

A novel therapeutic strategy targeting multiple members of the γ c-family cytokines; principles, relevance and potentially broad clinical applications.

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Abstract

Cytokines play diverse roles in normal and abnormal immunity. Dysregulated production of cytokines underlies a variety of disorders in humans by causing improper immune responses. Intervention to such abnormal cytokine action has been utilized in the clinical field as "anti-cytokine therapy" and proven effective as treatments. However the current anti-cytokine approaches lack effective and safe options for treating human diseases involving more than two cytokines as the pathogenic reason. This is an issue as the list of multi-cytokine diseases is expanding. To address this, a novel technology was developed by generating a novel class of multi-cytokine inhibitors (MCI). In short, cytokine-mimetic peptides were rationally designed each of which contains a motif shared only by the cytokines of the target and tested using specific biological assay. Peptides showing desired antagonistic activity were screened further for target specificity. The lead MCI, BNZ 132-1, specifically inhibits IL-2, -9 and -15 while it does not affect other γ c-cytokines (IL-4, -7, and -21) or non- γ c cytokines. This design ensures safety of the peptide upon clinical use by limiting off-target effects to the minimum, unlike small-molecule Jak kinase inhibitors which suppress a diverse array of cytokines and cause multiple adverse effects. Currently, we are conducting clinical trials involving BNZ 132-1 and saw expected transient decrease of select subsets of lymphocytes with minimum toxicity. BNZ 132-1 may provide a novel opportunity for treating many diseases such as myelopathy caused by human T-cell leukemia virus-1 (HTLV-1), non-viral human T-cell malignancies, autoimmunity (Alopecia areata, rheumatoid arthritis), graft-versus-host disease (GvH) upon transplant and cytokine release syndromes post microbial infections.

Introduction.

In the following sections, we will:

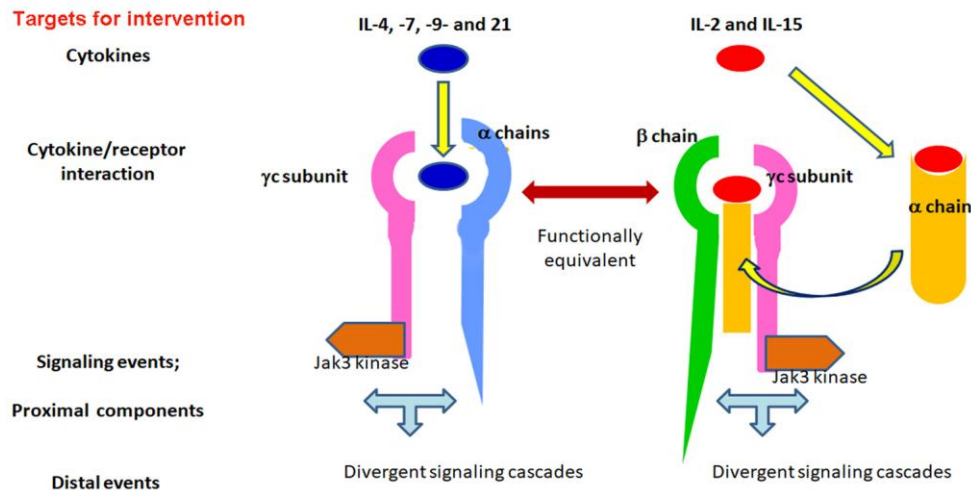
1. describe the basic nature of the γ c-cytokines and their receptor which are primary therapeutic targets for us.
2. discuss about an emerging concept of "multi-cytokine diseases" in which functionally redundant cytokines from a single family are pathogenically involved.
3. overview the currently available anti-cytokine strategy in the context of treating multi-cytokine diseases, which involves the special discussion on the limitations of current anti-cytokine strategy.

4. present our new BNZ technology and elaborate on the underlying concept, in vitro proof-of-concept data, its potential applications, preclinical data and on the on-going clinical trial involving the BNZ 132-1 peptide, the lead compound from our new technology.

1. The γ c-family cytokines.

Since these are the intervention targets of our interest, we will quickly review the intriguing characteristics of this family of cytokines, as the function and receptor configuration of these cytokines directly connect with the objective, challenges and the uniqueness of our endeavor.

Figure 1. Anatomy of the γ c-receptor system for the rational design of therapeutic interventions



The composition of multi-subunit receptors for γ c-cytokines is depicted. In this family, IL-2 and IL-15 form a subfamily of their own and the receptor configuration (three subunits instead of two) illustrates the uniqueness of the subfamily. The α chains for IL-2 and IL-15 are not a signaling subunit, but mediate the unique trans-presentation for IL-15 (66). In the cascade of ligand to intracellular signaling components, the γ c and Jak3 are the point of conversion and targeting these molecules is the logical choice to block multiple γ c-cytokines by a single agent.

Cytokines are small polypeptides (Molecular weight ranging from 5 to 40kDa) that induce activation and differentiation of various hematopoietic cells (1, 2). Most, if not all, cytokines assume a distinct topology called "the four-helix bundles" structure (1) and the γ c-family cytokines, namely IL-2, -4, -7, -9, -15 and -21, belong to this group. All γ c-cytokines use the common gamma (γ c, CD132) as a subunit in their multi-component receptor systems. Since γ c is a signal transducing molecule, these cytokines naturally possess overlapping signaling pathways and display functional redundancy. The physiologic roles of each cytokine have been elucidated in detail using gene-targeting (Knock-out or KO) and transgenesis technologies in mice (3-19). Basically all 6 γ c-cytokines are required for preparing various types of immune cells and participate in dictating the proper direction of the immune response upon microbial invasion and from other causes, as their biological relevance is well demonstrated by the phenotype of mice defective of the shared γ c-receptor which have no T, B, NK cells and manifest severe immunodeficiency (20, 21). We have a long history of studying the mechanisms, receptors, and biological functions of γ c-cytokines because of our interest in the activation and development of T and NK cells.

2. Why blocking multiple-cytokines is a unique and worthy scientific goal?

2-a. Human diseases caused by more than two cytokines (multi-cytokine diseases); Though cytokines belonging to the same family show functional redundancy, each cytokine has diverse unique functions (called the pleiotropic nature). Uncontrolled production or the lack of a cytokine leads to pathogenic conditions as demonstrated by the phenotypes of cytokine knock-out or transgenic mice. In humans, there are many examples of diseases caused by the

abnormal functioning of cytokine (cytokine diseases). Anti-cytokine therapy has been long developed to treat cytokine diseases. However, we conducted a literature search and noted that quite a few cytokine diseases involve more than two cytokines (and very likely those belonging to the same family) (22) and that the current anti-cytokine strategy has limitations in dealing with cytokine diseases involving multiple cytokines (22), which prompted us to design and develop a unique group of cytokine inhibitors which can block more than two cytokines for treating such multi-cytokine diseases. One example of multi-cytokine diseases is a myelopathy associated with HTLV-1 (human T-cell leukemia virus 1) called HAM/TSP (HTLV-1 associated myelopathy/tropical spastic paraparesis) (23, 24). HAM/TSP is a progressive and chronic disease of the spinal cord seen in 0.3~0.5% of HTLV-1 infected individuals, manifesting painful stiffness and weakness of the legs. In HAM/TSP patients, T lymphocytes which cause pathogenesis cross the blood-brain barrier and damage the spinal cord by the production of proinflammatory factors and by the direct cytotoxic action. As we showed in our publications, T cells are constitutively activated in vivo in these patients and spontaneously proliferate when they were transferred to ex vivo culture without the addition of mitogens or cytokines (25, 26). We later discovered that two γ c-cytokines, namely IL-2 and IL-15 are transcriptionally activated by the HTLV-1, and cause the autocrine proliferation and activation of CD8 T cells, a process likely leading to the tissue damage in HAM/TSP. We also demonstrated that both IL-2 and IL-15 need to be blocked in order to subside the pathogenic activation of T cells from HAM/TSP patients (22, 27-30). Examples of multi-cytokine disease are increasing in number. There are more examples of multi-cytokine diseases. To name a few,

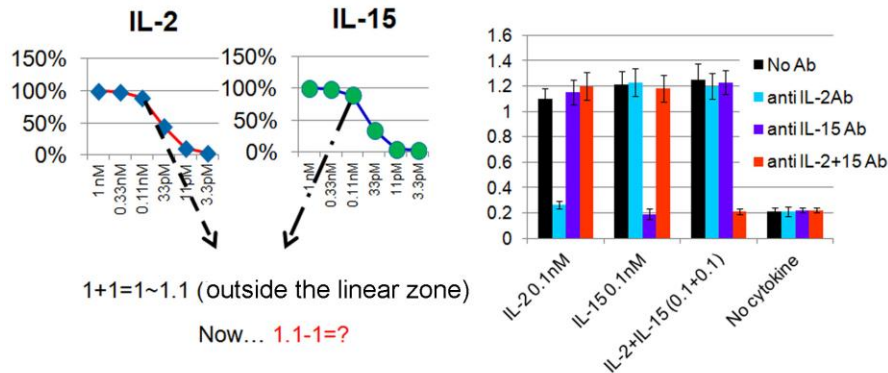
rheumatoid arthritis which may involve IL-15, -21, -6, and -7 as pathogenic factors (31-33) and Celiac Disease which may involve IL-2, -10, -15 and -21 (34-41). Co-production of cytokines belonging to a family may not be too rare. It is reasonable to assume that cytokines belonging to a family evolved through gene duplication (of the cytokine and of the receptor subunits) and thus the regulatory mechanisms still share common elements. We use IL-2 and IL-15 as an example to look into this. IL-15 and IL-2 form a subfamily in the γ c-cytokine family because they share another signaling subunit, IL-2R/IL-15R β (CD122) in addition to the γ c (Figure 1). Despite the functional similarities between IL-2 and IL-15, their production seems strictly separated from each other under normal circumstances. IL-2 is almost exclusively produced by T cells as was demonstrated in an IL-2/GFP knock-in mouse (42) whereas IL-15 is produced by activated monocytes, dendritic cells, and stromal cells (43, 44). Yet, their promoter regions share a few motifs including the one for NF κ B (45), accounting for the fact that Toll-like receptor stimulation in monocytes lead to the production of IL-15 by monocytes and T-cell activation results in production of IL-2 by T cells through the activation of the NF κ B pathway. Moreover, the co-production of IL-2 and -15 in HAM/TSP seems to involve the transcriptional activation of these two genes by the trans-activator Tax of HTLV-1 (45) through the activation of the NF κ B pathway. Therefore, it may not be surprising that IL-2 and IL-15 are simultaneously produced from different cells through an overlapping molecular mechanism, in particular under pathological conditions. This would apply to the co-production of multiple cytokines belonging to the same family.

2-b. A pitfall - neutralization of one of the factors in a multi-cytokine disease may not show visible effects even though the factor is pathogenically involved; After identifying that a human disease belongs to the category of cytokine disease, the logical strategy is to follow a reductionist approach; To gather information to identify the suspect cytokine and to test if neutralization of the target cytokine would effectively block the pathogenic signs associated with the disease. Even if the involvement of multiple factors was anticipated, the intuitive assumption is that the blockade of "a major factor" should have visible impacts. However this intuition may be misleading. We have shown this in our recent publication (22) but will discuss it here as it is a critical point of our argument; If two functionally redundant cytokines (IL-2 and IL-15 on T or NK cells is a perfect example) are stimulating the target cells as a 1:1 mixture, how much of the biological activity will be neutralized when only neutralizing anti-IL-2 or anti-IL-15 antibody (Ab) was added? The intuitive answer would be "a 50% inhibition". However, this is not true at some circumstances, because the biological response of a cell is not always linear to the dose of cytokine. The dose-response relationship follows a sigmoid-curve. At the near-saturating dose, additional increase of the stimulant no longer guarantees visible increase of the response. In reversal, anti-IL-2 antibody added to a mixture of 0.5 nM each of IL-2 and IL-15 to PHA-activated human T lymphocytes (which responds to both cytokines at high affinity; ~50 pM) would bring down the 100% response of the cells only to 90~95% (Figure 2). This was experimentally verified in our previous publication (22). The important take-home-message is that the "lack of a visible inhibition" using an antibody to one cytokine when more than two cytokines are cooperating to cause cellular activation does

not allow to eliminate the target cytokine from the equation. In other words, **no visible inhibition does not mean no participation of the target cytokine**. This may become more critical in vivo because it is difficult to know how many cytokines are in operation and how much is the contribution by each involved cytokine. For example, if cytokine A was highly suspected in the cause of a disease but antibody therapy targeting cytokine A failed. There might have been an example of this scenario in the actual clinical trials involving Asthma. Asthma has been connected to multiple cytokines

including IL-4, IL-13, IL-9, and IL-5 in previous studies (16, 46-54). Accordingly, clinical trials involving anti-IL-4 and anti-IL-9 therapy have been approved but separately conducted (47, 48, 55-57). Unfortunately, neither gave rise to a satisfactory therapeutic result. This may simply mean that other cytokines are equally participating in the cause of the disease thus a co-inhibition of both factors could be a "simple" solution. However, the implementation of this strategy may not be as simple as it may look like.

Figure 2 Co-inhibition is the only way to effectively block cytokine-cooperation



On the left shown is a typical proliferative response of mitogen-activated T cells to various doses of IL-2 or IL-15. They respond to low-concentrations of cytokines because they express high-affinity (K_d ; ~ 50 pM) IL-2 and IL-15 receptors. Importantly, when a cytokine doses pass over half-saturation concentration (which is equal to the K_d value), cellular responses are no longer linear to the increase of the dose of the cytokine. When the saturating doses of IL-2 and IL-15 were combined, they no longer create a response over that attained by either cytokine ($1+1=1$ in this non-linear world). Inversion of the equation ($1-1=1$) illustrates neutralization efforts on only one of the two cytokines (leaving the other cytokine at a saturation dose) which leads to almost zero inhibition of the cellular response. The other side of the coin is that an antibody to a cytokine may fail in vivo to show any visible inhibition even though the target cytokine is relevantly involved in the activation of the target cell. This observation quite often misleads us into a conclusion that "the cytokine we studied was not involved in the pathogenic activation".

On the right shown is an actual experimental result that we conducted in support of our argument. Note that the same dose of anti-IL-2 (or anti-IL-15) mAb effectively neutralized the single target cytokine, but shows negligible effect when two cytokines are combined (the third group from left).

2-c. Technical limitations of the current anti-cytokine therapy; as demonstrated above, monotherapy may not only be rather ineffective but also the failure of it may mislead the researchers to a wrong conclusion when dealing with multiple-cytokine diseases. It is particularly true for cytokine diseases because of the unique nature of the cytokine that most of them belong to families and have multiple siblings/relatives sharing biological functions. So what about combining antibodies (the most preferred modality in anti-cytokine therapy) to neutralizing multiple cytokines? This is very simple with in vitro experiments, though we wish to point out that a random combination of antibodies may be a waste of resource, energy and time so that one has to build up a rational strategy before proceeding to the execution. However, a clinical application of combinatorial use of antibodies for therapeutic purposes seems infeasible, due to the expensive nature of the antibody therapy. The goal can still be attained by using other modalities than antibodies. We will discuss it below.

3. Rational approach to design multi-cytokine inhibitors

We will start from a general discussion on how to rationally design a cytokine intervention, and shift the emphasis to γ c-cytokines which are of our interest.

3-a. Targeting the ligand (cytokine). As shown above in Figure 1, there are multiple-levels of targets when intercepting cytokine function. Neutralizing antibodies are available for most cytokines. However, as discussed above, a monoclonal anti-cytokine antibody (mAb) is not suited for treating multi-cytokine diseases, unless the antibody is bispecific (58). However, generating a bispecific antibody is a

laborious work and is still in the early developmental stage for clinical applications.

3-b-1. Targeting a shared receptor component. The next level target is the cytokine receptor. More to the point, the aim is **targeting the shared receptor component**. Many cytokines share receptor components with other cytokines due to their evolution through gene duplication. This goal can be accomplished in general by using anti-cytokine receptor antibodies against the shared component. A soluble version of the shared receptor component can be used as well. Alternatively, a ligand analog that lacks agonistic function (antagonistic cytokine mimetics, (59)) can be a potent inhibitor. With the γ c-cytokines, however, no neutralizing anti- γ c antibody exists. This is perhaps because the binding surface of the γ c is highly structurally conserved across species so the production of antibodies recognizing this portion may be blocked by the immune tolerance mechanism. As mentioned above, IL-2 and IL-15 share another signaling receptor subunit, IL-2R/IL-15R β (CD122). Heterodimerization of γ c and β are needed for signal transduction by IL-2 or IL-15 (60) so inhibiting the β -cytokine interaction can block the signaling triggered by IL-2 or IL-15. There exist a few neutralizing anti- β antibodies including Mik β 1 (61). The humanized Mik β 1 has been tested in clinical trials involving T-cell large granular lymphocyte(LGL)-leukemia (62) and proposed for a diverse array of disorders including Celiac disease and other autoimmunity in which IL-2 and IL-15 might allegedly be pathogenic. Nevertheless, there is a setback for this antibody as well. However, we need to know a little more about the dual mode of IL-15 receptor configuration in order to understand this issue.

3-b-2. The *trans*- and *cis*-mode of IL-15 action. As mentioned above, the β/γ hetero-dimer is the functional core of the IL-2/IL-15 receptor, However, these two are not sufficient for the "in vivo" function of these cytokines because they only bind target cytokines at relatively low affinity (Kd; 0.7 nM) and thus unable to bind them at their physiological concentrations (10~100 pM). For the IL-2, a ligand-specific component IL-2R α (CD25) joins and converts the β/γ receptor complex to a high-affinity IL-2 receptor (Kd; 10 pM). The in vivo functional IL-2R assumes an IL-2R α -IL-2R/IL-15R β - γ c hetero-trimeric configuration. However, IL-15R is structurally different. Like the IL-2R α (CD25), there is an IL-15-specific binding partner called IL-15R α . However, this molecule behaves as a component of a hetero-dimeric single cytokine (like IL-12 and its siblings) which consists of a complex of IL-15 and IL-15R α (hence calling the latter molecule "a receptor" may not be entirely accurate though it is a binding protein to IL-15) (63-65). They can be formed as a membrane-bound form on activated dendritic cells (DCs) and monocytes and then be presented in *trans* to T or NK cells which only expresses the β - γ c molecules (the *trans*-presentation mechanism, originally shown by us and validated by others (63-83). Several lines of studies indicated that this is the major physiologic form of IL-15 in vivo. In other words, the β - γ c complex expressed on the surface of resting T or NK cells in the absence of IL-2R α /CD25, is a natural receptor for IL-15 which can be brought to T/NK cells by monocytes or dendritic cells expressing the IL-15/IL-15R α complexes on the surface. IL-15R α can be induced on the surface of T/NK cells and converts the β - γ c complex to respond to soluble IL-15 monomer (IL-15R in the *cis*-configuration), but we have observed that CD8 T cells constitutively expressing IL-15R α are more

prone to developing into leukemia (84), a plausible reason why the *trans*-configuration of the IL-15R might be preferred by nature. Now let us go back to the issue of cytokine-neutralizing anti-IL-2R/IL-15R β antibody. The setback is that the Mik β 1 mAb only effectively blocks the action of IL-2 or IL-15 when the private α is not present on the same cells that express the β/γ . This means IL-2 signaling in vivo cannot be efficiently blocked by this antibody (since the only physiological IL-2R consists of all 3 chains expressed by the same cell). Interestingly, the *trans*-action of IL-15 can be efficiently blocked by Mik β 1, but the *cis*-action is not (62). As we and others have indicated in the past, *trans*-IL-15 action is preferred upon normal immune responses, therefore Mik β 1 is effective in blocking such IL-15 actions in vivo. However, pathologic IL-15 action such as the one seen with CD8 T-cell leukemia seems to require the *cis*-IL-15R configuration by target cells (84) and the Mik β 1 mAb cannot effectively block this. Thus, Mik β 1 may have limitations for blocking IL-2, IL-15 or both under disease conditions.

3-b-3. Cytokine mimetics. The cytokine-mimetics are recent developments. These are fragments from a cytokine or mutated cytokines with altered functions. Small peptides containing the active site (one that is used for interacting with receptor subunit) of cytokines can mimic the cytokine action and trigger a signal (agnostic mimetics). Alternatively, they can block the active binding site on the cytokine receptor and inhibit the action of cytokines (antagonistic mimetics). For therapeutic purposes, antagonistic mimetics are perhaps more beneficial. Examples include IL-4 mimetics (85) which have been tested in clinical trials targeting Asthma with limited success (86, 87). Importantly, an IL-4 mutant (tyrosine at residue 124 mutated to

aspartic acid, Y124D) which has good binding affinity to IL-4R α with no signaling capability can block both IL-4 and IL-13 actions due to the sharing of IL-4R α between these two cytokines (88). In a way, this IL-4 mutant (hIL-4Y124D) is a prototype multi-cytokine inhibitor. Agonistic cytokine mimetics have been tested to substitute the cytokine, but they are not the major focus for this review. Examples include those for erythropoietin (89), thrombopoietin (90, 91), interferon (92), TGF- β (93), hepatocyte growth factor (94), and chemokines (95), which are being tested in clinical applications. Our BNZ peptides (BNZ-1~3) belong to antagonistic small-molecule γ c-cytokine mimetics.

3-b-4 Soluble receptors. There have been attempts to generate soluble cytokine receptors for therapeutic purposes (96). Because soluble cytokine receptors lack the signal transducing elements, but still bind to the cytokine with similar affinity as do membrane-bound counterparts, they can compete with the membrane-bound receptors for the cytokine binding and function as inhibitors. Related to the γ c-cytokine world, soluble IL-4R presents an intriguing example of multi-cytokine inhibition, as it blocks both IL-4 and IL-13 actions due to the IL-4R α sharing between IL-4 and IL-13 as mentioned above (88, 97). Though IL-13 is not a γ c-cytokine, IL-4 and IL-13 form a subfamily of their own, which has been implicated in Asthma. Thus, modalities (e.g., antibodies, mutant IL-4, and soluble IL-4R) neutralizing these cytokines have been tested in clinical trials with some success (47, 48, 98). However, only monotherapy targeting each cytokine has been tested so it is not yet clear if a co-inhibition of IL-4 and IL-13 has superior therapeutic effects to those attained by the single cytokine targeting.

3-c. Targeting Signaling molecules.

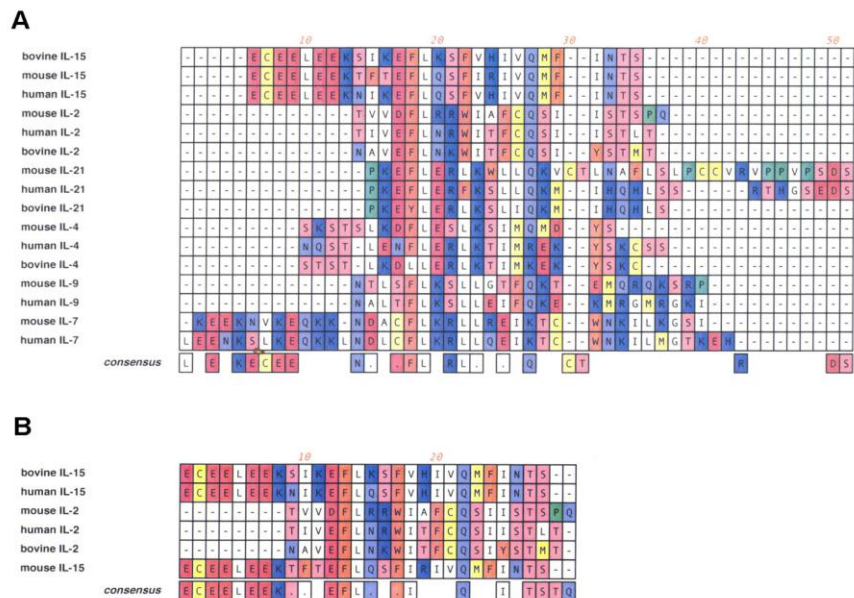
Another class of targets for intervening cytokine action includes signaling molecules. Unlike many hormone receptors (e.g., those for insulin, EGF, and Growth Hormones), cytokine receptor molecules generally lack an intrinsic kinase activity. Instead, members of the Janus kinase family (Jak1, Jak2, Jak3, and Tyk2, reviewed in (99)) specifically interact with cytokine receptors and help transducing the signal. The membrane proximal events are probably easier to target because this is where the entire cascade of cytokine signaling becomes convergent. In particular, the γ c-cytokine family presents a unique case because not only all γ c-cytokines use Jak3, but also the Jak3 usage seems exclusively limited to γ c-cytokines (100-107). This is underscored by the observation that Jak3 knockout mice (108) or humans with loss-of-function Jak3 mutations (105) show identical phenotypes to γ c-knockout mice (20, 109) or humans with defective γ c-molecule (X-linked SCID, (110)). A brilliant idea was proposed by John O'shea's group, based on their own discovery of Jak3 and of its exclusive use by γ c-cytokines, to target Jak3 for blocking actions by a single or multiple γ c-cytokines, and they developed a Jak3 inhibitor (CP690, 550 (111)) in collaboration with Pfizer for this purpose. The compound has been approved by the FDA as a clinical drug 'Tofacitinib' (29, 84, 112-120). Currently Tofacitinib is being tested in clinical trials for the treatment of rheumatoid arthritis (114), graft-versus-host diseases associated with organ transplantation (111, 115, 119), inflammatory bowel disease/celiac disease (117, 120) and T-cell leukemia/lymphoma (22, 29, 84). We will further evaluate the efficacy and shortcomings of Tofacitinib as a clinical drug below.

4. Can we selectively target multiple γ c-cytokines?

4-a. Safety concerns for the Jak3 inhibitor. So a question may arise as to why we needed to develop a new technology to block multiple γ c-cytokines? Why are we not satisfied with the Jak3 inhibitor(s)? Here is our answer: though a Jak3 inhibitor would comprehensively block γ c-cytokines, it does not allow us to selectively choose the target cytokine(s). What we want was to specifically choose target cytokines within the γ c-family and block only them, but not other cytokines. In addition, it became clear that the inhibition by Tofacitinib is not limited to Jak3 (121) because it inhibits the kinase activity of Jak2 and Jak1 at high concentrations (122). Since Jak2 is critically involved in erythropoiesis and thrombopoiesis, a continued use of Tofacitinib might induce anemia and bleeding tendency (118, 123-126). Other concerns also cropped up as represented by the action of the European Commission which initially declined to approve Tofacitinib citrate (Xeljanz) for treating rheumatoid arthritis due to potential carcinogenesis, mutagenesis and infertility observed in animal studies. Eventually, it was recently approved in Europe as well, and there are on-going efforts to develop purely-Jak3 specific inhibitors (127) which will undoubtedly improve the safety of this strategy. The use of Jak3 inhibitor as a clinical drug will thus keep advancing. Nevertheless, we saw in our own hands that its clinical application may require further sophistication of the technology. First, Tofacitinib is a very potent inhibitor to all γ c-cytokine in vitro. However, it does not seem to show similarly potent inhibitory effects in vivo. In our mouse model of IL-15-induced leukemogenesis in which the continued production of IL-15 by

transgenesis leads to the development of CD8 T cell leukemia, anti-IL-15 modalities such as neutralizing mAb to IL-15 or mAb to IL-2R/IL-15R β perfectly protected those mice from leukemic death (22, 29, 84). So did BNZ 132-1 (22). Tofacitinib was administered using subcutaneous pump, but failed to protect host mice from leukemic death, only prolonged their survival (22, 29, 84). This may be due to the hydrophobic chemical nature of Tofacitinib, which makes it difficult to maintain the compound at effective dose all the time in vivo. Furthermore, the strategy of blocking Jak3 kinase itself may present another setback, because it blocks IL-7. IL-7 is a uniquely important member of the γ c-family because IL-7 broadly maintains the homeostasis and development of multiple lymphocyte lineages. Genetic perturbations or defects of the IL-7/IL-7R system lead to a profound loss of T and B cells in mice (9, 128), or a loss of T cells in humans (129). Normal numbers of B cells are present in the absence of the IL-7 signaling in human neonates, but a further study indicated that these individuals will develop B-cell dysfunction later in life (130). Thus, the impact of IL-7/IL-7R dysfunction is almost close to that of the loss of γ c function itself in magnitude (SCID in mice (20, 109), and X-linked SCID in humans (110)). The only difference in IL-7- and γ c-dysfunctions is the presence or absence of NK cells which are exclusively maintained by IL-15 (5, 131). These studies suggest that blocking IL-7 for a continued period of time may invite unfavorable outcomes as it could create severe immune suppression. We propose that a prolonged inhibition of IL-7 for therapeutic purposes should be avoided or kept at a minimum, unless IL-7 itself is the cause of the disease as reported in subgroups of T-cell or B-cell acute lymphocytic leukemia (T-ALL/B-ALL) (132, 133).

Figure 3, Alingment of D-helices of γ c-cytokines from various species



Alignment was performed using the Clustal Omega algorithm by the EMBL-EBI (151). The consensus sequence (A) contains many amino acids that participate in direct physical contact between the cytokine and γ c-molecule, providing evidence that the sequence conservation is underscored by a structural reason.

Thus, we saw a need to develop a new technology to block IL-2 and IL-15 (or other combinations of multiple γ c-cytokines, except IL-7) using a single agent. In a way, we tried to "shoot many γ c-cytokines by a single stone with a precise target guidance system". We hypothesized that such blocking is possible by interrupting the interaction of γ c-cytokines and the shared γ c subunit. Though all γ c-cytokines use a broad overlapping interface, there should be minor spots that are only occupied by select γ c-cytokines, but not by others. By targeting such unique spots, we should block select γ c-cytokine combinations.

4-b. The BNZ technology. The past two decades have seen a tremendous advancement in our structural knowledge on the interaction between cytokines and

receptors, owing to dedicated works by structural biologists. The IL-2/IL-15R system was no exception. The Garcia group (Stanford University) has conducted extensive studies on these receptor systems and mapped the amino acid residues by which various cytokines interact with their receptors. They have also delineated the binding interface utilized by IL-2 or IL-15 when they are binding to the γ c-molecule (134-139). These studies taught us that the D-helix (the last of the four helical bundles, a structure shared by many cytokine molecules) of the γ c-cytokines primarily interacts physically with the γ c-subunit. Naturally, it makes sense that D-helix is the most conserved among the all four α -helices of a γ c-cytokine across species (degree of similarities, D helix>B>A>C (22)). We noticed a moderately conserved motif

(which we named "the γ c-box", (22)) in this region of 6- γ c cytokines from many species (Figure 3A). We thus hypothesized that a peptide which mimics the sequence (and hence the 3D structure) of this region of γ c-cytokines may have the capacity to inhibit multiple γ c-cytokines. Since the amino acid sequences of the D-helices are moderately conserved, we hypothesized it would be possible to design a peptide which only resembles IL-2/IL-15, but not IL-4, -7, -9, or -21. However, this was not a simple task. We first synthesized a peptide corresponding to the D-helix of the human IL-15 protein, hoping this might be a specific inhibitor to IL-15, but not to other γ c-cytokines. This peptide did not inhibit IL-15 (determined by the inhibition of the T/NK cell proliferation in vitro) (Figure 4). It later turned out that the IL-15's D-helix peptide has a very weak binding affinity to the γ c so that it was easily competed off from the binding pocket by the natural ligand. We learned from this failure that we have to leverage the affinity of the peptide- γ c interaction and specificity of the peptide in interfering with the binding of each cytokine to γ c. So, we first carefully examined the published structural data concerning the binding of IL-15 to γ c (138). This study identified several crucial amino acids of the D-helix of IL-2 and IL-15 which form the binding interface with the γ c. We made these amino acids as the "backbone frame" of our peptides to ensure stable and tight binding of the peptides to the γ c. We then compared the sequences of IL-2 and IL-15 across species and generated "a consensus sequence" which consists of "conserved" and "wobble (variable)" positions (Figure 3B, (22)). To the wobble positions, we assigned amino acids based on

the usages by mammalian IL-2 and IL-15 molecules. We then ran rigorous computer docking simulation to rule out peptides that would not stably bind to the γ c at the predicted binding surface. We then synthesized over 30 peptides which passed the computer testing and examined their inhibitory nature using a specific bioassay and came up with the BNZ 132-1 peptide (a 19-mer synthetic peptide with the sequence of "IKEFLQRFIHIVQSIINTS", (22, 30)).

5. Proof-of-concept of the BNZ technology

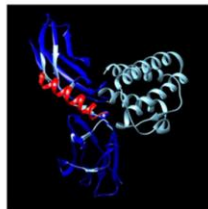
As shown in our recent publication, BNZ132-1 peptide efficiently blocks the in vitro function of IL-2 and IL-15 and to a lesser extent IL-9 as well (22, 30). We note that BNZ 132-1 has tight target specificity as it does not inhibit other γ c-cytokines (e.g., IL-4, IL-7, IL-21) or non- γ c cytokines (22). Our analysis indicated that the major signaling pathways downstream of the IL-2/IL-15/IL-9R are equally and completely blocked by BNZ 132-1 (22). To our knowledge this is the first example of a short peptide which blocks more than two cytokines from a family. We extended this technology to generate additional novel antagonistic cytokine mimetics which inhibit IL-15 and IL-21 (BNZ 132-2) and IL-4 and IL-9 (BNZ 132-3). It is also noteworthy that BNZ technology generates multi-cytokine inhibiting cytokine-mimetics without targeting IL-7, as we can exclude peptides which block IL-7 during the biological selection process. Though not yet published, we confirmed that BNZ 132-2, our second antagonistic mimetic peptide which blocks IL-15 and IL-21, does not block IL-7 function.

Figure 4 Failure of IL-15 inhibition by the D-helix peptide derived from human IL-15.

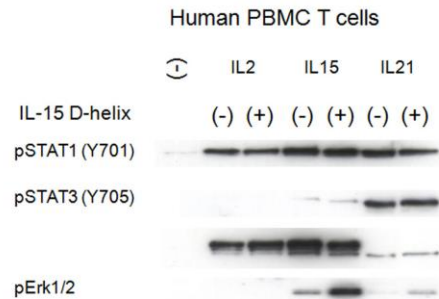
I-K-E-F-L-Q-S-F-V-H-I-V-Q-M-F-I-N-T-S; A 19 mer-D-helix of human IL-15

No binding to the γ c-pocket

Lack of inhibition on the IL-15 signaling



Computer-simulated docking;
 3D-structure of
 IL-15-D-helix/IL-15 vs. γ c



We first conducted a simple experiment to test our initial hypothesis that the D-helix peptide can be an antagonistic cytokine-mimetic agent. We ran a docking simulation (PatchDock algorithm, (152, 153)) which predicted that this peptide does not form stable binding with the γ c-molecule at the reported cytokine binding interface of this molecule (22). This highly suggests that the peptide may not work as an antagonist to γ c-cytokines. We then conducted a biological assay in which we incubated mitogen-activated human T cells with low doses of γ c-cytokines (IL-2; 50 pM, IL-15; 50 pM, IL-21; 10 pM) to induce tyrosine-phosphorylation of signaling molecules including STAT1, STAT3 and Erk1/2 as shown in the figure. We saw no inhibition of STAT1 and Erk1/2 which are signature signaling events associated with IL-15 activation of T-lymphocytes. The signaling by IL-2 or IL-21 was not inhibited, either. This experiment taught us that the D-helix sequence needs to be optimized so that the peptide functions as an antagonist by competitively inhibiting the binding of target cytokines to the γ c-molecule at the right interface.

6. Target diseases for the BNZ technology

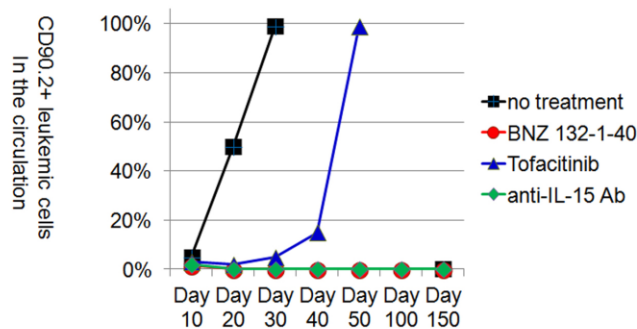
When we chose IL-2 and IL-15 as the target and started designing the inhibitory peptide, we already had in mind that such a peptide can be used for treating HAM/TSP caused by human T-cell leukemia virus-1 (HTLV-1) (23, 24). The constitutive in vivo activation of T-lymphocytes in HAM/TSP patients is initiated by the Tax-1 protein (140) of this virus and maintained by the combination of IL-2 and IL-15 produced as a consequence of HTLV-1 activation of CD4 T cells as we and others have previously demonstrated (25-28, 45, 141). In

addition to IL-2 and IL-15, a recent report suggests that the IL-9/IL-9R system is also playing a pertinent role in this disease (142). Incidentally, IL-9 is another target of BNZ 132-1 which was not originally anticipated by us when we designed BNZ132-1. As shown in our recent publication, BNZ132-1 alone can very efficiently block the spontaneous proliferation of T cells from multiple HAM/TSP patients (30). Furthermore, BNZ 132-1 also down-regulated various hallmark activation events associated with HAM/TSP T cells, including the tyrosine-phosphorylation of signaling molecule STAT5, and the survival of

HTLV-1 specific CD8 T cells (as determined by a Tax-tetramer staining) (30). Thus, BNZ 132-1 is a prime candidate by which to treat HAM/TSP patients. We expect that the PEGylated form of this peptide (see below) will stably circulate in the patients, subside the constitutive activation of T cells and revert the

lymphokine-activated killer (LAK) cells and CTLs to resting state so that they will stop damaging neuronal tissues/cells in HAM/TSP patients. Subsequent studies made us convinced, however, that there are more diseases that can be efficiently treated by the BNZ 132-1 peptide. We will describe our recent progress below.

Figure 5 Expansion of IL-15 leukemic cells was completely suppressed by BNZ 132-1 in syngeneic mice



The IL-15 leukemic cells line was established from the IL-15 Tg mouse (C57B6, CD90.2 background) we generated (6, 84). These cells grow in syngeneic normal mice (C57B6 strains) and form secondary leukemia upon grafting. Without treatment, the recipient mice will die of CD8 T cell leukemia within 30 days of the injection of leukemic cells. The graph shows the preferential expansion of CD90.2⁺ CD8 leukemic T cells in the CD90.1⁺ congenic C57B6 mice as determined by flow cytometry (Black line). The only CD90.2⁺ cells in these mice are from grafted IL-15 leukemic cells. Since these leukemic cells are dependent on IL-15 for their growth, injection of a neutralizing anti-IL-15 mAb (Mab247, R & D Systems, 150 μ g /intraperitoneal injection, twice per week for two weeks) eradicates leukemic cells thus no CD90.2⁺ leukemic T cells are found in the blood or spleen of recipient mice (Green line). Treatment by BNZ132-1-40 (PEGylated form, 1 mg/kg per dose, intraperitoneal injections twice per week) shows similar protective effects (Red line) as anti-IL-15mAb treatment. Curiously, Tofacitinib, infused by a subcutaneous pump, only delayed the onset of explosive expansion of leukemic cells, but failed to show a complete remission (Blue line).

7. BNZ 132-1 as a therapeutic drug.

7-a. Stable Pharmacokinetic nature of BNZ 132-1. Since BNZ 132-1 is a short (19-mer) peptide, it will quickly be lost from the body as the result of rapid renal clearance. Thus, we tried a few

modifications of the peptide to elongate the biological half life and PEGylation, a commonly used modifying technique for proteins and peptides, to enhance the pharmacokinetic (PK) properties of the peptide.

7-b. Pharmacodynamics of BNZ 132-1; The IL-15 leukemia model as the proving ground that BNZ 132-1 can be a therapeutic compound for LGL-type leukemia in humans.

After the confirmation that the PEGylated version (BNZ 132-1-40) has similar in vitro inhibiting activity on IL-2 or IL-15-gated cellular proliferation as the original peptide (22), Pharmacokinetics (PK) and pharmacodynamics (PD) of the drug were determined using mouse models of cytokine challenge. The over-expression of IL-2 leads to the preferential expansion of CD4 regulatory T cells (T-regs) (CD4⁺CD25^{hi}Foxp3⁺ T cells with suppressive function, reviewed in (143, 144)) as we have previously shown in the IL-2 transgenic (Tg) mouse that we generated (4). This can be imitated by administering recombinant human IL-2 into mice. Likewise, the injection of recombinant IL-15 into mice causes a temporal expansion of mouse NK cells (84). As reported (22), two bolus injections of recombinant human IL-2 or human IL-15 (150 μ g/shot) led to a temporal expansion of T-regs or NK cells, respectively. This transient expansion of T-regs or NK cells were almost completely blocked by the concomitant administration of BNZ 132-1-40 (1 mg/kg, two injections over a week). These experiments convinced us that BNZ 132-1-40 is stable in vivo and can be a drug-candidate for humans. PK studies in the currently on-going phase I-a clinical trials validated that the $T_{1/2}$ of BNZ 132-1-40 is around 72 hr in normal individuals, thus we could maintain it above minimum effective dose (0.2~0.4 mg/kg) by injecting a single dose of 1~2 mg/kg BNZ 132-1-40 over the next 14 days, reaffirming that BNZ 132-1-40 is a stable compound suited for clinical use.

Next, we tested the in vivo efficacy of the PEGylated form of BNZ 132-1 in a

clinically relevant model involving the over-production of IL-15. We and others had generated IL-15 transgenic (Tg) mice previously (6), and observed that they develop a fatal CD8 T cell leukemia (7, 8, 84). This experimental leukemia resembles human T-cell malignancies developing from CD8/LGL cells (7, 8). Using the IL-15 transgenic mouse they generated, the Caligiuri group (Ohio State University) demonstrated that this model recapitulates characteristics associated with human LGL leukemia and cutaneous T-cell leukemia (CTCL). In addition, they recently showed that the epigenetic modification which occurs at the *il15* locus in human LGL-leukemia patients can be observed in IL-15 Tg mice as well (145-147). Based on these observations, they proposed to treat patients with LGL leukemia and CTCL by anti-IL-15 therapies (147). Likewise, we explored our IL-15 Tg mouse model to test similar therapeutic possibility. As reported (84), we have established leukemic CD8 T cell clones from our IL-15 Tg mice. These cells can be expanded in culture without any addition of cytokines. These leukemic clones are still dependent on the IL-15 which they produce because the inclusion of anti-IL-15 antibody to the culture immediately stops cellular proliferation and induces apoptotic death (84). Importantly, these cells can be grafted into a syngeneic host mouse (the C57B6 strain) and kill the host within 4-6 weeks time by forming a secondary leukemia/lymphoma (29, 84). This model was used to test if BNZ 132-1 suppresses the leukemic growth of IL-15-producing CD8 T-cells and protects host mice from leukemic death. As control and comparison, anti-IL-15 neutralizing antibody and Tofacitinib were included in the experiment. Control mice without any treatment died within 30 days following the injection of IL-15 Tg leukemic cells (K2) as reported before (29, 84). Anti-IL-15, or anti-mouse IL-

2R/IL-15R β (anti-mouse CD122, TM β 1) mAb protected mice as expected. BNZ 132-1-40 (the PEGylated form), administered intravenously (IV) 4 times (on days 1, 4, 8, and 11 after the injection of leukemic cells) at 1 mg/kg dose, prevented the mouse as efficiently as anti-cytokine/cytokine-receptor antibodies. Moreover, experiments were conducted to allow distinguishing grafted leukemic cells and host lymphocytes by congenic markers (i.e., leukemic cells are from CD90.2 C57B6 mice whereas the recipient mice are CD90.1 C57B6). A thorough flow-cytometric examination of the blood and secondary lymphoid organs revealed that there were no traceable CD90.2⁺ leukemic cells remaining in host mice after the BNZ132-1-40 treatment. Autopsy and histological analyses validated the assessment. These data strongly suggest that BNZ132-1-40 has completely eradicated IL-15-dependent CD8 leukemic T cells from leukemic mice. Curiously, Tofacitinib, which was administered using a cutaneous pump showed only marginal therapeutic effects as described above (22, 29). These results suggest that BNZ132-1-40 may be a superior "in vivo blocker" of IL-2 or IL-15 than Tofacitinib.

7-c. IND approval and on-going clinical trials involving BNZ132-1. As described above, BNZ132-1 has received an IND approval from the FDA to conduct a phase I-a clinical trial targeting normal volunteers to evaluate its drug candidacy. In preclinical studies that we conducted before an IND submission, we saw reduction of NK and some other types of T cells in the mice and in the cynomolgus macaques, as expected based on the known homeostatic roles of IL-15 to maintain these cells and those of IL-2 on T-regs, respectively. In the Phase I-a clinical trial that has just been completed, a significant decrease of NK cells and of other relevant lymphocyte

subsets was observed following a single administration of BNZ 132-1-40 (0.2, 0.4, 0.8, 1.6, 3.2 and 6.4 mg/kg). The detailed results will be published elsewhere in the near future (manuscript in preparation). To apply BNZ132-1 to wider diseases, studies are in progress to test if BNZ132-1 effectively suppresses graft-versus-host (GvH) responses upon transplantation. Previous studies suggested that this process involves IL-15, but not IL-2 (148-150). However, in our experimental system we saw that only co-inhibition of IL-2 and IL-15 (by combining anti-IL-2 and anti-IL-15 antibodies, or by BNZ132-1-40), but not the blockade of a single cytokine, ameliorated the GvH symptoms (manuscript in preparation). This represents another compelling example in which monotherapy (targeting a single cytokine) fails to display therapeutic effects and multi-cytokine inhibition is needed for attaining the goal. Thus, we are continuing our efforts to obtain additional IND approvals on BNZ132-1 for treating other hematologic and immunologic diseases including GvH disease and autoimmunity.

Summary

Years ago, we observed that the ex vivo spontaneous proliferation of HAM/TSP T cells which is driven by a combination of almost equal amounts of IL-2 and IL-15 cannot be efficiently blocked by a single neutralizing mAb either to IL-2 or IL-15. This contrasted with our intuitive prediction in which we expected to see a near 50% neutralization by the individual antibody, which made us realize the fraud in our logic and a related gap in the existing anti-cytokine approach, and triggered subsequent research to develop a new tool (multi-cytokine inhibitor) that we have now. Our recent publications represent the proof-of-concept of our novel technology to rationally design multi-cytokine inhibitors.

Our research shows that an increasing number of human diseases may be efficiently treated by our multi-cytokine inhibitors (MCIs). Currently our compounds are still limited to blocking the γ c-family cytokines, but we plan to develop new class of inhibitors that block cytokines belonging to different cytokine families..

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BNZ 132-1 and other BNZ peptides were protected by the US patent (08455449) and owned by the BIONIZ therapeutics.

Animal experiments were conducted at the BioQual Inc. (Rockville, MD) under protocols approved by the IACUC committee.

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