# The role of G-protein-coupled receptors in the biological activity of $\gamma\delta$ T lymphocytes

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#### **Abbreviations**

αβ T cells alpha-beta T cells

BSA bovine serum albumin

Ca<sup>2+</sup> calcium

cAMP cyclic adenosine-3',5'-monophosphate

CI chemotactic index Ctx. cholera toxin

C5a complement factor 5a

DAG diacylglycerol

dimaprit S-(3-dimethylaminopropyl)isopthiourea 2HCl (dimaprit-2HCl) EGTA ethyleneglycol-bis(beta-aminoethylether)-N,N'-tetraacetic acid

ERK extracellular signal regulated kinase

FACS fluorescence cytometry

FCS fetal calf serum
FITC fluoro-isothiocyanate
fMLP N-formyl-Met-Leu-Phe

FPR fMLP-receptor

FPRL1, FPRL2 fMLP receptor-like 1 and 2

γδ T cells gamma-delta T cells
GCPR gamma-delta T cells

H1R, H2R, H3R, H4R histamine receptor subyptes (1-4)

HTMT 6-[2-(-imidazolyl)ethylamino]-N-(4-trifluoromethylphenyl)

heptanecarboxamide dimaltate

IL- interleunkin-IFN- interferon

IP<sub>3</sub> inositol-1,4,5-trisphosphate NK cell natural killer cells

NFκB nuclear factor 'kappa – light – chain –enhancer'

PBMC peripheral blood mononuclear cells

PBS phosphate-buffered saline

PE phycoerythrin

PIP<sub>2</sub> phosphatidylinositol (4,5)-bisphosphate PIP<sub>3</sub> phosphatidylinositol (3,4,5)-trisphosphate

PI3K phosphatidylinositol 3-kinase

Ptx. pertussis toxin

Raji human Burkitt's lymphoma cell line

RANTES regulated on activation-normal T cell expressed and secreted thioperamide N-cyclohexyl-4-(1H-imidazol-4-yl)-1-piperidinecarbothioamide

maleate salt (thioperamide)

### **Summary**

The data collected in this work represent the first evidence that G-protein-coupled signalling exist in γδ T cells via histamine and fMLP receptors. The bigenic amine histamine and the bacterial peptide fMLP have been demonstrated to be novel chemoattractant factors for circulating human γδ T cells, which are critical members of the immunological tumor surveillance machinery. Here, we analyzed the influence of histamine on the interaction of human γδ T cells with tumor cells such as the A2058 human melanoma cell line, the human Burkitt's non-Hodgkins lymphoma cell line Raji, the T-lymphoblastic lymphoma cell line Jurkat, and the NK cell-sensitive erythroleukaemia line K562. We found that histamine inhibits the spontaneous cytolytic activity of  $\gamma\delta$  T cells in response to these cell lines. The downregulation of  $\gamma\delta$  T cell mediated cytotoxicity involves the histamine receptor subtype 2 (H<sub>2</sub>R), the activation of G<sub>s</sub> proteins and increased cAMP intracellular levels. On the other hand, histamine activates the common signalling pathways of chemotaxins such as G<sub>i</sub>-protein-dependent actin reorganization, the increase of intracellular Ca<sup>2+</sup> and the induction of migratory response in  $\gamma\delta$  T lymphocytes. Our data indicate that histamine contributes to the mechanism by which tumor cells escape immunological surveillance.

The bacterial-cell-wall-derived peptide N-formyl-Met-Leu-Phe (fMLP) is a well characterized chemotactic factor for phagocytes such as neutrophils, monocytes and dendritic cells. Here, we analyzed the influence of fMLP on isolated human peripheral blood  $\gamma\delta$  T cells. We found that fMLP induces intracellular calcium transients, actin reorganization, CD11b upregulation and the migration of  $\gamma\delta$  T cells. Pretreating  $\gamma\delta$  T cells with pertussis toxin inhibited all fMLP-stimulated cell responses, implicating the involvement of Gi proteins in the induced signalling cascade. The present data suggest that, in addition to phagocytes, N-formyl peptides also regulate the trafficking and activation of  $\gamma\delta$  T cells.

#### Zusammenfassung

Die Kommunikation zwischen Zellen ist eine essentielle biologische Funktion in multizellulären Organismen. Externe Signale können durch spezifische Rezeptoren der Zielzellen wahrgenommen werden und lösen Protein-Protein-Interaktionen, Protein-Phosphorylierungen und die Bildung sekundärer Botenstoffen aus und führen schließlich zu spezifischen, zellulären Reaktionen.

G-Protein gekoppelte Rezeptoren gehören der Superfamilie der sogennanten Serpentin Rezeptoren an, deren charakteristisches Merkmal sieben transmembranäre Helices sind. G-Proteine selbst sind aus drei Unterinheiten ( $\alpha$ , $\beta$  und  $\gamma$ ) aufgebaut. Bindet ein Botstoff, beispielsweise Histamin oder fMLP an seinen Rezeptor, so wird durch eine Konformationswandlung die Bindungsstelle für G-Proteine freigelegt. Innerhalb des Immunsystems sind G-Proteine Regulatoren von zellulärer Reaktionen. Die G-Protein gekoppelte Rezeptoren sind an die Einrichtung von verschiedenen Zellfunktionen beteiligt. Beispielsweise induzieren Chemotaxine wie fMLP in neutrophilen Granulozyten G-Protein-abhängig die gerichtete Migration und die ROS-Produktion der Zellen. Mittels RT-PCR, Western blot Analyse und Flow Zytometrie wurden Histamin und fMLP Rezeptoren an der Oberfläche von aus dem peripheralen Blut isolierten humane γδ T Zellen identifiziert. Histamin (2-(4-Imidazolyl)-ethylamin), das biogene Amin wird durch Abspaltung von Kohlendioxid aus der Aminosäure Histidin gebildet und insbesondere in Mastzellen, basophilen Granulozyten und Nervenzellen gespeichert. N-formyl-Met-Leu-Phe (fMLP) ist ein in den Bakterien vorkommender chemotaktische Faktor. Es wurde gezeigt, dass Histamin und fMLP als chemotaktische Faktoren für humane γδ T Zellen funktionieren. Es wurde weiterhin demonstriert, dass die Migration der humanen γδ T Zellen zu den beiden Faktoren eine G<sub>i</sub>-Proteinabhängige Reaktion der Zellen ist. Wir haben nachgewiesen, dass es in humane γδ T Zellen für Histamin drei unterschiedlichen Rezeptoren-Untereinheiten (H<sub>1</sub>, H<sub>2</sub>, H<sub>4</sub>) existieren. Von denen der G<sub>s</sub>-Protein gekoppelte H<sub>2</sub>-Rezeptor für die Regulierung der Zytotoxizität Reaktion und der G<sub>i</sub>-Protein gekoppelte H<sub>4</sub>-Rezeptor für die chemotaktische Antwort der humane γδ T Zellen auf Histamin verantwortlich ist.

# 1. Introduction

# 1.1. The immune system

The immune system is an organization of cells and molecules with specialized roles in defense against infection. There are two fundamentally different types of responses to invading microbes. Innate responses occur to the same extent however many times the infectious agent is encountered, whereas adaptive responses become more exact on repeated exposure to a given infection [1]. Innate and acquired responses usually work together to eliminate pathogens. The innate responses use neutrophils, monocytes, macrophages, basophils, mast cells, and eosinophils, natural killer cells, and  $\gamma\delta$  T cells. The molecular components of innate responses include complement, acute-phase proteins, and cytokines [1]. Acquired responses involve the proliferation of antigenspecific B and T cells. Specialized cells, called antigen presenting cells, present the antigen to lymphocytes and collaborate with them in the response to the antigen, to initiate the target-specific response. The response of the immune system to pathogens, immune complexes, and injured tissues depends in large part on the functional relationships between the different immune cells.

The accumulation of leucocytes at sites of inflammation and microbial infection in response to locally produced chemotactic factors is an important feature of the immune response [2]. The accumulation of selected populations of leukocytes is an orchestrated event that probably involves the coordinated expression of a variety of adhesion molecules, target-cell-specific chemokines, cytokines, growth factors and enzymes that contribute to the ensuing tissue remodelling, as defined by the degradation of existing and the deposition of new extracellular matrix proteins and proteoglycans, and angiogensis [3]. A large group of chemotactic factors such as all chemokines, histamine

and N-formyl peptides, e.g N-formyl-Met-Leu-Phe (fMLP) engage seven-transmembrane G-protein-coupled-receptors (GPCRs).

# 1.2.1. G-protein-coupled receptors (GPCRs)

With nearly 1000 members, G-protein-coupled receptors (GPCRs) constitute the largest group of cell-surface proteins that are involved in signal transduction. They account for more than 2% of the proteins that are encoded by the human genome [4]. GPCRs are often referred as serpentine receptors because they span the plasma membrane seven times with α-helical segments separated by alternating intracellular and extracellular loop regions. The N-terminus is located extracellularly, whereas the C-terminus is found intracellularly [5]. These receptors are activated by a diverse array of ligands, from peptide and non-peptide neurotransmitters, to chemokines and nucleotides [6.7]. Consistent with their ability to recognize many antigens, GPCRs participate in a wide variety of physiological functions. GPCRs are also involved in a number of human diseases: this involvement is reflected in the fact that they are targets directly or indirectly of 50-60% of all known therapeutic agents to date.

When receptors activated by agonists, the conformation of the transmembrane  $\alpha$ -helices of GPCRs changes profoundly and previously masked G-protein-binding sites appear. This, in turn, promotes GDP-GTP exchange on the  $\alpha$ -subunit, which results in their activation. Consequently, G $\beta\gamma$ - and GTP-bound G $\alpha$  proteins stimulate effector molecules. The biochemical changes that are induced are highly dependent on the individual receptor-coupling specificity for each of the four families of mammalian G-protein subunits:  $\alpha_s$   $\alpha_i$   $\alpha_q$  or  $\alpha_{12/13}$  [8]. Receptors that are coupled to G $\alpha_s$  activate adenylyl cyclases, thereby increasing cyclic-AMP levels; whereas those that stimulate G $\alpha_i$ -family members inhibit adenylyl cyclases and reduce cAMP levels as well as

activating phospholipases and phosphodiesterases [8]. Each G protein has a unique biological function. The G<sub>s</sub> protein activates adenylyl cyclise, resulting in the accumulation of cyclic adenosine monophosphat (cAMP) in the cells. G<sub>i</sub> proteins are highly homologous members of the family of heterotrimeric signal-transducing guanine nucleotide-binding proteins (G proteins) involved in coupling numerous receptors for extracellular signalling molecules to the inhibition of adenylyl cyclase, the stimulation of phospholipase C, or the opening of potassium channels [9, 10, 11]. The activation of phospholipase C results in the hydrolysis of phosphatidylinositol bisphosphate (PIP<sub>2</sub>) and the generation of diacylglyceride (DAG), an activator of protein kinase C (PKC), and inositol(1,4,5)trisphosphate . [12]. The termination of GPCR signalling cascades after agonist activation is highly regulated and can occur through several mechanisms [8].

# 1.2.1.1. Toxins influencing G-protein coupled receptor pathways

There are two main subgroups of G proteins, the cholera-toxin-sensitive G proteins and the pertussis-toxin-sensitive ones. In the cholera-toxin-sensitive G proteins, the arginine residue present in the carboxy terminal of the  $\alpha$  subunit of  $G_s$  will be ADP-ribosylated, whereas the pertussis toxin ADP ribosylates the cysteine residue in the carboxy terminal of the  $\alpha$  subunit of  $G_i$  and  $G_o$  proteins while it inhibits their interaction with agonist-occupied receptors [13].

#### 1.2.2. The role of adhesion molecules

Leukocytes must engage several sequential adhesion pathways to leave the circulation. Initially, tethers are formed by adhesion receptors that are specialized to engage rapidly and with high tensile strength. The most important initiators of adhesion are the three

selectins expressed on leukocytes L-selectin, endothelial cells (P- and E-selectin), and activated platelets (P-selectin) [14]. Selectin-mediated bonds are too impermanent to arrest cells at the vessel wall. As the blood flow exerts pressure, adhesion bonds dissociate at the cell's upstream end and new bonds form downstream. This results in a rolling motion that is much slower than that of free-flowing cells. To stop rolling, cells must engage additional (secondary) receptor molecules, which belong to the integrin family. Integrins are obligate heterodimers containing two distinct chains, called the  $\alpha$ - and  $\beta$ -subunits. They are expressed on the most of the cells and are responsible for anchoring the cells to the extracellular matrix. Only a limited number of integrins are expressed in leukocytes, e.g. the  $\beta_2$ -integrin CD11b/CD18 [15, 16, 17].

# 1.2. Chemotaxis of leukocytes and G-protein-coupled signalling

Chemotaxis is the ability of cells to detect and move towards the source of a chemoattractant signal. Such directional migration is essential for a variety of cellular processes, including cell movement during development, immune responses and wound healing, in addition to the metastasis of tumor cells. In response to chemoattractants, leukocytes rapidly polarize in the direction of the signal, forming a pseudopod on the side exposed to the highest chemoattractant concentration, and a uropod or a posterior domain on the opposite side of the cell. These structures become the leading and trailing edges of the migrating cell [18, 19]. The formation of the leading edge results from localized chemoattractant-induced F-actin assembly, whereas the cell's side and posterior become enriched in assembled myosin II. The regulation of these cytoskeletal components during chemotaxis provides the driving and contracting forces required for cell motility [20]. Cells are able to detect and respond to chemoattractant concentrations that differ by as little as 2-5% between the front and the back of the cell. These

differences are then translated by the cell into a steep intracellular gradient of signalling components, leading to an asymmetric cellular response [21]. One of the first asymmetric responses to chemoattractant stimulation in leukocytes is the localized accumulation of phosphatidylinositol-3 kinases [PI(3,4,5)P<sub>3</sub>], the product of class I phosphatidylinositol-3 kinases (PI3Ks), at the site of the new leading edge [21, 22]. PI3K, which is made up of multiple isoforms, comprises a regulatory p85 subunit and a catalytic p110 subunit, of which in  $\alpha\beta$ T lymphocytes p110 $\alpha$  and p110 $\delta$  are expressed; both participate in the activity of the enzyme [23]. The three catalytic isoforms p110 $\alpha$ ,  $\beta$ ,  $\delta$  are classified as class IA PI3Ks; the only member of the class of IB PI3Ks is p110 $\gamma$ [24, 25]. The pharmacological inhibitors of PI3Ks are wortmannin [26] or LY294002 [27]. PI3K activation results in binding of p85 to adapters and in turn the enzyme to the plasma membrane and subsequently generates PI(3,4,5)P<sub>3</sub> and PI(3,4)P<sub>2</sub> by activating the catalytic subunit p110 [28]. This localized enrichment of [PI(3,4,5)P<sub>3</sub>] at the leading-edge membrane of the cell recruits and activates proteins that preferentially bind to PI(3,4,5)P<sub>3</sub>, including proteins that contain a plextrin-homology (PH) domain (e.g. protein kinase B [Akt/PKB], guanine nucleotide exchange factors [GEFs]) [29].

# 1.3. The role of $\gamma\delta$ T cells

The immune system has maintained two different types of T cell receptors (TCRs) throughout evolution,  $\alpha\beta$  T cells and  $\gamma\delta$  T cells.  $\gamma\delta$  T cells form an independent population of lymphocytes expressing the  $\gamma\delta$  TCR genes [30, 31, 32] and representing a complex system of developmentally and functionally differentiated subsets [33].  $\gamma\delta$  T cells are an ancient type of lymphocytes derived from haematopoetic stem cells; they share different characteristics with other immune cells such as the capacity to present antigens [34, 35] and the cytolitic effect [34, 36].  $\gamma\delta$  T cells participate in the

polarization of Th<sub>1</sub>- and Th<sub>2</sub>- immune responses by releasing various cytokines and helping  $\alpha\beta$  T lymphocytes mature [37] and B cells [38].  $\gamma\delta$  T cells, unlike  $\alpha\beta$  T lymphocytes, do not need MHC presentation by antigen presenting cells to recognize antigens. They are able to directly recognize natural phosphoantigens derived from plants [39], bacteria [40, 41, 42] protozoa [43], endogenous ligands from some tumors [47, 48, 49], and viruses [45, 46]. Early events in an immune response stimulate the production of cytokines; these direct the subsequent development of T cell subsets with discrete patterns of cytokine production. Cytokines secreted by human  $\gamma\delta$  T cells are able to activate both innate host defenses such as IFN $\gamma$ , or IL-2 [34, 50, 48, 35, 49, 51] and humoral host defenses with IL-4 and IL-10 [50, 52, 53].

# 1.3.1. γδ T cell subtypes: Circulating and resident γδ T cells

Two main subsets of  $\gamma\delta$  T cells are distinguished on the basis of their location. Resident  $\gamma\delta$  T cells are found in the skin, uterine and epithelium, whereas the circulating/systemic  $\gamma\delta$  T cells are found in the peripheral blood or lymphoid tissues [54, 55]. The majority of circulating human  $\gamma\delta$  T cells express a TCR comprising the variable segments  $\gamma9$   $\delta2$  (V $\gamma9V\delta2$  TCR). The functional characteristics of these cells are MHC-unrestricted cytolysis, production of Th1 type cytokines such as IFN- $\gamma$  and TNF- $\alpha$ , and expansion in a variety of bacterial or parasite infections [56, 57, 58, 34]. In the peripheral blood of most healthy individuals, T cells expressing the V $\delta2$  gene paired with one particular V $\gamma$  chain (V $\gamma9$ ) account for 50% to 90% of the  $\gamma\delta$  T cell population [34]. The major population of V $\gamma9V\delta2$  T peripheral  $\gamma\delta$  T cells is strongly activated by a variety of bacteria and parasites [59, 36, 35]. Sequencing the rhesus monkey V $\gamma9V\delta2$  TCR revealed a strong sequence homology to the human V $\gamma9V\delta2$  TCR with important

preserved sequence motifs [60] macaque and human  $V\gamma9V\delta2$  TCR share several conserved motifs [61]. Intestinal intraepithelial  $\gamma\delta$  T cells frequently express the V $\delta$ 1 gene, which can associate with different  $V\gamma$  elements [54].

#### 1.3.2. γδ T cell antigen recognition

 $\gamma\delta$  T cells have been shown to recognize classical MHC molecules (MHC I/II) as well as nonclassical MHC molecules, such as nonpolymorphic MHC-related CD1 protein and the closely related MHC class I b molecules T22 by TCR. MHC molecules are recognized as antigens, whereas the loaded peptides on the membrane do not play any role as ligands. In addition,  $\gamma\delta$  T cells can recognize non-MHC molecules, for examples, isopentenyl pyrophosphate, heat shock protein (Hsp), MICA/B, ULBP3, virus protein directly *via*  $\gamma\delta$  TCR or cooperatively through antigen specific receptor, such as NKG2D without antigen processing and presentation requirements [62, 63, 64, 65].

#### 1.3.2.1. Bacterial antigens

Early reports on the reactivity of  $\gamma\delta$  T cells suggested that heat-shock proteins were the main stimulatory fraction of mycobacteria [66, 67]. However, later studies have shown that stimulation of  $\gamma\delta$  T cells by M. tuberculosis appears to occur mainly through the recognition of the phosphorylated nucleotides TUBag1-4 [68].

#### 1.3.2.2. Amino-bisphosphonates

The antitumor effect of amino-bisphonates mainly results from their capacity to activate  $\gamma\delta$  T cells, as they have been found to induce V $\gamma$ 9V $\delta$ 2 T cell proliferation in vitro [69] and in vivo [70] and to trigger the V $\gamma$ 9V $\delta$ 2 TCR-mediated lysis of tumor lines [69].

 $V\gamma 9V\delta 2$  T cells are stimulated by cognate recognition of human cells pulsed with aminobisphosphates, e.g., pamidronate or zoledronate, probably because their pharmacological action leads them to accumulate of mevalonate metabolites [69, 71]. Amino-bisphosphonates induce  $V\gamma 9V\delta 2$  T cell proliferation in vitro and in vivo, and trigger the  $V\gamma 9V\delta 2$  TCR-mediated lysis of many tumor cell lines and reduce the survival of autologous myeloma cells [36].

### 1.3.2.3. Phosphoantigens

The most potent natural phosphoantigen is (E)-4hydroxy-3-methyl-but-enyl pyrophosphate, a metabolite of the nonmevalonate pathway of isoprenoidsynthesis, which is found in plants, bacteria and protozoa [43]. Less potent phosphoantigens are ubiquitous metabolites of isoprenoid synthesis, such as isopentenyl pyrophosphate (IPP) [72]. IPP-mediated  $\gamma\delta$  T cell activation does not require accessory cells as does the activation of the aminobisphosphonate pamidronate [73].

#### 1.3.3. Receptors expressed on the surface of $\gamma\delta$ T cells

#### 1.3.3.1. TCR receptor

# 1.3.3.1.1. The classical TCR signalling pathway in T lymphocytes

The physiological ligand for T cell activation is a foreign peptide bound to a major histocompatibility complex (MHC)-encoded molecules presented on the surface of professional antigen presenting cells (APCs) such as dendritic cells (DC) [15]. Biochemical signals initiated by the TCR receptor make up  $\alpha/\beta$  subunits and signal-transducing subunits  $\epsilon$ ,  $\gamma$ ,  $\delta$ , and  $\zeta$ ; these determine the specifity of the TCR receptor

activation and events initiated by other membrane proteins such as MHC receptors CD4/CD8; costimulators such as CD2, CD28 or integrins modulate the intracellular signaling thresholds required to initiate a T cell immune reaction [74]. The activation of tyrosine kinases initiates a cascade of signalling pathways in T cells resulting in the activation of small GTPases-signalling networks and the regulation of inositol phospholipid metabolism: this regulation is responsible for controlling intracellular calcium (Ca<sup>2+</sup>) and the activity of diverse serin/threonine kinases, including members of the PKC family and phosphatidyl inositol-3 kinase (PI3K)-controlled serin kinases [29]. Unlike  $\alpha\beta$  T,  $\gamma\delta$  T cells are endowed with the capacity to interact with low-molecular-mass-phosphate-containing nonprocessed antigens, such as pyrophosphomonoesters [72] and alkylamines [39]. Recognition of these antigenic compounds is mediated by TCR but does not require presentation by conventional MHC molecules [75, 76].

# 1.3.3.1.2. NKG2D receptor

NKG2D belongs to a sub-family of C-type lectin-like receptors, which have lost the ability to bind both calcium and carbohydrate. In addition to TCR ligation, the activation of  $V\gamma9V\delta2$  T cells is modulated by a range of inhibitory and activating NK-cell receptors [77, 78].

NKG2D ligation is known to provide a powerful co-stimulus for the antigen-mediated activation of both CD8<sup>+</sup> and V $\gamma$ 9V $\delta$ 2 and intraepithelial V $\gamma$ 1V $\delta$ 1 T cells; however, the signalling pathways leading to the activation of these cell types are poorly understood. In V $\gamma$ 9V $\delta$ 2 T cells, activation through NKG2D receptors happens independently of the antigen, as also happens in the NKG2D-mediated activation of NK cells [36]. Several lines of evidence show that both  $\gamma\delta$  T cell populations have antitumor reactivity [79, 48, 80], which they perform with their production of perforin [54, 50, 80] and granzymes

[34, 81]. Certain tumors and lymphomas activate  $\gamma\delta$  T cells via endogenous ligands in  $\gamma\delta$  TCR-dependent fashion [75, 48]; other ligands of tumors or transformated cells bind the NK cell receptor NKG2D on the surface of  $\gamma\delta$  T cells [79].

#### 1.3.3.2. The role of $\gamma\delta$ T cells as effector cells: the role of chemokines and cytokines

Early during the course of infection, cytokines produced by  $\gamma\delta$  T cells may help eliminate pathogens and contribute to effective cellular immunity against intracellular bacteria and parasites. V $\delta$ 2 T cells display a range of innate effector functions, including the rapid secretion of chemokines MIP- $\alpha$ , MIP- $\beta$ , or RANTES [82, 83] and cytokines TNF- $\alpha$  and INF- $\gamma$  [84, 85].

# 1.3.3.2.1. Response to chemokines

Chemokines or chemotactic cytokines are a family of small (7-15 kDa), inducible proteins, which can be classified into four families: C-X-C chemokines or  $\alpha$ -chemokines, CC or  $\beta$ -chemokines C or  $\gamma$ -chemokines, and CX3C or  $\delta$ -chemokines. Chemokines are implicated in multiple pathological processes, such as allergic disorders, autoimmune diseases and ischemia [12]. In addition to chemotaxis, chemokines have been shown to modulate a number of biological responses in leukocytes, including cellular activation, adhesion, enzyme release, degranulation, tumor cell lysis, and intracellular microbe death [86].

MCP-1/CCL2 [87], RANTES/CCL5, MIP-1 $\alpha$ /CCL3 and MIP-1 $\beta$ /CCL4 were found to chemically attract  $\alpha\beta$  T lymphocytes and circulating  $\gamma\delta$  T cells, whereas neither  $\alpha\beta$  nor  $\gamma\delta$  showed response to IL-8/ CXCL8 or IFN-inducible protein 10 (IP-10/CXCL10) [88, 89]. The corresponding chemokine receptors - seven-transmembrane-spanning G

protein-coupled receptors - share structural features, and some of which are involved in intrathymic [90, 91] and memory T cell development, in orchestrating T and B cell interactions and in differentiating effector T cells [87, 92].

# 1.3.3.3. The role of $\gamma\delta$ T cells as effector cells: the way of killing, cytolytic activity

 $\gamma\delta$  T cells not only inhibit the early stages of tumor development but also limit the progression of neoplastic cells; combined, these functions suggest there are two or more different antitumor mechanisms in these cells. In summary, mouse studies have elucidated three different pathways by which  $\gamma\delta$  T cells may provide anticancer activities; direct killing of transformed cells, early IFN- $\gamma$  production and critical immunoregulatory function.  $\gamma\delta$  T cell lines have been shown to efficiently kill extraand intracellular *M. tuberculosis* bacilli through the release of the cytotoxic molecule granulysin, although granulysin kills both extra- and intracellular *M. tuberculosis*, eliminating the latter requires perforin [59]. Activated  $\gamma\delta$  T cells are shown to lyse target cells also by producting granzymes and perforin [36].

#### 1.3.3.3.1. $\gamma\delta$ T cells in the tumor microenvironment

The broad antigen-specific recognition of hematopoietic tumors by  $V\gamma9V\delta2$  T lymphocytes results in cytotoxic activity, inducing Th1 cytokine production and proliferation [77, 93]. Like NK cells,  $V\gamma9V\delta2$  T lymphocytes exert a cytotoxic activity that is controlled at the effector level by the expression of killer Ig-like receptors (KIR) [94] which interact with major histocompatibility complex class I (MHCI) molecules at the surface of the target [78, 95]. Tumor cell lines which are sensitive to this NK-like

cytolytic activity, such as the chronic myelogenous leukaemia cell line K562, often lack the ability to express MHC-class I molecules [96, 97, 98].

Some tumors and lymphomas, such as the B cell line Daudi, express endogenous ligands, which in turn activate  $\gamma\delta$  T cells in a V $\gamma$ 9V $\delta$ 2 T cell-dependent fashion [75]. Moreover, both the percentage and the absolute number of circulating  $\gamma\delta$  T cells are known to be significantly reduced in patients with cutaneous primary melanoma, thus suggesting a specific deficit for this lymphocyte population. The reduction of  $\gamma\delta$  T cells correlated with the decrease of the V $\delta$ 2 T cell subset, which is involved in the reactivity toward microbial antigens and tumor cell antigens. The in vitro expansion of  $\gamma\delta$  T cells, particularly the V $\delta$ 2 T subset isolated from peripheral blood of patients with cutaneous primary melanoma, was significantly reduced; in summary, not only the proliferative capacity but the percentage of  $\gamma\delta$  T cells producing either TNF- $\alpha$  or IFN- $\gamma$  is significantly reduced in melanoma patients [99].

#### 1.3.3.3.1. γδ T cells in immunosuppression and cancer immunotherapy

Human  $\gamma\delta$  T cells are prime effector cell candidates for immune therapy against malignancy, since there is a growing evidence that V $\gamma$ 9V $\delta$ 2 T cells have cytotoxic antitumor activitiy against a large range of tumor types [100, 101]. Activated V $\gamma$ 9V $\delta$ 2 T cells provide alternative mechanisms for tumor-targeted recognition and killing, increasing the possibility that treatment with other modalities may enhance antitumor effects. Pre-treating tumor cells with low concentrations of chemotherapeutic agents may lead to their isoprenoid-mediated recognition and perforin following TCR-dependent rapid killing by V $\gamma$ 9V $\delta$ 2 T cells [102].

#### 1.4. Characterization of histamine and fMLP as chemoattractant factors

Since circulating  $\gamma\delta$  T cells have a pivotal role in host defense against invading pathogens, bacteria, and viruses as well as in the tumor microenvironment, and the existence of G-protein-coupled receptors on the surface of these cells has been not thoroughly examined, in this work we investigated the signal transduction pathways for histamine and fMLP in circulating  $\gamma\delta$  T cells. All the chemoattractants mentioned above are ligands of GPCRs and influence the immune response of leukocytes.

#### 1.4.3. Histamine

Histamine ( $\beta$ -imidazolylethylamine) is an endogenous mediator involved in various responses of immune and non-immune cells; it is stored in the granules of tissue mast cells and blood basophlis and is released upon activation by IgE cross-linking [103]. The half-life of histamine in plasma is < 20 minutes, whereas the suppression of T-cells and phagocytes lasted  $\geq$  4 hours after subcutane histamine injection [104].

Beyond its physiological functions, histamine was described as an autocrine and paracrine/exogene growth factor for malignant melanomas and leukemic cells. Histamine and histidine decarboxylase (HDC) [105], the only enzyme that catalyzes histamine production, has been proven to be present in elevated concentrations in proliferating tissues, including tumor cells [106]. In the case of the myeloproliferative disease of chronic myeloid leukaemia (CML), this gene was the first known leukaemia-specific oncogene reported to be involved in the regulation of histamine production in neoplastic cells [107]. Malignant haematopoetic diseases characterized by hyperhistaminism in peripheral blood respond favorably to INF- $\alpha$ , an NK- and T-cell-activating cytokine [107, 108]. Histamine also interacts directly with H<sub>2</sub> receptors (H<sub>2</sub>R) on T-cells to inhibit T-cell proliferation and cytokine production [109]; however, the

inhibition of T-cell function may not be expressed if histamine is combined with IL-2 [110, 111].

# 1.4.3.1. Histamine receptor subtypes

Histamine receptors are expressed on the surface of human monocyte-derived dendritic cells [112], eosinophils [113], mast cells [114], T cells, keratinocytes [115], and NK cells [116]. The effects of the numerous physiological activities of histamine are mediated by several histamine receptors. To date, four subtypes (H<sub>1</sub>R, H<sub>2</sub>R, H<sub>3</sub>R and H<sub>4</sub>R) of histamine receptors have been identified and cloned. All belong to the seven transmembrane domain G-protein- coupled receptor family, and coupled G proteins, and subsequently activated intracellular signals have been characterized [117]. In vitro studies made in human peripheral blood lymphocytes have demonstrate that histamine and the histamine receptor 2 (HR<sub>2</sub>) agonists, impromidine and dimaprit, were effective in suppressing cell proliferation induced by low rather than high doses of PHA [118, 119, 120].

#### 1.4.2. Formyl-methyl-leucyl-phosphate (fMLP)

N-formyl-methionine-leucine-phenylalanine (fMLP) is the prototype for N-formylated bacterial peptides and also a chemoattracting tripeptid. Its receptor in humans was pharmacologically defined as a high affinity binding site on the surface of neutrophils [121]. fMLP stimulates glucose uptake in CHO cells expressing chemoattractant-receptors which is due to the translocation of GLUT1 mediated by the  $G\alpha$  subunit through unknown pathways [122]. The fMLP-stimulated 2-DG uptake of human peripheral blood cells is inhibited by the PKC inhibitor calphostin C and tyrosine kinase inhibitor genistein [123]. In phagocytes, the chemoattractant fMLP activates the FPR to

stimulate diverse biological processes including chemotaxis, transcriptional activation, and actin reorganization [124].

# 1.4.2.1. The formyl peptide receptor (FPR)

First reports of the cloning and functional expression of human fMLP receptor, published in the early 1990s [125, 126], found the fMLPR to be constitutively active [127]. The formyl peptide receptor (FPR) genes of non human primates are 95-99% identical of their human ortholog [128]. Expression of the receptor has been described in phagocytic leukocytes, monocytes and neutrophils [129, 130], and is observed in immature dendritic cells, microglial platelets, spleen and bone marrow, nonhaematopoetic cells and tissues such as hepatocytes, fibroblasts, astrocytes, neurons of autonomic nervous system, lung and lung carcinoma cells, thyroid, adrenals, heart, the tunica media of coronary arteries, uterus, ovary, placenta, kidney, stomach and colon [131]. The first ligand described for the FPR was the prototypical formylated peptide fMLP derived from E. coli, which binds and activate FPR with high affinity [131]. Because the formylation of peptides also occurs in mitochondria; the secondary release of formylated peptides secondary to cell death, might allow for the attraction of phagocytic leukocytes through FPR. Indeed disrupted mitochondria purified bovine mitochondrial N-formylatmethionyl-proteins, but not non-formylated mitochondrial proteins; the latter are chemoattractants for neutrophils in vitro [132]. The fMLP receptor is coupled to G<sub>i</sub> proteins, as indicated by the total loss of cell response to their respective agonists upon exposure to pertussis toxin [121, 133, 134]. More specifically, the fMLP receptor is coupled to  $G_{i\alpha 1}$ ,  $G_{i\alpha 2}$  [135, 136, 137],  $G_{i\alpha 3}$  but coupling to  $G_o$ ,  $G_z$ ,  $G_{\alpha 16}$  [138] and in xenopus oocytes to  $G_q$  [139] has also been described [140, 141]. After ligation of the FPR, the heterodimeric G protein is activated, which dissociates into a and  $\beta\gamma$  subunits; these subunits then activate the phospholipase C (PLC). The hydrolysis of phosphatidylinositol 4,5-bisphoshate (PIP<sub>2</sub>) by PLC generates IP<sub>3</sub> [142], which releases intracellular calcium ([Ca<sup>2+</sup>]<sub>i</sub>) from endoplasmatic reticulum stores, and diacylglycerol (DAG); DAG, which activates protein kinase C (PKC) isofoms. The phosphoinositide 3-kinase (PI3K)-Akt/PKB pathway is also activated, in particular PI3K $\gamma$ , which is the isoform coupled to most chemoattractant receptors [143, 141]. Since fMLP such as IL-8 induce Ca<sup>2+</sup> efflux and leukocyte chemotaxis, which are sensitive to PTX treatment, the fMLP receptor may use the same signal transduction pathways as the IL-8 receptors [144]. Co-transfection assays have been demonstrated that the fMLP receptor can specifically activate PLC  $\beta$ 2 but not PLC  $\beta$ 3 in a PTX-sensitive manner; thus, the interaction of the receptor with the G protein causes the release of the G $\beta\gamma$  subunits, which in turn specifically activate PLC  $\beta$ 2 [138].

#### 1.5. The aim of the study

The biogenic amine, histamine, has been described as playing a pathophysiological regulatory role in the cellular processes of various immune cells. This work analyzes the biological activity of histamine-influencing human peripheral blood  $\gamma\delta$  T lymphocytes, which are critical members of the immunological tumor surveillance. The bacterial cell-wall-derived peptide N-formyl-Met-Leu-Phe (fMLP) is a well characterized chemotaxin for phagocytes such as neutrophils, monocytes and dendritic cells. Both fMLP and histamine receptors belong to the family of G-protein-coupled seven-transmembrane receptors (GPCRs). Receptor ligand interactions are known to activate intracellular signal transduction pathways in neutrophils and monocytes, leading to directed motility, phagocytosis, superoxide anion generation, and degranulation responses culminating in death and elimination of invading microorganisms.

In this regard, the present study addresses the biological questions below:

- 1) What is the role of fMLP and histamine in the chemtaxis of human circulating  $\gamma\delta$  T lymphocytes?
- 2) What is the role of histamine in the cytolytic activity of human circulating  $\gamma\delta$  T lymphocytes?

# 2. Materials and methods

# 2.1. Materials

# 2.1.1. Biological Material

#### 2.1.1.1. Cell lines

The human metastatic melanoma line A2058, from a brain metastasis in a 43-year-old man [145], the human Burkitt's non-Hodgkin lymphoma cell line Raji [146], the T-lymphoblastic lymphoma cell line Jurkat, established from the peripheral blood of a 14-year-old boy with acute lymphoblastic leukemia (ALL) at first relapse in 1976 [147] and the erythroleukaemia cell line K562 originating from patients with chronic myeloid leukaemia and blast crisis [148] were maintained at 37°C in a 5% CO<sub>2</sub> incubator in RPMI 1640 supplemented with 10% fetal bovine serum, 10 U/ml penicillin, 10 U/ml streptomycin, and 1 mM L-glutamine [149].

#### 2.1.1.2. Human peripheral blood $\gamma\delta$ T lymphocytes

The use of human cells was approved by the Research Ethics Board of the Friedrich Schiller University of Jena. Human peripheral blood  $\gamma\delta$  T lymphocytes used in the work were isolated from the buffy coats of healthy volunteers produced by the Institute of Transfusion Medicine of Friedrich Schiller University of Jena.

#### 2.1.2. Cell culture medium

Isolated human peripheral blood  $\gamma\delta$  T lymphocytes were cultured in RPMI medium supplemented with 10% fetal calf serum (FCS), 10 U/ml penicillin, 10 U/ml streptomycin, and 1 mM L-glutamine.

# 2.1.3. Reagents

Albumin from bovine serum (BSA) Sigma-Aldrich, Taufkirchen

Cholera toxin (Ctx.) Sigma-Aldrich, Taufkirchen

DNA Molecular Weight Marker Roche Diagnostics, Mannheim

Dimethylsulfoxid (DMSO) Sigma-Aldrich, Taufkirchen

FastStart Taq DNA Polymerase Kit Roche Diagnostics, Mannheim

Fetal bovine serum, 10 U/ml Promocell, Heidelberg

N-formyl-Met-Leu-Phe (fMLP) Sigma-Aldrich, Taufkirchen,

Fura/2AM MERK, Darmstadt

L-glutamine Promocell, Heidelberg

Hank's balanced salt solution (Hank's BSS)

Promocell, Heidelberg

High Pure RNA Kit Roche Diagnostics, Mannheim

Histamine Sigma-Aldrich, Taufkirchen

HTMT dimaleate TOCRIS (BIOZOL), Eiching

rHu interleukine-2 (Proleukine) Chiron, Ratingen

imetit TOCRIS (BIOZOL), Eiching

MicroScint-40 cocktail PerkinElmer, Jügesheim

 $Na_2^{51}CrO_4$  (100 $\mu$ Ci) Amersham, Freiburg

NBD-phallacidin Invitrogen GmbH, Karlsruhe

Phosphate buffered saline (PBS)

Promocell, Heidelberg

Penicillin (10 U/ml) Promocell, Heidelberg

Phaseolus vulgaris phytohemagglutinin (PHA) Sigma-Aldrich, Taufkirchen

Pertusis toxin Sigma-Aldrich, Taufkirchen

Primers Invitrogen GmbH, Karlsruhe

RANTES Promokine, Heidelberg

RPMI 1640 culture medium

SeaKem LE agarose

Streptomycin, and 1 mM

Thioperamide malate salt (T123)

Triton-X-100

Wortmannin

Promocell, Heidelberg

Cambrex, Tuafkirchen

Promocell, Heidelberg

Sigma-Aldrich, Taufkirchen

Sigma-Aldrich, Taufkirchen

Sigma-Aldrich, Taufkirchen

#### 2.1.4. Antibodies

# 2.1.4.1. Flow cytometry analysis antibodies

anti-TCR  $\gamma\delta$  hapten-antibody anti-hapten microbeads-FITC antibody goat anti-mouse-FITC (sec. AB) gouse anti-human cAMP antibody mouse anti-human CD88-FITC antibody mouse anti-human fMLPR-PE antibody mouse anti-human V $\gamma$ 9 TCR antibody mouse anti-human V $\delta$ 2 TCR antibody mouse anti-human-TCR- $\gamma/\delta$ -1 antibody mouse anti-human-TCR- $\gamma/\delta$ -1 antibody

Miltenyi, Bergisch Gladbach
AL-Immunotools, Friesoythe
Abcam, Cambridge
Acris Antibodies, Hiddenhausen
BD Biosciences, Heidelberg
BD Biosciences, Heidelberg
BD Biosciences, Heidelberg
BD Biosciences, Heidelberg

BD Biosciences, Heidelberg

Miltenyi, Bergisch Gladbach

# 2.1.4.2. Western blot analysis antibodies

goat anti-human actin [(I-19:sc-1616)] goat anti-human H<sub>1</sub>R (A20) antibody goat anti-human H<sub>3</sub>R (P-15) antibody

Santa Cruz Biotechnology, Heidelberg Santa Cruz Biotechnology, Heidelberg Santa Cruz Biotechnology, Heidelberg goat anti-human H<sub>4</sub>R (IL-15) antibody Santa Cruz Biotechnology, Heidelberg

rabbit anti-human H<sub>2</sub>R (H-70) antibody Santa Cruz Biotechnology, Heidelberg

2.1.5. Consumables

Cell culture flasks 75 cm<sup>2</sup> Greiner, Frickenhausen

Cell culture plates (12-, 24-, 96-well) CellStar,

LS-Separation columns Miltenyi, Bergisch Gladbach

Minifac tubes Costar, Jena

Pipettes (5 ml, 10 ml, 25 ml) Greiner, Frickenhausen

Pipette tips (10 μl, 100 μl and 1000 μl) Greiner, Frickenhausen

Plastic tubes (500 µl, 1000 µl, 2000 µl) Eppendorf, Hamburg

Syringes 60 ml Braun Braun, Melsungen

Tubes (15ml, 50 ml) Greiner, Frickenhausen

2.1.6. Buffers and solutions

Ammonium-chloride solution: 41.5 g NH<sub>4</sub>Cl, 0.5 g KHCO<sub>3</sub>, 0.179 g EDTA, 500 ml

 $H_2O_{(dest.)}$ 

PBS-BSA-EDTA solution: PBS pH 7.2, 0.5% BSA, 2 mM EDTA

EDTA solution: 0.5M, pH8.0

Ethidium bromide: 10 mg/ml

TBE buffer (5x), 1000 ml:

54 g Tris-base, 27.5 g boric acid, 20 ml 0.5 M EDTA (pH 8.0)

*NET-G buffer (1x)*:

Tris-HCl pH 8.0 10 mM, NaCl 150 mM, EDTA 5 mM, Tween 20 0.05 % (v/v),

Gelatine 0.05 % (v/v)

Running buffer (4x):

Tris-base, SDS 10% (v/v)

Stacking buffer (4x):

Tris-HCl pH 6.8 0.5 M, SDS 10 % (v/v)

Running gel (10%):

H<sub>2</sub>O<sub>(dest.)</sub> 6.0 ml, Running buffer (4x) 3.6 ml, Acrylamide Stock 4.8 ml

Stacking gel (5%):

H<sub>2</sub>O<sub>(dest.)</sub> 2.8 ml, Stacking buffer (4x)

1.2 ml, Acrylamide Stock 0.8 ml

*Protein electrophoresis buffer* (10x):

Tris-HCl pH 8.3 250 mM, Glycin 2 M, SDS 1 % (v/v)

*Transfer buffer*:

Tris 25 mM, Glycin 192 mM, Methanol 20 % (v/v)

Stripping buffer:

Tris-HCl pH 6.7 62.5 mM, 2-mercaptoethanol 100 mM, SDS 2% (v/v)

2.1.7. Equipment

Autoclave Varioklav, Freiburg

Balance Sartorius, Göttingen

Centrifuge 5804R Eppendorf, Hamburg

Centrifuge RC-3B Sorvall Heraeus, Langenselbold

Digital fluorescence microscope unit Atofluor Zeiss, Jena

Electrophoresis gel chambers Gibco BRL Life Technologies Inc., USA

Flow cytometer FACSCalibur BD Biosciences, Heidelberg

Freezer -20°C Siemens, Berlin

Freezer Hera -80°C Heraeus, Langenselbold

Gamma counter TopcountTM Packard Instruments,

Heating block HB-130 Unitek, Freiburg

Incubator CO<sub>2</sub> Sanyo, Sakata, Japan

Laminar-Air HB 2472 workbench Heraeus, Langenselbold

Microplate Reader Emax Molecular Devices, Freiburg

Microscope Telaval 3 Carl Zeiss, Jena

Pipettes Easypet Eppendorf, Hamburg

Pipettes (20ml, 200ml, 1000ml) Gilson, Hamburg

Power supply, PowerPac 300 Bio-Rad, USA

Thermomixer comfort Eppendorf, Hamburg

Vortex Bender&Hobein, Freiburg

#### 2.2. Methods

# 2.2.1. Separation of blood leukocytes using the Ficoll-Paque gradient

A 50 ml buffy coat was diluted 1:1 with PBS (v/v). Carefully, 25 ml of the cell suspension was layered over 15 ml Ficoll-Paque PLUS solution in 50 ml conical tubes and centrifuged at 1800 rpm for 30 min at 20°C, without braking. Differential migration during centrifugation results in the formation of layers containing different cell types (Figure 2.2.1). The bottom layer contains aggregated erythrocytes. The layer above the erythrocyte layer contains mostly granulocytes. Because of their low density, the lymphocytes are found at the interface between the plasma and the Ficoll-Paque gradient with other slowly sedimenting particles (platelets and monocytes). The interphase cells (lymphocytes, monocytes and thrombocytes) were transferred by

Pasteur pipette to a new conical tube and subjected to short washing steps, with a balanced salt solution to remove platelets, Ficoll-Paque and plasma.

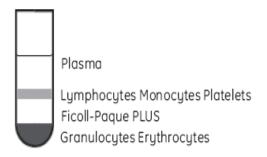


Figure 2.2.1. Differential migration during centrifugation results in the formation of layers containing different cell types.

# 2.2.1.1. Preparation of γδ T cells

The use of human cells was approved by the Research Ethics Board of the Friedrich Schiller University of Jena. Peripheral blood mononuclear cells (PBMCs) were isolated using the Ficoll separation protocol [33]. Briefly, a density gradient centrifugation of buffy coats was performed. The leukocyte-containing pellet was resuspended in phosphate-buffered saline (PBS) pH 7.2, supplemented with 0.5 % BSA and 2 mM EDTA, and the cells were labelled with an anti-TCR  $\gamma\delta$  hapten-antibody and anti-hapten micro-beads-FITC antibody. Labelled cells were separated with magnetic separation columns. Positive selected  $\gamma\delta$  T cells were cultured for 7-10 days in the presence of phaseolus vulgaris phytohemagglutinin (PHA) (2  $\mu$ g/ml) for 3 days and IL-2 (100 IU/ml) until day 7 [80, 79, 99].

# 2.2.2. Flow cytometer

The flow cytometer enables a cell population to be acquired on the basis of its cell size and granularity. Additionally, surface molecules, or the purity or vitality of a cell population can also be examined by fluorescence. Purified cells and cells being cultured for 7-10 days were incubated with FITC-conjugated mouse anti-human-TCR- $\gamma$ / $\delta$ -1 (1:200, BD Pharmingen) monoclonal antibodies for 30 min at 4°C. Isotype matched antibodies (IgG1 control, Immunotools) were used as negative control. Analysis was done with a FACSCalibur flow cytometer (BD Biosciences). All experiments were done with a purity of 96-99% of  $\gamma\delta$ TCR positive T cell population. The cell viability was measured with the trypan blue exclusion test of cell viability and we accepted populations with a cell viability of min. 95%.

2.2.3. mRNA isolation, reverse transcription and polymerase chain reaction (RT-PCR) analysis

mRNA was isolated from  $1x10^6$  human peripheral blood  $\gamma\delta$  T cells using a High Pure RNA isolation kit. A Fast Start Taq cDNA Polymerase Kit and a Fast Start Taq DNA Polymerase kit were used to obtain cDNA and PCR products. The primers were designed to recognize sequences specific for each target cDNA:

- H<sub>1</sub>R, (403 bp): sense 5'-CATTCTGGGGGCCTGGTTTCTCT-3'; antisense 5'-CTTGGGGGTTTTGGGATGGTGACT-3'
- H<sub>2</sub>R, (497 bp): sense 5'- CCCGGCTCCGCAACCTGA-3' antisense 5'-CTGATCCCGGGCGACCTTGA-3'
- H<sub>3</sub>R, (589 bp): sense 5'-CAGCTACGACCGCTTCTTGTC-3' antisense 5'-GGACCCTTCTTTGAGTGAGC-3'

H<sub>4</sub>R, (396 bp): sense 5'-GGTACATCCTTGCCATCACATCAT-3'

antisense 5'-ACTTGGCTAATCTCCTGGCTCTAA-3'

fMLPR (bp): sense 5' TTC CGG ATG ACA CAC ACA GT

antisense 5' TCC GAA CAA GTT GAT GTC CA

The phases of the PCR process were:

The initial phase: the denaturation of the DNA-double strand at 95°C for 3 minutes

Main cycles: the amplification of the DNA with variable number of cycles (in three

steps):

a. denaturation of the DNA-double strand (94°C, 1 minute)

b. hybridization of the specific primers to the single-strand DNA (58 °C, 60 °C)

c. the Taq-polymerase chain reaction (amplification) (72 °C, 1 minute)

The final phase: Taq-polymerase chain reaction (End-polymerization) (8 minutes)

2.2.4. Mobilization of Ca<sup>2+</sup> transients

Intracellular free Ca<sup>2+</sup> was measured in Fura-2-labeled γδ T cells using the digital

fluorescence microscope unit Attofluor (Zeiss, Oberkochen, Germany). The purified

circulating γδ T cells were resuspended in Hank's 1% BSA solution and were loaded

with 4 µM Fura-2/AM. The suspension was then incubated in dark, 37°C for 30

minutes. After the incubation cells were washed two times and were resuspended in

buffer and measured in dark. After the stimulation the fluorescence was measured and

the ratio was calculated from the absorption between the wavelength 340 nm and 380

nm [150].

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#### 2.2.5. Filamentous (f) actin measurements

Probes of stimulated  $\gamma\delta$  T cells ( $10^6$  /ml; 50ul/probe) were fixed in a 7.4 % formaldehyde buffer and mixed with the staining mixture containing 7.4 % formaldehyde, 0.33  $\mu$ M NBD-phallacidin and 1 mg/ml lysophosphatidylcholine. The fluorescence intensity was measured by flow cytometry. The relative f-actin content was compared to unstimulated controls [151].

# 2.2.6. Migration assay

The chemotaxis of isolated human peripheral blood  $\gamma\delta$  T cells was performed in 48-well-microtechnic chambers from Neuro Probe (Gaithersburg, US). Wells in the bottom of the chamber were filled with 29 µl medium containing the indicated concentration of stimulus. Over this filled chamber, a polycarbonate membrane (thickness 10 µm, diameter of the pores 8 µm) and a gasket made of silicone was fixed. The device was screwed on the top of the gasket and the cells were added in 29 µl/well (resuspended in a concentration of  $1x10^5$ /ml) in the upper wells of the chamber. For the assay, the chamber was incubated for 240 minutes at 37°C. The non-migrating cells from the wells of the upper chamber were removed after the incubation period. The filter and gasket were then removed and the cells from the bottom chamber were collected, fixed in formaldehyde (3.7 %) and counted by flow cytometry. A chemotactic index (CI) was calculated as the ratio between stimulated and random migration.

#### 2.2.7. In vitro cytotoxicity assay

Cytotoxicity was determined with a standard  $^{51}$ Cr release assay. Target cells were labeled at  $37^{\circ}$ C for 1 h with  $100 \,\mu$ Ci  $Na_2^{51}$ CrO<sub>4</sub>. Cells were washed and resuspended at

a cell density of 1×10<sup>6</sup> cells/ml in RPMI 1640 culture medium supplemented with 2 % fetal calf serum. Effector and target cells at different ratios (10:1; 5:1 and 2.5:1) were placed into individual wells of 96-well plates in a total volume of 200 μl at 37°C for 4 h. After incubation, 100 μl culture supernatant was collected from each well, mixed with a MicroScint-40 cocktail and analyzed with a gamma counter (TopcountTM, Packard Instruments). To obtain the value of total lysis, target cells were incubated with 2 % Triton-X. The percentage of specific lysis was calculated using the following formula:

#### 2.2.8. Measurement of cAMP levels

 $\gamma\delta$  T cells (1 x 10<sup>6</sup>/ml) were fixed and made permeable before the intracellular staining was performed. The amount of intracellular cAMP in the  $\gamma\delta$  T cell preparation was determined by flow cytometry [152].

#### 2.2.9. Western blot analysis

# 2.2.9.1. Lysis of the cells

Equal numbers of cells were washed and resuspended in PBS in 1.5 ml Eppendorf tubes and stimulated at 37°C. Supernatants were removed and cell pellets were transferred to ice and resuspended in an appropriate volume of lysis buffer containing 0.5 % of the detergent NP-40. Cell membranes were dissolved, while the cell cores and remaining cell compartments remained intact and were separated by centrifugation. The different protease inhibitors were added to the lysis buffer directly before use.

Lysis buffer: 20 mM Tris-HCl pH 8.0; 137 mM NaCl; 2.7 mM KCl; 1 mM MgCl<sub>2</sub>; 1 mM CaCl<sub>2</sub>; 0.5% NP-40; 0.5 mg/ml BSA; 1 mM Na<sub>3</sub>VO<sub>4</sub>; 1 mM PMSF; 21 μg/ml Aprotinin; 12 μg/ml AEBSF; 5 μg/ml Leupeptin; 5 μg/ml Pepstatin A.

After 10 min on ice, the lysates were centrifuged (15 min, 14000 rpm, 4 °C) and the NP-40 soluble fraction containing membranes and cytosolic proteins were collected and transferred into 1.5 ml eppendorf tubes.

## 2.2.9.2. SDS-Polyacrylamide-Gel electrophoresis (SDS-PAGE)

After cells were lysed, the NP40-soluble fraction contained in the supernatant was removed and diluted with 5x Laemmli sample buffer [153] and heated to 95°C for 5 minutes. Samples (20 μg protein/ lane) were electrophoresed in 10% SDS polyacrylamide gels (80 mA). Proteins separated by SDS-PAGE were transferred from the gel to PVDF membranes (HybondTM ECL<sup>TM</sup>, Amersham Biosciences) by electrophoresis (Western blotting) 50V for 90 min. The primary antibody was diluted 1:2000 with NET-G buffer and incubated 12 hr at 4°C. After being washed with NET-G buffer, the membrane was incubated with the secondary antibody (1:10000 in NET-G) for 1 hour at room temperature. After being washed, the protein of interest was detected by enhanced chemical luminescence (ECL) utilizing the reaction of Luminol and H<sub>2</sub>O<sub>2</sub>. For this purpose, the membrane was incubated with the ECL (Perbio, Bonn, Germany) reagents for 1 min and the chemiluminescence signal was visualized with the Bio Imaging System MF-ChemiBIS 3.2 (Biostep, Germany).

To reprobe the membrane with different antibodies, the previously applied antibodies first had to be removed. For this purpose, the membrane was incubated with 100-200 ml stripping buffer for 45 min at 56°C. The membrane was then washed several times with

NET-G (1x), until no residual 2-mercaptoethanol could be detected. The blot was again blocked in NET-G (10x) for 1 h, and a new primary antibody applied as described.

# 2.2.10. Statistical analysis

Significant values (p<0.05) were determined using the nonparametric two-tailed Student's t-test.

### 3. Results

# 3.1. Histamine and $\gamma\delta$ T cells: the involvement of pertussis-toxin- and choleratoxin-sensitive functions of isolated human peripheral blood $\gamma\delta$ T lymphocytes

### 3.1.1. γδ T cells expressed histamine receptors: H<sub>1</sub>R<sub>1</sub> H<sub>2</sub>R, and H<sub>4</sub>R mRNAs

In order to characterize the biological activity of histamine in  $\gamma\delta$  T cells, the expression of different histamine receptor subtypes was analyzed. Using reverse transcriptase and polymerase chain reaction (RT-PCR) analysis, the expected products for the  $H_1R$ ,  $H_2R$  and  $H_4R$  subtypes were detected in isolated human-peripheral-blood  $\gamma\delta$  T cells. In contrast, the  $H_3R$  was undetectable (Fig. 1). Moreover,  $H_1R$ ,  $H_2R$  and  $H_4R$  expression in  $\gamma\delta$  T cells at the protein level has been shown by western blot analysis (Fig. 2).

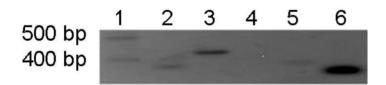


Fig. 1: Expression of mRNA of  $H_1R$ ,  $H_2R$  and  $H_4R$  in isolated human peripheral blood  $\gamma\delta$  T lymphocytes. (A)  $\gamma\delta$  T cells were isolated from human peripheral blood and the expression of mRNA for histamine receptors was analyzed. Lane 1) DNA molecular weight marker XIV (100-1500bp); lane 2)  $H_1R$  (403 bp); lane 3)  $H_2R$  (497 bp); lane 4)  $H_3R$  (589 bp); lane 5)  $H_4R$  (396 bp); lane 6)  $\beta_2$ -microglobulin (259 bp). Experiments were repeated three times with identical results.

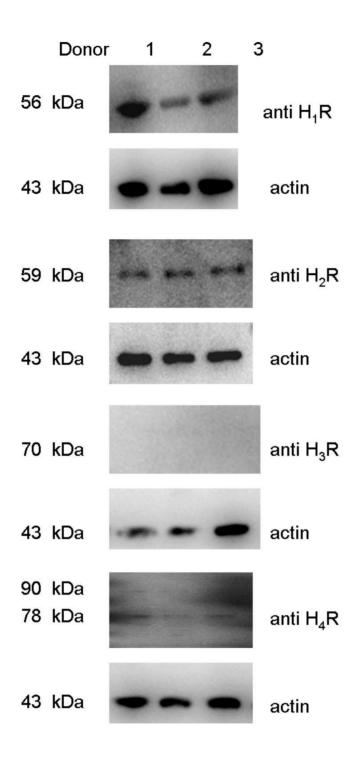
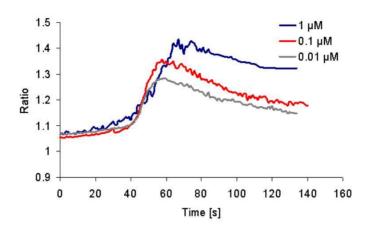


Fig. 2: Expression of  $H_1R$ ,  $H_2R$  and  $H_4R$  at a protein level in isolated human peripheral blood  $\gamma\delta$  T lymphocytes.  $\gamma\delta$  T cells were isolated from human peripheral blood and the expression of different histamine receptor subtypes was analyzed by western blot analysis. Block 1)  $H_1R$  (56 kDa), actin (43 kDa); block 2)  $H_2R$  (59 kDa), actin (43 kDa); block 3)  $H_3R$  (70k Da), actin (43 kDa); block 4)  $H_4R$  (78/90 kDa), actin (43 kDa). Experiments were repeated three times with identical results.

3.1.2. Histamine induces actin polymerization, intracellular  $Ca^{2+}$  mobilization, and chemotaxis in  $\gamma\delta$  T cells through pertussis toxin-sensitive  $G_i$  proteins

Histamine induces Ca<sup>2+</sup> transients in different types of leukocytes [154]. In accordance with the literature histamine also induced a rapid and concentration-dependent intracellular response in human  $\gamma\delta$  T cells isolated from peripheral blood (Fig. 3A). Ca<sup>2+</sup> transients are mainly induced by mobilization from intracellular stores or by influx across the plasma membrane from the medium. In order to discriminate between these experiments in the presence of ethyleneglycol-bis(betamechanisms. aminoethylether)-N,N'-tetraacetic acid (EGTA) were performed. Preincubating of γδ T cells with EGTA (4 mM) did not influence the histamine-initiated Ca<sup>2+</sup> intracellular rise, which implicated the mobilization of Ca2+ from intracellular stores (data not shown). To investigate the involvement of G<sub>i</sub> proteins in this response we took advantage of pertussis toxin. This toxin uncouples G<sub>i</sub> proteins from serpentine receptors by ADP-ribosylation. Pretreating  $\gamma\delta$  T cells for 1 h with pertussis toxin (100 ng/ml) strongly inhibited the histamine-induced Ca<sup>2+</sup> increase in γδ T cells, which in turn implicated the involvement of G<sub>i</sub> proteins in this histamine response (Fig. 3B). To check the responsiveness of pertussis-toxin-treated cells, experiments with ionomycin were performed. Ca<sup>2+</sup> transients induced by ionomycin were not influenced by pertussis toxin (data not shown).

A)



B)

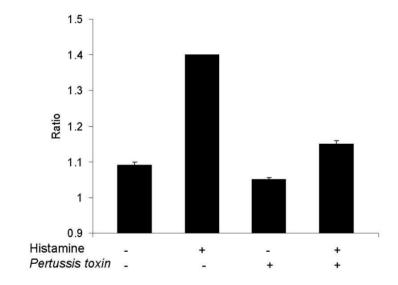
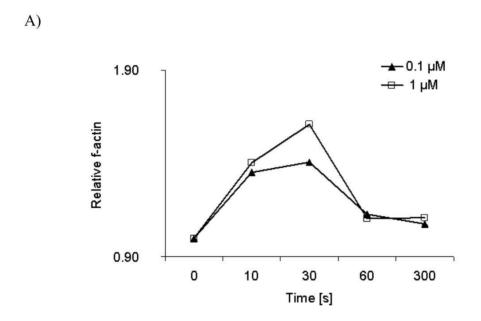


Fig. 3: Histamine induces calcium transients in isolated human peripheral  $\gamma\delta$  T lymphocytes. (A) Cells were loaded with Fura-2/AM and stimulated with 0.01  $\mu$ M, 0.1  $\mu$ M, and 1  $\mu$ M histamine. Intracellular Ca<sup>2+</sup> transients were followed by digital fluorescence microscopy and a ratio between 340 nm and 380 nm was calculated. Representative data from one experiment are shown. Experiments were repeated three times with identical results. (B) Cells were pre-incubated with or without pertussis toxin (100 ng/ml) for 90 min at 37°C and stimulated with 1  $\mu$ M histamine. The ratio calculated after stimulation with and without histamine is shown. Data are means of three different experiments of three different donors  $\pm$  SEM.

Next, actin reorganization was analyzed by flow cytometry. A rapid increase in f-actin content (circa 60%) was induced when  $\gamma\delta$  T cells were stimulated with histamine (Fig. 4A). The response was transient and reversible, with maximal values reaching within 30

seconds. To test the participation of  $G_i$  proteins in this response,  $\gamma\delta$  T cells were also pre-incubated with pertussis toxin before they were exposed to histamine (Fig. 4B). Pretreating  $\gamma\delta$  T cells with pertussis toxin almost completely abolished the effect of histamine on actin polymerization. In contrast pretreating them with cholera toxin did not result in a significant difference compared to control cells (Fig. 4C).



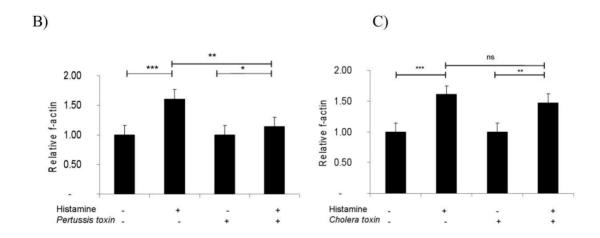
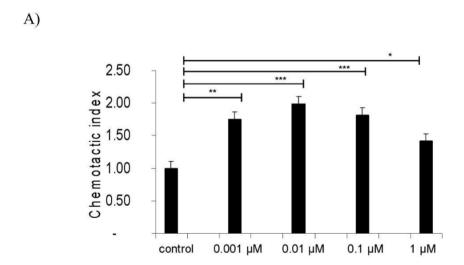


Fig. 4: Effects of histamine on actin polymerization in isolated human-peripheral-blood  $\gamma\delta$  T cells. (A) Cells were exposed to 0.1  $\mu$ M - 1  $\mu$ M histamine and f-actin content was measured by flow cytometry. (B) Cells were pre-incubated with or without pertussis toxin (100 ng/ml) for 90 min at 37°C and stimulated with 1  $\mu$ M histamine for 30 seconds, and the increase in f-actin content was analyzed. (C) Cells were pre-incubated with or without cholera toxin (0.5  $\mu$ g/ml) for 90 min at 37°C and stimulated with 1  $\mu$ M histamine for 30 seconds and the f-actin content was analyzed. All data are means of three

different experiments using three different donors ± SEM, (\*\*\*: p<0.0001; \*\*: p>0.005; \*: p>0.05; ns:

non-significant).

Intracellular  $Ca^{2+}$  transients and actin reorganization are prerequisites for cell migration. Therefore, human peripheral blood-isolated  $\gamma\delta$  T cells were exposed to different concentrations of histamine (0.01  $\mu$ M - 1  $\mu$ M), and migration across the porousmembrane was evaluated. Histamine induced a typical bell-shaped concentration dependent chemotactic response in  $\gamma\delta$  T lymphocytes (Fig. 5A). Maximal chemotactic responses were observed when cells were stimulated with 0.01  $\mu$ M histamine. Moreover, histamine-stimulated migration was abolished by pretreating  $\gamma\delta$  T cells with pertussis toxin (100 ng/ml) (Fig. 5B).



B)

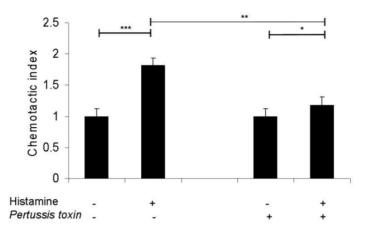
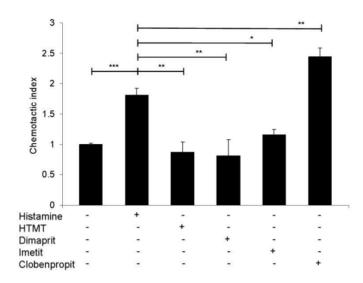


Fig. 5: Effects of histamine on chemotaxis in  $\gamma\delta$  T cells. Cells were exposed to different histamine concentrations (0.001 μM - 1 μM) in Boyden chambers. Migrated cells in the bottom wells of the Boyden chamber were fixed with formalin and counted by flow cytometry. (B)  $\gamma\delta$  T cells were pre-treated with pertussis toxin for 1h at 37°C, and migration in response to 0.1 μM histamine was analyzed. All data are means ±SEM, (\*\*\*: p<0.0001; \*\*: p>0.005; \*: p>0.05).

To specify the subtype of histamine receptor involved in the histamine-induced chemotactic response in human peripheral blood  $\gamma\delta$  T cells, receptor-selective agonists and antagonists were used. The H<sub>1</sub>R agonist 6-[2-(-imidazolyl)ethylamino]-N-(4-trifluoromethylphenyl) heptanecarboxamide dimaltate (HTMT), the H<sub>2</sub>R agonist S-(3-dimethylaminopropyl) isopthiourea 2HCl (dimaprit-2HCl) and the H<sub>3</sub>R agonist S-[2-(imidazol-4-yl)ethyl]isothiourea dihydrobromide (imetit) did not induce any significant change in chemotactic activity (Fig. 6A), whereas pretreating  $\gamma\delta$  T cells with the H<sub>4</sub>R antagonist N-cyclohexyl-4-(1H-imidazol-4-yl)-1-piperidinecarbothioamide maleate salt (thioperamide) prevented histamine-induced migration (Fig. 6B), suggesting that in  $\gamma\delta$  T cells, migration in response to histamine occurs through H<sub>4</sub>R.





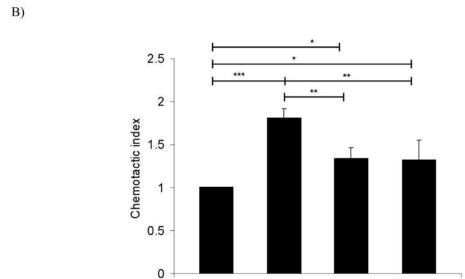


Fig. 6: Effects of histamine receptor agonists and antagonists on chemotaxis in  $\gamma\delta$  T cells. (A)  $\gamma\delta$  T cells isolated from healthy donors were exposed to histamine and isoform-selective agonists HTMT (H<sub>1</sub>R), dimaprit (H<sub>2</sub>R), and imetit (H<sub>3</sub>R), and migration was analyzed. (B)  $\gamma\delta$  T cells were pre-treated with the H<sub>4</sub>R antagonist thioperamide, and the migration assay was performed. All data are means ±SEM, (n=3); (\*\*\*: p<0.0001; \*\*: p>0.005; \*: p>0.05).

### 3.1.3. Histamine-induced intracellular cAMP levels in γδ T cells

Histamine Thioperamide

Histamine is known to affect the intracellular cAMP levels in human dendritic cells via  $H_2R$  and  $G_s$  proteins [112]. In order to characterize the functional expression of  $H_2R$  in

isolated human peripheral blood  $\gamma\delta$  T cells, intracellular cAMP levels after stimulation with histamine were determined by flow cytometry. A significant increase (p<0.0001) in cAMP levels, as reflected by increased in mean fluorescence intensity (MFI) was observed within 1 minute of histamine treatment (Fig. 7A). Moreover, cAMP levels reached a maximum after 4 minutes and remained high 30 minutes after histamine stimulation (Fig. 7B).

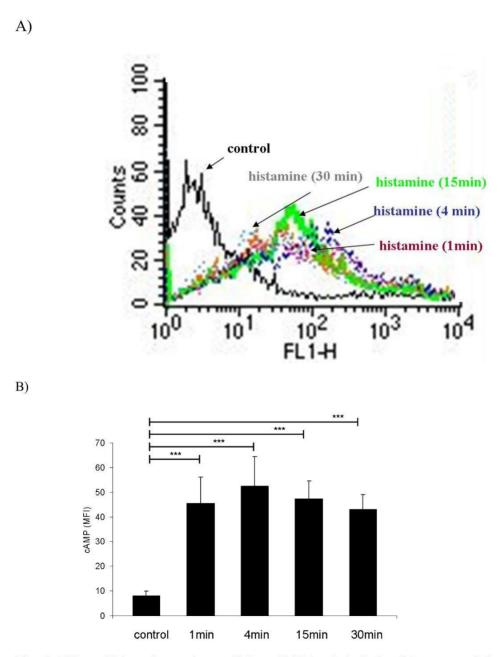


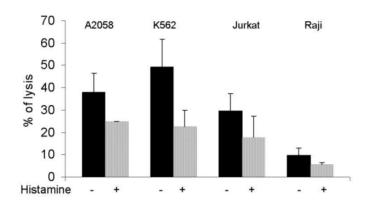
Fig. 7: Effect of histamine on intracellular cAMP levels in isolated human-peripheral-blood  $\gamma\delta$  T cells. (A) Distribution of fluorescence intensity in control cells and  $\gamma\delta$  T cells stimulated with 1  $\mu$ M

histamine for 15 min is shown. Aliquots of cells were fixed and stained as described in Materials and Methods. Fluorescence intensity was measured by flow cytometry. Representative data from one experiment are shown, performed three times in triplicates. (B) Time course of cAMP levels after stimulation with 1  $\mu$ M histamine. Experiments were repeated five times with  $\gamma\delta$  T cells isolated from different donors. Data are means  $\pm$ SEM, (n=5); (\*\*\*: p<0.0001).

### 3.1.4. Histamine affects the cytotoxic activity of $\gamma\delta$ T cells in response to tumor cells

We have shown that the activation of  $G_s$ -protein-coupled receptors and the upregulation of cAMP lead to the downregulation of cytotoxic responses in NK cells [151]. In addition,  $\gamma\delta$  T cells are known to process cytolytic activity towards different human tumor cell lines such as the myeloid leukaemia cell line (K562), cutaneous malignant melanoma cells and the non-Hodgkins T cell line Jurkat [99, 155, 156]. To better characterize the cytolytic activity of  $\gamma\delta$  T cells, in vitro radioactive assays were performed using different cell lines. Target cells were labelled for 1 hour with chromium (100  $\mu$ Ci/10<sup>6</sup> cells) and co-cultured for 4 hours at 37°C with  $\gamma\delta$  T cells to allow spontaneous cytotoxicity. As shown in (Fig. 8A), although  $\gamma\delta$  T cells displayed cytolitic activity in response all cell lines tested, their lytic capacity was greatest in response to K562 cells. Therefore, this cell line was chosen for further experiments to analyze the influence of histamine on on the cytolytic activity of  $\gamma\delta$  T cells. Histamine significantly reduced the cytolytic capacity of  $\gamma\delta$  T cells in response to K562 cells at all effector:target (E:T) ratios tested (Fig. 8B).

A)



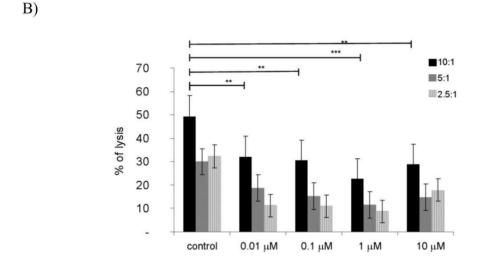
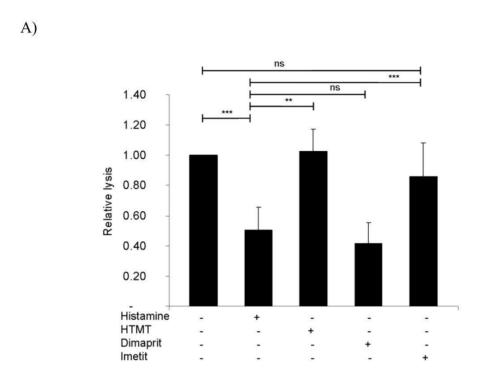


Fig. 8: Cytotoxic activity of  $\gamma\delta T$  cells from healthy donors in response to tumor cell lines: (A)  $\gamma\delta$  T cells isolated from healthy donors were co-cultured with different tumor cell lines and their spontaneous cytolytic capacity was determined. (B)  $\gamma\delta$  T cells were stimulated with the indicated concentrations of histamine (0.01  $\mu$ M - 10  $\mu$ M) and cytotoxicity in response to the chronic myeloid tumor cell line K562 was analyzed. Cells were co-cultured in different effector: target (E:T) ratios as indicated. Data are means  $\pm$ SEM, (n=3); (\*\*\*: p<0.0001; \*\*: p>0.005; \*: p>0.05).

In order to specify which subtype of histamine receptor modulates cytotoxicity in  $\gamma\delta$  T cells, experiments with receptor-specific agonists and antagonists were performed. Isoform-specific  $H_1R$ - and  $H_3R$ -agonists did not affect the spontaneous lysis of K562 cells by  $\gamma\delta$  T cells, whereas the  $H_2R$ -agonist dimaprit significantly reduced the spontaneous lysis capacity of  $\gamma\delta$  T cells in response to K562 cells (Fig. 9A). Moreover, the  $H_4R$  antagonist did not prevent the histamine-induced effect on cytotoxicity while

the  $H_2R$ -antagonist prevented the histamine-induced effect (Fig. 9B). These experiments suggest that the modulatory effect of histamine on  $\gamma\delta$  T-cell-mediated cytotoxicity requires the activation of the  $H_2R$  isoform.



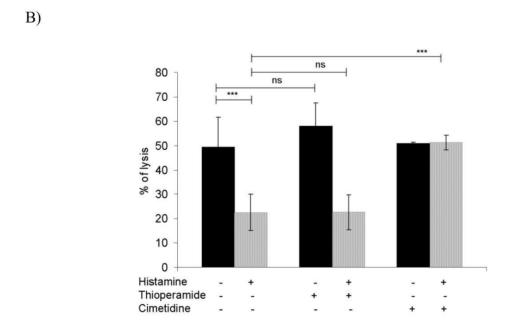


Fig. 9: Dimaprit inhibits the cytotoxic activity of  $\gamma\delta$  T cells. (A)  $\gamma\delta$  T cells were stimulated with 1  $\mu$ M histamine, 10  $\mu$ M H<sub>1</sub>R-agonist HTMT, 10  $\mu$ M H<sub>2</sub>R-agonist dimaprit and 0.01  $\mu$ M H<sub>3</sub>R-agonist imetit and co-cultures with human myeloid cell line K562 were performed in order to analyze the level of cytolytic

activity. (B)  $\gamma\delta T$  cells were stimulated with 1  $\mu M$  histamine in the presence or absence of 10  $\mu M$  H<sub>4</sub>R-antagonist thioperamide, or the H<sub>2</sub>R-antagonist, cimetidine and co-cultures with the human myeloid cell line K562 were performed in order to analyze cytolytic activity. Data are means  $\pm$  SEM, (n=3); (\*\*\*: p<0.0001; \*\*: p>0.005; ns: non-significant).

We next determined the involvement of different G proteins on histamine-modulated cytotoxicity. Thus,  $\gamma\delta$  T cells were pretreated with  $G_i$  protein inhibitor pertussis toxin and the  $G_s$  activator cholera toxin, and their cytotoxic activity in response to K562 cells was examined (Fig. 10). Pertussis toxin did not significantly alter the effect of histamine on cytotoxicity. On the contrary, pretreating  $\gamma\delta$  T cells with cholera toxin alone inhibited cytotoxicity by more than 50% with respect to control cells untreated with histamine. This inhibitory effect of the cholera toxin was further enhanced by histamine.

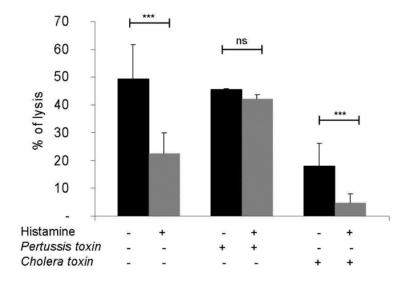
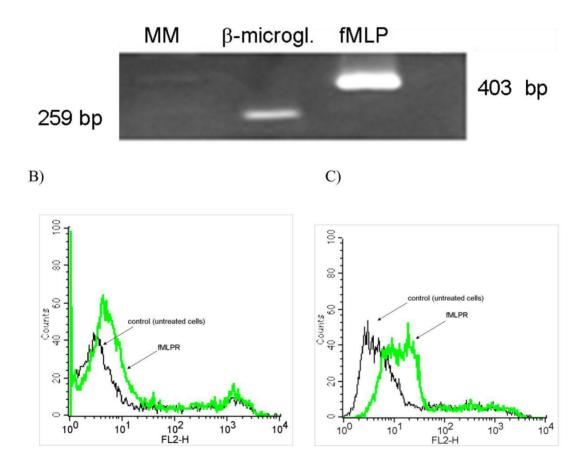


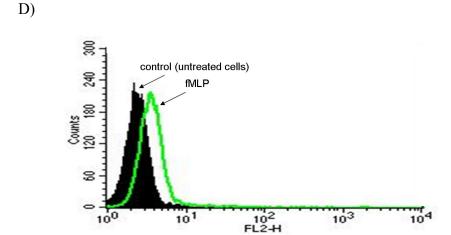
Fig. 10: Cytotoxic activity of  $\gamma\delta$  T cells in resonse to tumor cell line K562 is dependent on  $G_s$  proteins  $\gamma\delta$  T cells were pre-incubated with or without cholera toxin (0.5 µg/ml) or pertussis toxin (100 ng/ml) for 1h at 37°C. Thereafter,  $\gamma\delta$  T cells were stimulated or not with histamine and co-cultured with K562 cells. Data are means  $\pm$  SEM, (n=3); (\*\*\*: p<0.0001; ns: non-significant).

# 3.2. fMLP and $\gamma\delta$ T cells: involvement of pertussis-toxin-sensitive functions of isolated human-peripheral-blood $\gamma\delta$ T lymphocytes

The expression of the mRNA of the fMLP receptor was analyzed using reverse transcriptase and polymerase chain reaction (RT-PCR). The expected products for the fMLP receptor were detected in isolated human-peripheral-blood  $\gamma\delta$  T cells by gel electrophoresis (Fig. 11A). No PCR-product could be obtained by omitting reverse transcription (data not shown). In addition, the expression of fMLP proteins on the surface was analyzed. The fMLP expressions on  $\gamma\delta$  T cells are shown by flow cytometry (Fig. 11B+C). Neutrophils were used as control cells (Fig. 11D+E).

A)





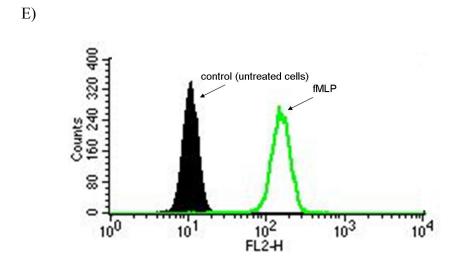


Fig. 11: Expression of fMLP mRNA by RT-PCR and surface receptors detected by flow cytometry staining in isolated human peripheral blood  $\gamma\delta$ T lymphocytes. (A)  $\gamma\delta$  T lymphocytes isolated from human peripheral blood express the mRNA for fMLP receptors. (B;C;D;E)  $\gamma\delta$  T lymphocytes express the cell surface receptors for fMLPR.  $\gamma\delta$  T lymphocytes were isolated and cultured as detailed in Materials and Methods. Cells were incubated for 30 min at 4°C with PE-conjugated anti-fMLPR mAb (B:  $\gamma\delta$  T lymphocytes extracellular staining; C:  $\gamma\delta$  T lymphocytes intracellular staining D: neutrophils extracellular staining; E: neutrophils intracellular staining). Fluorescence intensity of 10 000 cells/sample was analyzed by flow cytometry. Abscissa: fluorescence intensity FL-2H, PE; FL-1H, FITC; ordinate: number of cells.

fMLP is able to induce  $Ca^{2+}$  transients in neutrophils [157]. Here, the influence of fMLP on  $Ca^{2+}$  in  $\gamma\delta$  T cells was measured by digital fluorescence microscopy after loading the cells with Fura-2. fMLP induced a rapid and concentration-dependent intracellular response in human  $\gamma\delta$  T cells isolated from peripheral blood (Fig. 12A).  $Ca^{2+}$  transients

are mainly created by their mobilization from intracellular stores or by their influx across the plasma membrane from the medium. In order to distinguish between these two mechanisms, experiments in the presence of EGTA in the medium were performed. The preincubation of  $\gamma\delta$  T cells with EGTA (4 mM) did not influence the fMLP-initiated intracellular Ca<sup>2+</sup> rise, implicating the mobilization of Ca<sup>2+</sup> from intracellular stores (data not shown). To investigate the involvement of  $G_i$  proteins in this response, we took advantage of pertussis toxin. This toxin uncouples  $G_i$  proteins from serpentine receptors by ADP-ribosylation. Pretreating  $\gamma\delta$  T cells for 1h with pertussis toxin (100 ng/ml) inhibited the fMLP-induced Ca<sup>2+</sup> increase in  $\gamma\delta$  T cells (Fig. 12B). To check the responsiveness of pertussis-toxin-treated cells, experiments with ionomycin were performed. Ca<sup>2+</sup> transients induced by ionomycin were not influenced by pertussis toxin (data not shown).

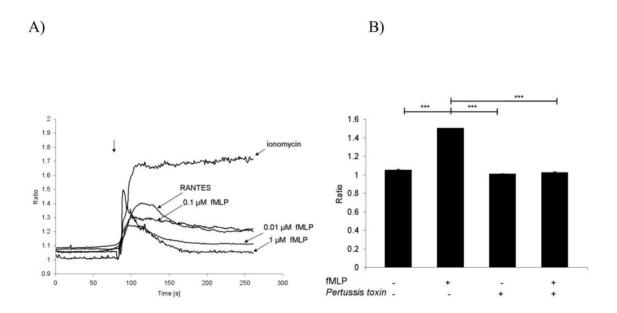
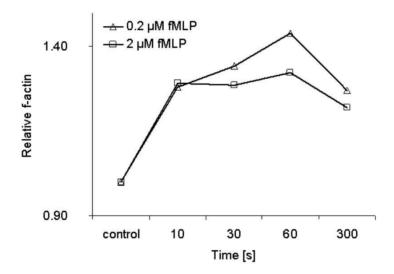


Fig. 12: fMLP activation induces intracellular  $Ca^{2+}$  transients in human peripheral  $\gamma\delta T$  lymphocytes. (A) Cells were loaded with Fura-2/AM and stimulated with different concentrations of fMLP (0.01  $\mu$ M -1  $\mu$ M). Intracellular  $Ca^{2+}$  transients were followed in fura2-labelled human peripheral blood  $\gamma\delta$  T lymphocytes. fMLP induces a rapid and concentration-dependent increase of intracellular calcium  $Ca^{2+}$ . Ionomycin and RANTES were used as positive controls. Changes in fluorescence intensity were followed spectrofluorometrically after stimulation of the cells with fMLP, and the ratio between 340

nm and 380 nm was calculated. (B) Cells were preincubated with pertussis toxin (100 ng/ml) for 90 min at 37°C and stimulated with fMLP (1  $\mu$ M). Representative data from three experiments using cells isolated from different donors are shown.

The reorganization of f-actin is a prerequisite for the migration of different types of leukocytes [158, 159]. Next, the influence of fMLP on the f-actin content in human peripheral blood  $\gamma\delta$  T cells was analyzed by flow cytometry. Rapid f-actin formation was induced when  $\gamma\delta$  T cells were stimulated with fMLP (Fig. 13A). The increase in f-actin content upon fMLP stimulation was transient, but extended for at least 5 min. Pretreating  $\gamma\delta$  T cells with pertussis toxin completely abolished the effect of fMLP on f-actin polymerization (Fig. 13B).

A)



B)

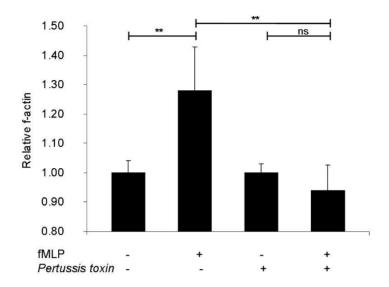
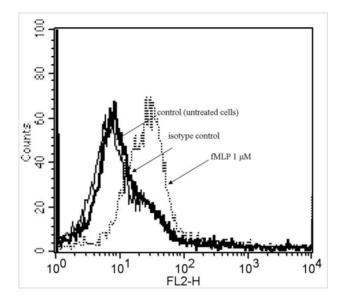


Fig. 13: Effects of fMLP on actin polymerization in isolated human-peripheral-blood  $\gamma\delta T$  cells. (A) fMLP induced f-actin polymerization in  $\gamma\delta T$  cells. (B) Cells were pre-incubated without or with pertussis toxin (100ng/ml) for 90 min at 37°C and stimulated (10s) with a concentration of 0.2  $\mu$ M fMLP. The relative f-actin content after 0s, 10s, 30s, and 300s was measured by flow cytometry. A significant increase in f-actin polymerization was observed 30s after fMLP stimulation compared to the untreated control. Data are means  $\pm$  SEM, (n=3); (\*\*: p<0.0001; ns: non-significant).

CD11b is known to be the key  $\beta_2$ -integrin involved in cell adhesion and leukocyte chemotaxis [160]. Thus we measured the influence of fMLP on CD11b expression by flow cytometry. fMLP induced a concentration-dependent upregulation of CD11b on the surface of human  $\gamma\delta$  T lymphocytes (Fig. 14A-B).

A)



B)

35
30
25
W
20
W
20
15
10
5

0.01 µM

control

Fig. 14: fMLP-induced CD11b expression in  $\gamma\delta T$  cells.  $\gamma\delta$  T cells were stimulated with the indicated concentrations of fMLP and then incubated for 30 minutes at 37°C. The reaction was stopped by diluting the sample to 100 times the original concentration with ice-cold buffer. Samples were incubated for 40 minutes on ice with PE-conjugated anti-CD11b mAbs and then analyzed by flow cytometry. Representative histograms of stimulated (0.01 μM - 10 μM fMLP) (A-B) and unstimulated anti-CD11b-stained  $\gamma\delta$  T cells are shown. Representative data from three experiments using cells isolated from different donors are shown.

 $0.1 \mu M$ 

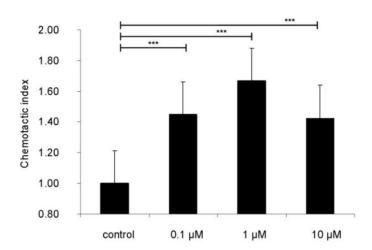
1 µM

10 µM

Intracellular  $Ca^{2+}$  transients, actin reorganization and CD11b expression are prerequisites for cell migration in various types of leukocytes. Therefore, isolated human-peripheral-blood  $\gamma\delta$  T cells were exposed to different concentrations of fMLP

 $(0.01~\mu M$  - 1  $\mu M)$ , and migration across a porous membrane was evaluated. Both fMLP induced a typical bell-shaped concentration-dependent chemotactic response in  $\gamma\delta$  T lymphocytes (Fig. 15A). Again, fMLP-stimulated migration was abolished by pretreating  $\gamma\delta$  T cells with pertussis toxin (Fig. 15B).







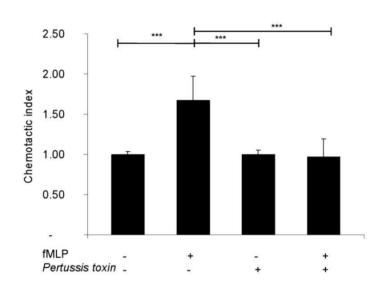


Fig. 15: fMLP-induced chemotaxis of circulating γδT cells. (A) Cells were exposed to different fMLP concentrations (0.1  $\mu$ M-10  $\mu$ M) in Boyden chambers. Migrated cells collected from the bottom wells were fixed with formalin and counted by flow cytometry. (B) γδ T cells were pre-treated with the  $G_i$  protein inhibitor pertussis toxin. fMLP-induced migration of isolated human-peripheral-blood γδ T lymphocytes is significantly inhibited by pertussis toxin. Data are means  $\pm$ SEM, (\*\*\*: p<0.0001; \*\*: p>0.005; ns: non-significant).

### 4. Discussion

### 4.1. γδ T cells

γδ T cells are thought to be major effector cells in a number of infectious diseases, e.g. tuberculosis [66, 67], brucellosis [161], listeriosis [162], leishmaniasis [163], malaria [164], and toxoplasmosis [165]. The recruitment and activation of y\delta T lymphocytes to sites of inflammation are caused by various agents and natural phosphoantigens derived from plants [39], bacteria [40, 41, 166], protozoa [43], endogenous ligands [32, 48, 35, 49, 167] and viruses [167, 168]. Early events in an immune response involving  $\gamma\delta$  T cells stimulate the production of cytokines that direct innate host defenses and the subsequent development of T cell subsets with discrete patterns of cytokine production.  $\gamma\delta$  T cells are a population of lymphocytes expressing functional  $\gamma\delta$  T cell receptor genes [30]. This subtype of T cells, presumably an ancient type of lymphocyte, is derived from haematopoietic stem cells that share certain characteristics with other immune cells, e.g. the capacity for antigen presentation, immune modulatory properties and cytolytic activity [34, 80]. Two main subsets of γδ T cells are distinguished according to their location. Resident γδ T cells are found in skin, uterine and epithelial tissues, whereas circulating/systemic γδ T cells can be isolated from peripheral blood or lymphoid tissues [54, 55]. In contrast to αβ T lymphocytes, γδ T cells do not need antigens presenting on classical MHC molecules for recognition [55]. Instead, they recognize antigens bound to CD1 molecules. [45, 169]. Several lines of evidence implicate  $\gamma\delta$  T cells in primary host defenses as well as in tumor surveillance. They are also known to attack bacterial and virus-bearing cells as well as transformed cells [79]. This cytotoxic activity is mediated by the production and release of perforin and granzymes [79, 34]. The antitumoral activity of γδ T cells is either mediated via endogenous ligands in  $\gamma\delta$  TCR-dependent fashion or depends on the interaction of the cells with NK cell receptor NKG2D [75, 79].

Different subpopulations of immune cells can infiltrate tumor lesions, although there is usually a limited number of T lymphocytes detectable even in those neoplasias (such as melanoma) in which the lymphocyte infiltrate appears to correlate with disease course [169]. The limited number of T lymphocytes in the microenvironment of tumors suggests that mechanisms acting in down-modulating antitumor immune responses are likely to be at work [170]. Two lymphocyte subsets T cells and NK cells are frequently found within malignant tumors and in the surrounding inflammatory tissue. Fractions of these intra- or peritumoral lymphocytes are frequently impaired compared to T and NK cells in peripheral blood or in adjacent non-malignant tissue, a phenomenon referred to as tumor-induced immunosuppression [171, 172]. Tumor and tumor-associated cells actively and directly inhibit immune responses by secreting of immunosuppressive cytokines such as interleukin (IL)-10 [173] and transforming growth factor (TGF)- $\beta$  [173, 174], as well as by skewing CD4<sup>+</sup> T cell responses into  $T_{\rm H}2$  rather than  $T_{\rm H}1$ -dominated immunoreactions [175, 176].

MCP-1/CCL2 [87], RANTES/CCL5, MIP-1 $\alpha$ /CCL3 and MIP-1 $\beta$ /CCL4 were found to be chemically attractive for  $\alpha\beta$  T lymphocytes and circulating  $\gamma\delta$  T cells, whereas neither  $\alpha\beta$  nor  $\gamma\delta$  responded to IL-8/ CXCL8 or IFN-inducible protein 10 (IP-10/CXCL10) [89]. The corresponding chemokine receptors are seven-transmembrane-spanning G-protein-coupled receptors. They share structural features and some of them are involved in intrathymic [90], and memory T cell development, others in the orchestration of T and B cell interactions and the differentiation of effector T cells [87].

# 4.2. G protein-coupled receptor pathways in circulating $\gamma\delta$ T cells: histamine and fMLP receptor pathways

Our data suggest that γδ T cells' responses to the stimulation of histamine or fMLP are dependent on G protein-coupled receptor pathways. The heterotrimeric G proteins are composed of three subunits:  $\alpha$ ,  $\beta$  and  $\gamma$ . About twenty  $\alpha$  subunits have been reported, belonging to four families. Two distinct G protein subtypes are sensitive to bacterial toxins: cholera toxin ADP-ribosylates the arginine residue present in the carboxy terminal of the  $\alpha$  subunit of  $G_s$ , whereas pertussis toxin ADP-ribosylates the cystein residue in the carboxy terminal of the  $\alpha$  subunit of  $G_i$  or  $G_o$  proteins [177]. Each G protein has a unique biological function. The G<sub>s</sub> protein activates adenylyl cyclise, resulting in the accumulation of cyclic adenosine monophosphat (cAMP) in the cells. G<sub>i</sub> proteins are highly homologous members of the family of heterotrimeric signaltransducing guanine nucleotide-binding proteins (G proteins) and by coupling numerous receptors for extracellular signalling molecules to inhibit adenylyl cyclase, stimulate phospholipase C, or open potassium channels [9,10]. The activation of phospholipase C results in the hydrolysis of phosphatidylinositol bisphosphate (PIP<sub>2</sub>) and the generation of diacylglyceride (DAG), an activator of protein kinase C (PKC), and inositol(1,4,5)trisphosphate [12].

4.2.1. Histamine affects  $\gamma\delta$  T cell effector functions: the involvement of pertussis-toxinand cholera-toxin-sensitive functions of isolated human peripheral blood  $\gamma\delta$  T lymphocytes

Since its discovery in 1910, histamine has been regarded as one of the most important mediators in allergy and inflammation and is known to be involved in smooth muscle-stimulating, vasodepressor action and is involved in reactions during anaphylaxis [178].

Although histamine is located in most body tissues, it is highly concentrated in the lungs, skin and gastrointestinal tract [179], where it has been shown to regulate both the secretion of gastric acid in the stomach and the functions of several neurological transmitters [180, 181]. In the central nervous system, histamine is involved in regulating drinking, body temperature, and blood pressure and in perceiving pain [182, 183]. Histamine has also been described as an autocrine/paracrine or exogenous growth factor for cancer cells, e.g. malignant melanomas and leukaemia cells. In the case of chronic myeloid leukaemia, the secretion of histamine is the consequence of a leukaemia-specific oncogene [107].

Histamine is known to regulate humoral and cellular immunity by controlling the production of pro-inflammatory cytokines, the expression of adhesion molecules and the migration of inflammatory cells such as eosinophils, dendritic cells, NK cells and αβ T cells [184, 185]. Histamine and histidine decarboxylase (HDC) [105], the only enzyme that catalyzes histamine production, has been proven to be present in elevated concentrations in proliferating tissues including tumor cells [157]. In the case of the myeloproliferative disease chronic myeloid leukaemia (CML), this gene was the first known leukaemia-specific oncogene reported to be involved in the regulation of histamine production in neoplastic cells [107]. By mediating the immune response histamine also interacts directly with H<sub>2</sub> receptors (H<sub>2</sub>R) on T cells to inhibit T cell proliferation and cytokine production [109]. H<sub>2</sub>R antagonists increase the likelihood that patients with gastric and colorectal cancer will survive [186, 187]. Histamine mediates the dose-dependent inhibition of human IL-12 and stimulates IL-10 via H<sub>2</sub>R receptors [188]; moreover, it inhibits the IFN-  $\gamma$  production of T<sub>H</sub>1 cells but has no effect on IL-4 production from T<sub>H</sub>2 clones [189]. Furthermore, histamine directly enhances the production of human IgE from B cells [190], while it inhibits T cell cytotoxicity. Available data suggest that histamine induces a  $T_H2$  shift at the level of monocytes [191].

It has been previously shown that  $\gamma\delta$  T cells exposed to histamine do not expand [39]; however, our data show that  $\gamma\delta$  T cells express three distinct histamine receptor subtypes and functionally respond to two of them. To better understand the role of histamine in the cross-talk between immune cells and tumor cells, we performed cell studies and co-culture experiments with γδ T cells. Here, we demonstrated that histamine stimulates actin polymerization and Ca2+ transients in a concentrationdependent manner and migration in a typical bell-shaped concentration curve. Moreover, we showed that the increase in intracellular Ca<sup>2+</sup> is due to mobilization from intracellular stores, inasmuch as it is insensitive to the chelation of extracellular Ca<sup>2+</sup>. A long-lasting migration response requires continuous interaction between migrationinducing ligands and the cell surface receptors that can be activated via actin polymerization to induce the "cell motor". Therefore, a gradient of ligands, continuously available cell surface receptors and a very sensitive signal transduction mechanism are necessary to transmit the external signal to the internal cellular movement machinery. Thus, a low concentration of chemotactic ligands can activate and direct the cell over a long period of time. At high ligand concentrations, the receptors at the cell surface are very quickly occupied and consequently desensitized as well as internalized via endocytosis. In this case, the chemotactic index is low because ligands find no form of receptors at the cell surface that can be activated and are not able to induce movement until either novel transcriptionally regulated receptors are synthesized or the internalized receptors are recycled [192].

Histamine is able to act by binding to different G protein-coupled receptors. In order to demonstrate the participation of  $G_i$  proteins in these stimulated cell responses, experiments with pertussis toxin were performed. This toxin selectively uncouples  $G_i$ 

proteins from the intracellular sites of receptors by ADP-ribosylation. Pretreating  $\gamma\delta$  T cells with pertussis toxin blocked histamine-induced actin polymerization, Ca<sup>2+</sup> transients and migration in  $\gamma\delta$  T cells; these consequences implicate the involvement of  $G_i$  proteins in these cell responses. Principally, histamine is able to act by binding to different receptor subtypes,  $H_1R$ ,  $H_2R$ ,  $H_3R$ , and  $H_4R$ .

In the present work, reverse transcription-polymerase chain reactions revealed the mRNA expression of H<sub>1</sub>R, H<sub>2</sub>R and H<sub>4</sub>R in γδ T cells. No mRNA expression for H<sub>3</sub>R was detected. Moreover, H<sub>1</sub>R, H<sub>2</sub>R and H<sub>4</sub>R expression at the protein level has been shown by Western blot analysis. The involvement of the different receptors in these cell responses was dissected using isoform-specific receptor agonists and antagonists. Our experiments revealed that histamine regulates actin polymerization, Ca<sup>2+</sup> transients and chemotaxis via H<sub>4</sub>R, but provided no evidence for the involvement of H<sub>1</sub>R, H<sub>2</sub>R and H<sub>3</sub>R in these cell responses. This receptor-isoform specific cell regulation is consistent with reports in eosinophils, mast cells and NK cells [185, 114]. Therefore, one can assume that H<sub>4</sub>R in γδ T cells activates pertussis-toxin-sensitive heterotrimeric G<sub>i</sub> proteins, which in turn dissociate into the guanosine triphosphate- $\alpha$  subunit and free  $\beta \gamma$ dimers. The latter activates phospholipase  $C\beta_2$  [193]. This enzyme cleaves phosphatidylinositol (4,5)-bisphosphate into diacylglycerol and trisphosphate, which mobilizes Ca2+ from intracellular stores [194]. In leukocytes, Gi proteins regulate the reorganization of the actin cytoskeleton independently of activated phospholipase C [195]. These G<sub>i</sub>-protein-coupled signalling pathways are essential components of the migration response in different subtypes of leukocytes [196].

Unlike classical chemotaxin receptors, such as chemokine receptors or the complement C5a receptor, the coupling of different types of histamine receptors is pleiotropic, including the interactions of  $H_2R$  with  $G_s$  proteins with the consequent activation of adenylyl cyclase.

Our cell studies combining histamine and isoform-specific receptor agonists or antagonists showed enhanced cAMP levels and  $H_2R$  activation in  $\gamma\delta$  T cells. In different subtypes of leukocytes, e.g. NK cells and CD8<sup>+</sup> T cells, the cytotoxicity response by cAMP has been reported to be inhibited [197]. Our data show that the spontaneous cytolytic activity of isolated human peripheral blood  $\gamma\delta$  T cells is prevented by histamine. Neither HTMT nor imetit nor thioperamide altered the spontaneous cytolytic capacity of  $\gamma\delta$  T cells, but it was inhibited by dimaprit, suggesting that  $H_2R$  may be involved in the inhibitory effect of histamine on the cytotoxicity of  $\gamma\delta$  T cells in human peripheral blood. Moreover, it has been shown that cholera toxin impairs cytotoxicity in  $\alpha\beta$  T lymphocytes and NK cells [198].

As the literature suggests [198], we found that the  $G_s$ -protein activator cholera toxin inhibited the spontaneous cytotoxicity of  $\gamma\delta$  T cells, enhancing cAMP levels. Infiltration by lymphocytes, macrophages, mast cells, and neutrophils is a hallmark of inflammatory, defense and tissue repair reactions, which are often present in tumors. Various types of tumor-infiltrating macrophages and lymphocytes are considered to be potential effectors of anti-tumor immunity and may interfere with tumor expression [199]. In this work, we showed that histamine, which is present in the inflammatory and neoplastic microenvironment, induces the migration of isolated human peripheral blood  $\gamma\delta$  T cells; in contrast, the spontaneous cytolytic effect of  $\gamma\delta$  T cells is prevented by histamine [108, 200]. Neither HTMT nor imetit nor thioperamide altered the spontaneous cytolytic effect of  $\gamma\delta$  T cells, but it was inhibited by dimaprit, suggesting that H<sub>2</sub>R may be involved in the inhibitory effect of histamine on human peripheral

blood  $\gamma\delta$  T cell-mediated cytotoxicity. Our data suggest that histamine contributes to the ability of tumor cells to escape the immunological surveillance.

4.2.2. fMLP affects effector functions of  $\gamma\delta$  T cells: the involvement of pertussis-toxinsensitive functions of isolated human peripheral blood  $\gamma\delta$  T lymphocytes

Gram-negative bacteria-derived and synthetic N-formyl peptides play key chemotactic roles for phagocytic cells. N-formyl-Met-Leu-Phe (fMLP) is the prototype for Nformylated peptides. Its receptor in humans has been pharmacologically defined as a high affinity binding site on the surface of phagocytes such as neutrophils [121]. Natural fMLP was subsequently purified and identified in supernatants of gram negative bacteria. Nevertheless, over the past five years, data from several groups have indicated that fMLP receptors might act in a more complex manner. A large number of nonformylated peptide ligands for the fMLP receptor, including derivates of mitochondrial proteins, have been identified. Moreover, physiological ligands for the fMLP receptor also presumably derive also from viral envelope proteins and the fibriolytic cascade [133]. However, it is currently thought that the main effector cells activated by fMLP represent phagocytes such as neutrophils, dendritic cells and monocytes. Recently, the reduced resistance of fMLP receptor knockout mice to infection by Listeria monocytogenes has been reported [201]. Since there is good evidence that  $\gamma\delta$  T cells are major effector cells in listeriosis, the direct effect of fMLP on cell adhesion and migration was analyzed in these cells.

fMLP exerts its biological functions through three different receptor subtypes: fMLP-receptor (FPR), and fMLP receptor-like 1 and 2 (FPRL1, FPRL2). Human FPR was first defined biochemically as a high affinity binding site for the prototype fMLP on the surface of neutrophils. Two additional human genes, designated FPRL1 and FPRL2

(FPR-like), were subsequently isolated. [202, 203, 204]. It is generally accepted that FPRL1 shares signal transduction features with FPR. Both receptors are sensitive to pertussis toxin and possess a high degree of amino acid identity in the cytoplasmic signalling domains [205].

Here, we demonstrated the expression of the fMLP receptor (FPR) surface protein on  $\gamma\delta$  T cells. Moreover, we showed that fMLP stimulates  $Ca^{2+}$  transients in  $\gamma\delta$  T cells. The increase in intracellular  $Ca^{2+}$  is due to mobilization from intracellular stores, inasmuch as it is insensitive to extracellular chelation by EGTA. Pretreating  $\gamma\delta$  T cells with pertussis toxin inhibited both fMLP-induced  $Ca^{2+}$  transients. This toxin selectively uncouples  $G_i$  proteins from the intracellular sites of receptors by ADP ribosylation. These data suggest that fMLP in  $\gamma\delta$  T cells induces similar or identical signal pathways as in phagocytes. Binding fMLP to their receptors might induce GDP/GTP exchange in the  $G\alpha_i$  subunit of heterotrimeric G proteins. The activated heterotrimeric G protein complex (GTPform) is unstable and dissociates into  $G\alpha_i$  subunits and free  $\beta\gamma$  dimers, which activate phospholipase  $C\beta_2$  [194]. This enzyme cleaves phosphatidylinositol (4,5)-bisphosphate into diacylglycerol and the inositol trisphosphate, which in turn mobilizes  $Ca^{2+}$  from intracellular stores [195].

In leukocytes,  $G_i$  proteins regulate the reorganization of the actin cytoskeleton independently of activated phospholipase C [196]. These  $G_i$ -protein-coupled signalling pathways are essential components of the migratory response in different subtypes of leukocytes [197]. To determine whether fMLP-treated cells can initiate the signalling required for actin reorganization  $\gamma\delta$  T cells were activated with different concentrations of fMLP (0.1  $\mu$ M - 10  $\mu$ M) labelled with NBD and analyzed by flow cytometry. Rolling of the cells is mediated by the selectin family of adhesion molecules and their counterstructural carbonhydrate ligands, such as sialyl Lewis X, on leukocytes and sulphated

polysaccharides, such as fucoidan, on endothelial cells. The more stable adhesion, sticking, to the vessel wall is mediated by CD11/CD18 leukocyte adhesion molecules ( $\beta_2$  integrins) on leukocytes and intercellular adhesion molecules (ICAMs) on endothelial cells. After attaching firmly, leukocytes start to migrate across the endothelium via intercellular junctions into the subendothelial space. Finally, leukocytes become attached to the inflammatory sites a process known as chemotaxis through the production of exogenous and endogenous chemoattractant mediators. Exogenous chemotaxins include bacterial oligopeptides of the fMLP type, lectins, denatured proteins, some lipids and lipopolysaccharides. Endogenous chemotaxins are produced by the host organism and are humoral (complement fragment C5a and C3a, fibrinopeptides, kallikrein and plasminogen activators) or cellular (from different cells: leukotrine B4, platelet activated factor, chemotactic cytokines) [206, 207].

We demonstrated that fMLP induces actin polymerization and CD11b upregulation in a concentration dependent manner and migration in a typical bell-shaped concentration curve. The requirements of these cell responses were discussed above already in the chapter for histamine.

#### 4.3. Concusion

The data collected in this work represent the first evidence that G-protein-coupled signalling exist in γδ T cells via histamine and fMLP receptors. The bigenic amine histamine and the bacterial peptide fMLP have been demonstrated to be novel chemoattractant factors for circulating human γδ T cells, which are critical members of the immunological tumor surveillance machinery. Here, we analyzed the influence of histamine on the interaction of human γδ T cells with tumor cells such as the A2058 human melanoma cell line, the human Burkitt's non-Hodgkins lymphoma cell line Raji, the T-lymphoblastic lymphoma cell line Jurkat, and the NK cell-sensitive erythroleukaemia line K562. We found that histamine inhibits the spontaneous cytolytic activity of  $\gamma\delta$  T cells in response to these cell lines. The downregulation of  $\gamma\delta$  T cell mediated cytotoxicity involves the histamine receptor subtype 2 (H<sub>2</sub>R), the activation of G<sub>s</sub> proteins and increased cAMP intracellular levels. On the other hand, histamine activates the common signalling pathways of chemotaxins such as G<sub>i</sub>-protein-dependent actin reorganization, the increase of intracellular Ca<sup>2+</sup> and the induction of migratory response in γδ T lymphocytes. Our data indicate that histamine contributes to the mechanism by which tumor cells escape immunological surveillance.

The bacterial-cell-wall-derived peptide N-formyl-Met-Leu-Phe (fMLP) is a well characterized chemotactic factor for phagocytes such as neutrophils, monocytes and dendritic cells. Here, we analyzed the influence of fMLP on isolated human peripheral blood  $\gamma\delta$  T cells. We found that fMLP induces intracellular calcium transients, actin reorganization, CD11b upregulation and the migration of  $\gamma\delta$  T cells. Pretreating  $\gamma\delta$  T cells with pertussis toxin inhibited all fMLP-stimulated cell responses, implicating the involvement of Gi proteins in the induced signalling cascade. The present data suggest that,

in addition to phagocytes, N-formyl peptides also regulate the trafficking and activation of  $\gamma\delta$  T cells.

### 4.4. Outlook

Our data suggest that the responses of circulating human  $\gamma\delta$  T cells to histamine are mediated by two different G-protein-coupled pathways. Data suggest that the downstream pathways of cAMP is involved in the responses of  $\gamma\delta$  T cells to histamine stimulation. Further analysis of the involvement of phospholipase C (PLC), protein kinas B (Akt) and protein kinase A is required.

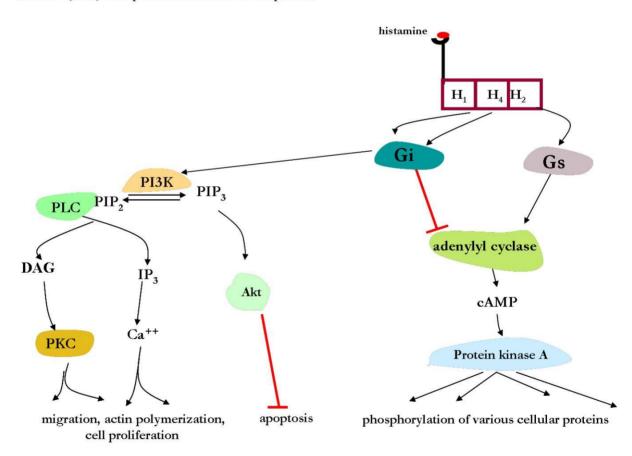
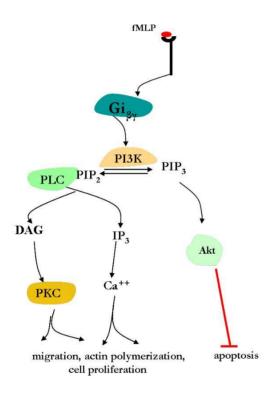


Figure 4.4.1. The possible G-protein-coupled receptor signalling in  $\gamma\delta$  T cells via the histamine receptor subtypes H<sub>1</sub>R, H<sub>2</sub>R, H<sub>4</sub>R. Histamine may recruit the  $\gamma\delta$  T cells to the site of tumors and inhibit their cytolytic functions. Once there,  $\gamma\delta$  T cells secrete different chemokines, e.g. RANTES, that may provide tumor development. (Abbreviations: PI3K: phosphatidylinositol 3-kinase; PIP<sub>2</sub>: phosphatidylinositol 4,5-bisphosphate; PIP<sub>3</sub>: phosphatidylinositol (3,4,5)-trisphosphate; PLC: phospholipase C; DAG: diacylglycerol; PKC: protein kinase C; IP<sub>3</sub>: inositol-1,4,5-trisphosphate; Ca<sup>++</sup>: calcium; Akt: proteinkinase B; cAMP: cyclic adenosine-3',5'-monophosphate; H<sub>1</sub>R, H<sub>2</sub>R and H<sub>4</sub>R: histamine receptor subtypes 1,2 and 4; G<sub>i</sub>: G<sub>i</sub> protein; G<sub>s</sub>: G<sub>s</sub> protein).

Furthermore, in the case of the fMLP receptor pathway in circulating human  $\gamma\delta$  T cells, further studies analyzing the different subtypes of the fMLP receptor family are required to describe the exact receptor pathway involved in the responses of  $\gamma\delta$  T cells to fMLP.



**Figure 4.4.2.** The possible G-protein-coupled signalling pathway in  $\gamma\delta$  T cells via the fMLP receptor. (Abbreviations: **PI3K**: phosphatidylinositol 3-kinase; **PIP<sub>2</sub>**: phosphatidylinositol 4,5-bisphosphate; **PIP<sub>3</sub>**: phosphatidylinositol (3,4,5)-trisphosphate; **PLC**: phospholipase C; **DAG**: diacylglycerol; **PKC**: protein kinase C; **IP<sub>3</sub>**: inositol-1,4,5-trisphosphate; **Ca**<sup>++</sup>: calcium; **Akt**: proteinkinase B; **fMLP**: N-formyl-Met-Leu-Phe).

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# 7. Selbständigkeitserklärung / Declaration

Hiermit erkläre ich, dass ich die vorliegende Arbeit selbständig und nur unter Verwendung der angegebenen Hilfsmittel, Literatur und persönlichen Mitteilungen angefertigt habe.

Ich versichere, dass ich diese Arbeit noch an keiner anderen Hochschule eingereicht habe, um ein Promotionsverfahren eröffnen zu lassen. Die Hilfe eines Promotionsberaters wurde von mir nicht in Anspruch genommen. Dritte haben von mir weder unmittelbar noch mittelbar geldwerte Leistungen für Arbeiten erhalten, die im Zusammenhang mit der vorgelegten Dissertation stehen. Die Promotionsordnung der Biologisch-Pharmazeutischen Fakultät ist mir bekannt.

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- 2. Bagi É., K. Tóth, **K. Truta-Feles**, L. Lénárd: A sóéhség központi idegrendszeri angiotenzinerg szabályozása patkány zona incertában. A MÉT LXIX. Vándorgyűlése, E55, p.: 49, 2005.42
- 3. Bagi É., K. Tóth, **K. Truta-Feles**, L. Lénárd: Angiotensinergic regulation of salt hunger in the zona incerta of rat brain. *Acta Physiologica Hungarica*, 92(3-4): 237, 2005.
- 4. Bagi É, Tóth K, **Truta-Feles K**, Lénárd L: Angiotensinergic mechanism in the central nervous system regulating salt intake and fluid balance. *Clin. Neurosci.*, 58:(Suppl 1) 10- p. (2005)
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- 6. Lagadari M, **Truta-Feles K**, Lehmann K, Berod L, Ziemer M, Idzko M, Barz D, Kamradt T, Maghazachi AA, Norgauer J.: Lysophosphatidic acid inhibits the cytotoxic activity of NK cells: involvement of G<sub>s</sub> protein-mediated signaling. *Int. Immunol.* 2009 Jun;21(6):667-77. Epub 2009 May 21.
- 7. **K. Truta-Feles**, M. Lagadari, K. Lehmann, L. Berod, S. Cubillos, S. Piehler, Y. Herouy, D. Barz, T. Kamradt, A.A. Maghazachi, and J. Norgauer: A novel role for histamine as a modulator of  $\gamma\delta$  T cell migration and cytotoxicity: Involvement of  $G_i$  and  $G_s$  protein-coupled signalling pathways. (in press)
- 8. **K.** Truta-Feles, K. Lehmann, M. Lagadari, L. Berod, S. Cubillos, S. Piehler, D. Barz, A.A. Maghazachi, T. Kamradt, and J. Norgauer: A novel role for fMLP as a modulator of human  $\gamma\delta$  T cell migration: Involvement of  $G_i$  protein-coupled signalling pathways. (submitted)
- 9. **K. Truta-Feles**, M. Lagadari, K. Lehmann, L. Berod, S. Cubillos, S. Piehler, D. Barz, A.A. Maghazachi, T. Kamradt, and J. Norgauer: A novel role for C5a as modulators of human  $\gamma\delta$  T cell migration: Involvement of  $G_i$  protein-coupled signalling pathways. (submitted)