

Imatinib Therapy in Chronic Myeloid Leukaemia: Review of Mechanisms of Resistance and Therapeutic Options in Imatinib Failure

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ABSTRACT

Chronic myeloid leukaemia is a triphasic, clonal malignancy, arising from the haemopoietic stem cell. It is characterised by the presence of Philadelphia chromosome, which results from the reciprocal translocation between chromosome 9 and 22. The resulting oncogene- bcr-abl has proliferative activity and survival advantage against normal cell and this account for the clinical and laboratory manifestation of this myeloproliferative disorder. Imatinib, a tyrosine kinase inhibitor (TKIs) is currently the first line treatment, however, one third of patients develop resistance to it, thus necessitating alternative TKIs. Many factors are associated with the development of resistance to imatinib, such as mutation in the bcr-abl gene, increased production of the mutant protein and activation of alternative pathways amongst other causes. The aim of this review is to explore these factors, and to also evaluate current TKIs that are used as alternative in Imatinib resistant cases.

KEYWORDS: Imatinib Therapy, Resistance, Alternative.

Introduction

Chronic myeloid leukaemia (CML) is a clonal malignancy of the haemopoietic stem cell (HSC), characterized by the presence of Philadelphia chromosome (Ph). This arises from reciprocal translocation between the Abelson gene (Abl) on the long arm of chromosome 9 and the breakpoint cluster region (bcr) on the long arm of chromosome 22 [t(9,22)(q34;q11.1)]¹.

The resultant fusion oncogene (bcr-abl) codes for an abnormal protein with tyrosine kinase activity in excess of the wild type, consequently multiple signal transduction pathways are activated, such as, the STAT5/JAK, phosphatidylinositol 3 (PI3), the Ras, Raf and MAPK family of kinases. The result is the uncontrolled proliferation and

gain of survival advantage of the malignant clones over normal cells².

In Ph-positive CML, the breakpoint in the BCR gene occurs in a small 5.8 Kb major breakpoint cluster region (M-bcr). This hybrid bcr-abl gene is transcribed into a novel 8.5 kb mRNA with a b3a2 and/or b2a2 junction that encodes a 210-kD fusion protein with enhanced protein tyrosine kinase activity³. The cellular form of Abelson leukaemia virus tyrosine kinase (c-Abl) is related to the src family of tyrosine kinases. It consists of about 1150 residues, with an "N" terminal cap of about 80 residues involved in autoinhibition. The fusion of the gene encoding c-Abl with the breakpoint cluster region (BCR) results in the formation of a chimeric protein with all of the c-Abl preserved except its N-terminal region. This fusion distorts the control mechanism that keeps the c-Abl in an inactive form and the enhanced tyrosine kinase activity results in chronic myeloid leukaemia⁴⁻⁷.

The clinical course of CML is characterized by a chronic phase (CP), an accelerated phase

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(AP) and a blast crisis (BC). The last two stages are difficult to treat and associated with poor outcome¹. However, the advent of Imatinib (IM), a tyrosine kinase inhibitor (TKI) two decades ago revolutionised the treatment of CML, and is currently the first line treatment for all phases of the disease. Given that the Bcr-Abl enzyme is necessary and sufficient for the initiation and propagation of CML, inhibition of the Abl kinase activity was an ideal target for a molecular-based therapy⁸. Imatinib is a potent and selective inhibitor of the abnormal kinase domain (KD), it competitively inhibits the binding of ATP onto its binding site on the bcr-abl domain

(figure 1). Importantly, it binds to the KD in the inactive form, this makes Imatinib highly selective for the bcr-abl kinase compared to other agents (NB – the kinase are similar in their active form, but dissimilar when inactive). Imatinib displaces ATP and traps an inactive conformation of the kinase domain⁹. Unfortunately, about 1/3rd of patients may not respond, or may need to discontinue treatment due to intolerable side effects². Multiple factors have been implicated in the mechanism of resistance to this agent. The aim of this review is to explore these factors, and to also evaluate current TKIs that are used as alternative in Imatinib resistant cases.

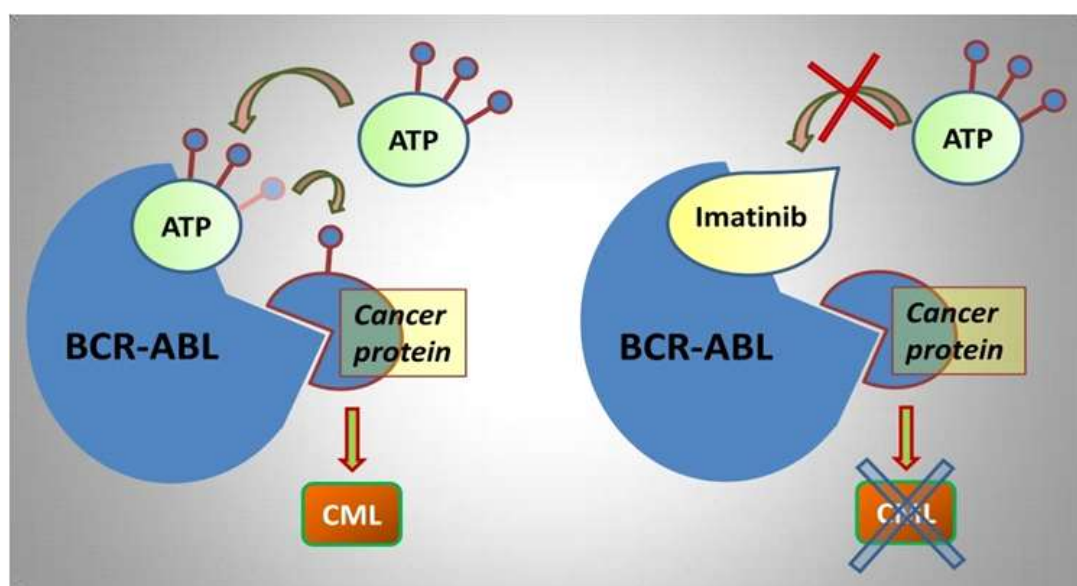


Figure 1: The mechanism of action of Imatinib. It binds to the bcr-abl ATP binding site and thereby prevents phosphorylation of the leukaemic protein (*adapted from www.intechopen.com*)

Response to imatinib treatment:

Ninety percent of patients with chronic phase CML achieve haematologic responses to imatinib, and about 50% of these patients achieve cytogenetic response using standard cytogenetic assays. Approximately 60% of patients with advanced-stage CML (blast crisis) respond to imatinib, but nearly all these patients relapse despite continued therapy¹⁰. The landmark 'IRIS' (international randomized study of interferon vs ST1571)

trial, where newly diagnosed patients with CML were randomized to treatment with Imatinib (ST1517) or interferon, established Imatinib 400mg as the first line drug for patients with CP CML¹¹. Certain criteria are used to assess if there is optimal response following treatment with imatinib. Complete haematological response (CHR) indicates a normal peripheral blood count after 3 month of therapy. Cytogenetic response can either be partial cytogenetic response (pCyR) (Ph

chromosome levels less than 35% at 6 month) or complete cytogenetic response (CCyR) (absence of Ph⁺ positive metaphases within the bone marrow at 12month)². Cytogenetic response to treatment can be monitored using either conventional cytogenetic assessment of bone marrow metaphase, or FISH using peripheral blood or marrow metaphase or interphase cells. TKIs are highly effective at reducing the number of tumour cells in the bone marrow and should be monitored by RT-PCR analysis for BCR-ABL1 transcripts in marrow or blood and/or karyotypic analysis of the bone marrow, typically at 3, 6 and 12 months¹⁰. Molecular response is assessed as a ratio of BCR-ABL1 transcripts to ABL1 transcripts and it is expressed as BCR-ABL1% on a log scale, where 10%, 1%, 0.1%, 0.01%, and 0.001% correspond to a decrease of 1, 2, 3, 4, and 5 logs respectively below the baseline. A major molecular response (MMR) is defined as BCR-ABL1 transcript level of < 0.1% or less than pre-treatment levels¹¹.

Resistance to imatinib treatment:

Failure to respond to imatinib can be classify as follow^{2,12}:

- Primary resistance: defined as incomplete haematological response at 3/12, no CyR at 6/12 months, or less than partial or complete CyR in 12 and 18 month respectively.
- Secondary resistance (acquired): this is loss of the initial responsiveness to treatment. The risk factors include older patients, advanced disease or a high sokal score at presentation, previous treatment with interferon or new clonal evolution.
- Suboptimal response: patients respond to Imatinib treatment at a lower than expected rate
- Intolerance: this group of patients require the discontinuation of treatment due to intolerable side effects.

Mechanism of resistance to Imatinib

Various mechanisms have been implicated in imatinib resistance and these can be broadly divided into BCR-ABL dependent or independent causes as follow:

BCR-ABL dependent mechanisms include:

Mutations: Numerous mutation have been implicated in resistance to Imatinib (Figure 2); this can occur in various locations in the abl sequence, including Imatinib binding sites (T3151), ATP binding sites (p-loop), catalytic and activation sites^{10,12}. Although the total number of known point mutations continues to grow, their clinical significance in an individual patient may vary, in addition, the frequency of mutation increases with disease progression. P-loop mutations account for majority of the mutation (48%), causes resistance by altering the conformation form required by imatinib to bind, associated with poor prognosis, and increase risk of disease transformation¹³. In samples obtained from imatinib-resistant patients, it was observed that a single nucleotide substitution (C→T) at position 944 of the Abl gene results in the substitution of amino acid threonine with isoleucine at position 315 of the abl protein sequence (Th³¹⁵→Ile³¹⁵; T315I); resulting in the elimination of a critical oxygen molecule required in the formation of hydrogen bonding between imatinib and the kinase domain; as well as steric hindrance to the binding of imatinib. T3151 mutation confers resistance to all TKIs in clinical use¹⁴.

Increase production or expression of BCR-ABL transcript:

Associated with secondary resistance and account for a minority of cases. The mechanism is not fully explained, but cells with increased transcript are less sensitive to imatinib in vitro¹⁵.

Presence of alpha 1 acid glycoprotein (AGP):

mediate resistance by decreasing the concentration of imatinib and thus inhibiting it at a dose dependent manner¹².



WHAT ARE THE OPTIONS IN THE FACE OF RESISTANCE

Treatment options in cases of resistance to Imatinib include

1. Imatinib dose escalation: the use of higher doses of imatinib may overcome primary resistance arising from leukemia cells exposure to sub therapeutic levels of Imatinib¹⁷. An increase from the standard dose of 400mg has been suggested to be beneficial in such resistant cases however; this may not be suitable in patient intolerant to standard dose of Imatinib. There is paucity of studies in favour of dose escalation^{18,19}.
2. Individualized therapy: Patients that fail to respond to imatinib should be screened for compliance as well as an assessment of imatinib drug levels. Additionally, mutational assessments should be made to identify possible mechanisms of resistance. Studies have indicated that patients failing imatinib can often respond to a second or third-line TKI-based therapy, including those with and without identifiable KD mutations^{19,20,21}.
3. The second generation TKIs are the next line of treatment in cases of resistance to imatinib¹⁹.
Example of some second line TKIs:
 - **Dasatinib:** inhibits both bcr-abl and src family of kinases. Shown to have more activity against the bcr-abl (>350) compared with imatinib. Effective against all phases of CML, and has a good tolerability profile. It has been extensively evaluated in the START trial, a large multicentre that included both open and randomized studies; it showed sustained response against all subgroup of patients; cross-intolerance with imatinib has not been documented, thus beneficial in

patients who are intolerable to Imatinib²². Not effective against T3151 mutation.

- **Nilotinib:** an Imatinib analogue and has been shown to be effective in patients resistant or intolerant to imatinib. Active against all phases of the disease. Side effects include elevated lipase level and hyperglycaemia, dose modification required in patients with diabetes, liver or kidney disease¹².
4. The third generation TKI bosutinib, a dual Abl/Src inhibitor, with limited activity against c-Kit and the platelet-derived growth factor receptors (PDGFRs) has been used primarily in imatinib-resistant CP-CML patients, and after a median follow-up of 8 months, 79% have achieved a CHR, with 40% achieving a MCyR, and 29% a CCyR. Ninety-one percent of patients have maintained their MCyR over this brief interval¹⁹

Unfortunately, few options are available for patients who fail to respond to both first and second generation TKIs. In addition, none of the TKIs is effective against the T3151 mutation (Table 1). Stem cell transplantation is reserved as salvage treatment in patients who relapse or resistant to second generation TKIs, and currently the only potential cure for CML, however, the procedure is associated with high morbidity and mortality, availability of suitable donor also limits its use. Indicated in patients with primary resistance or advanced disease²³.



Table 1: Summary of the various agents targeting the resistant bcr-abl clone:

Drug class	Mechanism	Effect on T3151	Comment	Trials	Reference (s)
2 nd generation TKIs: Bosotunib	Dual kinase inhibitor	No	>200fold activity compared with IM	Phase 1	Burke et al ²⁴
Batetinib (INNO-406)	Non-ATP mediated kinase Inhibition.	No	>55 potency compared with imatinib Good toxicity profile.	Phase 1	Khoury et al ²⁵
3 rd generation TKI: Ponatinib (AP24534)	Multiple kinase inhibitor-flt, src and bcr-abl	Yes	Good toxicity profile.	Phase 1/11	Bixby&Talpe ^{z17}
Aurora KI (AT9238)- Danusertib	Multiple kinase inhibitor	Yes	Nil	Phase 1	Burke et al ²⁴
Heat shock protein inhibitor (hsp).eg 17AAG	Inhibition of hsp results in down regulation of bcr-abl activities& apoptosis	No	Synergistic effect with imatinib	Nil	Burke et al ²⁴
Arsenic trioxide	Induces apoptosis and proliferation of CML cells	No	Effective against M351T,Y253F mutation	Nil	Roy-Chowdhery and Talpaz ²⁶
Homoharringtonine. (HHT) Eg: omacetaxine, chemaxenine.	By product of plant alkaloid. Induces apoptosis	No	Nil	Phase 11	Meso&Chua ^{h27}



Imatinib therapy in chronic myeloid leukaemia

Histone deacetylase inhibitors.eg Verinostat, SAHA.	Hyper acetylation of histone residues, leading to ↑ expression of p21&p27 and apoptosis	No	Increases sensitivity of mutant cells to imatinib	Nil	Meso&Chuah, ²⁷
Proteasome inhibitors.eg bortezomib	Cell cycle inhibition and induces apoptosis	No	Combination of low dose bortezomib and IM lead to anticancer	Nil	Meso&Chuah, ²⁷
Flavone and flavopiridol	Cyclin dependent kinase inhibitors	No	Additive effect with IM in resistant cells	Phase 11	Meso&Chuah, ²⁷
Farnesyltransferase inhibitor.eg Tipitarnib,lonatarnib	Ras/MAPK inhibitors	Partial	CHR achieved in IM resistant cells	Phase 11	Burke et al ²⁴
Raf -1- inhibitors	Multiple kinase inhibitor	No	Effective against IM resistance	Phase1/11	Burke et al ²⁴
MEK inhibitor.eg CI-1040	Nil	Partial-with arsenic and hsp)	Nil		Burke et al ²⁴
MTOR inhibitors.eg Rapamycin, sirolimus	Nil	Partial	Poor biochemical property	Phase1/11 (everolimus)	Burke et al ²⁴



Monitoring Response to Treatment

Detection of mutation at an early stage can allow for modification of patients' treatment. In the last decade, various techniques that can detect the bcr-abl mutations have been developed, which has allowed for the ongoing monitoring of patients to help identify those at risk of disease progression²⁸.

Current guidelines recommend that patient monitoring should include a baseline profile and regular monitoring as outlined below^{29,30}.

- Baseline assessment: these should include a blood bcr-abl transcript detection (RQ-PCR) to establish the diagnosis of CML; type of transcript present [e.g of common transcript are e13a2 (b2a2) and e14a2 (b3a2)], this allow for identification of any new transcript due to disease progression; bone marrow cytogenetic to detect presence of extra cytogenetic abnormalities or translocations.
- Response to treatment: for patients responding to treatment, monitoring entails twice monthly blood count, bcr-abl transcript every 3/12 and marrow for cytogenetic analysis every 6/12.

- Establishment of CCyR: monitoring of bcr-abl transcript, importantly, to characterize the pattern of change, such as, whether it is declining, steady or rising. FISH can also be used to assess for CCyR.

- Mutation analysis: this is indicated in patients with rising levels of transcript, advance disease at presentation, loss of initial response or suboptimal response in patients with CP disease. Also important in deciding second line TKIs to use or if SCT should be offered.

Conclusion

Chronic myeloid leukaemia resulting from Philadelphia chromosome positivity can be reliably managed with imatinib, however, the malignant clone are unstable and prone to developing resistance. Patient with resistance can be offered treatment with second generation or third generation TKIs or salvage treatment with SCT, however, the outcome is poor with the latter. Various agents targeting the resistant bcr-abl clone are under evaluation, with the aurora kinase family showing positive result against the T3151 mutation.

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Cited this article as: A. I. Ladu, Aisha M. Abba.

Imatinib Therapy in Chronic Myeloid Leukaemia: Review of Mechanisms of Resistance and Therapeutic Options in Imatinib Failure. *Bo Med J* 2017; 14(1): 18-27.

Source of Support: Nil, **Conflict of Interest:** None declared.

