model	Recruiting	Skin cancer	NCT03136783
3D Organoid model	Recruiting	Vaginal cancer	NCT04278326
Patient-derived organoid model	Recruiting	Breast cancer	NCT03544047
Patient-derived organoid model	Recruiting	Esophageal cancer	NCT03283527
Organoid model	Recruiting	Kidney cancer	NCT04342286
Develop <i>in-vitro</i> organoid model	Recruiting	Liver and Pancreatic cancer	NCT02436564
Xenograft model	Phase I	Liver cancer	NCT04690972
Patient-derived xenograft model	Completed	Triple negative breast cancer	NCT02247037
Patient-derived xenograft model	Recruiting	Prostate cancer	NCT03786848
Patient-derived xenograft model	Completed	Breast cancer	NCT02752893

TABLE 6. Some patents on use of different *in vitro-in vivo* models for cancer research.

Title	Year	Patent number	Inventors/Assignee
Mice models of human prostate cancer progression	2002	US6365797B1	The Regents of the University of California (USA)
<i>In-vitro</i> model for a tumor Microenvironment	2017	US9617521B2	HemoShear, LLC, Charlottesville (VA, USA)
Rodent mammary window for Intravital microscopy of orthotopic breast cancer and related method	2004	US20040151666A1	Dewhirst MW, Shan S
Method and a kit for the <i>in-vitro</i> diagnosis of Pancreatic ductal Adenocarcinoma or for determining the predisposition to pancreatic ductal adenocarcinoma	2015	US20150177249A1	Natimab Therapeutics, SRL, Colletterto Giacosa, Torino
PIK3CA H1047R knock-in non-human animal breast cancer model	2013	WO2013015833A3	Seshagiri S (USA)
Spontaneously immortalized prostate cancer cell line	2014	US20140017721 A1	The Research Foundation of State University of New York
<i>In-vitro</i> tumor metastasis model	2012	WO2013017282A1	Roche diagnostics GMBH
Model of colorectal cancer	2014	WO2015013432A1	Erica J, Kevin L (USA)
An <i>in-vitro</i> 3d cell culture model-based tumor relapse assay	2016	WO2017029414A1	Insphero AG
Methods for the <i>in-vivo</i> detection and treatment of patient-centric tumor dependencies	2017	WO2017197290A1	Board of Regents, The University of Texas System (USA)
Animal models of prostate cancer and methods for their use	2003	WO2004000010A2	The Regents of the University of California (USA)
Mice model of human prostate cancer	2002	US6828471B2	The Regents of the University of California (USA)
Genetically modified mice expressing humanized PD-1	2017	US20190387724	Biocytogen Pharmaceuticals (Beijing) Co., Ltd
MMTV-SV40-spy1A and spy1A-pTRE transgenic mouse models	2013	US20140109244	University of Windsor
Animal model of human cancer and methods of use	2012	US20140047570	Altiok S
Transgenic Non-human animal model of lung tumorigenesis	2006	US20080313759A1	Dartmouth college
Knock-in mouse prostate cancer model	2003	WO2004044197	Procyon Biopharm Inc.
Three-dimensional bioprinted pancreatic tumor model	2015	WO2016022830A1	Oregon Health and Science University, US
Patient-derived xenograft model in gastric cancer and a use thereof	2015	KR101743340B1	Jeong J, Moon-Hee S, Lee J, Yoon-young C
Xenograft model of human bone metastatic prostate cancer	2012	US20140304844A1	The Regents of the University of California, (US)
A liver cancer PDX standardization model base	2016	CN105684989A	West China Hospital Sichuan University

in vitro and *in vivo* systems because they reflect similar cellular, molecular, and phenotypic features that closely generalize the characteristics of human malignant neoplasm. Varied animal (rodent) models have been developed for the cancer drug discovery process, which closely mimics human cancer environment and comfort to study the etiology of many types of cancer, assessment of drug toxicity, and the therapeutic potential of drugs. The zebrafish model can be used as an alternative *in vivo* model for investigating the cancer biology due to its cost-effectiveness, resembling a high homology to humans, rapid development, optical transparency, and the ease of

genetic manipulation. The choice of a preclinical model is crucial to the drug discovery process. Therefore, there is no single model that fits all drug discovery programs, and no single type of model is universally the best. There is still a lot that needs to be done to fill the gap in translating the results of the animal model into the clinic. The development of animal models that fully epitomize the evolution, complex and heterogeneous biological process dynamics, including molecular, cellular, and histological changes, as well as the development of tumors in humans, are a prerequisite for the effective progress of translational research.



CONFLICT OF INTEREST:

The Authors declare that they have no conflict of interests.

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