



Mixing signals: molecular turn ons and turn offs for innate $\gamma\delta$ T-cells

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Lymphocytes of the gamma delta ($\gamma\delta$) T-cell lineage are evolutionary conserved and although they express rearranged antigen-specific receptors, a large proportion respond as innate effectors. $\gamma\delta$ T-cells are poised to combat infection by responding rapidly to cytokine stimuli similar to innate lymphoid cells. This potential to initiate strong inflammatory responses necessitates that inhibitory signals are balanced with activation signals. Here, we discuss some of the key mechanisms that regulate the development, activation, and inhibition of innate $\gamma\delta$ T-cells in light of recent evidence that the inhibitory immunoglobulin-superfamily member B and T lymphocyte attenuator restricts their differentiation and effector function.

Keywords: BTLA, dermatitis, $\gamma\delta$ T-cell, IL-7, lymphotoxin, ROR γ t

INTRODUCTION

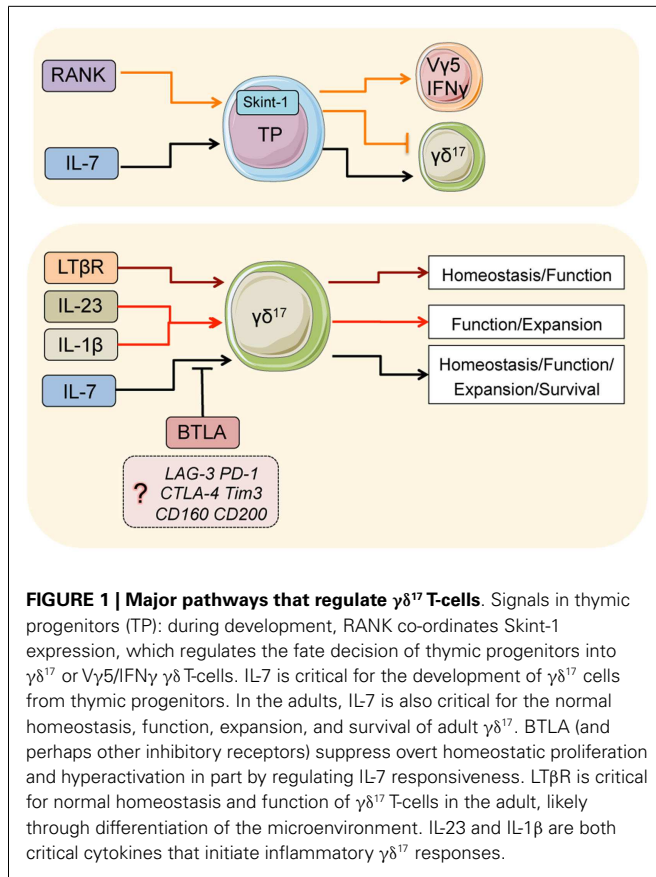
The ability to generate antigen receptor diversity by somatic recombination evolved approximately 500 million years ago (1) and became the founding biological property of what we now know as adaptive immunity. This evolutionary milestone provided our immune system with an innate and an adaptive arm that synergized for the fight against infection and the recognition of oncogenesis. Lymphocytes of the gamma delta ($\gamma\delta$) T-cell lineage are evolutionary conserved among species (2) and although they express rearranged antigen-specific receptors, a large proportion display innate properties. In the mouse, where innate $\gamma\delta$ T-cells have been mostly studied, approximately 25% of lymph node $\gamma\delta$ T-cells respond rapidly to cytokine stimuli similar to innate lymphoid cells (ILCs) and appear to have reduced T-cell receptor (TCR) signaling capacity (3). Innate $\gamma\delta$ T-cells are characterized by the spontaneous and high expression of interleukin (IL)-17 ($\gamma\delta^{17}$) as well as IL-22 and express functional Toll-like receptors (TLR) (4, 5). Importantly, IL-17 and IL-23 receptor (IL-23R) expression, which is critical for IL-22 induction, are turned on during embryonic development in the thymus strongly pointing toward a *bona fide* innate nature (6–8). Although a new interferon gamma (IFN γ)-producing innate $\gamma\delta$ T-cell subset with no IL-17 potential has recently been described (3), this review will discuss briefly some of the key cytokines, cytokine receptors, and transcription factors (TFs) that regulate the development, activation, and inhibition of mouse innate $\gamma\delta^{17}$ cells (Figure 1).

IL-23 AND IL-1 β : KEY PROINFLAMMATORY AND ANTI-BACTERIAL MEDIATORS

Innate $\gamma\delta^{17}$ cells localize mainly at barrier and mucosal surfaces such as the skin, gut, and lung (9) and within the lymph nodes, they position themselves in close proximity to the subcapsular sinus and interfollicular regions both of which specialize in the capture of antigen (10). Therefore, infectious and inflammatory stimuli can readily activate $\gamma\delta^{17}$ cells either directly through TLR ligation or through cytokines such as IL-23 and IL-1 β that are produced by local innate sensors.

IL-23 induces the expression of IL-17 and IL-22 as well as the transcription factor retinoid-related orphan receptor gamma-t (ROR γ t) in T-helper 17 (T_H17) cells while at the same time promoting survival and cell proliferation (11). $\gamma\delta^{17}$ cells express functional IL-23R as early as embryonic day E18 in the thymus (7), in contrast to CD4⁺ T-cells that upregulate the IL-23R upon T_H17 differentiation (12). Although IL-23 or IL-23R has not been reported to be important for $\gamma\delta$ T-cell development, they enhance the production of IL-17 and IL-22 and can promote cellular proliferation (3, 13). *In vivo* infectious and inflammatory models have shown that IL-23 can be important for the activation of the $\gamma\delta$ T-cell response.

During imiquimod (IMQ)-induced psoriasis, genetic ablation of IL-23 or IL-23R results in a significant reduction of IL-17 production by $\gamma\delta^{17}$ cells, diminished accumulation of these cells in the skin, and a subsequent decrease in inflammatory symptoms (14–16). In this model, IL-23 is produced locally in the skin by resident



macrophage and dendritic cell (DC) populations that receive a combination of TLR and neuronal signals (15, 17, 18). The onset of experimental autoimmune encephalomyelitis (EAE), which is often used to model human multiple sclerosis, also depends to a certain extent on IL-23-driven IL-17 production by $\gamma\delta$ T-cells (5, 19). More specifically, it has been shown that IL-23-activated $\gamma\delta^{17}$ cells are important for optimal T_H17 polarization (5) and the suppression of regulatory T-cell responses (19). In a mouse model of brain ischemic injury, absence of IL-23 also abrogated $\gamma\delta^{17}$ -induced inflammation (20). In addition to regulating inflammatory reactions, $\gamma\delta^{17}$ cells and IL-23 have been linked with protection from a number of bacterial infections. Thus, cutaneous infection with *Staphylococcus aureus* triggers a $\gamma\delta$ T-cell orchestrated IL-17 response that depends on the combined effects of IL-23 and IL-1 β (21). Furthermore, infection with *Listeria monocytogenes* elicits an IL-23-driven $\gamma\delta^{17}$ response that is important for bacterial clearance (22, 23), and the IL-23 pathway appears also to operate during $\gamma\delta^{17}$ activation by *Mycobacterium tuberculosis* (24). Together, these data highlight the role of IL-23 in activating $\gamma\delta^{17}$ cell-induced inflammatory responses, both to pathogens and in driving autoimmune disease.

Similar to IL-23, IL-1 β has also been linked with IL-17-related immunity both in CD4⁺ T as well as in innate $\gamma\delta$ T-cells. $\gamma\delta^{17}$ cells constitutively express the IL-1 receptor and respond to *in vitro* IL-1 β stimulation by rapid proliferation and upregulation of IL-17 (3, 5, 13). Interestingly, IL-1 β appears to be important

for IL-23-mediated $\gamma\delta$ T-cell expansion and IL-17 production although the molecular mechanism is not yet understood (5, 13). Effective IL-1 β signaling was critical for $\gamma\delta$ T-cell activation and disease progression in the EAE model (5). However, during IMQ-induced psoriasis, usage of *Ilr1*^{-/-} mice has resulted in conflicting conclusions. Whereas an earlier report presented no impact of IL-1 β on either dermatitis or $\gamma\delta^{17}$ activation (25), a more recent study showed that *Ilr1*^{-/-} mice were consistently protected with severely compromised $\gamma\delta$ T-cell responses (13). A key difference in the two studies was the site of inflammation: ear (no IL-1 β effect) (25) versus dorsal epidermis (strong IL-1 β effect) (13), suggesting that IL-1 β may have site-specific regulatory roles, such as differential effects on resident stromal and epithelial cells or due to differences in lymphatic drainage.

IL-7: KEEPING THE BALANCE BETWEEN HOMEOSTASIS AND INFLAMMATION

IL-7 is one of the best-studied T-cell homeostatic cytokines. IL-7 deficiency is associated with lymphopenia and dysfunction of naive and memory T-cell subsets (26). IL-7 is essential for the development of $\gamma\delta$ T-cells (27, 28) by regulating the survival of early thymic progenitors and by inducing V(D)J recombination within the TCR- γ locus (29, 30). Further experiments have shown that in addition to its developmental role, IL-7 supports the homeostatic proliferation of $\gamma\delta$ T-cells (31). Although IL-7 is strongly associated with signaling via the signal transducer and activator of transcription 5 (STAT5) (32), it has been shown to induce STAT3 phosphorylation in diverse lymphocyte populations such as thymocytes (33), B-cell progenitors (34), and $\gamma\delta$ T-cells (35). STAT3 is a critical component of the IL-23 and IL-6 signaling pathways, which are important for the differentiation of CD4⁺ T-cells into the T_H17 lineage (11, 36), in part by antagonizing STAT5 (37). Of the $\gamma\delta$ T subsets, IL-7 was found to preferentially expand and activate innate $\gamma\delta^{17}$ cells in a STAT3-dependent manner (35), although it sustained survival of all $\gamma\delta$ T-cells (38).

We have recently demonstrated that in $\gamma\delta^{17}$ cells, STAT5-mediated IL-7 signaling induces surface expression of the checkpoint receptor B and T lymphocyte attenuator (BTLA), which is necessary for their normal homeostasis and activation during skin inflammation (38). Blockade of IL-7 signaling itself has been shown to acutely diminish $\gamma\delta^{17}$ -driven dermatitis (35) while during viral hepatitis IL-7 co-operates with IL-23 to rapidly activate intrahepatic $\gamma\delta^{17}$ cells and initiate inflammation (39). Whether IL-7-induced STAT5 and STAT3 phosphorylation operate in parallel, sequentially, or as mutually exclusive processes within the $\gamma\delta^{17}$ population is unknown. However, $\gamma\delta$ T-cells deficient in STAT3 display normal homeostatic responses (40) suggesting that at steady state STAT5 may have a dominant role.

In addition to its direct effects on $\gamma\delta$ T-cells, IL-7 indirectly influences innate $\gamma\delta$ T-cell development by promoting the generation of lymphoid tissues in part by inducing the expression of tumor necrosis factor (TNF) superfamily members. IL-7 is produced homeostatically in the developing thymus and lymph node anlagen (41) and has been shown to induce the expression of surface lymphotoxin- $\alpha\beta$ (LT $\alpha\beta$) on resident embryonic lymphoid tissue inducer (LTi) cells (42). LT $\alpha\beta$ expressed by LTi interacts with the LT β receptor (LT β R) in order to initiate lymph node

development and organization (43, 44). Genetic ablation of LT β R results in the absence of all secondary lymphoid tissues in addition to disorganized splenic and thymic architecture (45, 46). Several members of the TNF superfamily have been shown to directly regulate $\gamma\delta$ T-cell development, homeostasis, and function, as outlined below.

LYMPHOTOXIN AND THE TNF NETWORK: CRITICAL REGULATORS OF INNATE $\gamma\delta$ T-CELLS

Innate IL-17 producing $\gamma\delta$ T-cells as well as V γ 5 (V γ 3 in Garman nomenclature) expressing cells that colonize the skin as resident dendritic epidermal T-cells (DETCs) are strictly dependent on the embryonic microenvironment (8, 47). Thus, adult progenitors cannot reconstitute either of the aforementioned populations even if they are provided with a fetal thymus suggesting the need for embryonic-only progenitors (8). Thus, the fetal thymus contains fully functional $\gamma\delta^{17}$ cells that develop between E15–18 (8). The development of these cells is intimately associated with the TNF superfamily since as early as E15 V γ 5⁺ progenitors express the TNF ligand RANKL (receptor activator of NF- κ B ligand) and condition the thymic medulla to upregulate Skint-1 (48), an immunoglobulin (Ig) superfamily protein that is necessary for the development of V γ 5 cells (49–51). Interestingly, in Skint-1 deficient animals, V γ 5 cells are reprogrammed into a $\gamma\delta^{17}$ -like phenotype with severely reduced IFN γ production (52). This suggests that innate $\gamma\delta^{17}$ T-cells are likely to represent the default differentiation pathway of most $\gamma\delta$ T-cell progenitors pre-Skint-1 selection. This is in line with the evolutionary evidence that IL-17-producing $\gamma\delta$ T-cells are conserved between non-jawed vertebrates and human beings (2) while Skint-1 and related genes (e.g., Btn1a1) are highly restricted to mammals (www.ensembl.org).

In addition to RANK, LT β R has also been linked with the development and functional maturation of $\gamma\delta$ T-cells. Early reports showed that $\gamma\delta$ T-cells can acquire LT β R expression in the thymus, and that activation of these receptors by LT $\alpha\beta$ - and LIGHT-expressing double-positive (DP) thymocytes drives maturation of $\gamma\delta$ T-cells assessed by the production of IFN γ (53). However, the expression of IL-17 or other $\gamma\delta^{17}$ -related properties was not evaluated. The authors suggested that LT β R-induced maturation likely occurred at the late stages of thymic development when DP cells predominate. Given that $\gamma\delta^{17}$ T-cells develop during early embryonic life (8), one scenario to explain these findings is that during thymic development the LT β R pathway in part regulates the IFN γ potential of $\gamma\delta$ T-cells, presumably following Skint-1 selection. In agreement with this argument, the TNF receptor CD27 is required by thymic progenitors to induce the innate IFN γ -related differentiation program and to sustain expression of LT β R (7). Thus, while CD27 deficient animals retain an intact $\gamma\delta^{17}$ compartment, they showed a marked reduction in IFN γ and LT β R expression (54). These results predict that LT β R signaling is not absolutely necessary for $\gamma\delta^{17}$ development and function, although mice deficient in LT β R or its ligands had very few IL-17-producing $\gamma\delta$ T-cells in the spleen and thymus (55). Mice lacking the NF- κ B TFs RelA and RelB also showed reduced IL-17-producing $\gamma\delta$ T-cells (55). Since the NF- κ B pathway is central to TCR signaling and T-cell development (56), low

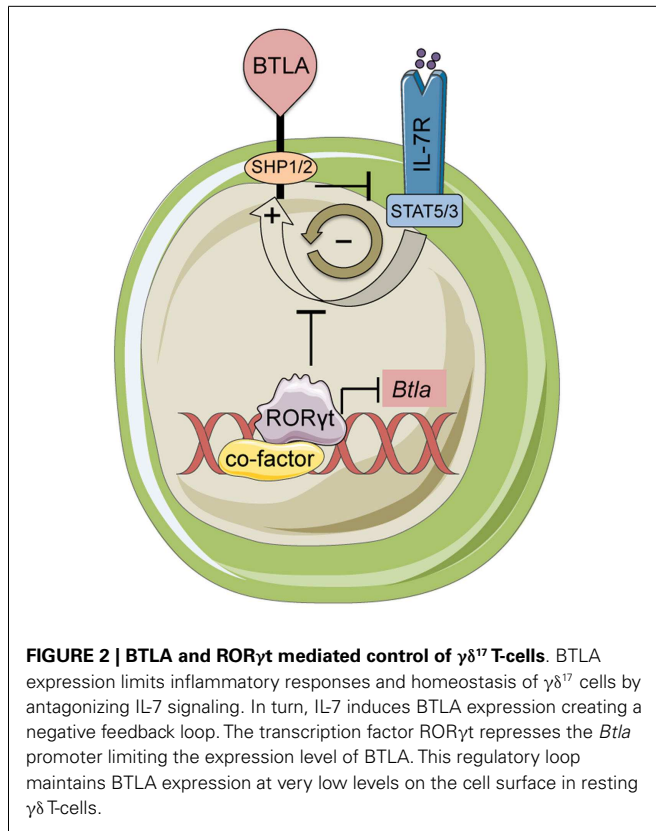
IL-17 production might be reflective of impaired TCR stimulation rather than loss of LT β R signals. Furthermore, lack of lymph nodes in LT β R deficient mice (45) may relocate $\gamma\delta^{17}$ cells to the skin or intestine and thus explain their reduced numbers in the spleens. Importantly, loss of LT β R results in abnormal thymic organization and maturation of the medullary epithelium (46, 57), which may negatively affect $\gamma\delta^{17}$ T-cell development. Alternatively, organized secondary lymphoid tissues may be important for the survival and steady-state turnover of $\gamma\delta^{17}$ cells. Of note, LT β R has been shown to participate in the production of IL-7 by fibroreticular stromal cells in the lymph node (58), which might explain why deficiency in LT β R can result in reduced $\gamma\delta^{17}$ responses.

In addition to its involvement in stromal cell development, LT β R is expressed on tissue resident DCs and macrophages (59) both of which have been linked with the IL-23-mediated activation of $\gamma\delta^{17}$ T-cells, whether this is in the context of skin (15, 17) or brain inflammation (5). Notably, LT β R regulates the homeostasis of DCs (60, 61) and can directly induce their production of IL-23 (62). Interestingly, an LT β R-LT $\alpha\beta$ interaction has been linked with the production of IL-22 by intestinal ILCs (63, 64) raising the possibility that a similar mechanism may be in place at sites where $\gamma\delta^{17}$ cells preferentially localize, such as the skin.

BTLA AND INHIBITORY RECEPTORS: PUTTING THE BRAKES ON

In human beings, herpesvirus entry mediator (HVEM) interacts with the two TNF ligands LIGHT (shared with LT β R) and soluble LT α , and the Ig superfamily members CD160 and BTLA. BTLA is an inhibitory receptor with an immunoreceptor tyrosine inhibitory motif (ITIM) that has been shown to interact with the Src homology 2 (SH2)-domain containing protein tyrosine phosphatase 1 (SHP1) and SHP2 and to inhibit T-cell activation (65–67) upon interacting with HVEM, its only identified ligand thus far (66, 68, 69). In addition to its inhibitory role in T-cell responses, BTLA was shown to prevent overt TLR stimulation in DCs (70) and to diminish cytokine production by natural killer T (NKT) (71) and follicular T-cells (72) suggesting a regulatory role both in adaptive and innate immunity.

BTLA and HVEM signal bi-directionally providing inhibitory signals in T-cells and survival signals in cells expressing HVEM (68). BTLA expression varies $\sim 10^3$ fold among hematopoietic lineages, and co-expressed with HVEM forming a complex *in cis* that may contribute to homeostatic signaling (73). Constitutive surface expression of BTLA (74) implicates a unique ability among inhibitory receptors to sustain the homeostatic balance of T-cells (75) and DCs (61). Similarly, our recent data showed that BTLA is necessary to inhibit homeostatic expansion and activation of lymph node and skin resident $\gamma\delta^{17}$ T-cells (38). $\gamma\delta^{17}$ but not other $\gamma\delta$ T-cell subsets deficient in *Btla* were hyperresponsive to IL-7 stimulation suggesting that BTLA diminishes IL-7 receptor (IL-7R) signaling. Interestingly, IL-7 increased surface BTLA on $\gamma\delta^{17}$ cells in a STAT5-dependent way revealing the presence of a negative feedback loop between IL-7 and BTLA (38) (Figure 2). Given the broad range of SHP1 and SHP2 targets (76), it is likely that these phosphatases can inactivate both STAT3 and STAT5 in response to IL-7. However, the exact molecular details of BTLA-mediated



suppression of IL-7R or other $\gamma\delta^{17}$ -expressed cytokine receptors are currently not known.

Although there are numerous functional inhibitory receptors that have been reported on the surface of lymphocyte subsets either at steady state or after activation, there is little information regarding their role on innate or non-innate $\gamma\delta$ T-cells. Several reports have mapped the expression of inhibitory molecules like programmed death-1 (PD-1) (77, 78), lymphocyte activation gene-3 (LAG-3) (79), CD200 (80), Tim-3 (81), CD160 (82), and cytotoxic T lymphocyte antigen-4 (CTLA-4) (83) on human or murine $\gamma\delta$ T-cells but the capacity to target these receptors using agonistic or antagonistic manipulation has in general not been addressed. Notably, we found that activating BTLA receptors using an agonistic antibody limited pathology in mice (38). Additionally, blockade of BTLA signaling enhanced activation of lymphoma-specific human V γ 9V δ 2 T-cells (84). Thorough investigation of the expression patterns and function of the different inhibitory receptors on innate $\gamma\delta$ T-cells may provide promising targets for intervening when these lymphocytes need to be turned on or off. Currently, and in combination with its suppressive activity, BTLA appears to be a key targetable pathway for regulating innate $\gamma\delta$ T-cells.

TRANSCRIPTIONAL CONTROL: IS EVERYTHING PRE-PROGRAMMED?

It is now well-appreciated that there is an extensive network of TFs that are expressed early in pre-committed progenitors and are necessary for the development, functional differentiation, and survival

of all innate cells including $\gamma\delta^{17}$ T-cells. A subset of these TFs control lineage specification, either through activating or repressing gene transcription. A number of TF mouse knockout lines result in the complete abolishment or severe reduction in the numbers of the $\gamma\delta^{17}$ subset in the periphery and in the thymus. Thus, mice deficient for the high-mobility group (HMG) box TFs Sox13 and Sox4 show severe reduction of IL-17-producing $\gamma\delta$ T-cells due to a differentiation block early on during development (85, 86), which correlates with high expression levels of Sox13 and Sox4 in $\gamma\delta^{17}$ -committed T-cell progenitors (86–88). Interestingly, the function of Sox13 can be counteracted embryonically by Egr3, which drives the DETC differentiation program and IFN γ expression (52), while TCF1, another HMG box TF, suppresses $\gamma\delta^{17}$ differentiation (86). Notch signaling turns on TCF1 (89), which can also induce expression of Hes1, another TF critical for the generation of $\gamma\delta^{17}$ cells during embryonic differentiation (40). Interestingly, a subset of innate $\gamma\delta$ T-cells has been shown to depend on the expression of promyelocytic zinc finger (PLZF), which is also required for the development of ILCs (90, 91). It remains to be seen whether PLZF is specifically required for the development of $\gamma\delta^{17}$ cells.

Although, ROR γ t is necessary for the differentiation of T_H17 cells (36), it is not essential for the development of $\gamma\delta^{17}$ progenitors in the fetal thymus (40). However, consistent with its ability to bind to and transactivate the *Il17* promoter (92), ROR γ t is important for optimal IL-17 production (40). Interestingly, despite being developed, ROR γ t deficient $\gamma\delta$ T-cells cannot persist in the periphery (40), suggesting a potentially critical role for ROR γ t in the homeostasis of adult $\gamma\delta^{17}$ T-cells. This could be either cell-extrinsic or cell-intrinsic. ROR γ t is necessary for the development of all secondary lymphoid tissues (93). Thus, upon export in the periphery, $\gamma\delta^{17}$ T-cells may not have the appropriate microenvironment in order to sustain homeostasis (cell-extrinsic). In the cell-intrinsic scenario, ROR γ t may be important for the survival of $\gamma\delta^{17}$ cells by regulating the levels of the anti-apoptotic protein Bcl-xL (93). Our data have demonstrated that via its interaction with LxxLL containing nuclear co-factors ROR γ t can function as a transcriptional repressor and suppress expression of BTLA (38) (Figure 2). Therefore, an alternative cell-intrinsic hypothesis is that loss of ROR γ t results in aberrant expression of BTLA and perhaps other co-inhibitory receptors (such as LAG-3; Bekiaris/Ware, unpublished observations) leading to a sustained inhibition of homeostatic expansion.

CONCLUSION

$\gamma\delta^{17}$ and other $\gamma\delta$ T-cell subsets comprise a unique family of lymphocytes that provides an innate powerhouse to the immune system. The innate nature of $\gamma\delta^{17}$ cells is demonstrable by a number of key biological properties including rapid response to cytokines, functional maturation during embryogenesis, largely TCR-independent responses, and TF-dependent lineage commitment. Resolving the complex and fascinating biology of these cells has been breaking the Frontiers of Immunology for a number of years and has taught us a great deal about how lymphocytes develop and function. The continued knowledge of how all innate $\gamma\delta$ T-cells work will certainly push forward these frontiers and perhaps allow us to develop tools in order to manipulate them for the treatment of human disease.

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