

REVIEW

Pharmacological targeting of the KIT growth factor receptor: a therapeutic consideration for mast cell disorders

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KIT is a member of the tyrosine kinase family of growth factor receptors which is expressed on a variety of haematopoietic cells including mast cells. Stem cell factor (SCF)-dependent activation of KIT is critical for mast cell homeostasis and function. However, when KIT is inappropriately activated, accumulation of mast cells in tissues results in mastocytosis. Such dysregulated KIT activation is a manifestation of specific activating point mutations within KIT, with the human D816V mutation considered as a hallmark of human systemic mastocytosis. A number of other activating mutations in KIT have recently been identified and these mutations may also contribute to aberrant mast cell growth. In addition to its role in mast cell growth, differentiation and survival, localized concentration gradients of SCF may control the targeting of mast cells to specific tissues and, once resident within these tissues, mast cell activation by antigen may also be amplified by SCF. Thus, KIT inhibitors may have potential application in multiple conditions linked to mast cells including systemic mastocytosis, anaphylaxis, and asthma. In this review, we discuss the role of KIT in the context of mast cells in these disease states and how recent advances in the development of inhibitors of KIT activity and function may offer novel therapies for the treatment of these disorders.

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Abbreviations: Btk, Bruton's tyrosine kinase; FcεRI, high-affinity receptor for IgE; FLT3, FMS-like tyrosine kinase 3; HMC, human mast cell line; IL, interleukin; M-CSF, macrophage colony stimulating factor; PDGFR, platelet-derived growth factor receptor; PI3K, phosphoinositide 3-kinase; SCF, stem cell factor; TNF-α, tumour necrosis factor-α; VEGFR, vascular endothelial growth factor receptor

Introduction

The human *c-KIT* oncogene, which is mapped to the *W* locus in the mouse, encodes for a protein, KIT (CD117), which is a member of the transmembrane receptors with tyrosine kinase activity superfamily. This family also includes other growth factor receptors, namely FMS-like tyrosine kinase 3 (FLT3), the platelet-derived growth factor receptor (PDGFR), and the macrophage colony stimulating factor (M-CSF) receptor (Broudy, 1997; Patnaik *et al.*, 2007). Expression of human KIT is primarily restricted to melanocytes, germ cells and cells of haematopoietic lineage, including bone marrow progenitor cells, mast cells, megakaryocytes (Kitamura *et al.*, 1995, 2006; Roskoski, 2005a,b; Alexeev and Yoon, 2006) and, to a lesser extent, basophils and eosinophils (Yuan *et al.*, 1997; Heinemann *et al.*, 2005). It has also been reported to be

expressed in other cell types such as epithelial cells and vascular smooth muscle cells (Al-Muhsen *et al.*, 2004; Hollenbeck *et al.*, 2004). KIT is expressed as several alternatively spliced isoforms, which, following glycosylation, give rise to proteins with multiple molecular weights around 145 kDa (Yarden *et al.*, 1987). The ligand for KIT, stem cell factor (SCF), also known as steel factor based on its generation by the Steel locus (*Sl*) in mouse (Yarden *et al.*, 1987; Huang *et al.*, 1990), exists in two forms that are produced by alternative splicing: a soluble form of approximately 31 kDa and a membrane-bound form of approximately 32 kDa, which lacks the proteolytic site for processing into the soluble form (Flanagan and Leder, 1990). In both human and mouse, there is evidence that KIT may respond differentially to these individual forms of SCF (Toksoz *et al.*, 1992; Kapur *et al.*, 1998; Trieselmann *et al.*, 2003). SCF is expressed in fibroblasts, thymus tissue, spleen, testes, placenta and mast cells (reviewed by Broudy, 1997; Reber *et al.*, 2006) and, under normal conditions, in humans it is present in the plasma at concentrations of approximately

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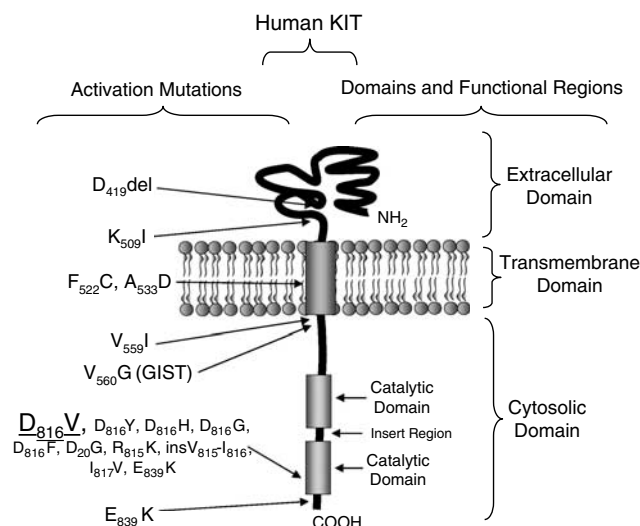


Figure 1 Structure of human KIT and selected mutations associated with specific human disease states.

1–3 ng mL⁻¹ (~0.2 nM) (Kojima *et al.*, 1997). However, local concentrations near the sites of origin and localized membrane-associated KIT concentrations are likely to be significantly higher.

KIT has five immunoglobulin-like regions within its extracellular domain that contain the binding site for SCF (Figure 1). The K_d for the binding of SCF to human KIT is reported to be 4.2 pM for high-affinity binding and 1.7 nM for lower-affinity binding (Broudy *et al.*, 1994). Thus, reasonable increases in localized concentrations of SCF would be sufficient for optimal activation of the receptor. As with other receptors with tyrosine kinase activity, KIT-mediated cellular responses follow ligand-induced dimerization of the receptor and resulting activation of its inherent tyrosine kinase activity. This activity is associated with a split catalytic domain contained within its cytosolic domain (Mol *et al.*, 2003) (Figure 1). The KIT kinase activity targets specific tyrosine residues also contained within the cytosolic domain of KIT. Thus, once ligated, KIT undergoes auto/transphosphorylation (Mol *et al.*, 2003). The Src kinase, Lyn, may also contribute to phosphorylation of specific sites within KIT (Shivakrupa and Linnekin, 2005). Following phosphorylation, sequences containing these phosphotyrosines become docking sites for associating critical signalling molecules. These molecules include the Src kinases Lyn and Fyn, phospholipase C γ , phosphoinositide 3-kinase (PI3K) and the adaptor molecules Grb2 and Shc (Lev *et al.*, 1992; Serve *et al.*, 1994; Herbst *et al.*, 1995; Linnekin *et al.*, 1997; Price *et al.*, 1997; Thommes *et al.*, 1999; Gilfillan and Tkaczyk, 2006; Samayawardhena *et al.*, 2006) (Figure 2). The specific residues responsible for these interactions have been mapped by site-directed mutagenesis, revealing that, in the case of human KIT, phospholipase C γ binds to pY936 (Herbst *et al.*, 1995), PI3K to pY721 (Herbst *et al.*, 1995), Grb2 to pY703 and pY936 (Herbst *et al.*, 1995; Thommes *et al.*, 1999), and Lyn and Fyn to pY568 and pY570 (Linnekin *et al.*, 1997; Samayawardhena *et al.*, 2006). By recruiting these molecules, a receptor-signalling molecule complex is established which coordinates the critical downstream signalling events

leading to the diverse cellular responses attributable to KIT. Also recruited are a number of signalling molecules whose function is to terminate an ongoing reaction. These include the phosphoinositide phosphatase SHIP (van Dijk *et al.*, 2000), which reverses the actions of PI3K, the protein phosphatase SHP1 (Kozlowski *et al.*, 1998), which reverses the phosphorylation events initiated by KIT, the tyrosine kinase Chk (Price *et al.*, 1997), which inactivates Src kinases, and SOCS6 (Bayle *et al.*, 2004), which downregulates signalling pathways leading to cytokine gene expression. As it is not the purpose of this review to discuss in length the downstream signalling processes that account for the biological responses mediated by KIT, readers are referred to several excellent reviews that contain in-depth discussions of these events (Broudy, 1997; Linnekin, 1999; Lennartsson *et al.*, 2005; Roskoski, 2005a,b; Reber *et al.*, 2006) and to Figure 2, which summarizes these processes.

Roles of KIT and normal and dysregulated control of mast cell function

Stem cell factor-dependent ligation of KIT, and the subsequent activation of KIT kinase activity, is essential for mast cell homeostasis. Mice that are defective for functional KIT expression (*W/W^v* and *W/W^{sh}*) (Kitamura *et al.*, 1978) and those defective for SCF (*Sl/SI^d*) (Kitamura and Go, 1979) are thus mast cell deficient. In humans, expansion and differentiation of mast cells from their CD34⁺/CD117⁺ progenitor cells is also dependent on SCF, as demonstrated in *ex vivo* culture systems (Kirshenbaum *et al.*, 1999). Once mature, the continued survival of human mast cells is also dependent on the presence of SCF, as cells deprived of SCF display signs of apoptosis within 24–48 h (Metcalf *et al.*, 1995). A number of activating mutations in KIT are associated with the dysregulated growth of mast cells associated with mastocytosis (Akin and Metcalfe, 2004). These activating mutations will be discussed later. Under experimental conditions, *in vitro*, SCF has been demonstrated to be a potent chemotactic agent for mast cells (Nilsson *et al.*, 1994; Dastyk *et al.*, 1998) and mast cell precursors (Taylor *et al.*, 2001), and to induce mast cell adhesion to basement membrane proteins (Dastyk and Metcalfe, 1994; Lorentz *et al.*, 2002). Thus, *in vivo*, in addition to its role in mast cell homeostasis, SCF likely contributes to the processes that regulate the homing of mast cells to their sites of residence within tissues/organs such as skin, intestinal mucosa and submucosa, alveolar walls, nasal mucosa, bronchial subepithelium and tonsils (Okayama and Kawakami, 2006). However, the extent to which SCF contributes to this process *in vivo* is currently unknown.

In addition to its ability to regulate mast cell homeostasis and tissue distribution, SCF is also recognized as a potent modifier of mast cell activation. When triggered, mast cells release an array of inflammatory mediators that contribute to the initiation of anaphylaxis and the inflammatory reactions associated with the asthmatic response (Metcalf *et al.*, 1997). These mediators include granule-associated bioactive amines such as histamine and 5-hydroxytryptamine (serotonin), eicosanoids such as prostaglandin D₂ and

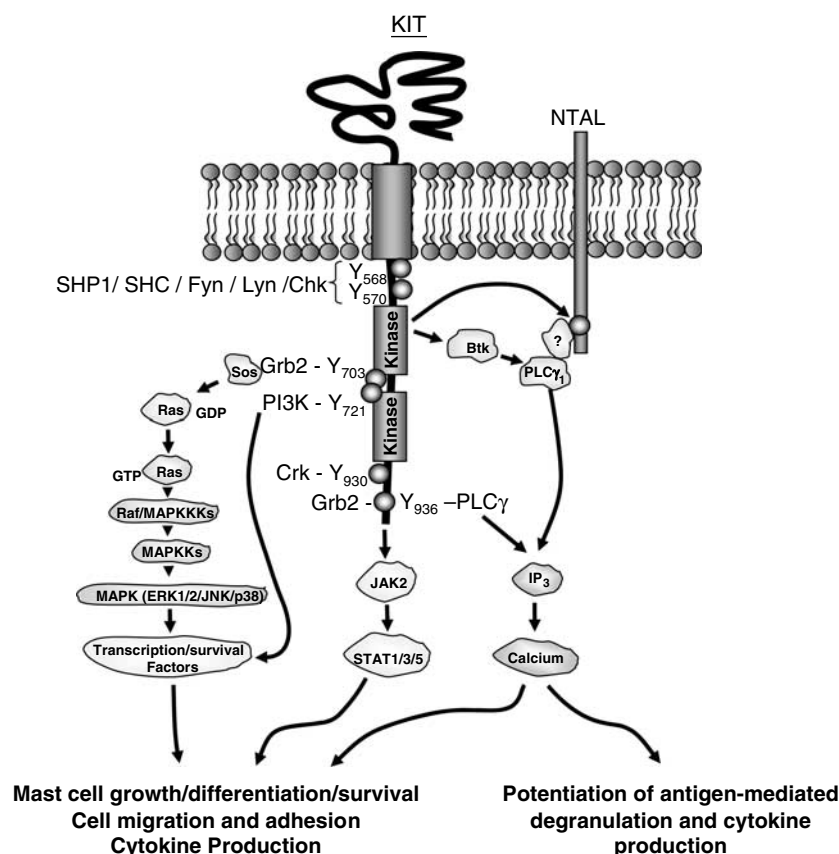


Figure 2 Binding of specific signalling proteins to phosphorylated tyrosine residues on KIT following receptor activation and the subsequent signalling events leading to mast cell responses. For clarity, only one receptor is shown in the figure. SCF-induced KIT dimerization enhances the tyrosine kinase activity associated with the split catalytic domain. This results in phosphorylation of specific tyrosine residues within the cytosolic domain of KIT. The tyrosine kinase, Lyn, may also contribute to this response. The consequential recruitment of critical signalling molecules such as the tyrosine kinases, Lyn and Fyn, the adaptor molecules, SHC and Grb2 and the signalling enzymes, PI3K and phospholipase $C\gamma$, results in the activation of the Ras–Raf–mitogen-activated protein kinase (MAPK) pathway, the enhancement of intracellular calcium levels, and the activation of transcription factors leading to mast cell growth, differentiation and survival; enhanced cell migration and chemotaxis; and cytokine production. Activation of the JAK2 and STAT1/3/5 likely contributes to these responses. Concurrent with these events, KIT induces activation of the Tec kinase Btk and the adaptor molecule NTAL (non-T-cell activation linker), resulting in an enhancement of antigen-mediated degranulation and cytokine production. (For further details, please refer to Broudy, 1997; Linnekin, 1999; Tkaczyk *et al.*, 2004; Iwaki *et al.*, 2005; Lennartsson *et al.*, 2005; Roskoski, 2005a, b; Gilfillan and Tkaczyk, 2006; Reber *et al.*, 2006.)

leukotriene C_4 , chemokines such as CCL2, CCL3, CCL5 and cytokines such as tumour necrosis factor- α (TNF- α), granulocyte macrophage colony-stimulating factor, interleukins (ILs) 3, 4, 5, 6, 10 (Galli *et al.*, 2005; Gilfillan and Tkaczyk, 2006). These events generally occur following antigen-dependent aggregation of IgE-occupied high-affinity receptors for IgE (Fc ϵ RI) on the mast cell surface (Metcalf *et al.*, 1997; Galli *et al.*, 2005; Rivera and Gilfillan, 2006; Tkaczyk *et al.*, 2006). Fc ϵ RI-mediated mast cell activation, however, can be dramatically enhanced by the concurrent activation of other receptors on mast cells, including KIT, and those for adenosine, sphingosine-1-phosphate, complement component C3a, PGE $_2$ and lysophosphatidic acid (Gilfillan and Tkaczyk, 2006; Kuehn and Gilfillan, 2007), although, in cultured human mast cells, adenosine and PGE $_2$ can also inhibit antigen-induced mast cell degranulation (Kuehn and Gilfillan, 2007). Owing to its unique influence on mast cell homeostasis (Kitamura *et al.*, 2006), however, of these receptors, KIT is likely the major modifier of antigen-mediated mast cell responses in a physiological setting. The evidence supporting the role of KIT as a co-activator of

mast cells, however, has primarily emerged from studies conducted in both rodent and human mast cell cultures where it has been demonstrated that the threshold for antigen-mediated degranulation is lowered and the magnitude of this response is synergistically enhanced by SCF (Gilfillan and Tkaczyk, 2006). These studies have also demonstrated that antigen-mediated cytokine production in mast cells is also dramatically enhanced by SCF. Thus, as with the role of KIT on mast cell homing, the precise contribution of KIT activation on mast cell activation *in vivo* has yet to be determined.

KIT and mast cell disorders

Although KIT is a critical molecule in haematopoiesis, gametogenesis, and mast cell development, activating mutations resulting in ligand-independent autophosphorylation may lead to dysregulated growth of the affected cells, thereby inducing tumorigenesis (Akin and Metcalfe, 2004). Diseases such as gastrointestinal stromal tumours, systemic

mastocytosis, germ cell tumours and core factor binding acute myeloid leukaemias carry mutations in KIT (Patnaik *et al.*, 2007). However, in terms of mast cell disorders, the two principal conditions where aberrant KIT activation may play a role are mastocytosis and anaphylaxis.

Mastocytosis

Pathologic constitutive activation of KIT is associated with the mast cell proliferative disorder, mastocytosis (Valent *et al.*, 2001, 2003b), as originally described by Nagata *et al.* (1995). More than 90% of patients with systemic mastocytosis have the D816V KIT point mutation, resulting in SCF-independent autophosphorylation (Akin, 2006; Garcia-Montero *et al.*, 2006). The aspartic acid produced by Codon 816 of KIT is located in the tyrosine kinase domain and is critically involved in ATP binding and subsequent phosphotransferase activity of the receptor (Mol *et al.*, 2003; Vendome *et al.*, 2005). Mutations resulting in replacement of this aspartic acid with valine stabilize the kinase in its active conformation (Mol *et al.*, 2004; Vendome *et al.*, 2005), thus obviating the need for binding SCF for autophosphorylation (Furitsu *et al.*, 1993). Although D816V is by far the most common mutation detected in mastocytosis, mutations involving other KIT domains (such as in the juxtamembrane, transmembrane and extracellular regions) have also been described (Figure 1).

The exact role of D816V KIT in pathogenesis of mastocytosis remains to be identified. *In vitro* studies showed that D816V KIT was able to confer growth factor independence to transformed haematopoietic cell lines (Kitayama *et al.*, 1995) as well as murine haematopoietic progenitor cells (Kitayama *et al.*, 1996). Human mast cells carrying D816V KIT were more resistant to apoptosis induced by SCF withdrawal (Akin *et al.*, 2003) and migrated more vigorously to SCF (Taylor *et al.*, 2004). Animal studies showed some, but not all (8 out of 28), transgenic mice carrying D816V c-KIT mutation under the control of chymase promoter develop a disease state resembling mastocytosis with infiltration of tissues such as skin and spleen (Zappulla *et al.*, 2005). On the other hand, mice transplanted with haematopoietic stem cells retrovirally transduced with D814V KIT, as well as mice transgenic for this mutation, develop acute leukaemias and lymphomas but not mastocytosis (Kitayama *et al.*, 1996). In humans, activating KIT mutations are seen not only in mastocytosis, but also in core factor binding acute leukaemias (Beghini *et al.*, 2000), gastrointestinal stromal tumours (Hirota *et al.*, 1998), germ cell tumours (Tian *et al.*, 1999) and some lymphoproliferative disorders (Hongyo *et al.*, 2000). The presence of KIT mutations do not appear to correlate with aggressiveness or prognosis of mastocytosis in humans. Overall, these observations suggest that, although KIT mutations (particularly D816V) are important pathogenic factors in mastocytosis, additional mutations or polymorphisms are needed to determine the final disease phenotype.

Anaphylaxis

In addition to a role for KIT mutations in the development of mastocytosis, it has been proposed that there is a link

between aberrant KIT activation and anaphylaxis, although this conclusion is somewhat more controversial. Certainly, based on *in vitro* studies, it would be logical to assume that an activating mutation in KIT would result in exaggerated antigen-mediated mast cell activation. Surprisingly, one study using mismatch amplification real-time PCR assay found a relatively high occurrence of the D816V mutation (2 out of 9, 22%) in subjects without a history of atopy or anaphylaxis (Lawley *et al.*, 2005). The same study identified the mutation in 9 out of 21 (43%) patients with anaphylaxis, although this apparently increased detection rate in anaphylaxis was not statistically significant (Lawley *et al.*, 2005).

Mice receiving chronic treatment of SCF do not have an increase in IgE-dependent anaphylaxis (Ando *et al.*, 1993). In contrast, there is evidence suggesting that the SCF-KIT axis is important in the development of non-IgE-dependent anaphylaxis. Anaphylactic-type dermal mast cell degranulation was observed in patients with advanced breast cancer who received subcutaneous injections of SCF in a Phase 1 clinical trial (Costa *et al.*, 1996). More recently, the D816V c-KIT mutation and other markers of clonal mast cell disease such as aberrant surface expression of CD25 by mast cells have been reported in a subgroup of patients with recurrent idiopathic anaphylaxis (Akin *et al.*, 2007). Some of these patients had a mild increase in mast cell numbers in bone marrow, which did not meet the diagnostic criteria for systemic mastocytosis. Such patients who experience anaphylaxis and carry a population of clonal mast cells without meeting the diagnostic criteria for systemic mastocytosis have been termed to have a monoclonal mast cell activation syndrome (Florian *et al.*, 2005; Akin *et al.*, 2007).

The multiple roles that SCF and KIT may play in dysregulated mast cell homeostasis and activation therefore provide a basis for considering inhibitors of KIT activity and function in the treatment of a number of mast cell related disorders including mastocytosis, atopic asthma, and anaphylaxis.

Pharmacological targeting of KIT

It is clear from mutational analysis and from studies conducted in knock out mice that KIT-induced phosphorylation of the tyrosines contained within the cytosolic tail and the subsequent recruitment of signalling molecules are essential events for the biological function of KIT (Broudy, 1997; Linnekin, 1999; Roskoski, 2005a, b; Akin *et al.*, 2007). Thus, pharmacological targeting of these processes, especially the KIT catalytic activity, has been a major strategy for blocking KIT-mediated responses. In the following sections, we will discuss the pharmacology of KIT inhibitors and how the disease states discussed above may be suitable target for potential targeting with KIT inhibitors.

The various tyrosine kinase inhibitors that have been described to inhibit KIT activity are listed in Table 1. The most widely recognized compound that blocks KIT catalytic activity is imatinib mesylate (imatinib) (also known as STI571, Gleevec and Glivec). Imatinib targets KIT at the ATP-binding site, thereby maintaining the receptor in a non-activated state. It is relatively selective as, in addition to KIT,

Table 1 KIT inhibitors and their targets

<i>Inhibitor</i>	<i>Additional names</i>	<i>Kit target</i>	<i>Additional targets</i>	<i>Reference</i>
Imatinib	Gleevec Glivec STI571	Wild type, V560G	Bcr-Abl, PDGFR	(Jensen <i>et al.</i> , 2007) (Levitzki <i>et al.</i> , 2006)
Nilotinib	AMN107	Wild type, V560G	Bcr-Abl, PDGFR	(Ma <i>et al.</i> , 2002) (Chow <i>et al.</i> , 2007)
PD180970		Wild type, V560G	Bcr-Abl, Src	(Gleixner <i>et al.</i> , 2006) (Corbin <i>et al.</i> , 2004)
Dasatinib	BMS-354825	Wild type, V560G, D816V	Src kinases, Tec, Btk	(Roskoski, 2005a, b) (Shah <i>et al.</i> , 2006) (Gleixner <i>et al.</i> , 2007) (Hantschel <i>et al.</i> , 2007)
Midostaurin	PKC412 N-benzoyl-staurosporine	Wild type, V560G, D816V	PKC, FLT3, VEGFR2, PDGFR, FGFR	(Patnaik <i>et al.</i> , 2007) (Fabbro <i>et al.</i> , 2000) (Gleixner <i>et al.</i> , 2006)
Hypothemycin		Wild type, D816V	^a	(Schirmer <i>et al.</i> , 2006)
EXEL-0862		Wild type, D816V	STAT3	(Pan <i>et al.</i> , 2007)
MLN518		Wild type, D816V	STAT3	(Corbin <i>et al.</i> , 2004)
AP23646/AP23848		Wild type, D816V	STAT3, Akt, ERK	(Patnaik <i>et al.</i> , 2007)
Semaxinib	SU5416	Wild type, D816V	STAT3, Akt, ERK	(Kosmider <i>et al.</i> , 2007)
Sunitinib	SU11248	Wild type, V559D, V645A, V559D/T670I, V670I	VEGFR, PDGFR, FLT3	(Chow <i>et al.</i> , 2007) (Prenen <i>et al.</i> , 2006)
Sorafenib	BAY 43-9006, Nexavar	Wild type	VEGFR 2,3, PDGFR, FLT3, Raf, MEK, ERK	(Liu <i>et al.</i> , 2006)
Pazapanib	GW786034	Wild type	VEGFR 1,3, PDGFRa,b	(Sonpavde and Hutson, 2007)
17-AAG		Wild type	HSP90, Akt, STAT3	(Ramanathan <i>et al.</i> , 2005)
MD-0354		V560G, D816V	NFkB	(Fumo <i>et al.</i> , 2004) (Tanaka <i>et al.</i> , 2005)

^aPKs with a conserved cysteine in the ATP-binding site.

it has been reported to inhibit only Bcr-Abl and the PDGFR. This may explain why imatinib induces relatively few side effects and is well tolerated (Levitzki and Mishani, 2006). Imatinib targets not only wild-type KIT but also KIT carrying the V560G mutation (Heinrich *et al.*, 2000). However, KIT carrying the D816V mutation associated with systemic mastocytosis is resistant to imatinib inhibition, due to the mutation changing the ATP binding site configuration, thereby blocking the binding of imatinib to KIT (Scheinfeld, 2006). Thus, although imatinib can prevent the growth of human mast cells that express wild-type KIT, the dysregulated growth of tumour mast cells linked to the D816V mutation is resistant to imatinib treatment (Zermati *et al.*, 2003). A similar pharmacological profile has been reported for the imatinib mimetics, nilotinib (AMN107) and PD180970, which can inhibit both wild-type KIT and KIT carrying the V560G mutation, but not KIT containing the D816V mutation (Corbin *et al.*, 2004; Verstovsek *et al.*, 2006a; Chow *et al.*, 2007). Nilotinib, in addition to targeting, KIT, Bcr-Abl and the PDGFR, has also been described to be cytotoxic to B cells, due to caspase activation, independently of kinase inhibition (Gleixner *et al.*, 2006). Besides KIT, PD180970 has been described to inhibit only Bcr-Abl and Src (Dorsey *et al.*, 2000). There has therefore been a focus on the development of KIT kinase inhibitors that overcome the drug-resistance associated with the D816V mutation.

Recently, several compounds have been identified that inhibit the catalytic activity associated with KIT carrying the D816V mutation. These include dasatinib (BMS-354825), midostaurin (PKC412, N-benzoyl-staurosporine), hypothemycin, EXEL-0862, MLN518, AP23646/AP23848 and

semaxinib (SU5416). These compounds are all multikinase inhibitors and therefore less specific than imatinib, nilotinib and PD18070. Dasatinib inhibits the growth of both human mast cell line (HMC)-1.1 and HMC-1.2 human mast cell lines, which express the V560G mutation or the V560G and V816D mutations, respectively. The growth inhibitory effect correlates with the compound's inhibitory effect on KIT autophosphorylation found in both of these cell lines (Shah *et al.*, 2006; Gleixner *et al.*, 2007). Dasatinib has also been reported to cooperate with midostaurin, nilotinib, imatinib and 2CdA (2-chloro-deoxy-adenosine) to produce an enhanced inhibition of growth, and the induction of apoptosis, of neoplastic mast cells (Gleixner *et al.*, 2007). In addition to KIT inhibition, however, dasatinib is also described to target Src kinases and the related Tec kinases Bruton's tyrosine kinase (Btk) and Tec. These properties would explain the reported ability of dasatinib to block histamine release from mast cells and basophils (Hantschel *et al.*, 2007): a process dependent on the activation of both Src kinases and Btk (Gilfillan and Tkaczyk, 2006). Midostaurin has a broad antiproliferative activity against various normal and tumour cell lines, including HMC-1.1 and HMC-1.2 cells, where it is also associated with induction of apoptosis (Fabbro *et al.*, 2000; Gleixner *et al.*, 2006). The many different cell types affected by midostaurin may be due to its several targets besides KIT, as this protein kinase inhibitor also targets PKC, FLT3, vascular endothelial growth factor receptor (VEGFR)-2, PDGFR and FGFR (Fabbro *et al.*, 2000).

In addition to inhibiting wild-type and D816V KIT, hypothemycin has also been described to inhibit other protein kinases with a conserved cysteine in the ATP-binding

site. These include several of the mitogen-activated protein kinases (Schirmer *et al.*, 2006). Studies conducted in mast cells also suggest that hypothemycin can inhibit an upstream regulator of Btk (Jensen *et al.*, 2008); however, hypothemycin does not appear to inhibit Btk directly as assessed in an *in vitro* kinase assay (Schirmer *et al.*, 2006). The ability of hypothemycin to inhibit the activation of Btk in mast cells would again account for the ability of the compound to inhibit antigen-induced and KIT-enhanced mast cell degranulation and cytokine production (Jensen *et al.*, 2008). EXEL-0862, MLN518 AP23646/AP23848 and semaxinib have been reported to inhibit STAT3 phosphorylation in addition to their effects on wild-type and D816V KIT (Corbin *et al.*, 2004; Kosmider *et al.*, 2007; Pan *et al.*, 2007; Patnaik *et al.*, 2007). Furthermore, AP23646/AP23848 and semaxinib have also been shown to affect Akt and ERK activities (Kosmider *et al.*, 2007; Patnaik *et al.*, 2007).

Certain VEGFR inhibitors have been found to also target KIT; however, their ability to inhibit KIT containing the D816V mutation is unreported. Sunitinib (SU11248), for example, inhibits VEGFR, PDGFR, KIT, FLT3 and has been reported to inhibit KIT containing V559D and V559D/T670I mutations in systemic mastocytosis and imatinib-resistant KIT mutations in gastrointestinal stromal tumours (V645A and T670I) (Prenen *et al.*, 2006; Chow and Eckhardt, 2007; Patnaik *et al.*, 2007). Sorafenib (BAY 43-9006, Nexavar), which targets VEGFR2 + 3, PDGFR, FLT3 and Raf (Liu *et al.*, 2006), inhibits MAPK/ERK kinase and ERK phosphorylation and downregulates the anti-apoptotic protein Mcl-1 in an MAPK/ERK kinase/ERK independent way and also has been demonstrated to inhibit wild-type KIT. Furthermore, pazopanib (GW786034), which targets VEGFR1, 2 and 3 and PDGFR- α and PDGFR- β , also inhibits wild-type KIT (Sonpavde and Hutson, 2007).

A number of compounds, not primarily considered to be tyrosine kinase inhibitors, have also been reported to inhibit KIT activity or KIT-mediated responses in mast cells. These include the following: (1) The anti-tumour antibiotic 17-allylamino-17-demethoxygeldanamycin, which, in addition to targeting the cytosolic heat shock protein 90 (Ramanathan *et al.*, 2005), was also reported to decrease the phosphorylation of KIT, Akt and STAT3 in both HMC-1 subclones (Fumo *et al.*, 2004); (2) IMD-0354, an nuclear factor- κ B inhibitor that has been shown to suppress proliferation of the subtype HMC-1.2 cells but not in cord blood-derived human mast cells expressing wild-type KIT (Tanaka *et al.*, 2005). HMC-1.2 cells have constitutively activated nuclear factor- κ B, which mediates upregulation of cyclin D3, resulting in cell cycle progression. Targeting nuclear factor- κ B and inhibiting its activation pathway therefore results in suppression of cell proliferation; (3) Rapamycin targets the serine/threonine protein kinase mTOR, which is activated by KIT in a PI3K-dependent manner in mast cells (Kim *et al.*, 2008). The mTOR pathway is critical for optimal mast cell growth, and tumour mast cells expressing the V560G or D816V mutation have constitutively activated mTOR. Rapamycin has been reported to only induce apoptosis in cells carrying D816V mutation, but not in V560G-mutated cells or wild-type mast cells (Gabillot-Carre *et al.*, 2006). However, in a later study,

rapamycin was reported to inhibit mast cell growth/survival, irrespective of the presence of a specific mutation (Kim *et al.*, 2008). These results are somewhat in contrast with imatinib, which only affects wild type and mast cells expressing V569G KIT.

The potential use of KIT inhibitors in the treatment of systemic mastocytosis

Cytoreductive therapy of systemic mastocytosis is generally reserved for patients with advanced forms of mastocytosis associated with decreased life expectancy (Valent *et al.*, 2003a). This is because of the concerns about potential mutagenicity, teratogenicity and lack of knowledge about the long-term toxicity of tyrosine kinase inhibitors. Inhibition of KIT as a potential cytoreductive approach in mast cell disease is based on several lines of evidence, perhaps the strongest of which is the high (greater than 90%) detection rate of the presence of the activating D816V KIT mutation in mastocytosis. Transfection of this mutation in haematopoietic cells renders growth factor independence (Kitayama *et al.*, 1995, 1996), thereby confirming the role of the mutation in neoplastic transformation. Moreover, there is a strong correlation between the ability of a given compound to inhibit KIT autophosphorylation and its cytotoxic effects on mast cell lines. *Ex vivo* experiments performed with mononuclear cells obtained from the bone marrow of patients with systemic mastocytosis showed that tyrosine kinase inhibitors effective against D816V KIT cause preferential cytotoxicity of mast cells carrying this mutation over other mononuclear cells (Akin *et al.*, 2003; Shah *et al.*, 2006). On the other hand, it is not known whether neoplastic mast cells have other survival pathways in addition to KIT. This is especially plausible in patients with advanced forms of mastocytosis such as aggressive systemic mastocytosis. Moreover, the concentrations required to inhibit mast cell growth may be difficult to achieve *in vivo* without toxicity to other tissues. This problem may be particularly pertinent for non-specific inhibitors with multiple targets.

The proof of concept that a KIT inhibitor can indeed cause regression of mastocytosis driven by a KIT mutation resulted from a study in which a patient with an unusual activating KIT mutation was treated with imatinib, and achieved complete remission (Akin *et al.*, 2004). This patient had a transmembrane mutation involving codon 522 (F522C), which was sensitive to imatinib. Treatment of the patient with imatinib doses up to 400 mg day⁻¹ resulted in a dramatic decrease in serum levels of the mast cell marker, tryptase, and mast cell burden. Imatinib is also effective in the treatment of rare cases of systemic mastocytosis associated with chronic eosinophilic leukaemia, which is characterized by the presence of the FIP1L1-PDGFR α rearrangement (Pardanani *et al.*, 2003a, b). This rearrangement occurs due to an approximately 800 kb interstitial deletion in chromosome 4 (detected by deletion of CHIC2 locus in 4q12 by immunofluorescence *in situ* hybridization or by reverse transcription-PCR), leading to constitutive activation of the intrinsic tyrosine kinase activity of PDGFR α . These patients, who have a multilineage

myeloproliferative disorder that involves mast cell as well as eosinophil progenitors, are generally male, display organ pathology due to eosinophilia and do not have the characteristic D816V KIT mutation observed in other categories of mastocytosis (Klion *et al.*, 2003, 2004; Pardanani *et al.*, 2003c; Maric *et al.*, 2007). In addition, a patient with a rare variant of mastocytosis associated with chronic basophilic leukaemia and a PDGFRB fusion has been shown to respond to imatinib (Lahortiga *et al.*, 2008). The therapeutic effect of imatinib in these disorders is due to its inhibition of PDGFR and not KIT.

Imatinib is currently approved by the US Food and Drug Administration for the indication of treatment of adult patients with aggressive systemic mastocytosis without the D816V c-Kit mutation or with c-Kit mutational status unknown. The bulk of the clinical evidence forming the basis of this recommendation appears to have come from patients with chronic eosinophilic leukaemia associated with systemic mastocytosis and FIP1L1–PDGFRA fusion gene as discussed above. However, as opposed to its beneficial effects in rare variants of systemic mastocytosis without codon 816 KIT mutations, or those with PDGFR rearrangements, imatinib is not effective in inhibiting codon 816 KIT mutations carried by the great majority of patients with systemic mastocytosis (Ma *et al.*, 2002; Akin *et al.*, 2003; Zermati *et al.*, 2003). As discussed earlier, this is thought to be due to the inability of the drug to bind the enzymatic pocket of KIT, whose structure is altered by the point mutation in codon 816. In apparent contrast to these findings, a recent study on 14 patients with mastocytosis reported a modest beneficial effect of imatinib on patients with D816V KIT who received the drug, as measured by decreased mast cell mediator levels, bone marrow mast cell numbers and symptomatic improvement (Droogendijk *et al.*, 2006). However, imatinib in this study was administered with glucocorticoids, complicating the interpretation of the response to imatinib. Interestingly, imatinib has been shown to decrease spontaneous histamine release in HMC-1.2 human leukaemic mast cells carrying the D816V mutation (Lober *et al.*, 2008), although it is not cytotoxic to this cell line, which may partially explain the symptomatic improvement observed in the study. However, because imatinib preferentially inhibits wild-type KIT, long-term use of this drug can potentially confer a growth advantage to the neoplastic clones with D816V KIT and therefore theoretically worsen the disease course. On the basis of *in vitro* data mentioned above as well as our personal experience (Cem Akin), our current practice is not to treat patients with codon 816 KIT mutations with imatinib.

Clinical trials with other KIT inhibitors have largely yielded disappointing results. In a phase-2 trial of nilotinib, among 60 patients (83% positive for D816V KIT) treated with 400 mg twice daily dose of nilotinib, only 2 showed complete remission (Hochhaus *et al.*, 2006). The overall response rate was 20% and two patients died due to disease progression. Nausea and headaches were the most frequently reported side effects. These results are consistent with the lack of *in vitro* activity of nilotinib on D816V KIT (Verstovsek *et al.*, 2006a). Dasatinib has been evaluated for its therapeutic effect on mastocytosis in a recent clinical trial. Thirty

patients with systemic mastocytosis received dasatinib at a dose of 70 mg twice daily (Verstovsek *et al.*, 2006b). There were two complete remissions, both observed in patients without D816V KIT mutation and low tryptase levels. There was a high incidence of systemic toxicity: the drug was stopped in 10 patients and dose reductions were necessary in 12. Six patients developed pleural effusions. These results may be due to the non-specific targeting of KIT by dasatinib, resulting in toxicity in concentrations required to inhibit D816V KIT.

An ongoing clinical trial with midostaurine (100 mg twice daily) has yielded more promising results in an interim analysis (Gotlib *et al.*, 2007). Eleven of 15 (73%) patients with advanced systemic mastocytosis have shown a response. Although there were no complete remissions, 5 patients displayed a major response and 6 had a partial response. Nausea and vomiting were the most frequently observed non-haematologic side effects. Midostaurine has also shown a partial, although temporary, mast cell cyto-reductive effect in a patient with mast cell leukaemia with associated myelodysplasia (Gotlib *et al.*, 2005).

Combination therapy of mastocytosis with regimens incorporating tyrosine kinase inhibitors is yet to be explored in clinical trials, although *in vitro* data obtained so far appear promising. Combination of dasatinib with midostaurine or cladribine yielded synergistic effects in HMC-1.2 mast cells carrying the D816V c-KIT mutation (Gleixner *et al.*, 2007). Another study employing antisense mcl-1 oligonucleotides in combination with midostaurine similarly showed a synergistic growth inhibition in the same cell line (Aichberger *et al.*, 2007).

Potential use of KIT inhibitors in other disorders

As discussed earlier, mast cells play a central role in the allergic reactions following antigen-dependent aggregation of IgE-occupied Fc ϵ RI, a response that, at least under experimental conditions in both mouse and human mast cells, can be strongly potentiated by SCF-induced KIT activation. This can occur even at antigen concentrations that produce minimal degranulation (Tkaczyk *et al.*, 2004). Thus, in addition to the primary targeting of the IgE-response as a therapeutic strategy for the treatment of allergic reactions, as discussed in a recent review article, targeting SCF and/or KIT in addition to Fc ϵ RI may have therapeutic relevance in the treatment of these disorders (Jensen *et al.*, 2007). However, use of KIT inhibitors in non-neoplastic conditions should be contemplated in clinical trials only after a careful consideration of risk-to-benefit ratio of the compounds to be investigated, including demonstration that such compounds do not increase the risk for developing secondary malignancies.

The concept of concurrent inhibition has been explored using hypothemycin, as this molecule in both human and mouse mast cells targets Fc ϵ RI signalling in addition to KIT (Jensen *et al.*, 2008). In addition to its ability to block Fc ϵ RI-mediated and KIT-enhanced degranulation and cytokine production in mast cells in culture, hypothemycin significantly reduced *in vivo* passive cutaneous anaphylaxis

in mice. These results demonstrate the potential of a combined KIT/Fc ϵ RI inhibition in the treatment of allergic diseases.

Mast cells have been implicated in the pathogenesis of rheumatoid arthritis (Malone *et al.*, 1986, 1987; Malone and Metcalfe, 1988), and a recent study (Juurikivi *et al.*, 2005) revealed that imatinib effectively induces mast cell apoptosis and reduces TNF- α production in human synovial tissue cultures. This could lead to attenuation of the inflammation in arthritic joints (Juurikivi *et al.*, 2005). Other diseases where mast cells have been proposed to play a role may also benefit from targeting KIT in a manner similar to that described above, although this has yet to be investigated. Such disorders include migraine headaches, inflammatory bowel disease, multiple sclerosis, atherosclerosis and angiogenesis associated with tumour progression. With regard to migraine headaches, degranulation of rat mast cells from the dura-mater induced by compound 48/80 was shown to result in excitation of meningeal nociceptors, which was accompanied by phosphorylation of the intracellular proteins ERK and c-fos, resulting in a long-lasting activation leading to the migraine headache (Levy *et al.*, 2007). Mast cell mediators have also been shown to be major regulators of the severity of inflammatory bowel disease. In this respect, histamine can induce the smooth muscle and endothelial cell contractions, increased vascular permeability, cytokine production and chemotaxis of inflammatory cells associated with this condition (Fogel *et al.*, 2005). In addition, patients with Crohn's disease have also been found to have an increased number of mast cells in the submucosal layer and muscularis propria of the gut (Oshitani *et al.*, 2006). Therefore, eliminating histamine release by silencing mast cells present in the gastrointestinal tract, in a KIT/SCF dependant way, could contribute to a better prognosis of these conditions.

A role for mast cells in multiple sclerosis has been suggested by their ability to be activated by the main multiple sclerosis protein, myelin basic protein, which results in the release of IL-8 and TNF- α . Furthermore, mast cells appear to markedly enhance T-cell activation by cell-to-cell contact and by TNF- α (Theoharides *et al.*, 2007). The ability of imatinib and hypothemycin to block antigen-mediated IL-8 and TNF- α release from human and mouse mast cells and to block the SCF enhancement of this response (Jensen *et al.*, 2008) may provide the basis for considering KIT inhibitors in the treatment of this disease. A role of mast cells in the progression of atherosclerosis may be due to release of the inflammatory cytokines IL-6 and interferon- γ following mast cell activation as demonstrated in the mouse (Sun *et al.*, 2007). Again, as imatinib and hypothemycin has been shown to inhibit IL-6 release from murine bone marrow-derived mast cells (Jensen *et al.*, 2008), these drugs may be able to dampen the role of mast cells in atherosclerosis. Finally, human tumours are often surrounded by mast cells, and this infiltration is associated with a poor prognosis (Theoharides and Conti, 2004). Using a oncogene mouse model carrying a mutation in the gene coding for RAS, it was shown that not only did these mice have more mast cells in dermis before development of skin tumours, compared with controls, but a decrease in KIT function inhibited the progress of carcinogenesis in the skin

(Muto *et al.*, 2007). This indicates that, even though an increase in mast cell number exist, if the KIT response is inhibited, a better prognosis might be obtained.

Conclusions

In summary, in this review, we have discussed how specific mast cell disorders are linked to dysregulated control of the kinase activity associated with KIT and how tyrosine kinase inhibitors targeting KIT show promise in the therapeutic management of such disease states. Future advances in the development of more selective tyrosine kinase inhibitors that would target both normal and mutated KIT may thus offer hope in the effective treatment of mastocytosis and potentially anaphylaxis, and other allergic responses. Furthermore, the potential role of mast cells in other clinical conditions may provide a basis for the investigation of the efficacy of KIT inhibitors in these disorders.

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Conflict of interest

The authors state no conflict of interest.

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