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# Systemic Review on Chronic Myeloid Leukemia: Therapeutic Targets, Pathways and Inhibitors

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#### **Abstract**

Chronic Myeloid Leukemia (CML) is a stem cell disorder, characterized by the translocation of 9<sup>th</sup> chromosome of Abelson (ABL) gene to the 22<sup>th</sup> chromosome of breakpoint cluster region (BCR) gene. Consequently, translocation results into the chimeric oncogene BCR-ABL which encodes the BCR-ABL oncoprotein. CML is mainly a disease of adults but it can occur in any stage of life and it accounts around 15% of the all the types of leukemia. Various methods have been used to combat this disease like Chemotherapy, Radiation therapy; tyrosine kinase inhibitors etc., Imatinib as a tyrosine kinase inhibitor has dramatically improved the survival rate of CML patients, hence can be referred as first generation drug against the CML. Later on, recurrence of the disease in some treated patients has also been seen probably due to mutation in oncogenes. Researchers have started to find out more efficient tyrosine kinase inhibitors which can work on mutated oncoprotein and which can be referred as second or third generation drugs. In this review, special emphasis have been given to the carcinogenic mechanism of abnormal fusion of the BCR-ABL genes, current therapeutic options to prevent this disease, and Systems Biology approach to explore the CML associated biochemical pathways. Various advantages and disadvantages of the all therapeutic options to combat CML have also been discussed.

Keywords: CML; BCR-ABL; Imatinib; Radiation; Chemotherapy

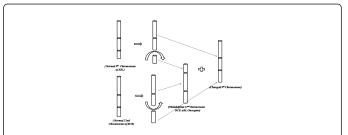
# Introduction

Chronic Myeloid Leukemia (CML) is probably the most studied human cancer because of its occurrence due to chromosomal abnormality and has been termed as Philadelphia Chromosome in 1960 [1]. In molecular investigation of the CML progression, it has been found that there is a translocation of Abelson (ABL) gene from 9<sup>th</sup> chromosome to the 22<sup>th</sup> chromosome of the Break Point Cluster genes and form a hybrid gene that is BCR-ABL oncogene [2]. The BCR-ABL gene encodes the oncoprotein namely BCR-ABL oncoprotein. Consequently this abnormal oncoprotein enhance the tyrosine kinase activity of the cells which are responsible for growth and replication by down regulating the associated pathways like Autophosphorylation, JAK/STAT, MAPK, RAS, RAF, JUN kinase, MYC etc. [3]. In continuation of the molecular analysis, ABL gene encodes a non receptor tyrosine kinase and homologous to the Abelosn murine leukemia virus .The ABL gene comprises of various structural domains, out of them three SRC homologous (SH-1, 2, and 3) are present towards the NH2 terminal [4]. Out of which, SH1 domain is responsible for tyrosine kinase activity, whereas SH2 and SH3 domains are involved in regulating the interaction activity of the BCR-ABL oncoprotein with other proteins [5].

It has been reported that ABL gene (which is expressed as 145-kd protein) is actively involved in the process of cell regulation and apoptosis with the help of extracellular and intracellular signals [5]. Likewise Break Point Cluster Region (BCR) gene encodes 160-kd oncoprotein and it consists of several structural motifs. The N-terminal exon encodes the serine-threonine kinase, whereas coiled domains of the exon region allowed BCR to dimerized [6]. The C-terminal region

of BCR consists of the GTPase which regulates the activity of Rac and consequently regulates the actin polymerization [7]. It has been noticed that, BCR can phosphorylate on various tyrosine residues especially T177 by binding with an adaptor Grb-2 and regulate the activation of Ras pathway [8]. Detail mechanism and domain orientation of the BCR-ABL protein is schematically represented into Figure 1.

This chimeric rearrangement of both the protein is also referred as the Philadelphia chromosome. The molecular consequence of this newly formed BCR-ABL fusion oncogene, which in turn translates into a Bcr-Abl oncoprotein and resulted into CML [6-9].



**Figure 1:** (A) Mechanism of BCR-ABL (breakpoint cluster region-Abelson) fusion and formation of Chronic Myeloid Leukemia causing agent.

# Stages of the CML

CML is a result of abnormal production of white blood cells by bone marrow cells. Proliferated leukemic cell starts affecting White blood cells, Red blood cells, and platelets [10]. CML usually occurs at adult

stage and rarely occurs at early stage of life with symptoms like fever, night sweats, and tiredness. There are three phases of chronic myeloid leukemia namely Chronic phase, Accelerated phase, and Blast phase [11]. Chronic is the most primitive stage of this disease, and it can be diagnosed by identifying simple symptoms like tiredness and fatigues, but sometimes symptoms mild and are not identifiable. If chronic phase passes away without getting noticed it ultimately enters into accelerated phase. In this stage CML cells becomes aggressive and is easily noticeable by analyzing symptoms consequently, Accelerated phase converted into Blast phase which is most aggressive and harmful to the patients. Most of the symptoms of Blast phase are similar to the Acute Myeloid Leukemia [10,11].

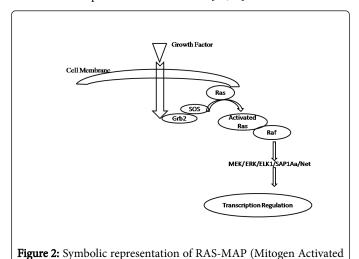
# Pathways Associated with Chronic Myeloid Leukemia

Multiple interlinked pathways have been reported that BCR-ABL influenced the PI3K/AKT/Mtor/JAK-STAT/RAF-RAS-MAPK pathways. Along with tyrosine kinase inhibition identification of common component of above said pathways can be a good therapeutic approach.

# Ras and MAP kinase pathways and targets

It's a signaling pathway by which cells receives the signal from receptor of the cell surface and communicate with the series of proteins in the cell and transmits signals to the nucleus. Signaling process starts by binding of the receptor molecule with the signaling molecule and ends with the binding of nucleus subsequently produces the changes into the systems like cell division [11-13]. Various reports had confirmed about the association between Ras and MAP kinase pathway with BCR-ABL protein.

The small ligand molecule Grb-2 and SOS autophosphorylate the reactions by binding with the tyrosine 177 docking site and activate the Ras shown in ref. [13]. Whereas other two adapter molecules like Shc and Crkl binds with SH2 and SH3 domains of BCR-ABL protein respectively and activates the Ras and MAPK pathways. Any abnormality in the chain reactions leads to the abnormal proliferation of the BCR-ABL protein and causes CML [14,15].



The activation of Ras by another ligand molecule like Crkl is still not clear because it is limited to the fibroblast only that's why interaction

between BCR-ABL and Crkl is not confirmed for transformation of myeloid cells [15-17]. Some evidence about involvement of Ras activity in pathogenesis of Philadelphia positive cells have been reported because of less common mutation presence even in Blast phase of the disease [18,19].

Raf initiates a signaling cascade through the serine-threonine kinases Mek1/Mek2 and Erk, which ultimately leads to the activation of gene transcription as shown in Figure 2 [20-22].

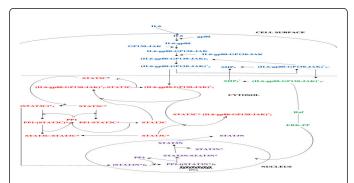
#### Jak-Stat pathway and targets

JAK-STAT pathway is highly expressed in white blood cells (WBC) and abnormal proliferation of the WBC is the cause of CML progression. This pathway is actively participating into the transcription process of the cells by transmitting the signal from outer environment of the cells through cell membrane and binds to the promoter of the nucleus. Phosphorylation of the transcription factors like STAT1-STAT5 is being reported in various CML literatures [20]. STAT5 significantly contributes during malignant cells transformation [23]. JAK/STAT signal transduction pathway is regulated by the two STAT3 phosphatases, PP1 and PP2.

Theses phosphates influence the activation of STAT component in the both cytosol and the nucleus. In this pathway as shown in Figure 3, IL-6 binds to its receptor gp80, which is not a part of the signaling process of the receptor. The receptor gp130 which is a part of the signaling process, attaches to the protein tyrosine kinases of the JAK family. Then IL-6-gp80 complex binds to the compound gp130/JAK to form the complex IL6-gp80-gp130-JAK. Formation of this complex results in the dimerization of the complex IL6-gp80-gp130-JAK to form (IL6-gp80-gp130-JAK)2.

SHP-2 binds to the phosphorylated dimer complex (IL6-gp80-gp130-JAK)2 to form (IL6-gp80-gp130-JAK)\*2-SHP-2 complex which act as the point of initiation for both the MAPK and the JAK/STAT pathway. In the JAK/STAT signaling pathway, the phosphorylated dimer, (IL6-gp80-gp130-JAK)\*2 recruits the transcription factor STAT3.

The phosphorylation of the component STAT3 leads to its dissociation from the (IL6-gp80-gp130-JAK)2 complex and further undergoes the process of dimerization. The dimerized complex of the STAT3 then enters into the nucleus to undergo dephosphorylation and is transferred back again into the cytosol for the cycle of phosphorylation-dephosphorylation cycle.



**Figure 3:** Symbolic representation of JAK-STAT and MAPK Pathway.

Protein) kinase pathway.

In Figure 3, Interaction between JAK-STAT and MAPK pathway have been described and can be seen that some of the molecules are involved in activation of other pathway's molecule.

# PI3 kinase pathway and targets

PI3 kinase pathway is directly related to the various cellular functions like cell division, cell differentiation, cell proliferation etc. It has been reported that PI3 kinase is highly expressed in BCR-ABL proliferation process. It forms complexes with Cbl, and adapter molecules like Crk and Crkl [24].

Membrane receptor has been catalyzed by the interaction with subunit of class I PI3Ks after receiving the extracellular signals. Catalyzed PI3Ks produces the PIP2 and PTEN, and deactivation of PI3Ks has been occurred due to dephosphorylation of PIP3 into PIP2 and acts as regulatory molecule of this pathway. PIP3 is activated by the binding of Pleckstrin Homology (PH) domains of the proteins and deposited to the cell membrane.

PIP3 is down regulated by the serine threonine kinase AKT/ PKB effectors. Binding of the PIP3 and PH domain activates the deposition of AKT to the membrane and activates the phosphorylation [25]. Schematic representation of the pathway has been shown in Figure 4.

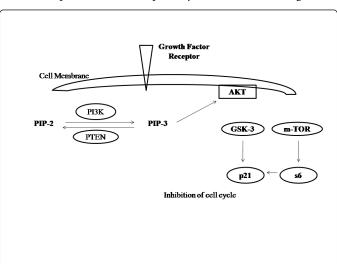


Figure 4: Symbolic representation of PI3 kinase pathway.

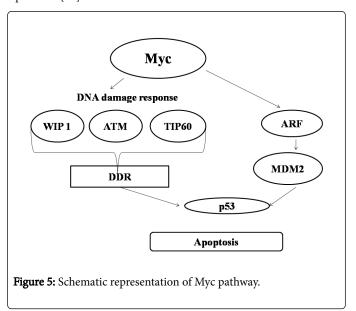
#### Myc pathway and targets

The Myc pathway is also highly expressed in various cellular processes like cell division, cell differentiation, proliferation, apoptosis etc. The mutation caused in pathways results into various types of malignancy [26]. It has been reported that Myc works as transcription factor, however its target is still unknown. Myc binds with the SH2 domain of the BCR-ABL protein and over expression of it causes the excessive proliferation of BCR-ABL [27].

Myc protein acts a transcription factor which activates the expression of genes by binding with enhancer box sequences (E-box) and histone acetyltransferases (HATs).

Myc protein is also known as transcriptional repressor because it inhibits the Miz-1target gene by binding with Miz-1 and replacement

of p300 co-activator. Myc pathway is directly associated with the DNA replication [28].



The direct association of Myc pathway with the BCR-ABL progression is still poorly understood, however some report suggests that transformation of v-abl is activated by the Ras/Raf, cdks, E2F transcription factors which ultimately affects the Myc promoter [28,29].

It has been noticed that effects of Myc on Philadelphia positive cells are almost similar to the other tumor cells like proliferation and apoptotic signals [30-32]. Schematic representation of the pathway has been shown in Figure 5.

# Important Components from the Various Pathways and their Targets

All significant components from the related pathways of chronic myeloid leukemia like Tozasertib, Bafetinib, Nilotinib, Dasatinib, Imatinib, Ponatinib, Bosutinib, Saracatinib etc. were collected and shown in Table 1. Respective targets of the each inhibitors are also been listed into the Table 1 with their PubChem ID.

S. No.	Reference Molecules	PubChem ID	Targets
1	Tozasertib	5494449	Aurora A,B, and C, FLT, Bcr-Abl
2	Bafetinib	24853523	Abl, Lyn
3	Nilotinib	644241	Bcr-Abl
4	Dasatinib	3062316	Abl, Src, c-Kit
5	Imatinib	5291	Abl, PDGFR, c-Kit
6	Ponatinib	24826799	Abl, PDGFRα, VEGFR2, FGFR1
7	Bosutinib	5328940	Abl, Src
8	Saracatinib	10302451	Src, EGFR, LCK, c-YES

**Table 1:** Reference molecules approved by FDA, their Pubchem ID, and their known targets.

# Therapeutic approaches to combat CML

There are three phase of the CML progression namely chronic, accelerated, and blast, out of these three phases chronic is a long lasting and primitive one. Whereas accelerated and blast are advanced and having reduced effect of the any treatment options like chemotherapy, stem cell transplantations, TKI etc. However for advanced stage patients stem cell transplantation option may be the preferred therapy if donor of the stem cells available. Specifically for the accelerated and blast phases, TKI like Dasatinib, Nilotinib etc are being reported significant.

However, after discovery of the Imatinib as a small molecule inhibitor for the targeting of causative agent BCR-ABL oncoprotein, survival rate of the patients have been remarkably improved specially in the case of chronic phase of the CML patients.

# Non-specific agent

Initial days, treatment process of the CML were completely based on non-specific agent like busulfan, hydroxyurea, and interferonalfa (INFa). These agents were able to improve the survival rate of the patients but their side effect like multitude of toxicities and probability of reoccurrence hinders the treatment process [33].

#### Stem cell transplantation

In continuation, Stem cell transplantation becomes second alternative of the CML treatment process, but its high risk of morbidity and mortality and lack of appropriate stem cell donor limits this technique [33].

#### Chemotherapy

Chemotherapy is useful in CML because through this technique patients have been treated by the certain dose of drug through vein or may be through mouth by which drugs enter in to the circulatory systems of the body and spread out into the whole body. Chemotherapy refers that the agent (hydroxyurea, busulfan, cyclophoshamide etc) which can specifically kill the abnormally growing cells. It was the extensively practiced treatment option decades ago [34].

Major limitations of this treatment are killing of non cancerous rapid dividing cells like bone marrow cells, the lining of the mouth and intestines, and the hair follicles etc which cause serious side effects like hair loss, mouth sores, loss of appetite, nausea and vomiting, Low white blood cell counts (leukopenia), which increases the risk of serious infection, Low blood platelet counts (thrombocytopenia), which can lead to easy bruising or bleeding, Low red blood cell counts

# Radiation therapy

Radiation therapy is used in a number of cancer cells treatments, especially when cancerous cells are localized in a particular area. It is used by bombarding t he high energy rays on a particular cancerous population and destroy them [36]. In Chronic Myeloid Leukemia, radiation therapy is not used as primary treatment, it is used in certain cases of the CML. Mainly it assists the chemotherapy by reducing the swollen internal organs like spleen. Sometimes low dose of radiations are given before the Stem cell transplantation treatment process of the CML.

Apart from its significance, there are some side effects of the radiation therapies which are as follows: Fatigue, skin changes (like redness, blistering, and peeling), mucositis, nausea, vomiting, diarrhea, low blood counts etc. [37].

# Tyrosine kinase inhibitors

After establishment of fact that the interaction of BCR and ABL protein is a causing agent, thereafter scientific community have started to find out the way through which interaction can be hampered and subsequently oncogenic activity or cancerous proliferation can be stopped. First success they have achieved after discovery of the Imatinib (STI-571), a small molecule that can inhibit the interaction by blocking the SH2 and SH3 domain of the ABL protein [38]. The FDA has approved Imatinib as first tyrosine kinase inhibitor drug in 2001, after discovery of Imatinib the entire arena of the CML treatment process has changed and patient's survival rate has increased remarkably) from 20 to 80-90% [39]. FDA has approved two more tyrosine kinase inhibitor like dasatinib, and nilotinib, after success of Imatinib. These drugs are referred as first generation of drug against the CML [40] shown in Figure 6.

#### **Imatinib**

Imatinib mesylate, (Gleevec, Novartis Pharmaceutical Corporation, NJ), a small molecule that can be used as a drug at chronic phase of the CML patients. It competitively inhibits the ATP binding site of the BCR-ABL protein and inhibits the phosphorylation process of cell signaling transduction. Clinical investigators have performed trial on 1106 patients by giving the dose of 400 mg/day till nineteenth months under the organization IRIS (International Randomized Study of Interferon and STI571).

In contrary around 30% patients didn't show anticipating response, which led the investigator to find out the second or third generation drugs against the CML. Many reasons have been drawn for the intolerance of Imatinib against the CML like point mutation of the specific residues like T315I, Y253H and F255K etc. Point mutation between Threonine and Isoleucine on 315 positions is also referred as Gatekeeper mutation which can be observed in 4-15 % Imatinib resistant patients [41].

Imatinib is a derivative of 2-phenyl amino pyrimidine and it accommodates with the active site of tyrosine kinase leads to reduction of activity. Imatinib is highly specific about binding with the receptor, however large numbers of tyrosine kinase receptors are available in the body but it acts against with ABL, c-kit, PDGFR etc only [42].

Tyrosine kinase receptor has ATP binding sites, where transfer of terminal phosphate group from ATP takes place and phosphorylation occurs. In case of BCR-ABL tyrosine kinase, Imatinib binds with the binding site by competitive inhibition with the ATP and blocks the space for ATP [43].

#### Nilotinib

In further continuation of drug discovery process of the Imatinib resistant CML cells, Nilotinib is a structural analogue of the Imatinib. It is produced as Tasigna, by Novartis Pharmaceutical Corporation and proven as 50 times better efficiency in terms of ATP binding affinity with BCR-ABL protein. Nilotinib gives the excellent outcomes in terms of survival rate [44].

# Multi Target Approach

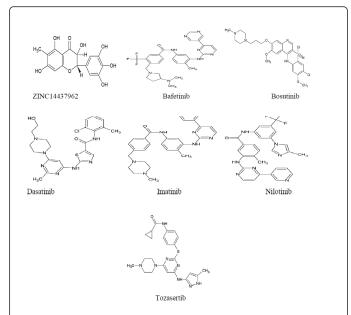
Above said first generation drugs like Imatinib, Nilotinib, and Dasatinib are invented for the inhibition of BCR-ABL fusion. Further it has been noticed that apart from BCR-ABL protein there are some other targets like SRC, STAT1, STAT3, STAT5, PDGFR etc., which are also involved in progression of this disease. Multi target inhibition by single inhibition has already been practiced in various severe disease combat strategies [40-44].

#### Dasatinib

Dasatinib is orally taken drug of CML and analogous to the Imatinib but in terms of efficiency it's 350 times efficient as compared to Imatinib. It is in market with the brand name Sprycel and produced by Bristol-Myers Squibb. Dasatinib is equally efficient against the BCR-ABL as well as SRC receptors [45]. It has been clinically approved that Dasatinib is a proven drug against chronic phase of the disease, Imatinib resistant cells, as well as newly treated patients [44,45].

# **Ponatinib**

Ponatinib has been discovered as multi-targeted tyrosine-kinase inhibitor, orally taken and marketed as Iclusig, and produced by the ARIAD Pharmaceuticals. It has been designed in a manner that it can efficiently inhibit the Imatinib resistant CML cells. This drug has been approved in 2012 by the FDA, but a ban has been imposed till Oct 2013 because some of the side effects like blood clot into the treated patients, but after rectification it is now widely used [46].



**Figure 6:** 2-D structures of the FDA approved inhibitors available for BCR-Abl and Src receptors.

After successful implementation of the Ponatinib as multi receptor inhibitors, many organizations have started to find out some more potent natural inhibitors which can bind to the multiple CML causing agent and efficiently inhibit the CML progression [45,46].

Some of the Bioinformatics approaches like virtual screening, molecular docking, molecular dynamics simulation, and ADMET

analysis become significant to get the small molecules which can be able to inhibit the multiple receptors. A recent study has been reported about the multi receptor inhibitors described in Table 2 [47]. In this study, small natural molecules have been screened out from the huge available natural databases. Reported molecules give the better result as compared to available single receptor inhibitor like Imatinib, Dasatinib, Nilotinib etc.

S. No.	Inhibitors	Docking Score with (BCR-ABL)	Docking Score With (SRC)
1	ZINC14437962	-9.493	-9.513
2	Bafetinib	-4.678	-7.085
3	Bosutinib	-4.686	-5.719
4	Dasatinib	-5.126	-8.794
5	Imatinib	-5.746	-7.831
6	Nilotinib	-5.590	-8.129
7	Tozasertib	-5.673	-7.957

**Table 2:** Comparison of the Docking scores of the inhibitors available for BCR-Abl and Src receptors.

#### Cytogenetic response

Ultimate goal of all the therapeutic options are to eliminate the leukemic stem cell populations from the patients for that cytogenetic analysis remains the standard options to monitor [3]. It has been reported that treatment through TKIs, patients are able to get complete molecular response. Through RTPCR, it has been observed the patients sample shows absence of the BCR-ABL transcripts. Second generation TKIs like Dasatinib and Nilotinib shows even better cytogenetic response [4]. However complete molecular responses against the leukemic cells are still matter of investigation.

# Conclusion

Successful trials of tyrosine kinase inhibitor like Imatinib has dramatically improved the survival rate of the CML patients and has drastically reduced the mortality rate to 1-2%. Moreover multiple therapeutic options involving various tyrosine kinase inhibitors like Imatinib, Nilotinib, Dasatinib, Bosutinib, and Ponatinib are being utilized extensively for treatment of CML patients. In view of the recent reports suggesting probable mutations in receptor sites the effectiveness of above said inhibitors are often questionable. Hence the multi target receptor inhibitors like Ponatinib became more potent alternative as compared to other single target inhibitors. The multi target inhibitors not only posses higher potency but also often shows low failure rate because of the tendency to avoid mutated active site. The effectiveness of dual inhibitors were being studied suggesting the ability of these inhibitors to interact with at least one of the reported drug targets for CML.

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