Suppression of autophagy by BCR/ABL

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1. ABSTRACT

Imatinib and second generation BCR/ABL tyrosine kinase inhibitors (TKIs) serve now as standard therapies for patients with chronic myelogenous leukemia (CML); however, CML stem cells are intrinsically insensitive to the cell death-inducing effects of TKIs, allowing the persistence of a "reservoir" of BCR/ABLexpressing CML-initiating cells potentially responsible for disease relapse and progression. Although it is still controversial whether the "insensitivity" of CML stem cells to treatment with TKI is due to BCR/ABL-dependent or independent mechanisms, treatment with IM appears to suppress BCR/ABL-dependent signaling in CML stem cells with no adverse effects on their survival. Recent evidence indicates that BCR/ABL suppresses and treatment of CML cells with IM/TKIs induces autophagy, a geneticallyregulated process of adaptation to metabolic stress which could allow tumor cells to become metabolically inert enabling their survival under conditions that may mimic growth factor/nutrients deprivation. Based on this hypothesis, TKI-induced autophagy may "antagonize" TKI-induced cell death and inhibition of autophagy may eliminate this survival mechanism by restoring "sensitivity" of CML stem cells to treatment with IM/TKI. Consistent with this, phenotypically and functionally defined CMLenriched stem cells insensitive to treatment with TKI are efficiently eliminated by the combination of TKI and chloroquine, an inhibitor of late stage autophagy. Thus, inhibition of autophagy may improve the potent and specific effects of TKIs by rendering CML stem cells sensitive to these targeted therapies.

2. INTRODUCTION

CML is a malignancy arising from transformation of the hematopoietic stem cell which typically progresses in three distinct disease stages: an indolent chronic phase (CP) characterized by the accumulation of mature granulocytes and myeloid precursors in the bone marrow and the peripheral blood; an accelerated phase (AP) characterized by an increase in disease burden and in the frequency of myeloid precursors: and an acute phase called blast crisis (BC) marked by increasing numbers of differentiation-arrested blast cells of either myeloid or lymphoid lineage (1-3). The hallmark of all phases is the Philadelphia chromosome (Ph1) which results from a reciprocal translocation of chromosomes 9 and 22 and generates BCR/ABL fusion genes that encodes a p210 KD protein with constitutive tyrosine kinase activity Transformation of hematopoietic stem cells by p210BCR/ABL requires its tyrosine kinase activity and depends on its ability to activate a multitude of intracellular signaling pathways which include the classical PI-3K/Akt and MAP kinase and the more recently described βcatenin/LEF-1 and Hedgehog pathways. Together, these pathways promote an expansion of the pool of committed myeloid progenitors and lead to their increased survival and proliferation and limited dependence on growth factors (5-10).

The generation of the BCR/ABL kinase ATP-competitive inhibitor imatinib mesylate (IM) has revolutionized the therapy of CML, since this drug is highly effective in the CP of the disease (11). However,

there are three major problems with IM-based therapy: i) the limited response of CML-BC or Ph¹ B-acute lymphoblastic leukemia (ALL) patients to IM (12-14); ii) the development of resistance caused in approximately 40% of cases by mutations in the BCR/ABL kinase domain which impair the ability of IM to interact with the protein (15-17); and iii) the relative insensitivity of Ph¹ stem cells to IM resulting in the persistence of residual leukemic cells (and possibly disease relapse) even in patients with complete cytogenetic response (CCyR) (18). For these reasons, more potent BCR/ABL inhibitors, also targeting IM-resistant mutants including BCR/ABLT315I, are being The insensitivity of developed and tested (19-22). primitive Ph^1 stem cells (which overexpress p210BCR/ABL) to treatment with IM and second generation (nilotinib, dasatinib, bosutinib) TKI (18, 23-27) is specially concerning because even new TKI capable of blocking the activity of the T315I mutant which is not bound by IM and second generation TKI (i.e. AP24534; reference 22) may not be able to eradicate Ph¹ stem cells from patients with the T315I mutation. The reasons for the insensitivity of CML stem cells to treatment with TKI are incompletely understood; several BCR/ABL-dependent and independent mechanisms including increased expression/activity of BCR/ABL, autocrine production of growth factors, down-regulation of drug transporters and persistent activation of proliferation/survival signaling pathways in spite of BCR/ABL inhibition have been proposed (28). Although it is still controversial whether TKI can suppress BCR/ABL activity in Ph¹ stem cells (25,26,29), a recent study provides strong evidence in support of IM being able to inhibit the tyrosine kinase activity of BCR/ABL in CML stem cells (29); however, this study also showed that cytokines support allowed Ph¹ stem cells to grow and survive like the normal counterpart (29), suggesting that CML stem cells may be "insensitive" to treatment with TKI via BCR/ABL-independent mechanisms or that the inhibition of BCR/ABL signaling in CML stem cells leads to "context-specific" metabolic changes that may antagonize its cell death-promoting effects. Thus, it is crucial to understand mechanistically why CML stem cells are insensitive to TKI and to develop novel therapeutic strategies that, in combination with TKI, might be effective in targeting the stem cell population.

3. REGULATION OF AUTOPHAGY BY BCR/ABL

3.1. Autophagy is an adaptation process that provides a survival mechanism for cancer cells

Macroautophagy (hereafter referred to as autophagy) is a degradative process in eukaryotic cells that results in the sequestration and the breakdown of intracellular organelles and proteins within specialized lysosomes called autolysosomes under homeostatic conditions or in response to stress (30, 31). This process allows cells to adapt to developmental changes and/or unfavorable environmental conditions (30, 31). Autophagy is a genetically controlled process which progresses through discrete steps which lead to the engulfment of long-lived proteins and whole organelles into multimembraned vacuoles called autophagosomes (30,31). Autophagosomes then fuse with lysosomes to produce

autolysosomes for final destruction and recycling (30.31). While in certain cellular contexts autophagy can serve as a cell death mechanism, named type II cell death (32, 33), it is becoming increasingly clear that this process can provide a cell survival mechanism which allows cells to adapt their metabolism to starvation caused by a decrease in metabolite concentrations or extracellular nutrients, consequences of loss of growth factor signaling (34-36). In this context, autophagy serves as an adaptation mechanism to evade programmed cell death. Consistent with a survival role played by autophagy, its genetic (knockdown of autophagy genes) or pharmacological (by use of drugs such as chloroquine, an inhibitor of autophagosome-lysosome fusion and lysosomal acidification (37)) inhibition results in cell death of growth factor-starved cells in which apoptosis has been genetically ablated (36,38). The prosurvival effect associated with induction of autophagy is not limited to growth factor-starved normal cells but was also observed in tumor cells: treatment of Myc-induced lymphomas with chemotherapy or drugs inducing p53 reactivation led to the appearance of morphological and biochemical markers of autophagy and pharmacological or genetic inhibition of autophagy enhanced the p53- or chemotherapy-dependent anti-tumor effects in vivo (37, 41). Together, these and other studies suggest that induction of autophagy provides a protective mechanism to tumor cells.

However, in tumors displaying defective apoptosis, inhibition of autophagy causes caspase-independent necrotic cell death, which, in turn, augments inflammation, leading to enhanced tumor burden (42). Autophagy can limit inflammation also in normal cells, thus counteracting tumorigenesis (42). Thus, the consequences of autophagy inhibition might be double-faced, as it can either promote or suppress tumorigenesis depending on the tumor type or inflammation status (37, 39, 42).

As shown by the study by Amaravadi *et al.* (39) in which mice with Myc-induced lymphomas were treated with alkylating drugs, the autophagic process can be also activated by chemotherapy and it provides a mechanism to chemotherapy-treated cells to evade cell death. Consistent with the findings of this study, autophagy inhibition sensitizes tumor cells to cell death induced by irradiation (43-45), alkylating agents (46), or arsenic trioxide (47), suggesting that cancer cells can react to chemotherapy by inducing autophagy as a self-defence mechanism (34). Such a mechanism might be particularly important in allowing persistence of drug-resistant dormant cancer cells (48, 49).

How an increase in intracellular lysosome-mediated catabolism results in reduced sensitivity to cell death is still unclear. Perhaps, clearance of mitochondria through autophagy (mitophagy) might sequester proapoptotic proteins (i.e. BH3-only proteins) and prevent the release into the cytosol of apoptosis activators such as cytochrome C (50). Alternatively, autophagy suppression may lead to reduced availability of metabolites for bioenergetic needs, thus impairing the survival capacity of

cancer cells (see also below). Finally, as specific proteins can be targeted for degradation via the p62 cargo pathway (51), autophagy suppression could result in reduced degradation of cell death inducers.

3.2. BCR/ABL suppresses autophagy by a tyrosine kinase-dependent mechanism

The oncogenic p210BCR/ABL protein has the ability to activate several survival pathways which mimic those activated by growth factor signaling (8,52), allowing transformed cells to acquire resistance to apoptosis induced by growth factor deprivation (8, 52). Inhibition of tyrosine kinase activity by IM results in the induction of cell death which might be caused by a sudden decrease of intracellular survival signals and a relative increase in proapoptotic signals (53). We have recently shown that BCR/ABL expression also prevents IL-3-deprivationinduced autophagy and that pharmacological inhibition of BCR/ABL tyrosine kinase activity, in addition to markers of apoptosis, leads to the rapid appearance of morphological and biochemical changes consistent with autophagy (54). Mechanistically, induction of autophagy by treatment with TKIs is not due to inhibition of c-Abl tyrosine kinase activity since expression of IM-resistant T315I c-Abl did not rescue IM-induced autophagy in hematopoietic cells expressing native BCR/ABL. Induction of autophagy was also independent from apoptosis as inhibition of caspase activity or Bcl-2 overexpression did not prevent IM-induced autophagy in BCR/ABL-expressing myeloid progenitor cells. Induction of autophagy did not require expression of wild type p53 since it was also induced by TKIs of p53-null K562 cells (54).

Although the precise mechanisms responsible for the ability of BCR/ABL to suppress autophagy remain unknown, it is likely that the PI-3K/Akt/mTOR is involved since treatment of BCR/ABL-expressing cells with Akt or mTOR inhibitors induced autophagy (54, data not shown). Moreover, the autophagic process activated by Imatinib treatment of BCR/ABL-expressing cells was preceded by the induction of endoplasmic reticulum (ER) stress and relied on intracellular calcium as autophagy was blocked by intracellular calcium chelators (54). However, it is unclear whether inhibition of Akt or mTOR is functionally linked to the ER stress response in the induction of autophagy.

Regardless of the mechanisms involved in the BCR/ABL-dependent suppression of autophagy, treatment with pharmacological inhibitors of autophagosomelysosome fusion (chloroquine or bafilomycin A) potentiates IM-induced cell death in CML cell lines and primary CML cells, including those carrying partially IM-resistant BCR/ABL mutants (52). Moreover, candidate CML stem cells, defined as colony forming CD34+CD38- cells and LTC-IC were also exquisitely susceptible to combination treatment (54). Genetic suppression of autophagy by knockdown of the autophagy genes Atg5 and Atg7 in K562 and primary CML cells also enhanced TKI-induced cell death, supporting the specificity of the effects induced by pharmacological inhibitors (54).

Together, these findings indicate that induction of autophagy provides a survival mechanism to IM-treated BCR/ABL-expressing cells, including the stem cell population, and suggest that inhibition of autophagy may improve the therapeutic efficacy of TKI in the treatment of CML by a more efficient elimination of stem cells. Based on this study, a randomized phase 2 clinical trial of IM versus hydrochloroquine (HCQ)/IM in CML patients with major cytogenetic response and residual disease by Q-PCR detection of bcr/abl transcripts has been approved and patients are presently being recruited in the UK at three centers (Glasgow, Liverpool and London) under the leadership of Dr Holyoake.

3.3. Is there a molecular basis for tki-induced autophagy versus apoptosis in cml stem cells?

The enhanced TKI-induced cell death associated with pharmacological inhibition of late stage autophagy (chloroquine treatment) in CML cells suggests that induction of autophagy by TKI has an "antagonistic effect" on TKI-induced cell death. This effect appears to be especially prominent in CML stem cells which were completely insensitive to apoptosis induction by TKI treatment unless autophagy was also suppressed (52). An explanation for these data is that TKI suppress BCR/ABL activity in CML stem cells but the consequence of this effect is a preferential induction of autophagy rather than apoptosis. Perhaps, inhibition of BCR/ABL tyrosine kinase activity in CML stem cells preferentially activates signal transduction pathway (s) leading to autophagy rather than apoptosis; although a number of pathways and genes regulating autophagy overlap with those involved in apoptosis (55), there are some (i.e. mTOR -dependent signaling) which appear to be more important for the regulation of autophagy than apoptosis (56), raising the possibility that TKI treatment of distinct CML progenitor/stem cell subsets elicits differences in the "intensity" of autophagy- versus apoptosis-regulatory signals.

One possibility is that IM treatment activates markers of autophagy (i.e., LC3-II expression and subcellular localization) rather than apoptosis (i.e., Bim levels) in stem cell-enriched (CD34+CD38-) versus progenitor-enriched (CD34+CD38+) CML subsets and this may reflect the preferential activation of signaling pathways which regulate autophagy versus apoptosis.

Although autophagy and apoptosis share many molecular regulators (i.e., the PI-3K/Akt pathway), another possibility is that different signal thresholds may be required in specific cell subsets (i.e. CD34+CD38- versus CD34+CD38+) to inhibit autophagy versus apoptosis (i.e. phosphorylation of Akt) and the consequence of suppressing Akt signaling (via IM treatment, or directly through an Akt inhibitor) may be the preferential induction of autophagy versus apoptosis (Figure 1).

Thus, there may be differences in the activation of autophagy- versus apoptosis- regulatory Akt-dependent signaling pathways in distinct progenitor subsets.

Tyrosine kinase inhibitors Cell death Autophagy Committed progenitors? Leukemic stem cells? Tumour killing Survival of cancer stem cells

Figure 1. Treatment with TKI may elicit different responses in distinct CML progenitor subsets. The response to TKI may be different in distinct CML progenitor subsets. One possibility is that in committed progenitors TKI preferentially activate cell death rather than autophagy. By contrast, more primitive progenitors (stem cells) would be more resistant to cell death due to their propensity to activate preferentially the process of autophagy. As a result, the debulking effect of TKIs would be accompanied by survival of resistant leukemic stem cells. Survival of these autophagy-prone stem cells would be responsible for promoting disease progression.

If such differences are not detected in spite of the fact that IM treatment induces autophagy but not apoptosis in the CD34+CD38- CML subset (54), alternative mechanisms (i.e., increased expression/activity of autophagy activators or apoptosis inhibitors) may explain the predominance of autophagy versus apoptosis in TKItreated CML stem cells. Experimental approaches to identify such alternative mechanisms may include screening of oligonucleotide RNA arrays, proteomic screening of pathway-specific or analyses and comprehensive phospho antibody arrays RNA/proteins of IM-treated CD34+CD38-CD34+CD38+ CML cells. It is also conceivable that CML stem cells are resistant to cell death following inhibition of BCR/ABL tyrosine kinase activity because of their quiescent state and limited bioenergetic needs. In this context, induction of autophagy would supply sufficient metabolites to provide energy to quiescent stem cells (Figure 2).

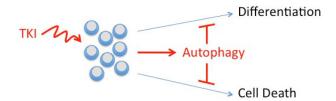
Induction of autophagy in TKI-treated CML stem cells may, mechanistically, prevent apoptotic cell death by autophagosome sequestration of regulators (i.e. BH3-only proteins) of the intrinsic, mitochondrial apoptotic pathway (Figure 2). In IM-treated K562 cells, inhibition of autophagosome formation by RNAi of Atg5 or Atg7

expression enhanced IM-induced cell death as effectively as inhibition of "autophagy flux" by chloroquine or bafilomycin A1 treatment (54). CML-enriched stem cells were sensitized to TKI by treatment with chloroquine, but genetic approaches are necessary to determine whether inhibition of autophagosome formation also enhances TKI-induced cell death of CML stem cells. The identification of the stage of the autophagic process which "antagonizes" TKI-induced cell death is not only biologically relevant but has also translational implications; the insensitivity of CML stem cells to TKI-induced cell death could be manipulated by harnessing mitochondria-independent cell death pathways or by restoring the expression/activity of cell death mediators lost/inactivated via enhanced autophagic flux.

4. CONCLUSION

Although autophagy appears to be induced earlier than apoptosis in CML cells (54), it is also possible that signals leading to either process could be activated simultaneously. In this regard, failure of TKI to induce apoptosis of CML stem cells may depend on the predominant activation of autophagy rather than apoptosis. Reversal of this "antagonistic" effect of autophagy seems capable of "sensitizing" CML stem cells to the apoptosis-

Role of autophagy in the cancer stem cell pool



HOW?

- •Selective degradation of proteins involved in cell death induction, cell cycle progression or differentiation
- •Generation of metabolism intermediates for energy production
- •Removal of dysfunctional mitochondria

Figure 2. Role of autophagy in the cancer stem cell pool. Induction of autophagy by TKI treatment may maintain CML stem cells by preventing cell death, inhibiting differentiation, or blocking cell cycle progression. Metabolism intermediates generated by autophagy may be also needed for maintenance of stem cells.

promoting effects of TKI, although *in vivo* studies have not been performed yet to assess the effects of autophagy inhibition on TKI-treated CML stem cells. Since failure to eliminate CML stem cells remains, probably, the major problem of TKI-based therapies (including second and third generation inhibitors targeting mutant BCR/ABL, including the T315I mutation), the ability of autophagy inhibition to "sensitize" CML stem cells to TKI would make unnecessary to target CML stem cells through BCR/ABL-independent, less specific approaches. At the same time, normal stem cells would be spared as pharmacological inhibition of autophagy alone has modest or no effects on normal or CML progenitors (54).

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Autophagy in CML stem cells

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