Novel anti-inflammatory targets and mechanisms of boswellic acids and celecoxib

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Meiner Familie!

Lieber einsam mit den Freien fallen, als mit den Massen im Triumphe ziehen!

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1 Abbreviations and reagents

1.1 Abbreviations

[³ H]-AA	tritium labelled arachidonic acid
AB	antibody
Α(β)-ΒΑ	3- <i>O</i> -acetyl-β-boswellic acid
(A)KBA	3-O-(acetyl-)-11- keto boswellic acid
ADP / ATP	adenosine diphosphate / adenosine triphosphate
Akt / PKB	protein kinase B
2-APB	2-aminoethyldiphenyl borinate
AUC	area under the curve
(β)-ΒΑ	β-boswellic acid
BAs	boswellic acids
BAPTA/AM	1,2-bis(aminophenoxy)ethane-N,N,N`,N`-tetraacetic acid tetrakis /
	acetoxymethylester
BS	Boswellia serrata
BSA	bovine serum albumin
B.spec.	Boswellia species
$[Ca^{2^+}]_i$	intracellular Ca ²⁺ concentration
CaCl ₂	calcium chloride
CatG	cathepsin G
CD62	cell surface protein 62
CDC	cinnamyl-3,4-dihydroxy-alpha-cyanocinnamate
CGI	cathepsin G inhibitor
CLP	coactosine-like protein
COPD	chronic obstructive pulmonary disease
COX	cyclooxygenase
CpG-DNA/-ODN	cytosine/guanine-rich DNA fragments / -oligodesoxynucleotides
cPLA ₂	cytosolic phospholipase A ₂
CytB	cytochalasin B
DAG	diacylglycerol
DMEM	Dulbecco's modified Eagle medium
DMSO	dimethylsulfoxide

DNA-PK(I)	DNA-dependent protein kinase (inhibitor)
DTT	dithiothreitol
EC ₅₀	effective concentration (50% of stimulation or inhibition)
EDTA	ethylenediaminetetraacetate
EGF	endothelial growth factor
eNOS	endothelial nitric oxide synthase
ERK	extracellular signal-regulated kinase
ETP	endogenous thrombin potential
FCS	fetal calf serum
FLAP	5-LO-activating protein
fMLP	N-formyl-methionyl-leucyl-phenylalanine
FPP(s)	farnesyl-pyrophosphate (synthase)
Fura-2/AM	[1-[2-(5-carboxyoxazol-2-yl)-6-aminobenzofuran-5-oxyl]-2-(2`-
	amino-5`-methyl-phenoxy)ethane-N,N,N`,N`-tetraacetic acid,
	pentaacetoxymethylester]
Glut-(K)BA	3-O-glutaroyl-11-(keto)-boswellic acid
GPCRs	G protein-coupled receptors
GSH	glutathione
HL-60	human leukemia cell line
HLE	human leukocyte elastase
H(P)ETE	hydro(per)oxyeicosatetraenoic acid
HPLC	high performance liquid chromatography
IC ₅₀	inhibiting concentration (50% of inhibition)
IGF-1	insulin-like growth factor
IgG / IgM	immunoglobuline G / M
ΙκΒ	inhibiting factor of kappa B
IL-6	interleukin 6
IKK	IkB-kinase
IMAC	ion metal affinity chromatography
IP ₃	inositoltrisphosphate
IPP	isopentenylpyrophosphate
IPTG	isopropyl-β-D-thiogalactopyranoside
(K)BA-Seph	11-(keto) boswellic acid coupled to Sepharose
LB-medium	Luria Broth base - medium

LNCaP	human prostate cancer cell line
5/12-LO	5/12-lipoxygenase
LPS	lipopolysaccharide
LTs	leukotrienes
M9	minimal medium 9
MALDI-TOF-MS	matrix-assisted laser desorption ionisation - time of flight - mass
	spectrometry
МАРК	mitogen-activated protein kinase
MCF-7	human mamma carcinoma cell line
11-me-BAs	11-methylene-boswellic acids
MFI	mean channel fluorescence intensity
MM6	human monocytic cell line
MQ	Milli Q water
MS	mass spectrometry
mTOR	mammalian target of rapamycin (kinase)
MWCO	mol weight cut off
ΝΓκΒ	nuclear factor k B
NMR	nuclear magnetic resonance spectroscopy
NSAIDs	non-steroidal anti-inflammatory drugs
OD	optical density
p12-LO	platelet-type 12-lipoxygenase
PAC-1	cell surface protein
PAF	platelet-activating factor
PAR-4	protease-activated receptor 4
PBS	phosphate-buffered saline
PDGF	platelet-derived growth factor
PDK-1	3-phosphoinositol-dependent kinase
PGC buffer	PBS plus 1 mg/ml glucose plus 1 mM CaCl ₂
PG	prostaglandin
PI3K	phosphatidylinositol 3 kinase
PIP ₃	phosphatidylinositol-trisphosphate
PKA, PKB, PKC	protein kinase A, B, C
PLC	phospholipase C
PMNL	polymorphonuclear leukocytes

PMSF	phenylmethylsulfonylfluoride
PRP	platelet rich plasma
pTyr	phosphotyrosine
r12-LO / his12-LO	recombinant 12-lipoxygenase / 6-his-tagged 12-lipoxygenase
Rap 1b	Ras-related protein 1b
RBL-1	rat basophilic leukemia cell line
RFU	relative fluorescence units
ROS	reactive oxygen species
RT	room temperature
RTK	receptor tyrosine kinases
SDS	sodium dodecylsulfate
SDS-b	2x SDS loading buffer
SDS-PAGE	SDS-polyacrylamide gel electrophoresis
Seph	EAH-Sepharose 4B
Suc-AAPF-pNA	N-succinyl-alanyl-alanyl-prolin-phenylalanine-p-nitroanilide
TBS	tris-buffered saline
ТСМ	traditional Chinese medicine
TFA	trifluoro acetic acid
TG	thapsigargin
TGFβ	transforming growth factor beta
ΤΝFα	tumour necrosis factor alpha
TRAP	thrombin receptor-activating peptide
TXA_2	thromboxane A ₂
WB	western blot
w/o	without

1.2 Reagents

- Boswellic acids (BAs) were prepared by Dr. J. Jauch¹ (Saarbrücken)
- *Boswellia serrata* (BS) extracts were generous gifts from Burg-Apotheke, Königstein, Germany and Pharmasan GmbH, Freiburg, Germany
- Celecoxib, etoricoxib, rofecoxib were sythesized by WITEGA Laboratories Berlin-Adlerhof GmbH, Germany
- Insulin was a generous gift from Aventis Deutschland GmbH, Frankfurt, Germany
- 5-LO antibody was provided by Dr. O.Rådmark, Stockholm, Sweden
- 12-LO antibody was provided by Dr. C.Funk, Kingston, Canada
- TGF β was prepared from human platelets as described ²
- BAPTA/AM, Fura-2/AM were from Alexis Corp, Lausen, Switzerland
- Ampicillin, Coomassie brilliant blue G250, IPTG, leupeptin, oxalacetic acid, were from Applichem, Darmstadt, Germany
- Antibodies against CD41-PC7, CD62-PE, and PE-labelled isotype IgG1 control were purchased from Beckman Coulter, Krefeld, Germany
- Antibodies against PAC1-FITC and FITC-isotype IgM and matrigel were from Becton Dickinson Biosciences, Freiburg, Germany
- CatG antibody, CDC and fluorescence dye secondary antibodies (Rockland Inc.) were from Biomol GmbH, Hamburg, Germany
- Bradford and Lowry reagents were from Biorad, München, Germany
- α-Cyano-4-hydroxycinnamic acid was purchased from Bruker Daltonics Inc.,
 Manning Park Billerica, MA, USA
- Alendronate, cathepsin G inhibitor I, chymotrypsin, DNA-PK inhibitor, elastase, PP2, PP3; Protein kinase assay kit (non-radioactive); staurosporine, SU6656 and tryptase were from Calbiochem, Schwalbach/Ts., Germany
- 5-HETE, LTB₄ and zileuton were from Cayman chemical company, Ann Arbor, MI, USA
- Antibodies against Akt and pAkt, pBad, DNA-PK, Ras; Akt1 kinase and Akt kinase assay kit were from Cell Signaling / New England Biolabs, Frankfurt, Germany
- Boc-Ala-Ala-Nva-SBzl, Cathepsin G, elastin-sepharose, N-Suc-AAPF-pNA, proteinase 3 were from Elastin Products Company Inc., Owensville, MS, USA
- α-Amyrin and ursolic acid were from Extrasynthèse, Genay, France

- [³H]-arachidonic acid, [³²P]-ATP, CM Sephadex C-50, EAH-Sepharose 4B, [¹⁴C]isopentenylpyrophosphate, TNFα release kit, nitrocellulose and PVDF membrane were from GE Healthcare, Amersham Bio-Sciences, Freiburg, Germany
- Plastic/PS-materials were from Greiner bio-one GmbH, Frickenhausen, Germany
- DMEM, glutamine, LB, LPS, penicillin/streptomycin, RPMI were from Invitrogen GmbH, Karlsruhe, Germany
- WB blocking buffer, fluorescence dye secondary antibodies were from LiCor Biosciences GmbH, Bad Hombug, Germany
- Grams violet, β-mercaptoethanol, methanol, trifluoracetic acid were from Merck Chemicals, Darmstadt, Germany
- Collagen reagent Horm was from Nycomed, Singen, Germany
- CpG ODN 1018 were from Operon biotechnologies GmbH, Köln, Germany
- Nycoprep and sodium pyruvate were from PAA, Pasching, Austria
- Optiphase HiSafe scintillation liquid was from Perkin-Elmer Deutschland, Rodgau-Jügelsheim, Germany
- DNA-PK, DNA-PK peptide substrate were from Promega Deutschland GmbH, Mannheim, Germany
- Ni²⁺-NTA resin was from Qiagen GmbH, Hilden, Germany
- Trypsin (sequencing grade) was from Roche, Mannheim, Germany
- Antibodies against β-actin, cPKC, pTyr were from Santa Cruz Biotechnology, Heidelberg, Germany
- ATP, glucose, GSH, PBS were from Serva, Heidelberg, Germany
- ADP, arachidonic acid, ASS, ATP-agarose, N-α-benzoyl-DL-arginine-pNA, bromophenolblue, BWA4C, chymase, cytochalasin B, dextran, diclofenac, DMSO, DTNB, DTT, farnesol, fatty acid free BSA, FCS, fMLP, geranylpyrophosphate, γglobulin, ionomycin, ionophore A-23187, isopentenylpyrophosphate, lysozyme, Nmethoxysuc-AAPV-pNA, m-3M3FBS, n-octyl-β-glucopyranoside, non essential amino acids, NP-40 (Igepal), PAF, PEG 4000, PGB₁, phosphatidylcholine, SDS, soybean trypsin inhibitor, sucrose, thapsigargin, thrombin, triethanolamine, trypan blue, Tween 20, Tx-100, U-73122, U-73343, wortmannin, zileuton and all other chemicals were purchased in analytical grade from Sigma-Aldrich Chemie, Steinheim, Germany
- Vivaspin cut-off columns were from Vivascience AG, Hannover, Germany

2 Introduction

2.1 Boswellic acids in traditional medicine and treatment of chronic diseases

Frankincense, the gum resin of the *Boswellia serrata* (BS) tree (fig.211), was known to ancient civilisations and is still used for ritual purposes in the Catholic Church and traditional ceremonies in Northern Africa. The first documentation of incense was the Papyrus Ebers written at the time of Pharao Amenophes I around 1500 BC³. The gum resin extracts of BS have traditionally been applied in folk medicine for centuries to treat various chronic inflammatory diseases, and experimental data from animal models and clinical studies on humans confirmed an anti-inflammatory potential of lipophilic BS extracts, for review see ⁴⁻⁷. Detailed analysis of the ingredients of these extracts revealed that the pentacyclic triterpenes boswellic acids (BAs) (fig.212) possess pharmacological activities which may be responsible for the respective anti-inflammatory properties ⁶. Among the pentacyclic triterpenes, derivatives of tirucallic acid, dehydro boswellic acid and lupeolic acid are major ingredients with pharmocological effects.

A number of relevant targets of BAs including 5-lipoxygenase (5-LO)⁸, platelet-type 12-LO (p12-LO)⁹, human leukocyte elastase (HLE)¹⁰, toposiomerase I and II¹¹ and I κ B kinases (IKK)¹² were identified. BAs induce apoptosis in tumour cells, ^{13, 14} accompanied by decreased



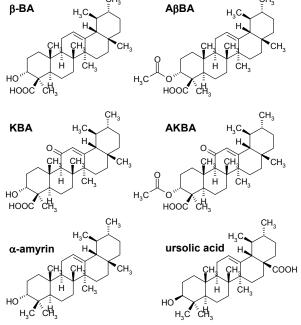


figure 211: Leaves and blossoms of the boswellia serrata tree ¹⁵

figure 212: Structures of the four major β -boswellic acids and two related pentacyclic triterpenes

extracellular signal-regulated kinase (ERK) phosphorylation ¹⁶ and enhanced caspase activity ¹⁴. BAs differentially interfere with signalling molecules and pathways related to inflammatory processes and tumour growth including Ca²⁺-influx, mitogen-activated protein kinases (MAPK) and Akt ^{17, 18}. Colon cancer proliferation is inhibited by BAs due to a p21-dependent pathway ¹⁹. All these effects are dependent on the structure of the BAs, since 11-keto-BAs have different targets than 11-me-BAs.

Several functional effects of BS extracts on animals and humans have been described; however, the underlying molecular mechanisms still seem to remain unclear. The extracts reduce carrageenan-induced oedema and chemical-induced colitis in rats and mice²⁰ and several studies on humans investigated the effectiveness of BS extracts in chronic inflammatory diseases like arthritis, Crohn's disease and ulcerative colitis. In a placebocontrolled (randomized, double-blind) study of patients with osteoarthritis, all patients (n =30) receiving drug treatment reported a decrease in knee pain, increased knee flexion and increased walking distance, which is statistically significant and clinically relevant ²¹. The effect of BS extract (3 x 350 mg/d) was compared to sulfasalazine treatment (3 x 1 g/d) in patients suffering from ulcerative colitis grade II and III. All clinical parameters improved and 82% of treated patients went into remission; in the case of sulfasalzine, the remission rate was 75%²². Studies in patients with Crohn's disease led to similar results: BS extracts were either as effective or even more effective than mesalazine or sulfasalazine ^{23, 24}. BS extracts (H15[®]) were successfully tested for cancer treatment in palliative therapy of patients with brain tumours and radiochemotherapy-related leukoencephalopathy. Oedema were reduced and all patients with leukoencephalopathy improved clinically for several months ²⁵. Due to these results, the medicinal product H15[®] (containing BS extract) was designated by the EMEA as an orphan drug and is approved in part of Switzerland for the treatment of peritumoral brain oedema.

It is generally believed that the anti-inflammatory properties of BAs are mainly due to interference with 5-LO product synthesis ^{4, 5} and also to interference with the nuclear factor κ B (NF κ B) signalling pathway ¹². Moreover, BAs modulate central signal transduction pathways such as Ca²⁺ signalling and MAPK activation ¹⁷, relevant for diverse functions of inflammatory cells. Despite these numerous putative target molecules of (11-keto-)BAs, the pharmacological relevance of the interactions with these targets is a matter of debate. Most of these interactions occur at relatively high concentrations that are far above the plasma levels of (11-keto-)BAs (<1 μ M) reached following oral application of a standard dosage of *B. spec*

extracts, and are considered not relevant *in vivo*^{26, 27}. In summary, direct interactions of BAs with high-affinity targets that are pharmacologically relevant *in vivo* still remain to be shown.

2.2 Cathepsin G; a neutrophil serine protease and its role in disease

Cathepsin G (CatG) is a neutral serine protease with a molecular weight (MW) of 25.8 kDa, mainly expressed in pro-inflammatory leukocytes (i.e. polymorphonuclear leukocytes (PMNL) and macrophages), where it is stored in azurophilic granules in their active form and released upon exposure to inflammatory stimuli by degranulation ^{28, 29}. Neutrophil infiltration, which is mediated by CatG, is a common pathological feature in acute inflammatory disorders and is necessary for phagocytosis and eradication of microorganisms ³⁰. It was demonstrated that CatG is crucial for resistance against infection with Staphylococcus aureus ³¹. In addition, CatG contributes to protection against certain fungal infections ^{32, 33}. However, neutrophils are also present in inflammatory diseases which are not caused by microorganisms, and active CatG is detectable at sites of inflammation ³⁰. It is known that ligation of cell-surface receptors, such as β_2 -integrins, activates a cascade of cytoplasmic signalling molecules and triggers an influx of calcium ions (Ca^{2+}) . The increase in the intracellular concentration of Ca^{2+} ([Ca^{2+}]_i) mediates the fusion of granules to the plasma membrane ³⁴. Once released, neutrophil serine proteases are potentially fully active because they function optimally in a neutral environment. Biochemical studies indicate that CatG preferentially hydrolyses peptide bonds after aromatic amino-acid residues³⁵. Following release into the plasma, it cleaves extracellular matrix proteins including laminin, proteoglycans, collagen, fibronectin and elastin, implying a role in local destruction of connective tissue at sites of injury ³⁶. Under normal circumstances, this extracellular protease activity is inhibited by serpins, which are endogenous serine protease inhibitors that form complexes with these extracellular proteases. Therefore, anti-chymotrypsin binds to and inhibits CatG^{35, 37}. In addition, neutrophil elastase and CatG are also inhibited by secretory leukocyte protease inhibitor ³⁷. It was recognized several years ago that serpins not only inhibit proteolysis but also prevent recruitment of inflammatory cells ^{38, 39}. Although the role of CatG in host defence is acknowledged, neutrophil serine proteases have also been implicated in various non-infectious, inflammatory processes. In support of this idea, inhibition of neutrophil serine proteases has been shown to reduce neutrophil infiltration and neutrophil-mediated injury ⁴⁰⁻⁴², and mice with a CatG

deficiency are resistant to experimental arthritis ^{43, 44}. Based on these properties, CatG is implicated in many inflammatory disorders including asthma, chronic obstructive pulmonary disease (COPD), emphysema, reperfusion injury, psoriasis and rheumatoid arthritis ³⁰. In addition, CatG may stimulate platelets via the protease-activated receptor (PAR)-4 for aggregation and secretion ⁴⁵. Truncation of chemokines and cytokines by CatG increased its monocyte chemotactic activity about 1,000-fold ⁴⁶. Another study by Sun et al. demonstrated that CatG is a chemotactic agonist for the G protein-coupled formyl peptide receptor (FPR) ⁴⁷. CatG binds FPR with low affinity, inducing a modest Ca²⁺ flux and weak activation of MAPK. However, the stimulation with CatG of cells that express FPR induced the translocation of protein kinase C ξ (PKC ξ) from the cytoplasm to the plasma membrane, which is essential for FPR-dependent chemotaxis. Therefore, extracellular CatG might contribute to leukocyte recruitment *in vivo* by binding to FPR on neutrophils and monocytes ⁴⁸. The agonistic activity of CatG on FPR might not depend on proteolysis, since no small and soluble molecules were released during the interaction of the enzyme with the receptor ^{48, 49}.

2.3 DNA-dependent protein kinase and Akt

Human DNA-dependent protein kinase (DNA-PK) and the protein kinase B (Akt/PKB) are important signal transducers for growth, survival and apoptosis⁵⁰⁻⁵². DNA-PK is a nuclear serine/threonine protein kinase that, when activated by DNA ⁵², phosphorylates several DNA-binding substrates, including the tumour suppressor protein p53 ⁵⁰. DNA-PK and Akt signalling pathway has emerged as a critical mediator of diverse cellular processes including metabolism, gene expression, migration, angiogenesis, proliferation and cell survival. Since Akt is tightly controlled, the consequences of its deregulation have been implicated in the development of cancers and diabetes ^{53, 54}. The activity of Akt is markedly stimulated in a phosphatidylinositol 3-kinase (PI3K)-dependent manner. Akt is phosphorylated at Thr-308 by the PI3K-activated 3-phosphoinositol-dependent kinase (PDK1). However, active Akt does not only depend on Thr-308 phosphorylation, but is also phosphorylated at Ser-473. The responsible kinase remained unclear for a long time, but was recently identified as DNA-PK (fig. 213) ⁵⁵.

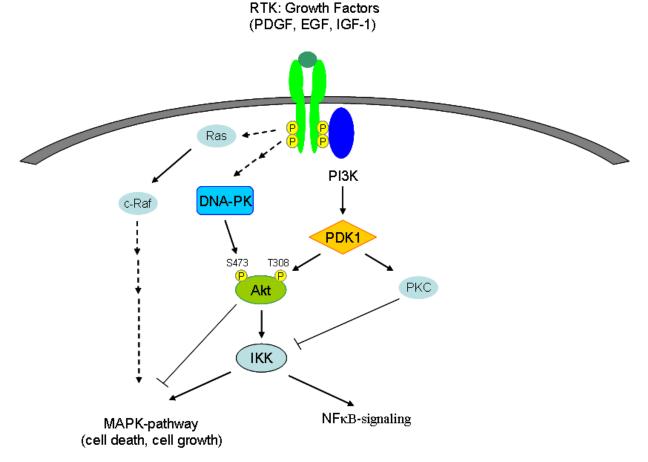


figure 213: Model for Akt (PKB) activation by upstream kinases. Upon stimulation by growth factors, PI3K is activated, which in turn generates the second messenger PIP_3 to recruit Akt and PDK1 to the membrane lipid rafts, where Akt is subsequently phosphorylated on Thr-308 and the hydrophobic motif Ser-473 by PDK1 and DNA-PK, respectively.

It has been shown that efficient DNA repair requires growth factor signalling ⁵⁶ and that this effect may be due to the physical association of DNA-PK with epidermal growth factor receptor ⁵⁷. DNA-PK is activated upon DNA damage by UV irradiation ⁵⁸ and bacterial products such as lipopolysaccharides (LPS) and CpG-DNA ⁵⁹, and these signals or factors also activate Akt ⁵⁹. Induction of apoptosis by cisplatin was explained by a decrease in DNA-PK activity through proteolytic degradation of DNA-PK ⁶⁰; in parallel, Akt activity and Ser-473 phosphorylation are also significantly inhibited by cisplatin treatment ⁶¹. A number of oncogenes and tumour suppressor genes that function upstream of Akt have been found to influence cancer progression by regulating Akt through increasing PI3K activity, including activated Ras and Bcr / Abl ⁶². PTEN has also been identified as a human tumour suppressor and loss of PTEN correlates with increased Akt activity ⁵⁴.

The survival factor Bad is a protein downstream of Akt which regulates (via co-recruitment of other proteins like Bcl-2 or 14-3-3) cell survival or apoptosis ⁵³. It is possible to determine Akt activity *in vivo* by measuring phosphorylation of Bad at residue Ser-136 (**fig. 214**). The pathways leading to DNA-PK/PI3K-Akt activation by LPS and CpG-DNA still remain poorly defined. Furthermore, DNA-PK and Akt are involved in activation of IKK and NF κ B ^{18, 63}, though cancer and inflammation are hardly influenced as well as bacterial infections. In conclusion, Akt is an important key mediator in survival / apoptosis mechanisms.

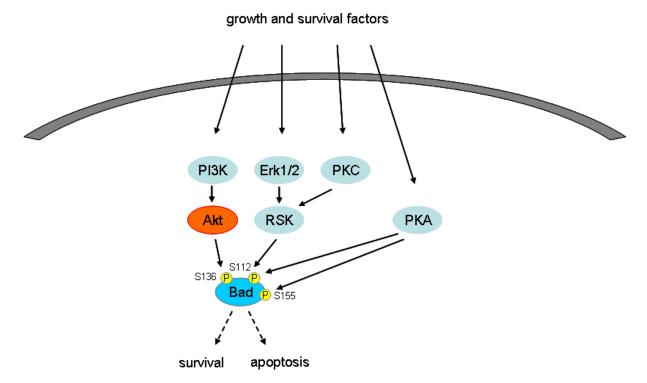


figure 214: **Downstream signalling of Akt**: Bad is phosphorylated by Akt on Ser-136 residue. Correcruitment of other proteins (14-3-3 or Bcl-xL) is related to cell survival or apoptosis.

2.4 Cellular signalling: Ca²⁺-homeostasis and MAP kinases

Many cellular processes are mediated by changes in $[Ca^{2+}]_i$. Stimulation of cells causes depolarisation of the plasma membrane and an opening of Ca^{2+} channels. The Ca^{2+} concentration between cytosol (10-100 nM) and extracellular space (1.5 – 2 mM) differs by factor 10⁴ to 10⁵ ⁶⁴. $[Ca^{2+}]_i$ does not only rise by opening plasma membrane channels, intracellular Ca^{2+} stores like sarcoplasmatic reticulum (SR) or endoplasmatic reticulum (ER) may also release Ca^{2+} , contributing to substantial elevations in $[Ca^{2+}]_i$. Rapid influx of Ca^{2+} acts as second messenger and influences important signalling cascades like the MAPK pathway. Because of cytotoxicity of high $[Ca^{2+}]_i$, Ca^{2+} pumps like Ca^{2+} -ATPase rapidly remove Ca^{2+} into intracellular storage sites or export it into extracellular lumen.

BAs influence Ca^{2+} -homeostasis, but the functional properties and the potencies of the BAs depend on their structure (**fig. 212**), in particular on the absence or presence of the 11-keto group. Indeed, 3-*O*-acetyl-11-keto- β -BA (AKBA) is frequently the most effective analogue ⁴, ^{9, 65, 66}, although in some instances the 11-keto-free BAs possess superior efficacy, depending on the target/effect but also on the respective cell type ^{11, 13, 17, 67} However, in PMNL, KBA and AKBA elevated [Ca²⁺]_i, whereas the 11-methylene BAs failed in this respect ⁶⁶. Finally, in monocytes, AKBA but not 11-methylene BAs suppressed intracellular Ca²⁺ levels ¹⁷.

Platelets play a critical role in the homeostasis of the cardiovasculature but also in the inflammatory pathophysiology ⁶⁸. Activation of platelets may lead to secretion of granular contents and release of arachidonic acid (AA), shape change, adhesion, and aggregation ⁶⁹. Thrombin or collagen are strong agonists of platelets, whereas platelet-activating factor (PAF), adenosine diphosphate (ADP), serotonin, or thromboxane (TX)A₂ require autocrine stimulation for the entire platelet response ⁶⁹. Soluble platelet agonists, such as thrombin, ADP, PAF, or TXA₂, typically activate specific G protein-coupled receptors (GPCRs), leading to the activation of certain phospholipase (PL)C isoforms that eventually results in elevation of [Ca²⁺]_i and activation of central signalling protein kinases (PK), including phosphatidylinositol 3-kinase (PI3K)/Akt, PKC isoforms and MAPK ^{68, 70-72}. Besides these effects in platelets, the four major BAs influence several signalling pathways in other cell types like MonoMac6 or HL-60 cells ¹⁷.

MAPK cascades are key signalling pathways involved in the regulation of normal cell proliferation, survival and differentiation. Aberrant regulation of MAPK cascades contributes to cancer and other human diseases. The ERK pathway in particular has been the subject of intense research for inhibitors for the treatment of cancer ⁷³. ERK is a downstream component of an evolutionary conserved signalling module that is activated by the Raf serine/threonine kinases (**fig. 213**) ⁷³. Raf, MEK and ERK are downstream effectors of Ras and receptor tyrosine kinases (RTK) like EGFR, which is mutationally activated and / or overexpressed in a wide variety of human cancers. RTK typically regulate cell growth and survival via several pathways, including Ras-Raf-MEK-ERK and PI3K-Akt-IKK cascades⁷⁴. Not only RTK inhibitors are important for the treatment of cancers, modulation and inactivation of downstream targets could also facilitate the cure of many diseases. BAs interact with MAPK, an interesting finding that could possibly lead to new approaches in (cancer) therapy.

2.5 COX and LOs: Key enzymes in eicosanoid biosynthesis

Inflammation is one of the most important processes in host defense; however, it often progresses to painful or chronically harmful diseases that require pharmacological treatment. The inflammatory response involves many effector mechanisms which produce a multiplicity of vascular and cellular reactions. One of the major findings since the discovery of the role of histamine in vascular processes of inflammation at the beginning of the twentieth century, is that many chemical mediators are involved in activating and coordinating various aspects of the inflammatory process ^{75, 76}. Now we know that vasodilatation, increased microvascular permeability, chemotaxis, cellular activation, pain and finally repair are mediated by the local production and release of several specific mediators. The arachidonic acid (AA) derivatives postaglandins (PGs), thromboxanes (TXs) and leukotriens (LTs), together with cytokines, oxygen and possibly nitrogen radicals, play a pivotal role. The non-steroidal anti-inflammatory drugs (NSAIDs), mainly the cyclooxygenase (COX) inhibitors, which have been used since the introduction of acetylsalicylic acid in 1899, share many pharmacological properties (and side effects) and are the main drugs used to reduce the unwanted consequences of inflammation ⁷⁶⁻⁷⁸.

As a result of proinflammatory stimuli or injuries, cytosolic phospholipase (cPL) A₂ releases AA from cell membrane phospholipids. AA is the substrate of either lipoxygenases (LOs) or COX. COX-1/2 form the cyclic peroxide PGH₂ as an intermediate product which is further converted to PGs and TXs. In addition to PGs derived from the COX pathways, LTs are formed by LOs from AA that exert pivotal biological functions. For LT biosynthesis, AA is first metabolized by 5-LO to LTA₄ (**fig. 215A**). The unstable LTA₄ is the precursor of the bioactive LTs such as LTB₄ and the cysteinyl-LTs C₄, D₄, and E₄. Interestingly, next to inflammatory and allergic diseases, 5-LO products play a crucial role in cancer development and atherosclerosis ⁷⁹. Polymorphonuclear leukocytes (PMNL) and monocytes/macrophages are the major cells capable of synthesizing LTs and express high amounts of 5-LO. These cells are major players in chronic inflammatory diseases. In the treatment of asthma, zileuton is the only 5-LO inhibitor that came into the market (USA). New potent and biocompatible inhibitors would open a new field in therapy of asthma, several cancers and atherosclerosis, because 5-LO is a major target in chronic inflammation. For example, 5-LO is overexpressed in prostate cancer, and other malignant (compared with benign) tumour tissues⁸⁰.

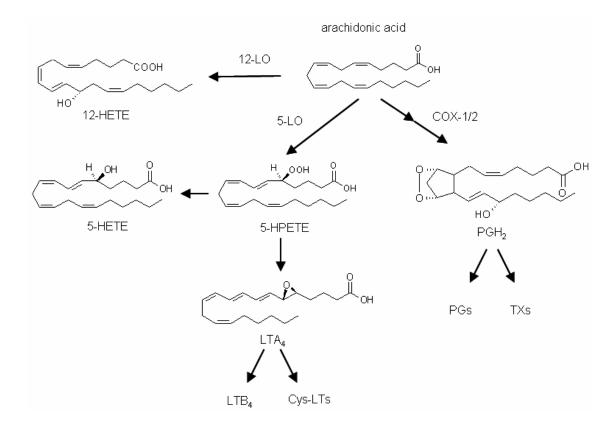
The activity of 5-LO in the cell is tightly controlled. The amount of free AA, released by cPLA₂, and its accessibility to 5-LO are determinants of LT biosynthesis. Thus, stimuli

capable of inducing leukotriene formation (e.g. fMLP, PAF and Ca^{2+} -ionophores) cause activation of 5-LO⁸¹.

The conversion of endogenously provided AA to LTA₄ occurs at the nuclear membrane, and it seems that leukotriene formation can be particularly prominent at this locus⁸². On cell stimulation, 5-LO migrates from a soluble compartment to the nuclear membrane. Membrane-bound FLAP might facilitate the transfer of AA to 5-LO. In cells lacking FLAP, or where FLAP is pharmacologically inhibited, transformation of endogenous AA by 5-LO is efficiently blocked ⁸². 5-LO is phosphorylated *in vitro* by the protein kinases p38 MAPK-regulated MAPKAPK-2/3, ERK1/2, CaMKII and PKA ⁸¹. Phosphorylation does not, however, seem to increase 5-LO activity *in vitro*, but PKA-mediated phosphorylation can inhibit 5-LO activity⁸³.

Because catalysis by 5-LO requires oxidation of Fe^{2+} to the active Fe^{3+} state by lipid hydroperoxides, the redox tone is an important parameter of cellular 5-LO activity. Conditions that promote lipid peroxidation, such as formation of reactive oxygen species by phorbol 12-myristate 13-acetate, addition of peroxides, inhibition of glutathione perxidase (GPx) enzymes and depletion of glutathione, upregulate 5-LO product synthesis, whereas reduction of peroxides by GPx-1 and GPx-4 suppresses 5-LO product formation⁸¹. Oxidative stress also activates p38 MAPK; thus, an increase in peroxide tone might activate 5-LO in the cell by promoting phosphorylation of 5-LO ^{84, 85}.

Leukotriene synthesis was first observed after cell stimulation by a Ca²⁺ ionophore ⁸⁶. Considerably lower concentrations of Ca²⁺ (~200 nM) seem to be sufficient for substantial activation of 5-LO in intact cells ⁸⁷, as compared with purified 5-LO (which requires 1–10 μ M Ca²⁺), indicating that cellular events and context modulate the Ca²⁺ requirement. Activation of 5-LO in intact cells, however, might not necessarily involve Ca²⁺. When exogenous AA was provided to intact human neutrophils, substantial 5-LO product formation occurred even when extra- and intracellular Ca²⁺ had been depleted by chelating agents such as EDTA and BAPTA/AM ^{79, 88}.



В

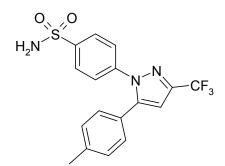


figure: 215: (A) Arachidonic acid metabolism: Arachidonic acid is released in inflammation and injuries by phospholipases and metabolized by lipoxygenases (5-LO; 12-LO; 15-LO) and cyclooxygenases (COX-1; COX-2) to leukotrienes (LTs) and prostaglandins (PGs), respectively. (B) Structure of celecoxib NSAIDs are used for the treatment of pain, fever and inflammation. Traditional NSAIDs inhibit both isoforms of COX which are key enzymes in the conversion of AA to PGs (**fig. 215A**). The housekeeping enzyme COX-1 is primarily responsible for maintenance of homeostasis (gastric and renal protection, platelet function), whereas COX-2 is mainly induced in response to growth factors and various cytokines at sites of inflammation and tumour growth ^{89, 90}. Thus, COX-2-selective drugs, such as celecoxib, etoricoxib, valdecoxib and rofecoxib, showed anti-inflammatory, analgesic and antipyretic efficacies similar to those of traditional NSAIDs, but displayed a lower incidence of symptomatic ulcers and ulcer complications ⁹¹.

Concomitantly with the introduction to the market, the COX-2-selective inhibitor, celecoxib (fig.215B) was shown to significantly reduce polyp formation and polyp size in patients with familial adenomatous polyposis at rather high doses $(2 \times 400 \text{ mg/d})^{92}$. Clear evidence that COX-2-inhibition alone is not responsible for the anticarcinogenic effects of celecoxib has been obtained from cell culture and animal studies showing antiproliferative effects of this drug in human colon carcinoma cells irrespective of COX-2 expression ^{93, 94}. Several COX-2independent molecular mechanisms with unclear in vivo relevance were identified and assumed to contribute to the anticarcinogenic effects of celecoxib, for instance 3phosphoinositide-dependent kinase-1 (PDK-1, $IC_{50} = 48 \ \mu M$ for celecoxib) and its downstream target Akt kinase. It was also shown that an inhibition of the endoplasmatic reticulum Ca²⁺-ATPase (IC₅₀ \approx 35 μ M) is associated with the induction of apoptosis ⁹⁵⁻⁹⁸. However, the impact of these findings was discussed controversially, since a significant discrepancy exists between the concentrations of celecoxib found in the plasma of patients under tumour therapy $(3-8 \mu M)^{99}$ and those concentrations needed to affect intracellular non-COX-2 targets (\geq 30 µM, *in vitro* assays). Hence, the exact molecular mechanisms responsible for the effects of celecoxib at higher doses are still a matter of intensive research.

2.6 Aim of this work

Extracts of B. spec. containing bioactive BAs have proven to be effective in the treatment of chronic inflammatory diseases such as rheumatoid arthritis and ulcerative colitis and have been applied in the Indian ayurvedic medicine and traditional Chinese medicine (TCM) for centuries. Due to a growing public interest in Germany in natural (plant-derived) compounds as safe and well-tolerable drugs during the last decades, the scientific community is increasingly focussing on the elucidation of the mechanisms by which these remedies act on the molecular and cellular level. This interest is substantiated by recent clinical trials demonstrating beneficial effects of BS extracts in several inflammatory diseases such as Crohn's disease, ulcerative colitis and certain types of cancer. However, the utilised extracts consist of a plethora of different compounds. Although BAs represent the largest fraction and are considered to be important pharmacological principles, application of this mixture in cellular experiments for the purpose of identifying molecular targets is not feasible, due to the heterogeneity of the extract and the diversity of the ingredients. Thus, investigations on the cellular level should start with the purified BAs. Research started 15 years ago, and has yielded a number of relevant target proteins, including 5-LO⁶. AKBA was regarded as the most potent BA but the other BAs are also active in several experimental settings. Modulation of LT formation, which are key mediators of inflammation with additional implications in carcinogenesis, was described by a series of studies. Other targets (e.g. HLE, IKK, NFkB), related to cell proliferation and apoptosis, were also identified ^{18, 100}. None of these findings, however, can presently provide a satisfying explanation for the anti-inflammatory effects of the extracts observed in clinical settings and in traditional folk medicine because of the high effective concentrations required. Hence, this thesis was designed to find pharmacologically relevant targets of BAs. Haematopoietic cells (leukocytes, platelets) and cancer cell lines (MCF-7, LNCaP, MM6, HL-60) were used as model systems. Investigations of the cellular and molecular effects of BAs were started in this group by Oliver Werz¹⁰¹. An initial thesis focussing on the effects of BAs on neutrophils was provided by Anja Altmann¹⁰². Based on these initial findings ⁶⁶, studies were continued and expanded toward other related cell types (monocytic cells, platelets) in Daniel Poeckel's thesis ¹⁰³. He examined a putative correlation of intracellular Ca²⁺ mobilisation and ERK1/2 activation by 11-keto-BAs ⁶⁶ to the generation of ROS, release of AA and formation of eicosanoids¹⁰⁴.

In this thesis, the antagonistic effects of BAs on Ca²⁺ mobilisation and functional responses in blood cells were investigated. The results with human platelets further underscored the interactions of BAs with platelet-type 12-LO. A construct of immobilised BAs was utilised to identify putative target proteins in cell lysate supernatants via a pulldown approach (protein fishing) and MS analysis. The functional consequences of the interactions of BAs with the identified proteins were studied. The most interesting proteins found were cathepsin G (CatG), DNA–dependend protein kinase (DNA-PK), Akt, PKC and farnesyl pyrophosphate synthase (FPPs).

Celecoxib is a well-known selective COX-2 inhibitor. Celecoxib was introduced to the European market in 1998 as Celebrex[®] and quickly captured the NSAIDs market. After rofecoxib (Vioxx[®]) was withdrawn, news surrounding the well-tolerated drug celecoxib quieted down. Several patients with colon carcinoma or strong polyp formation taking celecoxib strongly benefited from it and polyps or tumours regressed, but that occurred independently of COX inhibition ⁹⁴. Previous work suspected a role of leukotrienes in the formation of colon carcinoma, and this was the reason to test celecoxib for interference with 5-lipoxygenase *in vitro* and *in vivo* ⁹⁴.

3 Methods

3.1 Cells and cell culture

3.1.1 Cell culture

All cell lines were cultured in incubators (WTB Binder Labortechnik, Tuttlingen, Germany) at 37°C, 5% CO₂ and saturated humidity. The cultures were seeded at a density of 2 x 10^5 cells per ml. Cells were harvested by centrifugation (200 x g, 10 min at room temperature) and washed once in PBS, pH 7.4. To exclude toxic effects of BAs during various incubation periods, the viability of cells was analysed by means of trypan blue exclusion. Incubation with 30 μ M of BAs at 37°C for 30 min caused no significant change in the number of viable cells.

3.1.2 Mono Mac 6 cells

MM6 cells (monocyte like cell line) were obtained from Dr. Ziegler-Heitbrock (Munich, Germany) and maintained in RPMI 1640 medium with glutamine supplemented with 10% fetal calf serum, 100 μ g/ml streptomycin, 100 U/ml penicillin, 1 mM sodium pyruvate, 1x nonessential amino acids, 1 mM oxalacetic acid and 10 μ g/ml insulin. MM6 cells were treated with 2 ng/ml TGF β and 50 nM calcitriol for 4 days in order to obtain differentiated cells ¹⁰⁵.

3.1.3 LNCaP / HL-60

LNCaP (human prostate cancer cells) and HL-60 (human leukemic cells) were maintained in RPMI 1640 medium with glutamine supplemented with 10% fetal calf serum, 100 μ g/ml streptomycin, 100 U/ml penicillin.

3.1.4 MCF-7

MCF-7 (human mamma carcinoma cells) were maintained in DMEM high glucose medium supplemented with 10% fetal calf serum, 100 μ g/ml streptomycin, 100 U/ml penicillin and 1 mM sodium pyruvate.

3.1.5 RBL-1

RBL-1 (rat basophilic leukemia cells) were maintained in RPMI 1640 medium with glutamine supplemented with 10% fetal calf serum, 100 μ g/ml streptomycin, 100 U/ml penicillin, 1 mM sodium pyruvate, 10 mM HEPES and 1x nonessential amino acids.

3.2 Isolation of human PMNL (polymorphonuclear leukocytes) from venous blood

Human PMNL were freshly isolated from leukocyte concentrates obtained from St. Markus Hospital (Frankfurt, Germany). In brief, venous blood was taken from healthy adult donors and subjected to centrifugation at 4,000 x *g* for 20 min at 20°C for preparation of leukocyte concentrates. PMNL were promptly isolated by dextran sedimentation, centrifugation on Nycoprep cushions (PAA Laboratories, Linz, Austria), and hypotonic lysis of erythrocytes ^{85, 106, 107}. PMNL (7.5 x 10⁶ cells/ml; purity > 96–97 %) were finally resuspended in PBS pH 7.4 plus 1 mg/ml glucose and 1 mM CaCl₂ (PGC buffer).

3.3 Isolation of human platelets from venous blood

Platelets were isolated from supernatants (800 x g, 10 min, RT) after centrifugation of leukocyte concentrates on Nycoprep cushions (see above) to obtain platelet rich plasma (PRP). PRP was then mixed with PBS pH 5.9 (3:2, v/v), centrifuged (2,000 x g, 15 min, RT) and the pelleted platelets were resuspended in PBS pH 5.9 / 0.9% NaCl (1:1, v/v), washed by centrifugation (2,000 x g, 10 min, RT) and finally resuspended in PBS pH 5.9. For incubations with solubilized compounds, ethanol or DMSO was used as vehicle, never exceeding 1 % (v/v).

3.4 Human in vitro whole blood assay.

Aliquots of freshly heparinized human blood (450 µl) obtained from healthy male and female informed volunteers were pre-incubated with the drugs or vehicle (DMSO) for 30 min at 37°C. Formation of 5-LO products was initiated by the addition of Ca²⁺-ionophore dissolved in 50 µl autologous plasma to obtain a final concentration of 20 µM A23187 (final DMSO concentration was < 1 %, final volume 0.5 ml). The reaction was terminated after 15 min by rapid cooling of the plate on ice. Then, the samples were centrifuged at 1000 x g and 4°C for

15 min and eicosanoid concentrations (5-HETE, LTB₄, 12(S)-HETE, 15(S)-HETE and PGE₂) in the plasma supernatant were analyzed using LC/MS-MS. LC-MS/MS analysis was performed on a API 4000 triple quadrupole mass spectrometer (Applied Biosystems, Darmstadt, Germany). Linearity of the calibration curve was proven from 0.5 to 2500 ng/mL for each eicosanoid. Mean accuracy of the assay was found to be 99.9 +- 3.25% for LTB₄, 99.85 +- 4.8% for 5-HETE, 100.2 +- 4.8% for 12-HETE and 99.76 +- 4.4% for 15-HETE. This work was done in collaboration with Dr. TJ Maier and Dr. Geisslinger, Department of Clinical Phamracology, University of Frankfurt

3.5 Expression and purification of 5-LO from Escherichia coli

Human recombinant 5-LO protein was expressed in E. coli JM 109 cells, transfected with the plasmid pT3-5LO, and purification of 5-LO was performed via affinity chromatography as described ¹⁰⁸. Cells were grown overnight in LB medium supplemented with 100 µg/ml ampicillin, transferred to M9 minimal medium (48 mM Na₂HPO₄, 22 mM KH₂PO₄, 8.5 mM NaCl, 19 mM NH₄Cl, 6.3 mM NaOH, glycerol 2% and 100 µg/ml ampicillin, pH 7.4 casein 2 g/l) and expression of 5-LO was induced with 200 µM IPTG. Cells were harvested by centrifugation (1,500 x g, 15 min, 4°C) lysed by incubation in 50 mM triethanolamine/HCl 5 pН 8.0. mМ EDTA, soybean trypsin inhibitor (60 $\mu g/ml$), 1 mM phenylmethylsulfonylfluoride (PMSF), 1 mM DTT, and lysozyme (500 µg/ml), homogenized by sonification (Bandelin, Sonoplus HD 200) (3 x 15 s) on ice and centrifuged at 19,000 x g (Sorvall RC 5B plus) for 15 min at 4°C. The pellet was discarded. After centrifugation of the supernatant at 100,000 x g (Beckman Optima LE-80K) for 70 min at 4°C, the resulting supernatant was applied to an ATP-agarose column (Sigma-Aldrich, Munich, Germany), and the column was eluted with 20 mM ATP in PBS / EDTA ¹⁰⁵. Partially purified 5-LO was used immediately for in vitro activity assays.

3.6 Expression and purification of his-tagged platelet-type 12-LO from Escherichia coli

Human recombinant his-tagged platelet-type 12-LO (his12-LO) protein was expressed in E. coli JM 109 cells, transfected with the plasmid pT3-his12LO, and purification of 12-LO was performed via ion-metal affinity chromatography (IMAC). Cells were grown overnight in LB medium supplemented with 100 µg/ml ampicillin, transferred to M9 minimal medium (48 mM Na₂HPO₄, 22 mM KH₂PO₄, 8.5 mM NaCl, 19 mM NH₄Cl, 6.3 mM NaOH, glycerol 2% and 100 µg/ml ampicillin, pH 7.4, casein 2 g/l) and expression of his12-LO was induced with 200 µM IPTG. Cells were harvested by centrifugation (1,500 x g, 15 min, 4°C) lysed by incubation in lysis buffer (50 mM NaH₂PO₄, 300 mM NaCl, pH 8.0), 10 mM imidazol, soybean trypsin inhibitor (60 μ g/ml), 1 mM phenylmethylsulfonylfluoride (PMSF), and 10 mM β -mercaptoethanol, homogenized by sonification (3 x 15 s) on ice and centrifuged at 19,000 x g for 15 min at 4°C. The pellet was discarded. After centrifugation of the supernatant at 100,000 x g for 70 min at 4°C, the resulting supernatant was applied to a Ni²⁺-NTA-agarose column (Oiagen, Hilden, Germany), and the column was washed with 4 volumes of lysis buffer plus 20 mM imidazol and eluted with lysis buffer plus 250 mM imidazol (4 ml/l bacteria culture). Partially purified his12-LO was used immediately for in *vitro* activity assays ¹⁰⁵.

3.7 Determination of 5-LO product formation in intact cells

For assaying the 5-LO product formation of intact cells, 7.5 x 10^6 freshly isolated PMNL were resuspended in 1 ml PGC buffer. After pre-incubation with the test compounds or vehicle at 37°C for 10 min at the indicated concentrations, 5-LO product formation was started by the addition of the various stimuli (ionophore A23187, NaCl) as indicated. After 10 min at 37°C, the reaction was stopped by addition of 1 ml ice-cold methanol and 30 µl of 1 N HCl, 200 ng PGB₁, and 500 µl of PBS were added. 5-LO metabolites formed were extracted by solid phase extraction and analyzed by C-18 reversed phase HPLC (Waters, Eschborn) as described ¹⁰⁹. Mobile phase was 76% methanol, 24% H₂O and 0.007% trifluoracetic acid with a flow rate of 1.2 ml/min. PGB₁ (internal standard) and 5-LO products were detected and calculated by Empower Pro software (Waters, Eschborn).

3.8 Determination of 5-LO product formation in cell-free systems

For determination of 5-LO activity in cell homogenates, 7.5 x 10^6 freshly isolated PMNL were resuspended in PBS containing 1 mM EDTA, sonicated (3 x 10 s) at 4°C, and 1 mM ATP was added. For determination of the activity of recombinant 5-LO, partially purified 5-LO (0.5 µg in 5 µl PBS) was added to 1 ml of a 5-LO reaction mix (PBS, pH 7.4, 1 mM EDTA, 25 µg/ml phosphatidylcholine, 1 mM ATP, and 20 µg/ml γ -globulin). In some experiments DTT (1 mM) or GSH (5 mM) was added to the reaction mixtures as indicated. After incubation with test compounts or vehicle for 10 min at 4°C, samples were warmed up for 30 s at 37°C and 2 mM CaCl₂ and AA at the indicated concentrations were added to start 5-LO product formation. The reaction was stopped after 10 min by the addition of 1 ml ice-cold methanol and 30 µl of 1 N HCl, 200 ng prostaglandin B₁, and 500 µl of PBS were added ¹⁰⁸. Metabolites formed were analyzed by HPLC as described above.

3.9 Determination of 12-LO product formation

To determine p12-LO product formation in intact cells, freshly isolated platelets (10^8 /ml PG buffer) were supplemented with either 1 mM CaCl₂, 1 mM EDTA, or 1 mM EDTA plus 30 μ M BAPTA/AM. Platelets were preincubated with the indicated agents for 15 min at 37°C. After addition of stimuli and further incubation at 37°C for the times indicated, the reaction was stopped by the addition of 1 ml ice-cold methanol and 30 μ l of 1 N HCl, 200 ng prostaglandin B₁, and 500 μ l of PBS were added. p12-LO products (12(S)-hydro(pero)xy-6-trans-8,11,14-cis-eicosatetraenoic acid (12-H(P)ETE) were extracted and then analyzed by HPLC ¹⁰⁷. 12-HETE and 12-HPETE elute as one major peak, integration of this peak represents p12-LO product formation, expressed as ng metabolites per 10⁸ cells. For determination of p12-LO product formation in broken cell preparations, platelets (10^8 /ml PG buffer plus 1 mM EDTA and 1 mM PMSF) were sonicated (3 × 10 s) and lysates were

centrifuged (100,000 x g / 70 min / 4°C). To the resulting 100,000 x g supernatant or to partially purified his12-LO, BAs were added and samples were pre-warmed at 37°C for 30 sec. CaCl₂ (2 mM) was added as indicated and his- or p12-LO product formation was started by addition of AA (10 μ M). After 10 min at 37°C, the formation of 12-H(P)ETE was determined as described for intact cells.

3.10[³H]-Arachidonic acid release

PRP was labeled with 19.2 nM [3 H]-AA (1 µCi/ml, specific activity 200 Ci/mmol) for 2 h at 37°C in the presence of 100 µM aspirin to avoid clotting. Then, cells were washed twice with PBS pH 5.9 plus 1 mM MgCl₂, 11.5 mM NaHCO₃, 1 g/l glucose and 1 mg/ml fatty acid free BSA and finally resuspended in PG buffer (10⁸ cells /ml). Preparations of cells at pH 5.9 is thought to minimize temperature-induced activation ¹¹⁰. After 15 min at room temperature, 1 mM CaCl₂ was added 2.5 min prior stimulation with the indicated agents at 37°C. After 5 min, incubations were put on ice for 10 min, followed by centrifugation (5,000 x g, 15 min). Aliquots (300 µl) of the supernatants were measured (Micro Beta Trilux, Perkin Elmer) to detect the amounts of [3 H]-labeled AA released into the medium.

3.11 Immobilization of boswellic acids and protein pull-down assays

For immobilization of BAs at EAH Sepharose 4B beads, the free 3-OH group of the BAs was used (manuscript: Kather, N., Tausch, L., Poeckel, D., Werz, O., Herdtweck, E. and Jauch, J. (2008)). In brief, β -BA and KBA were treated with glutaric anhydride to form the half-esters Glut-BA and Glut-KBA, respectively, and analyzed by ¹H- and ¹³C-NMR as well as by MS. These substances were linked to EAH Sepharose 4B by standard amide coupling procedures. The carboxylic acid of the BA-core was unlikely to react under standard conditions due to steric crowding. The success of the coupling reaction was determined by two methods: a) Glut-BAs were used in defined excess (2 µmol of the Glut-BAs per 1 µmol NH₂-groups of the EAH Sepharose 4B). After the coupling reaction, the hypothetical excess of Glut-BAs (1 µmol) could be indeed recovered. b) Treatment of Glut-BAs with KOH in isopropanol under reflux for ca. 3 h cleaved the ester bond and gave BA and KBA respectively, analyzed by thin layer chromatography. Preparation of immobilized BAs was performed by Dr. J. Jauch, Saarbrücken.

For protein pull-down experiments, 3×10^7 PMNL, $5 \ge 10^7$ RBL-1, $3 \ge 10^9$ platelets, $1 \ge 10^7$ MM6, $3 \ge 10^7$ HL-60, $5 \ge 10^7$ LNCaP or $8 \ge 10^7$ MCF-7 were lysed in 1 ml lysis buffer (50 mM HEPES pH 7.4, 200 mM NaCl, 1 mM EDTA, 1% Triton X-100, 2 mM PMSF, 10 µg/ml leupeptin, 120 µg/ml soybean trypsin inhibitor). 50 µl of the sepharose slurries (50%, v/v) were added to the lysates and incubated at 4°C over night under continuous rotation. The Seph-beads were intensively washed 3 times with 5 volumes of binding buffer (HEPES pH

7.4, 200 mM NaCl, 1 mM EDTA) and precipitated proteins were finally separated and denatured by addition of SDS-b (20 mM Tris pH 8.0; 2 mM EDTA; 5% SDS; 10% β -mercaptoethanol). After boiling (95°C, 6 min), beads were removed by centrifugation and the supernatant containing proteins were separated by SDS-PAGE and visualized by WB, coomassie or silver staining, respectively.

3.12 SDS-PAGE

Freshly isolated PMNL, HL-60, or MM6 cells (5 x 10^6 each) were resuspended in PGC buffer; the final volume was 100 µl. After addition of the indicated stimuli (DMSO, final concentration < 1% (v/v)), samples were incubated at 37°C and the reaction was stopped by addition of 100 µl of ice-cold 2 x sodium dodecylsulphate–polyacrylamide gel electrophoresis (SDS–PAGE) sample-loading buffer (SDS-b; 20 mM Tris/HCl, pH 8, 2 mM EDTA, 5% SDS (w/v), 10% β-mercaptoethanol), vortexed and heated for 6 min at 95°C. Total cell lysates (20 µl) were mixed with 4 µl of glycerol/0.1% bromophenolblue (1:1, v/v) and analysed by SDS–PAGE using a Mini Protean III system (Bio-Rad, Hercules, CA, U.S.A.) on a 4-20% gel, unless stated otherwise.

3.13 Western Blot

After electroblot to nitrocellulose membrane (Amersham Pharmacia, Little Chalfont, UK), membranes were blocked with LiCor blocking buffer mixed with 50 mM Tris/HCl, pH 7.4, and 100 mM NaCl (Tris-buffered saline (TBS)) (1:1, v/v) for 1 h at RT. Correct loading of the gel and transfer of proteins to the nitrocellulose membrane was confirmed by Ponceau staining. Membranes were washed and then incubated with primary antibody (AB) overnight at 4°C. Antibodies were diluted 1 : 1000 in LiCor blocking buffer. The membranes were washed and incubated with 1 : 5000 dilution of fluorescence dye-conjugated immunoglobulin G (LiCor, Lincoln, NE, USA) for 45 min at RT. After washing (4 x with TBS plus 0.1% Tween 20 and TBS), visualisation of proteins was performed using a LiCor Odyssey 2-color Western detection system (LiCor, Lincoln, NE, USA), according to the manufacturer's instructions.

3.14 Colloidal Coomassie staining

Washed (in MQ) gels from SDS-PAGE were fixed and stained 6 - 12 h on a shaking table at room temperature in colloidal staining solution (0.08% Coomassie Brilliant Blue G250, 1.6% ortho-phosphoric acid, 8% ammonium sulfate, 20% methanol) and destained in MQ until background was clear. 1% acetic acid (v/v) was added as indicated ¹¹¹.

3.15 Silver staining

After SDS-PAGE, proteins in the gels were fixed in 5% acetic acid and 10% methanol (v/v, 4x for 30 min), washed in MQ and sensitized 2 min in freshly prepared 0.02% $Na_2S_2O_3$ solution. After washing in MQ and incubation in "silver solution" (0.1% AgNO₃) for 30 min the gel was developed several minutes under slow shaking in 2.5% Na_2CO_3 plus 300 µl/l formaline. The developed gel was fixed in 1% acetic acid and scanned in a scanner (AGFA Arcus II).

3.16 In-gel digestion

All gels were subjected to manual gel cutting using a razor blade. Dissected gel pieces were subjected to in-gel digestion protocols ^{112, 113} which were adapted for use on a Microlab Star digestion robot (Bonaduz, Switzerland) ¹¹⁴. Samples were reduced, alkylated and subsequently digested over night using bovine trypsin (sequencing grade, Roche, Mannheim, Germany). The gel pieces were extracted and the extracts were dried in a vacuum centrifuge and stored at -20°C until analysis.

3.17 MALDI-TOF-MS

MALDI-TOF-MS experiments were performed on an Ultraflex TOF/TOF mass spectrometer (Bruker Daltonics Inc., Manning Park Billerica, MA) in cooperation with Dr. M. Karas, Institute of Pharmaceutical Chemistry, University of Frankfurt. The samples were dissolved in 5 μ L of water/acetonitrile/TFA (29.5/70/0.5, v/v/v). α -cyano-4-hydroxycinnamic acid (3 mg/ml, Bruker Daltonics Inc., Manning Park Billerica, MA) in water/acetontrile/TFA (29.5/70/0.5, v/v/v) was used as matrix. Analyte and matrix were spotted consecutively in a 1:1 ratio on a stainless steel target and dried under ambient conditions. The dried sample was washed with ice-cold 5% formic acid to reduce salt contamination prior to analysis.

Spectra were externally calibrated with a SequazymeTM Peptide Mass Standard Kit (Applied Biosystems, Foster City, CA) and internally calibrated on a tryptic auto digestion peptide (m/z 2163.0564). The spectra were processed in flexAnalysis version 2.2 (Bruker Daltonics Inc., Manning Park Billerica, MA) using the SNAP algorithm (signal to noise threshold: 3, maximal number of peaks: 150, quality factor threshold: 40).

Proteins were identified by Mascot (www.matrixscience.com, Matrix Science, Boston, MA) (peptide mass tolerance: 100 ppm; maximum missed cleavages: 1) using the NCBInr database (2314886 sequences; 1066605192 date 26.01.2005). Proteins with a score of 76 or higher were considered significant (p < 0.05).

3.18 In vitro kinase assay for DNA-PK

To determine kinase activity of DNA-PK, 20 U of the kinase were incubated with the BAs for 10 min at 30°C in 50 μ l reaction buffer (50 mM HEPES pH 7.5, 100 mM KCl, 10 mM MgCl₂, 0.2 mM EGTA, 0.1 mM EDTA, 1 mM DTT) with 10 μ g/ml CpG-ODN 1018, 80 μ g/ml BSA, 200 μ M ATP, 3 μ Ci [³²P]-ATP and 100 μ g DNA-PK peptide substrate or 600 ng Akt-1 as substrate. Reaction was stopped by adding 20 μ l of 30% acetic acid, samples were separated by SDS-PAGE and bands were visualized by Coomassie-staining. The dried gel was exposed to a radioactivity sensitive-film and [³²P] incorporation was measured with a FLA-3000 reader (Fuji Film)⁶³.

3.19 In vivo phosphorylation

LNCaP or MCF-7 cells were cultured overnight in a 24-well cell culture plate (8 x 10^4 , 7 x 10^4 per well, respectively), and cells were starved in serum-free media for 24 h. After preincubation with the BAs for 1 h, cell activity was stimulated with 10 µg/ml CpG-ODN 1018 for 20 min. Cells were lysed with icecold lysis buffer (160 mM NaCl, 20 mM Tris pH 7.4, Tx-100 0.1%, NP-40 0.1%, 1 mM EDTA) and phosphorylation of Akt was detected by Western Blot using a phospho-specific antibody.

3.20 Akt activity assay

Active recombinant Akt-1 (50 ng) was incubated with eNOS (Ser-1177) biotinylated peptide in 50 μ l assay buffer (25 mM Tris pH 7.5, 10 mM MgCl₂, 5 mM β -glycerolphosphate, 0.1 mM Na₃VO₄, 2 mM DTT, 200 μ M ATP, 1.5 mM peptide) for 30 min at room temperature. Reaction was stopped with 50 mM EDTA, pH 8.0, and an aliquot was transferred to a streptavidin coated 96-well plate. After incubation with phospho-eNOS (Ser-1177) antibody and a fluorescence dye-conjugated secondary antibody, visualisation was performed using a LiCor Odyssey 2-color Western detection system (LiCor, Lincoln, NE, USA), according to the manufacturer's instructions. The amount of phosphorylated peptide is given as percent of maximum fluorescence¹¹⁵.

3.21 In vitro kinase assay of PKC

To preparate crude PKC containing samples, 6×10^7 HL-60 cells were harvested, washed and sonicated (5 x 10 s on ice) in 1 ml sample preparation buffer (50 mM Tris, 50 mM β -mercaptoethanol, 10 mM EGTA, 5 mM EDTA, 1 mM PMSF, 10 mM benzamidine, pH 7.5) and centrifuged (100,000 x g, 60 min, 4°C). Protein concentration was determined by Bradford-assay.

 $4.5 - 5.0 \ \mu$ g of the protein mixture were incubated (25 mM Tris, pH 7.0, 3 mM MgCl₂, 0.1 mM ATP, 2 mM CaCl₂, 50 μ g/ml phosphatidylserine, 0.5 mM EDTA, 1 mM EGTA, 5 mM β -mercaptoethanol) for 15 min at 25°C in a 96-well plate, coated with a PKC-pseudosubstrate (--RFARKGSLRQKNV). The reaction was stopped with 20% H₃PO₄, the wells were washed and treated with a biotinylated antibody against the pseudosubstrate. After incubation with a streptavidin-coated peroxidase and development with *o*-phenylenediamine, the OD 492 nm was measured in a plate reader.

3.22 Farnesyl pyrophosphate synthase activity assay

Farnesyl pyrophosphate synthase activity was determined in vitro. 100 mio HL-60 cells were homogenized in 5 ml ice-cold homogenisation buffer (0.3 M sucrose, 10 mM EDTA, 1.2 mM β -mercaptoethanol, pH 7.4) by sonication (3 x 8 s) on ice and centrifuged (900 x g, 5 min, 4°C) to remove cell debris. To obtain the peroxisomal fraction, the supernatant was centrifuged twice (8,700 x g, 10 min, 4°C and 10,000 x g, 10 min, 4°C). The resulting pellet was resuspended in ice-cold resuspension buffer (20 mM imidazol, 5 mM DTT, pH 7.4). Protein concentration was measured by Lowry-assay. The reaction was started by addition of 100 μ g of total protein in 150 μ l assay buffer (25 mM HEPES pH 7.4, 2 mM MgCl₂, 1 mM DTT, 5 mM KF, 1% n-octyl- β -glucopyranoside, 20 μ M geranylpyrophosphate, 13.3 μ M [¹⁴C]-IPP (55 mCi/mmol)) with the indicated BAs or alendronate. After 45 min at 37°C the reaction was stopped by addition of 150 μ l stop buffer (2.5 N HCl in 80% ethanol, 100 μ g/ml farnesol), the mixture was hydrolysed (30 min, 37°C) and pH normalized with 150 μ l NaOH 10%. The farnesyl pyrophosphate was extracted with 1 ml n-hexane and 200 μ l of the organic phase was pipetted into 3 ml scintillation liquid to measure (Micro Beta Trilux, Perkin Elmer) the produced [¹⁴C]-FPP .

3.23 Release of TNF α in MM6

Differentiated MM6 cells (10^6 / well) were preincubated in a 24-well plate for 30 min with the indicated compounds. After incubation with stimuli for 4 h, 50 µl of cell supernatants were transferred together with a biotinylated TNF α antibody (source: rabbit) into a 96-well plate (coated with an anti-rabbit antibody). After washing and incubation with a streptavidin-peroxidase, colour development of the ELISA-substrate was determined within 30 min at 450 nm.

3.24 Protease activity assays

Inhibition of the enzymatic activity of all proteases was determined in a 96-well plate format by mixing the protease with the test compounds or DMSO as vehicle (never exceeding 0.5%) as control in the respective assay buffer in a total volume of 200 μ l and subsequent incubation for 20 min at 25°C. The reaction was then started by addition of the respective chromogenic protease substrate and the proteolysis was monitored at 410 nm by spectrophotometrical measurement using a Victor² plate reader (PerkinElmer). Inhibition of the protease is given as the percentage of control without inhibitor (DMSO as vehicle).

For analysis of crude CatG, the enzyme was freshly prepared from PMNL $(2.5 \times 10^7 / \text{ml PGC})$ buffer) that were stimulated with 10 µM cytochalasin B and 2.5 µM fMLP for 5 min at 37°C ¹¹⁶. After centrifugation at 1,200 x g for 5 min at 4°C the resulting supernatant (containing approx. 10 µg CatG/ml) was immediately used for CatG activity assays. For analysis of isolated CatG, the enzyme was purified from PMNL as described below. The assay was composed of 100 ng crude CatG (approx. 10 µl supernatant) or alternatively 200 ng of

purified enzyme, diluted in 180 μ l HEPES 0.1 M, NaCl 0.5 M, pH 7.4. As substrate for CatG, N-Suc-Ala-Ala-Pro-Phe-pNA (Suc-AAPF-pNA) (1 mM final concentration) was used and the absorbance was measured at 410 nm at 25 or 37°C. Kinetic studies were performed with substrate concentrations from 0.1 to 4 mM and the K_m was determined from initial rate of hydrolysis by the Lineweaver-Burk method ¹¹⁷.

Inhibition of related proteases was performed in analogy to CatG adjusting the amount of protease, the assay buffer and the substrate individually to each type of protease as follows: Tryptase, 0.5 µg purified enzyme, Tris-HCl 0.1 M, pH 8.3 as assay buffer, and N-α-benzoyl-DL-arginine-pNA (1 mM) as substrate. Chymase, 0.1 µg purified enzyme, Tris-HCl 0.45 M, NaCl 1.8 M, pH 8.0, and 0.5 mM Suc-AAPF-pNA as substrate. Chymotrypsin, 0.1 µg purified enzyme, Tris-HCl 0.1 M, CaCl₂ 25 mM, pH 8.3 and 0.2 mM Suc-AAPF-pNA as substrate. Elastase, 0.15 µg purified enzyme, HEPES 0.1 M, NaCl 0.5 M, pH 7.4 and 0.2 mM N-methoxysuccinyl-Ala-Ala-Pro-Val-pNA as substrate. Proteinase 3, 0.5 µg purified enzyme, MOPS 0.1 M, NaCl 0.5 M, 5,5'-dithiobis-(2-nitro-benzoic acid) 0.1 mM, pH 7.5, and 1 mM Boc-Ala-Ala-Nva-SBzl as substrate.

3.25 Docking experiments

Rigid automated molecular docking was performed in collaboration with Dr. G. Schneider, Institute of Organic Chemistry and Chemical Biology, University of Frankfurt, using GOLD 3.01. (Cambridge Crystallographic Data Centre, Cambridge, UK, http://www.ccdc.cam.uk) which relies on a genetic algorithm 118 . We used the known crystal structure of CatG (1T32) from the database PDB. Hydrogens were added, and then energy minimized using the AMBER99 force field within the software MOE¹¹⁹. For the co-crystallized inhibitor (2-[3-{methyl[1-(2-naphthoyl)piperidin-4-yl]amino}carbonyl)-2-naphthyl]-1-(1-naphthyl)-2oxoethylphosphonic acid) hydrogen atoms were added, and energy minimization was performed using the MMFF94x force field ¹²⁰. The 3D structure of AKBA was recently determined (manuscript: Kather, N., Tausch, L., Poeckel, D., Werz, O., Herdtweck, E. and Jauch, J. (2008)). GOLD parameter settings for the genetic algorithm were: number of generations = 100000, population size = 100, selection pressure = 1.1, number of islands = 5, niche size = 2, migrate = 10, mutate = 95, crossover = 95. A 20 Å radius around the active site defined the binding pocket. The default Goldscore function ¹¹⁸ was employed for scoring the predicted receptor-ligand complexes. Larger positive score values indicate more favourable receptor-ligand complexes, negative values indicate unfavourable binding modes (nonbinding). Each docking run was repeated ten times to obtain average score values with standard deviations. The same method was used for the redocking of the co-crystallized inhibitor. RMSD values between the coordinates determent via x-ray crystallography and the docking solutions were computed, and a mean value with standard deviation was calculated. PyMOL was used for visualization of docking poses ¹²¹. This work was almost completely done by Lutz Franke, Dr. G. Schneider, Institute of Organic Chemistry and Chemical Biology, University of Frankfurt.

3.26 Measurement of platelet aggregation (turbidimetric)

Aggregation of platelets in (PRP or diluted PRP) washed platelets was determined using a turbidimetric light-transmittance device (ChronoLog 490 2D, Haverton, PA, USA). 30 ml whole blood was centrifuged (200 x g, 15 min, without brake) and 500 μ l of ACD (85 mM trisodium citrat, 65 mM citric acid, 100 mM dextrose) were added to 6 ml of PRP. After centrifugation for 10 min at 800 x g the pellet was resuspended carefully in Tyrode's buffer (129 mM NaCl, 8.9 mM NaHCO₃, 0.8 mM KH₂PO₄, 0.8 mM MgCl₂, 5.6 mM glucose, 10 mM HEPES, pH 7.4) without Ca²⁺ and 500 μ l of the suspension was gently prewarmed in temperated aggregation cuvettes for 2 min and 15 min incubated at 37°C with the indicated substances or DMSO as control, solvent concentration never exceeded 0.5%.

For aggregation, the response is given as percent of the maximal light transmission Amax. In Ca^{2+} -containing samples, 1 mM CaCl₂ was added right before the start of the measurement. Aggregation was recorded for 7 min.

3.27 Measurement of platelet activation markers CD62 and PAC-1 by flow cytometry

Whole blood samples (containing 3.13% sodium citrate), recalcified PRP, or washed platelets resuspended in PGC were incubated with β -BA, AKBA, thrombin receptor-activating peptide (TRAP), or vehicle (DMSO, control) for 2 or 15 min at RT. To measure CD62 and PAC-1, samples were diluted 1 : 1 in 20 mM HEPES, 137 mM NaCl, 2.7 mM KCl, 1 mM MgCl₂, 5.6 mM glucose, 1 mg/ml BSA, pH 7.4, and aliquots of 5 ml were incubated with saturating concentrations of CD41-PC7, CD62-PE, and PAC1-FITC at RT for 15 min in the dark. Samples were fixed with formaldehyde 1% (in PBS), washed twice (CellWash, Becton-Dickinson, Heidelberg, Germany), and resuspended in 300 µl PBS. Isotype-matched IgG and

IgM antibodies were used to correct for the nonspecific binding of the specific antibodies. P-selectin (represented by CD62) and PAC-1 antigen expression were quantified. Three-color flow cytometric analysis was used with logarithmic modes set for all channels. A gate was set around the platelet population (CD41), and 5000 events were acquired from each probe. The percentage of CD62- positive cells (%) as well as their mean channel fluorescence intensity (MFI) was determined at a level which yields a value of 1% for mouse IgG1-PE labelled sample. A histogram of PAC1-FITC against cell events was generated and MFI of the total platelet population was recorded. Acquisition of data was carried out using a FACSCalibur flow cytometer with CELLQuest software (Becton-Dickinson). The instrument calibration and compensation was assured daily with calibration beads (CaliBRITE Beads, Becton-Dickinson) and FACSCompt. Fluorescence-conjugated antibodies CD41-PC7, CD62-PE, and PE-labelled isotype IgG1 control were obtained from Beckman Coulter (Krefeld, Germany), PAC1-FITC and FITC-isotype IgM were from Becton-Dickinson.

3.28 Endogenous thrombin potential

Thrombin generation was assessed by using a fluorometric assay, based on the cleavage of a thrombin-specific fluorogenic substrate resulting from stimulation of recalcified or citratechelated PRP, yielding the so-called endogenous thrombin potential (ETP) ¹²². 80 μ l of PRP and 20 μ l of buffer containing the thrombin generation trigger were added to each well of a 96-well microtitre plate. The Fluoroskan Ascent Type 374 plate fluorometer (Labsystems; Finland) was used with excitation wavelength of 390 nm, emission wavelength of 460 nm, and a measurement integration time per well of 20 ms. The first derivative of the fluorescence–time curve reflects the course of thrombin activity in the sample. The parameter of interest in the samples using recalcified PRP was the maximal generation rate, which is the peak of the first derivative (ETP peak velocity, relative fluorescence units (RFU) per min) of the thrombin generation curve, or, due to low peak values in Ca²⁺-free samples, the ETP–area under the curve (AUC).

3.29 Intracellular Ca²⁺ measurement

Platelets (5×10^9 /ml PBS plus 1 mg/ml glucose) were incubated with 2 µM Fura-2/AM for 30 min at 37°C. After washing, 10^9 platelets were resuspended in 1 ml PBS plus 1 mg/ml glucose and 1 mM EDTA and incubated in a thermally controlled (37° C) fluorimeter cuvette in a spectrofluorometer (Aminco-Bowman series 2, Thermo Spectronic, Rochester, NY) with continuous stirring. After preincubation with the indicated inhibitors, stimuli and Ca²⁺ (2mM) were added and the fluorescence was measured, $[Ca^{2+}]_i$ was calculated according to Grynkiewicz et al. ¹²³. Fmax (maximal fluorescence) was obtained by lysing the cells with 1% Triton-X 100 and Fmin by chelating Ca²⁺ with 10 mM EDTA.

3.30 PMNL chemoinvasion assay

Freshly isolated PMNL (2×10^6) were resuspended in 1 ml HEPES-buffered RPMI 1640 medium with 10% (v/v) fetal calf serum (FCS) and preincubated with test compounds or with vehicle (DMSO, never exceeding 0.5% DMSO). 150 µl of the cell suspension were then placed on the upper chamber of two compartment Boyden chamber (8 µm pore-size filters) in a 24-well format. Cells were allowed to migrate through matrigel-coated pore-size filters for 40 minutes in the lower chamber containing buffer (negative control) or buffer plus fMLP (0.1 µM) as chemoattractant. Cells on the bottom of the wells were fixed with 3.7% formaldehyde, stained with grams violet, washed and the stain was solubilized using acetic acid. The absorption of the eluted stain was measured at 570 nm / 620 nm, respectively.

3.31 Purification of CatG from PMNL

CatG was purified as previously described ¹²⁴ with slight modifications. All purification steps were performed at 4°C. Human isolated PMNL (10^{10}) were suspended in ice-cold 0.15 M NaCl and sonicated (5 × 30 s, 65%). Remaining nuclei and cell debris were removed by centrifugation (600 x g, 10 min). The supernatant was again centrifuged (16,000 x g, 30 min) to separate the granular fraction (pellet) from the post-granular supernatant fraction. The granular fraction was resuspended in 1 M NaCl plus 0.05% Triton X-100, and stored for 36 h for protease extraction. The suspension was centrifuged at 16,000 x g for 30 min, and four volumes of water added to restore isotonicity. Proteins were precipitated by ammonium sulfate (60% saturation) and the pellet was resuspended in 40 ml 0.05 M Tris-HCl, pH 8.0. After centrifugation (16,000 x g, 30 min), the supernatant was subjected to elastin-Sepharose

affinity chromatography column (2.5×20 cm) equilibrated with 0.05 M Tris buffer, pH 8.0 (ÄKTA explore, GE Healthcare, Freiburg, Germany). The protein was loaded and the column was eluted with the equilibration buffer until the OD₂₈₀ returned to baseline. The column was then washed with two volumes of 0.05 M Na-acetate, 1 M NaCl, pH 5.0, and fractions containing CatG activity were eluted with 0.05 M Na-acetate, 1 M NaCl, 20% DMSO, pH 5.0. Active fractions were pooled, dialyzed in vivaspin cut-off columns (5,000 MWCO) versus 20 mM Na-acetate, 0.15 M NaCl, pH 5.5, and subjected to ion-exchange chromatography (CM Sephadex C-50) column, equilibrated in the same buffer. The sample was applied, washed with equilibration buffer and bound material was eluted by a linear NaCl gradient from 0.15 M to 1 M. The total elution volume was 300 ml and fractions of 6 ml were collected at a flow rate of 30 ml/h. The homogeneity of the purified material was checked by SDS-PAGE and CatG activity assays.

3.32 Determination of protein concentration

3.32.1 Lowry

BSA was diluted in Tris buffer (5 mM, pH 7.4) in the range 0.1 to 1 mg/ml. 25 μ l of Biorad Lowry reagent A (alk. coppertartrate) were mixed with 5 μ l standard or protein sample in triplicates in a 96-well plate. Under slow agitation, 200 μ l of Biorad Lowry reagent B (Folinciocalteau) was added and incubated for 15 min. The OD was determined at 620 nm in a microplate photometer (Digiscan, High Tech)^{125, 126}.

3.32.2 Bradford

Standard (BSA or γ -globulin) and protein of interest was diluted in appropriate buffer and pipetted in triplicates, 80 µl per well in a 96-well plate. After addition of 5x Biorad Protein assay Bradford reagent (20 µl) the plate was agitated for 15 min and measured in a microplate reader (Dynatech MR5000) at 620 nm^{127, 128}.

3.33 Crystallography and X-ray determination

Purified CatG from granules of human PMNL were used (see above). We screened for crystallization conditions with Sigma screen kits I and II (Sigma-Aldrich) and grew diffraction quality crystals by the hanging drop vapour diffusion method and at 20°C over 28 days. CatG (10 mg/ml) and Aβ-BA were mixed in a 1:2 molar ratio, and the crystals were grown from 30% poly(ethylene glycol) 4000, 0.2 M lithium sulfate, 20 mM zinc sulfate, and 0.1 M Tris, pH 8.5 ^{129, 130}. A second screen with Structure screen I & II HT 96 (Molecular Dimensions Limited, Suffolk, UK) with the sitting drop vapour diffusion method in a 96 well plate and CatG mixed with AKBA in a 1:2 molar ratio related to crystals grewing in 0.2 M sodium chloride, 0.1 M Na acetate pH 4.6, 30% (v/v) MPD and in 0.1 M Na citrate pH 5.6, 20% (v/v) 2-propanol, 20% (w/v) PEG 4000.

Crystals grew as long needles or cubes. For cryoprotection crystals were transferred into mother liquor containing 10% glycerol (v/v) and immediately frozen in liquid nitrogen. Data were collected at beamline ID23-1 at ESRF, in one degree oscillations. Diffraction was observed to a useable resolution of 2.7 Å processing and scaling were performed with HKL-2000. Crystallography and x-ray measurement was done in collaboration with Dr. J. Chen, Institute for Biophysical Chemistry, University Frankfurt.

3.34 Statistics

All data are presented as mean + s.e.m (standard error of the mean). For statistical analysis GraphPad Prism version 4.00 (GraphPad Software, San Diego, California, USA) was used. Data were subjected to Kolmogorov-Smirnov test to confirm Gaussian distribution followed by one-way ANOVA coupled with Bonferroni post hoc test for multiple comparisons (in celecoxib assays, ex vivo data were subjected by one-way ANOVA coupled with Dunnett's post t-test for multiple comparisons). All other data were subjected to paired, 2-sided t-tests. The IC₅₀ and ID₅₀ values were analysed using GraphPad Prism version 4.00 and a sigmoid curve fitting model.

4 Results

4.1 Target identification of BAs using a protein-fishing strategy

4.1.1 Pulldown experiments

An elegant technique for the identification of a high-affinity small molecule (ligand) - protein (target) interaction is the protein-fishing approach using an affinity matrix composed of the small molecule linked covalently to an insoluble resin ¹³¹. Prominent examples for a successful application of this strategy are the identification of human histone deacetylase as target for trapoxin ¹³² or of rapamycin binding protein (FKBP) as the target of the immunosuppressant FK-506 (rapamycin) ¹³³. A protein fishing strategy for the identification of molecular targets of BAs was developed in collaboration with Dr. J. Jauch, University of Saarland. Thus, KBA was linked at the C3-OH moiety to glutaric acid forming the half ester 3-*O*-glutaroyl-KBA (glut-KBA). The remaining free carboxylic group of glut-KBA was amide-coupled with the primary amine of EAH Sepharose 4B yielding KBA-Seph as an affinity resin (**fig. 411**). Freshly prepared lysates of Triton X-100 (1 %)-treated cells were incubated with KBA-Seph, EAH Sepharose 4B without glut-KBA (Seph) was used as negative control to exclude unspecific protein binding. The precipitates were intensely washed; proteins were detached from the resin using SDS-b.

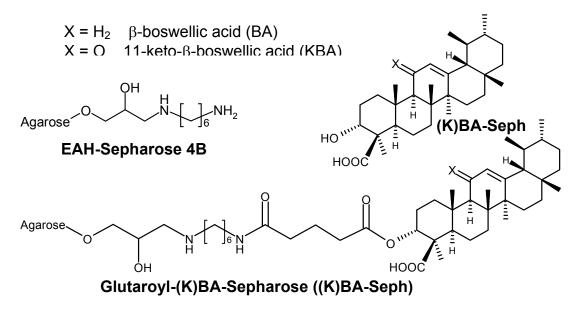


figure 411: Chemical structure of the KBA-Seph affinity matrix. KBA is linked to glutaric acid by esterification of the C3-OH group of KBA to EAH Sepharose 4B (Seph) via amide coupling, yielding the KBA-Seph affinity matrix.

4.1.2 Separation of precipitated proteins by SDS-PAGE

Proteins from pulldown experiments were separated by SDS-PAGE. The absolute amounts of precipitated proteins were comparable in pull-downs obtained with Seph and KBA-Seph. Proteins were visualized by Coomassie- and silver-staining (**fig. 412A-C**).

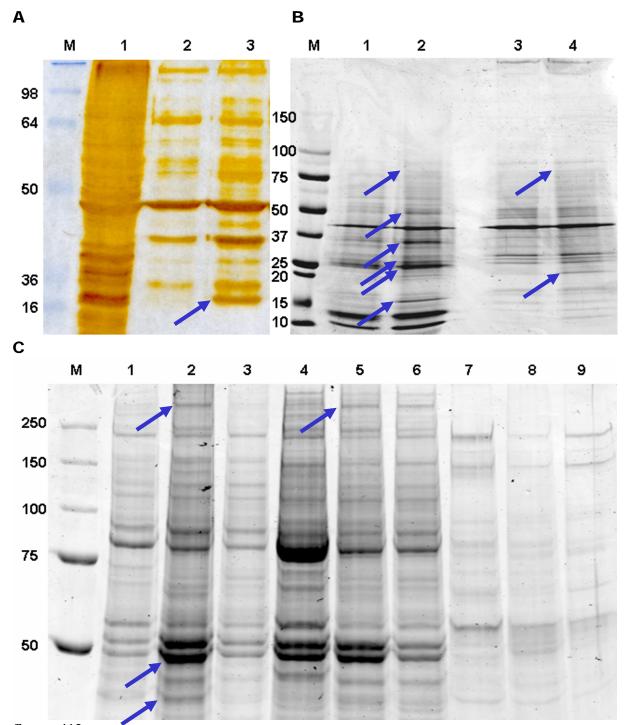


figure 412: SDS-PAGE of different fishing experiments. The gels were stained with silver or Coomassie, Biorad. All-blue protein marker (M) was used for estimation of the protein-size. Differential bands in KBA-Sepharose and Sepharose lanes were marked by arrows, results are given in table 411: (A) SDS-PAGE 10%, silver stain, PMNL-lysate, 1: lysate as control, 2: PMNL + Sepharose (Seph), 3: PMNL + KBA-Sepharose (KBA-Seph); (**B**) SDS-PAGE 4-20%, Coomassie, 1: PMNL + Seph, 2: PMNL + KBA-Seph, 3: platelets + Seph, 4: platelets + KBA-Seph; (**C**) SDS-PAGE 8%, Coomassie, 1-3: LNCaP-lysate (Seph, KBA-Seph, β -BA-Seph), 4-6: MCF-7-lysate, 7-9: PMNL-lysate

Control-lanes with samples of Seph-precipitated protein were compared with lanes of KBA-Seph-precipitated samples and bands visible in both lanes were excluded. Bands only existent in the KBA- or β -BA-Seph lane (arrows in **fig. 412A-C**) were analysed by mass spectrometry. These proteins bind specifically to KBA-Sepharose and could be a target for BAs.

4.1.3 Analysis of proteins by mass spectrometry

The protein bands of KBA-Seph pulldowns indicated by the arrows (fig. 412) were excised from the gel. Parallel to that, a corresponding piece of gel was excised from the neighboring lane, where proteins of the Seph-precipitates were separated. Proteins were in-gel-digested by trypsin and subjected to MALDI-TOF-MS in cooperation with Dr. M. Karas, Institute of Pharmaceutical Chemistry, University of Frankfurt. Analysis of the obtained peptide fragments using the peptide sequence data base www.matrixscience.com clearly indicated that the major proteins were significantly identified and could not be detected in the control sample (selection in table 411). Some proteins listed in table 411 were further analysed by Western blot. An example of database searching of protein fragments identified by MS is given in fig. 413. With a threshold of p<0.01, the analysed protein in the gel-band is considered as CatG.

MS data			found protain (atatiotically aignificant)	
cell type	species	size [kDa]	found protein (statistically significant)	
PMNL	homo sapiens	19.5	cathelicidin antimicrobial peptide CAP18	
platelets	homo sapiens	21.2	Ras-related protein Rap-1b	
			MS/MS: O-Krev precursor; Complexed With Gppnhp	
			And The Ras-Binding-Domain Of Human C-Raf1, Residues 51-131	
PMNL	homo sapiens	25.8	Cathepsin G	
PMNL	homo sapiens	26.1	proteinase 3	
LNCaP	homo sapiens	29.8	prohibitin	
RBL-1	Rattus norvegicus	41.1	Farnesyl pyrophosphate synthetase (FPP synthetase)	
PMNL	homo sapiens	41.6	vesicle amine transport protein 1 VAT1	
LNCaP	homo sapiens	51.1	ATP synthase beta subunit	
PMNL	homo sapiens	60	ATP synthase, H+ transporting	
PMNL	homo sapiens	76.4	UNC-112 related protein 2	
platelets	homo sapiens	76.5	UNC-112 related protein 2	
LNCaP	homo sapiens	470.1	DNA dependent protein kinase	

table 411: MS data pulldown experiments. KBA-Seph bound significant proteins in four different cell types. Proteins are assorted by size.

(MATRIX) Mascot Search Results Search title : Lars b : NCBInr 20050126 (2314886 sequences; 787107140 residues) Database : 4 Feb 2005 at 10:53:38 GMT Timestamp : 84 for gi 2392230, Chain A, Human Cathepsin G Top Score Probability Based Mowse Score Ions score is -10*Log(P), where P is the probability that the observed match is a random event. Protein scores greater than 76 are significant (p<0.05). Hits **Concise Protein Summary Report** ቼ <u>10</u> Number Significance threshold p< 0.05 Max. number of hits: 20 5 ð 70 60 Probability Based Mowse Score 1. gi|2392230 Mass: 25765 Score: 84 Expect: 0.0092 Queries matched: 14 Chain A, Human Cathepsin G Mass: 25767 Score: 84 Expect: 0.0092 Queries matched: 14 gi|3891975 Chain A, Human Cathepsin G gi|20664221 Mass: 27083 Score: 81 Expect: 0.02 Queries matched: 14 Chain B, Cathepsin-G gi|4503149 Mass: 29161 Score: 78 Expect: 0.033 Queries matched: 14 cathepsin G preproprotein [Homo sapiens] 2. gi|46324746 Mass: 45728 Score: 72 Expect: 0.15 Queries matched: 13 hypothetical protein Bucepa03000249 [Burkholderia cepacia R1808] gi|50355741 Mass: 26061 Score: 62 Expect: 1.3 Queries matched: 12 3. expressed protein [Oryza sativa (japonica cultivar-group)]

figure 413: Identification of CatG by MS and data base search

4.1.4 Immunodetection of proteins

The identity of selected proteins was analysed further by Western blot. A part of the results is shown in **fig. 414.** KBA-Sepharose significantly binds Ras, protein kinase C (cPKC β), platelet 12-LO, Akt, and several tyrosine-phosphorylated proteins with approximate sizes of 50, 60 and 80 kDa, whereas Seph as negative control does not bind these proteins. Because of unspecific protein-binding at the same size, proteins in **fig. 414** could not be detected selectively by MALDI-MS.

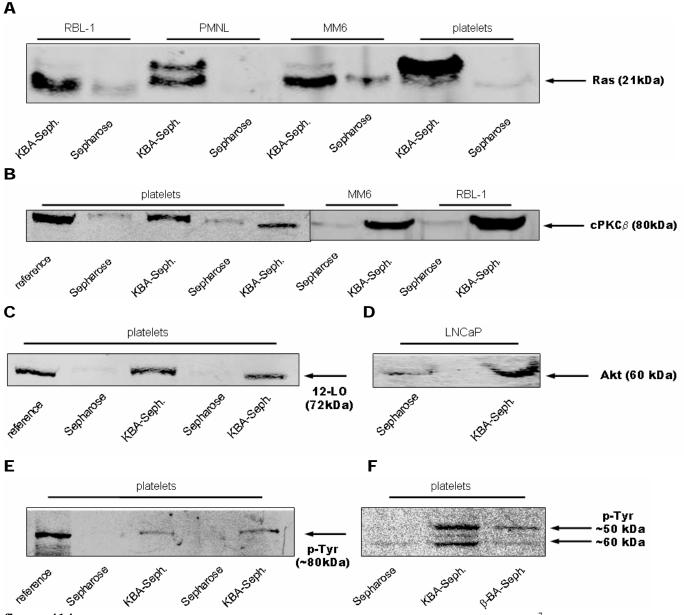


figure 414: Western blot analysis of selected proteins. Pulldowns of cell lysates (from 5×10^7 RBL-1, 3×10^7 PMNL, 1×10^7 MM6, 3×10^9 platelets or 5×10^7 LNCaP) were separated by SDS-PAGE, blotted and immunostained with the indicated antibody. Visualization was performed with an Odyssey reader. Reference = cell lysate (A) Ras (21kDa); (B) cPKC (80kDa); (C) 12-LO (72kDa); (D) Akt (60kDa); (E) pTyr (band at appr. 80kDa); (F) pTyr (bands at appr. 50/60kDa). Each blot is representative for at least three independent experiments.

4.1.5 FPPs activity

One of the identified proteins was farnesyl pyrophosphate synthase (FPPs). FPPs is one of the major proteins for terpene synthesis in humans and the target for bisphosphonates (like alendronate) in osteoporosis. As shown in table 411, it binds selectively to KBA-Sepharose but is not inhibited in vitro by BAs up to 30 μ M (**fig. 416**).

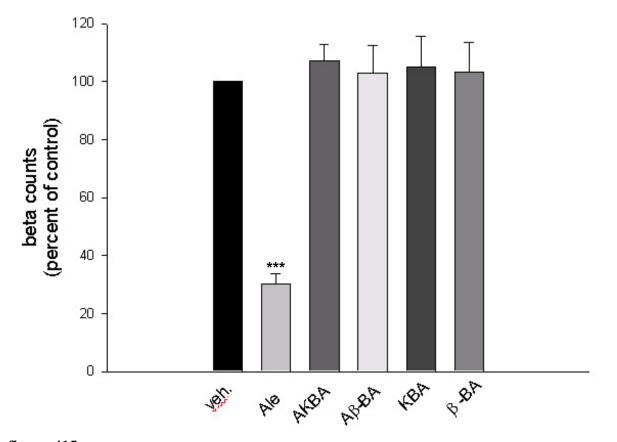


figure 415: Radioactive test system for FPPs activity in vitro. FPPs from the peroxisomal fraction of HL-60 cells was incubated with [¹⁴C]-isopentenylpyrophosphate and geranylpyrophosphate to form [¹⁴C]-FPP. Product was extracted and measured in a beta counter. DMSO was used as vehicle (veh.), alendronate 3 μ M (Ale) as positive control and BAs were used in concentrations of 30 μ M. Results are given as mean + s.e.m., n = 4-6, and analysed by one-way ANOVA followed by a Bonferroni post-hoc test: ***p<0.001.

4.1.6 PKC activity

Protein kinase C is involved in several diseases like cancer and diabetes via diverse signal cascades (see fig. 213). As shown in **fig. 414B**, PKC is bound to immobilized BAs whereas Sepharose as negative control does not bind PKC. The inhibition of the enzyme by BAs was tested in an activity assay in vitro (**fig. 417**). Only β -BA inhibits moderately at a concentration of 30 μ M, but is not statistically significant.

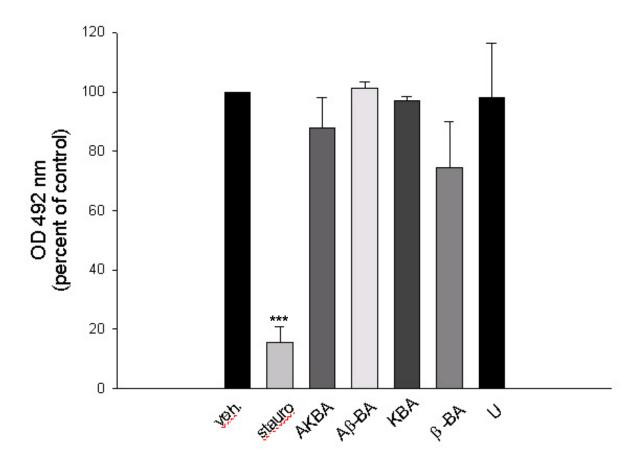


figure 416: PKC activity in vitro. In a 96-well plate, HL-60 lysate, preincubated with the indicated test compounds (U-73122 (U), BAs at 10 μ M, staurosporine (stauro) = 1 μ M) or vehicle (veh., DMSO) for 2 min at 25°C, converted a PKC-pseudo substrate and product was detected by ELISA as described in the *Methods* section. Data is given as mean + s.e.m., n = 3 and analysed by one-way ANOVA followed by a Bonferroni posthoc test: ***p<0.001.

4.1.7 Rap1b

The small guanine-nucleotide-binding protein Ras-related protein Rap1b is involved in signal transduction of apoptosis and cell survival in nearly all cells, but is essential for normal function of human platelets ¹³⁴. Rap1b efficiently activates various integrins including the platelet integrin $\alpha_{IIb}\beta_3$, which mediates fibrinogen binding required for platelet aggregation ¹³⁵. Rap1b is a common target of many different activation pathways in platelets. ADP, thrombin, collagen, epinephrine and platelet-activating factor induce Rap1-GTP formation. The pathways leading to Rap1 activation in platelets are not clearly defined ¹³⁶. For example, ADP has been suggested to activate Rap1 by Ca²⁺-dependent as well as Ca²⁺-independent mechanisms involving the Ga_i -coupled P₂Y₁₂ receptor ^{137, 138}. Recently, phosphatidylinositol 3-kinase (PI3K) was reported to mediate Rap1b activation by different platelet agonists ¹³⁹.

Rap1b was precipitated with KBA-Seph, therefore it was plausible to investigate the influence of BAs on Rap1b activity. AKBA but not β -BA inhibits ADP-induced Rap1 activation in human platelets (**fig 417**). This work was done in collaboration with Dr. Albert Smolenski, Institute of Biochemistry II, University of Frankfurt and will be continued in the thesis of Ulf Siemoneit, University of Tübingen.

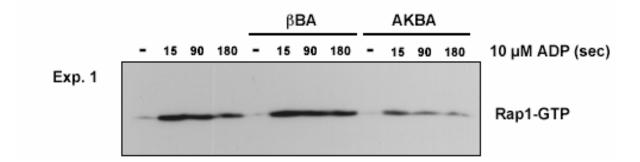


figure 417: Inhibition of ADP-induced Rap1 activation. ADP 10μ M, various time points, β -BA/AKBA 10μ M, preincubation 10 min. The blot is representative of at least three independent experiments.

4.2 Cathepsin G

4.2.1 Protein fishing with immobilized boswellic acids selectively precipitates cathepsin G

As shown in **fig. 421**, a number of proteins from PMNL were bound to the EAH-Sepharose 4B material and thus were detectable in both pull-downs. However, a 25.8 kDa protein was significantly enriched in KBA-Seph precipitates as compared to Seph lacking the ligand. No other protein selectively present in KBA-Seph but not in Seph precipitates was readily detectable. The sequence coverage using MALDI-TOF-MS was about 63% (**fig. 422A**). In order to confirm the identity of CatG as the select protein precipitated by KBA-Seph, the Seph and KBA-Seph pulldowns were subjected to SDS-PAGE and analysed by WB using CatG antibodies. Whereas no CatG was detected in the sample derived from the negative control (Seph precipitates), a strong immunoreactive band was apparent in the sample of KBA-Seph (**fig. 422B**). Moreover, purified CatG from human leukocytes was bound to KBA-Seph as detected by WB, but not to Seph, implying a specific and direct interaction with the BA (**fig. 422C**).

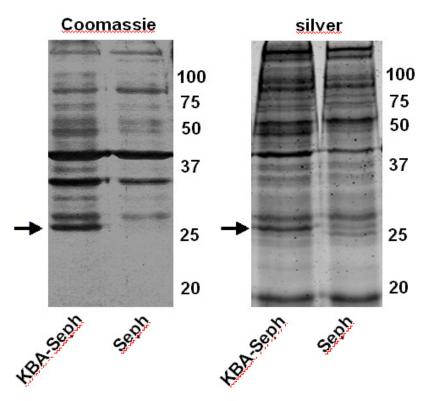


figure 421: 12,000 x g supernatants of PMNL lysates were incubated over night at 4°C with either KBA-Seph or crude Seph. Precipitates separated by SDS-PAGE and proteins were visualized by Coomassie- and by silver-staining. The band of interest at 25.8 kDa for KBA-Seph and a corresponding piece of gel from the Seph sample were excised, in-gel digested and analysed by MALDI-TOF-MS.

Α

1	IIGGR <mark>ESRPH</mark>	SRPYMAYLQI	QSPAGQSRCG	GFLVR EDFVL	TAAHCWGSNI
51	NVTLGAHNIQ	RR entqqhit	ARRAIRHPQY	NQRTIQNDIM	llqlsr rvrr
101	NR NVNPVALP	RAQEGLRPGT	LCTVAGWGRV	SMRRGTDTLR	EVQLRVQRDR
151	QCLR ifgsyd	PR RQICVGDR	RERK AAFKGD	SGGPLLCNNV	AHGIVSYGKS
201	SGVPPEVFTR	VSSFLPWIRT	TMRS		

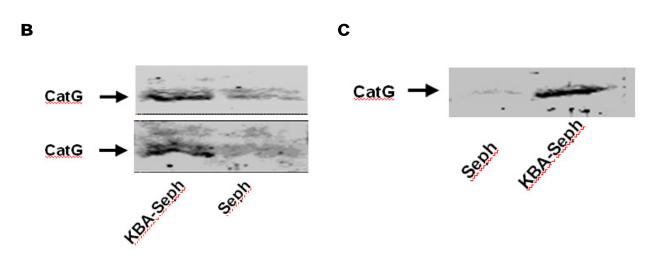


figure 422: (A) MALDI-TOF-MS data: Peptides matched to the protein sequence of human cathepsin G (chain A) (sequence coverage: 63%). The matched peptides are shown in bold (red) letters. Similar results were obtained in at least three additional experiments. (B) PMNL-lysate precipitates and (C) purified enzyme precipitates (Seph and KBA-Seph) analysed by SDS-PAGE and Western-blotting using specific antibodies against CatG.

In order to investigate whether or not the interaction of BAs with CatG may alter its enzymatic activity, the proteolytic activity of CatG, secreted from cytochalsin B/fMLP-stimulated human PMNL, was analyzed in the presence of structurally different BAs and the related triterpenes α -amyrin and ursolic acid (10 μ M each)¹¹⁶. As can be seen from **fig. 423A**, all BAs potently inhibited CatG activity, comparable to a synthetic CatG inhibitor (0.5 μ M, used as a reference compound)¹²⁹, whereas α -amyrin and ursolic acid caused no significant inhibition. Note that also the glutaroyl-linked-KBA (glut-KBA) was an effective inhibitor. Concentration-response curves (**fig. 423B**) for the BAs demonstrate concentration-dependent inhibition of CatG with IC₅₀ values of 0.6 μ M for AKBA, 0.8 μ M for β -BA, 1.1 μ M for A β -BA and 3.7 μ M for KBA. Variation of the peptide substrate concentration reduced the potency of BAs, and Lineweaver-Burke plots indicate a competitive inhibition of CatG (**fig. 423C**). Potent inhibition of 10 μ g CatG by 10 μ M AKBA (final concentration) obtained after incubation for 20 min at 37°C was reversed upon 50-fold dilution (**fig. 423D**), suggesting a reversible inhibition mode of CatG by BAs.

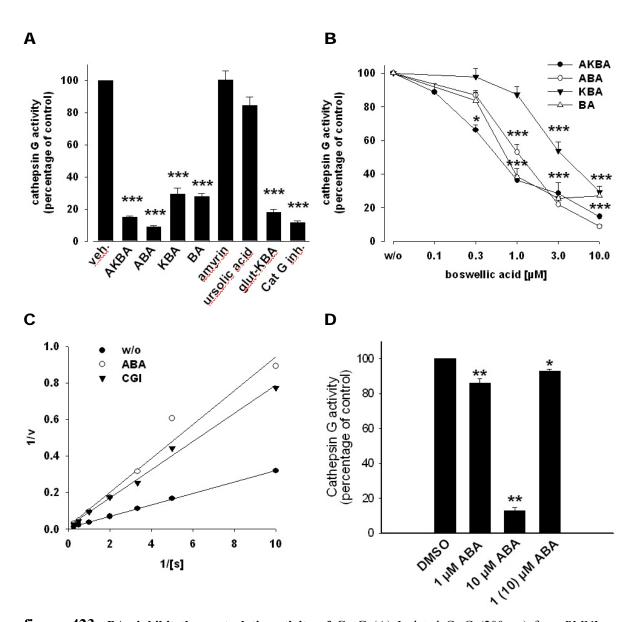
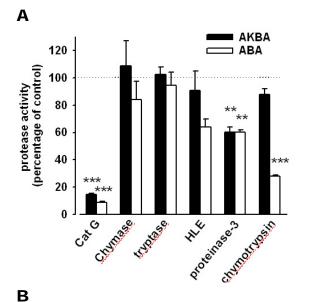
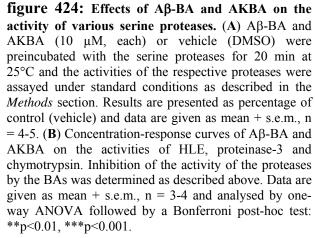


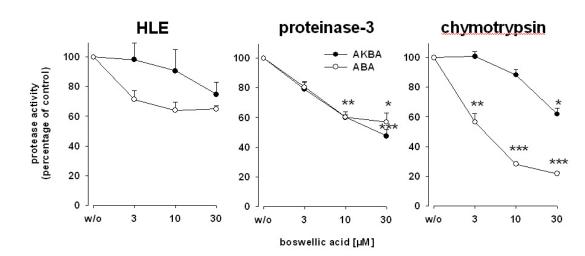
figure 423: BAs inhibit the proteolytic activity of CatG (A) Isolated CatG (200 ng) from PMNL was diluted in 180 µl HEPES 0.1 M, NaCl 0.5 M, pH 7.4 and preincubated with the indicated test compounds (all at 10 µM, except CGI = 0.5 µM) or vehicle (veh., DMSO) for 20 min at 25°C. Then, Suc-AAPF-pNA (1 mM final concentration in 200 µl final volume) was added to start the reaction and the absorbance was measured at 410 nm. The enzyme activity was determined by the progress curve method. Results are presented as percentage of control (veh.) and data are given mean + s.e.m., n = 4-6. (B) Concentration-response curves of BAs on CatG activity. Inhibition of the activity of CatG (200 ng isolated enzyme from PMNL) by BAs was determined as described above. Data are given as mean + s.e.m., n = 4. (C) Kinetic analysis of CatG inhibition by 10 µM Aβ-BA and 0.1 µM CGI. Data are given as mean of three independent experiments and results are presented as Lineweaver-Burke plots. The CatG substrate concentrations were 0.1, 0.2, 0.3, 0.5, 1, 2, and 4 mM. (D) Reversibility analysis of Cat G inhibition by Aβ-BA. Cat G is preincubated with 10 µM Aβ-BA for 5 min. After 1:10 dilution with incubation buffer, Cat G activity was determined as described above. Data are given as mean + s.e.m., n = 3. Data was further analysed by one-way ANOVA followed by a Bonferroni post-hoc (A, B) test or by a Dunnett post-hoc test (D): *p<0.05; **p<0.01, ***p<0.001.

4.2.3 Effects of boswellic acids on related serine proteases

In order to assess the selectivity of BAs for inhibition of CatG, the effects of BAs on the activity of the closely related serine proteases chymase, tryptase, HLE, proteinase-3 and chymotrypsin were determined under individually optimized, yet comparable experimental conditions. As shown in **fig. 424**, AKBA and Aβ-BA (10 μ M, each) failed to significantly inhibit chymase and tryptase. However, both BAs suppressed proteinase-3, and Aβ-BA (but not AKBA) also inhibited HLE and chymotrypsin, although less efficient as compared to CatG. Concentration response studies showed that for HLE and for proteinase-3 the IC₅₀ values of AKBA and Aβ-BA were \geq 30 μ M, whereas for chymotrypsin an IC₅₀ = 4.8 μ M for Aβ-BA was apparent. Together, AKBA and Aβ-BA are potent inhibitors of CatG, and apart from the inhibitory effect of Aβ-BA on chymotrypsin, these BAs are rather selective for CatG.







4.2.4 Docking of boswellic acids to cathepsin G

Automated molecular docking of AKBA in collaboration with Dr. G. Schneider, Institute for Biophysical Chemistry, University of Frankfurt using the protein data base (PDB) structure 1T32 as a reference for CatG, was performed for the purpose of finding a potential binding mode within the active site. To verify the validity of the GOLD software, we successfully redocked the 1T32 co-crystallized inhibitor (2-[3-{methyl[1-(2-naphthoyl)piperidin-4yl]amino}carbonyl)-2-naphthyl]-1-(1-naphthyl)- 2-oxoethylphosphonic acid) ¹²⁹, with an RMSD of 0.4 ± 0.06 . The acquired binding mode was identical to the X-ray structure and yielded a Goldscore of 102.6 ± 2.5 . Docking of AKBA into the same docking box resulted in an average docking score of 44.1 ± 1 . Both inhibitors partially occupy the same part of the active center (**fig. 425**).

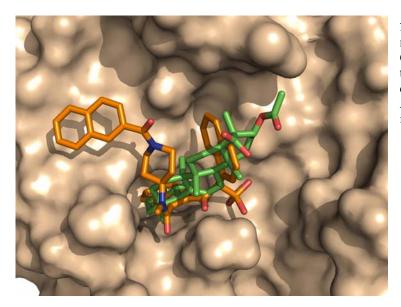


figure 425: Rigid automated molecular docking of AKBA into CatG. Active site of CatG (PDB: 1T32) together with the co-crystallized CGI displayed in orange. The orientation of AKBA in an overlapping binding mode is shown in green.

4.2.5 Boswellic acids inhibit (cathepsin G-mediated) PMNL chemoinvasion

Since CatG is assumed to contribute to invasion and migration of PMNL along chemoattractant gradients by degrading extracellular matrix proteins ³⁶, we determined the effects of AKBA and Aβ-BA on PMNL migration through matrigel towards the chemoattractant fMLP in a modified Boyden chamber assay. fMLP caused a 4.2-fold increase in the numbers of migrated cells and this was efficiently blocked by AKBA and Aβ-BA (IC₅₀ approx. 2.7 and 2.9 μ M, respectively). Comparison of the effect of CGI at 0.1 μ M demonstrated that AKBA and Aβ-BA at 8 to 9 μ M cause quantitatively the same inhibitory effect (approx. 75 % inhibition) on chemoinvasion of PMNL (**fig. 426**).

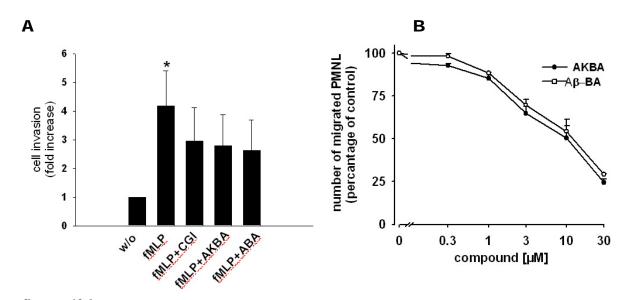


figure 426: BAs suppress CatG-mediated functional cellular responses. (A) A β -BA and AKBA inhibit PMNL chemoinvasion. PMNL were pretreated with the test compounds or with vehicle (DMSO) and placed on the upper chamber of a two compartment Boyden chamber. Cells that migrate through matrigel-coated pore-size filters in the lower chamber containing buffer (control; DMSO) or 0.1 μ M fMLP (positive control) within 40 minutes were fixed, stained with grams violet, and after washing the absorption of the solubilized stain was measured at 570 nm / 620 nm, respectively. CatG inhibitor 0.1 μ M (CGI) and AKBA 10 μ M were used as invasion inhibitors. Results are presented as fold-increase of the number of migrated cells where vehicle-treated cells were set to 1. (B) Concentration-response curves of β -BA and AKBA on the invasion inhibition of PMNL. Cells migrating toward the chemoattractant fMLP were set as 100% Data are given as mean + s.e.m., n = 5 and analysed by one-way ANOVA followed by a Bonferroni post-hoc test: *p<0.05.

4.2.6 Boswellic acids inhibit cathepsin G-mediated Ca²⁺ mobilisation in human platelets

CatG released from fMLP-stimulated PMNL can cleave the PAR-4 on platelets resulting in the mobilisation of Ca²⁺, suggesting that PAR-4 mediates platelet responses to CatG ⁴⁵. Accordingly, BAs should be able to block Ca²⁺ mobilisation in platelets evoked by fMLPactivated PMNL. Addition of fMLP to Fura-2-loaded platelets caused no Ca²⁺ mobilisation (not shown), but when PMNL were co-incubated then addition of fMLP resulted in a transient elevation of $[Ca^{2+}]_i$ in platelets (**fig. 427B**). In the presence of 3 μ M AKBA or 0.1 μ M CGI, this elevation of $[Ca^{2+}]_i$ was strongly suppressed. These data suggest that AKBA is able to block Ca²⁺ mobilisation in platelets due to inhibition of CatG (**fig. 427A**).

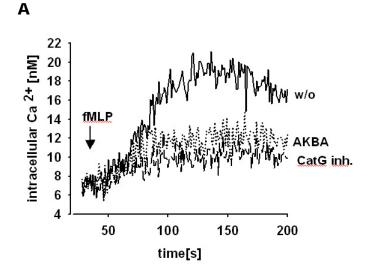
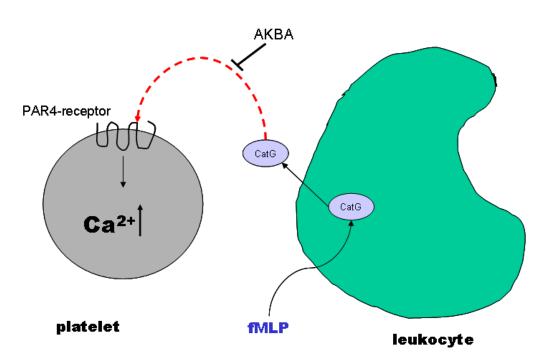


figure 427: (A) AKBA inhibits Ca^{2+} mobilisation in platelets induced by fMLPstimulated PMNL. Fura-2-loaded platelets (10^8 cells) were mixed with 10^7 unloaded PMNL in 1 ml PG buffer containing 0.1 mM EDTA and incubated with AKBA (3μ M), CGI (0.1 μ M) or vehicle (DMSO). After 5 min the measurement of $[Ca^{2+}]_i$ was started. After 30 sec, 100 nM fMLP was added and the fluorescence was recorded for another 100 sec. Curves are representative of at least 4 experiments. (**B**) Molecular mechanism of calcium influx in platelets via CatG-PAR4-receptor interaction

В



4.2.7 Inhibition of cathepsin G by Boswellia serrata extracts ex vivo

A multicentered, double-blind, randomized and placebo-controlled clinical trial aiming to protract an acute attack in patients with Crohn's disease was conducted by Pharmasan GmbH. Patients were in remission and use of aspirin, nonsteroidal anti-inflammatory drugs and glucocorticoids were not allowed in the 3 weeks before enrollment. The orally administered BS extract (Pharmasan GmbH, PS0201Bo, 400 mg) was taken twice daily in the morning/evening over a 12-month period and endpoint was an acute attack. Between October 2006 and October 2007, 13 healthy subjects were enrolled and randomization was computer-generated by using block randomization in two groups, but only one patient did not drop out until December 2007.

The effectiveness on CatG inhibition in plasma *ex vivo* was measured as part of this study. After screening and randomization, venous blood was first taken prior to medication (verum or placebo) and again after 4 weeks of treatment with 2 x 400 mg extracts/d (steady state). Citrated blood was stimulated with 10 μ M cytochalasin B and 2.5 μ M fMLP for 5 min and plasma was prepared for determination of CatG activity. As shown in **fig. 428A**, a mean reduction (47 ± 18 %) of CatG activity in the plasma of patients was measured after treatment as compared to plasma prior treatment. After partial deblinding in December 2007, data of seven patients could be analysed. CatG activity in patients who had taken verum (n=2) was reduced by 64 % ± 23 % and in patients who had taken placebo (n=5) by 15 % ± 25 % (**fig. 428B**). Plasma levels of BAs were 3380 / 2470 ng/ml (6.4 ± 1 μ M) for β-BA, 2190 / 2670 ng/ml (4.9 ± 1 μ M) for Aβ-BA, 188 / 135 ng/ml (0.34 ± 0.05 μ M) for KBA and 21.8 / 22.5 ng/ml (0.04 ± 0.01 μ M) for AKBA.

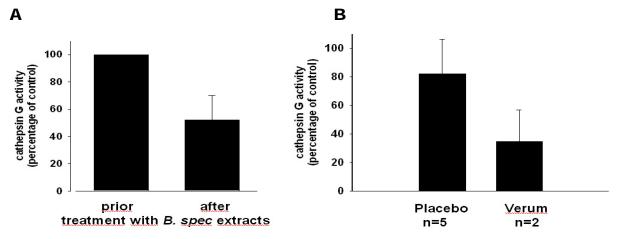


figure 428: BS extracts lower the CatG levels in human subjects *ex vivo*. Blood was taken from subjects prior medication as well as after four weeks of continuous administration of 2 x 400 mg/d BS extracts. After venipuncture, the blood was promptly stimulated with 10 μ M cytochalasin B and 2.5 μ M fMLP for 5 min, and plasma was prepared. Aliquots (20 μ l) of the plasma were immediately used for determination of CatG as described in the *Methods* section. (A) CatG activity prior and after treatment with BS extracts, n = 7. (B) CatG activity after 4 weeks treatment (steady state). Data are shown as mean + s.e.m.

In a second trial, an adult healthy male volunteer (28 years old) took three times a day three capsules with 400 mg BS extract (BS extract, acid fraction;Ch.B.: 2605010, BAs 91.37%; Euro OTC Pharma GmbH, Bönen; in total 3.6 g/d) over four weeks to demonstrate safety, tolerability, pharmacokinetics of BS extracts and to determine plasma levels of BAs in steady state. Co-medication was not allowed three weeks before and during the treatment. Blood was taken three times before treatment and three times during the fourth week under therapy and stimulated as described above to determine CatG activity *ex vivo*. CatG activity was reduced by intake of BS extract significantly by $24 \pm 1.5\%$ (fig. 429).

Plasma levels of BAs were 8013 ± 479 ng/ml ($17.6 \pm 1 \mu$ M) for β -BA, 6612 ± 82 ng/ml ($13.3 \pm 0.2 \mu$ M) for A β -BA, 355 ± 83 ng/ml ($0.76 \pm 0.2 \mu$ M) for KBA and 37 ± 6 ng/ml ($0.07 \pm 0.01 \mu$ M) for AKBA. BA plasma levels were determined by LC-MS/MS analysis ¹⁴⁰ in collaboration with Dr. Geisslinger, Clinical Pharmacology, University of Frankfurt. No side-effects appeared during intake, thus the BS extract was safe and tolerable.

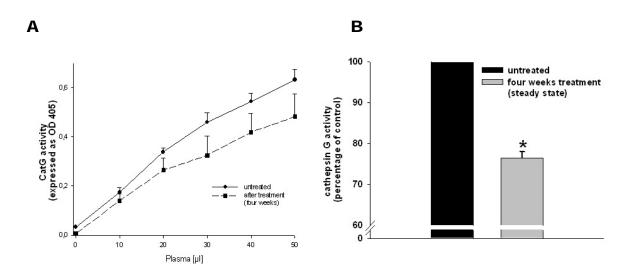


figure 429: CatG activity in human plasma before and after continuous treatment with 3.6 g extract per day. Blood was promptly stimulated with 10 μ M cytochalasin B and 2.5 μ M fMLP for 5 min, and plasma was prepared. Aliquots (20 μ I) of the plasma were immediately used for determination of CatG as described in the *Methods* section. (A) Dose response of treatment of the volunteer on CatG activity in plasma. (B) Activity of endogenous CatG ex-vivo before and after treatment with extract (3 x 1200 mg/d). Data are shown as mean + s.e.m, n = 3 and analysed by ANOVA followed by Bonferroni post-hoc test, *p<0.05.

In experiments with rats, animals were fed with gelatinous slurry as negative control or with three different extract slurries, containing BS extract, *Boswellia carterii* extract or BS solubilisate. Four hours after feeding, rats were sacrified; whole blood was prepared and stimulated with 10 μ M cytochalasin B and 2.5 μ M fMLP for 5 min. Then plasma was prepared for determination of CatG activity. Absolute activity measured in OD at 405 nm is shown in **fig. 4210A**. Only blood of rats treated with BS extract exhibited lower activity of CatG *ex vivo* as control (w/o), but not significantly because of high variations between individuals as well as the low number (n = 5). When exogenous Aβ-BA (10 μ M) was accessorily added to the plasma samples, CatG activity was only lowered in untreated animals and animals treated with the solubilisate (**fig. 4210B**), while CatG activity was already fully inhibited in extract-treated rats.

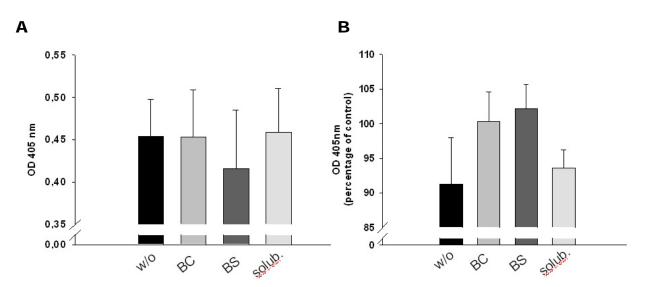


figure 4210: Cathepsin G activity *ex vivo* in rat plasma. Four hours after feeding with gelatinous slurry (w/o), BS extract (BS), *Boswellia carterii* extract (BC) or BS solubilisate (solub.), whole blood was prepared and stimulated with CytB and fMLP as described in the *Methods* section. Stimulated plasma was divided in two parts and CatG activity was measured with 1 µl DMSO as vehicle (A) or with 10 µM exogenous Aβ-BA (B). Results are shown as mean + s.e.m, n = 5 and analysed by ANOVA followed by Bonferroni post-hoc test.

4.2.8 Crystallography of CatG and Aβ-BA

CatG was purified as described in *Methods* (3.30). Concentrated protein (10 mg/ml) was mixed with Aβ-BA (1:2 molar ratio) and incubated at 22 °C for 15 min prior to crystallization trials ^{129, 141}. Crystals were grown by using the hanging drop method at 22 °C, with 0.2 M lithium sulfate, 20 mM zinc sulfate, 0.1 M Tris pH 7.4 and 30% (w/v) PEG 4000, as the precipitant. Crystals grew as long needles (**fig. 4210**) within 2 weeks to a maximum size. For cryoprotection, crystals were transferred into mother liquor containing 10% glycerol (v/v) and immediately frozen in liquid nitrogen. Data were collected at beamline ID23-1 at the European Synchrotron Radiation Facility in 1° oscillations. Diffraction was observed to a resolution of 3.0 Å. Aβ-BA was expected as inhibitor in the catalytic center. The crystals grew in space group P6(3), with a = 73.4 and b = 70.2. Because of the weak diffraction pattern and strong ice rings at 3 Å, indexing was only done to 3.5 Å. Solving the structure by molecular replacement appeared very problematic and no inhibitor could be observed at this resolution. Further condition screening and modifications in the freezing procedure should solve the problems.



figure 4211: CatG crystals under microscope. Crystals grew as long needles or in the form of a brush.

4.3 DNA-PK and Akt are influenced by boswellic acids

4.3.1 Akt phosphorylation is inhibited by BAs in vivo and in vitro

Downstream of receptor tyrosine kinases (RTK) (**fig. 213**), DNA-PK and Akt play critical roles in cell growth, apoptosis and the immune response ^{50, 51}. LNCaP were incubated for 1 hour with the indicated test compounds (AKBA, A β -BA, DNA-PK-inhibitor (DNA-PKI) or wortmannin), and then stimulated with CpG-ODNs for 20 min to activate DNA-PK ⁵². As shown in **fig. 431 and 432**, AKBA significantly inhibits phosphorylation of Akt at position Ser-473 in prostate cancer cells (LNCaP), acts synergistically with the PI3K-inhibitor wortmannin, and inhibition is concentration-dependent (**fig. 431B**). Bad is a downstream target of Akt, therefore, phosphorylation of Bad at residue Ser-136 is an indicator for the activity of Akt ⁵³. Beside Akt phosphorylation, Bad phosphorylation is also potently inhibited by AKBA (20 μ M) and the combinations of wortmannin (200 nM) and DNA-PKI (600 nM; **fig. 432**), but not by wortmannin or by DNA-PKI alone.

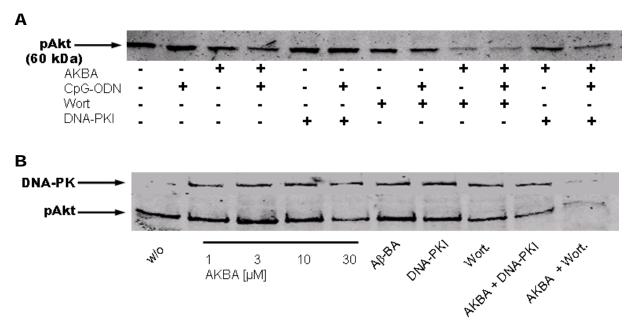


figure 431: Inhibition of Akt-phosphorylation in LNCaP cells by BAs. LNCaP were incubated for 1 hour with the test compounds prior to stimulation with CpG-ODNs (10 μ g/ml). Cells were lysed and immunoblotted against pAkt or pBad. (A) Akt-phosphorylation is inhibited by AKBA (20 μ M) and the control inhibitors wortmannin (200 nM) and DNA-PKI (600 nM); unstimulated or stimulated with CpG-ODNs. (B) Concentration-response of AKBA in Akt-phosphorylation, Aβ-BA 10 μ M, wortmannin 200 nM and DNA-PKI 600 nM and the combinations were used as controls. Blots are representative of at least four experiments.

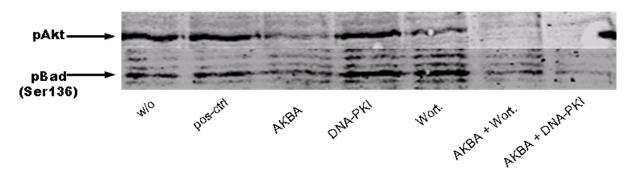


figure 432: Correlation of Akt- and Bad-phosphorylation. Bad is a downstream-target of Akt and an indicator of active Akt (used concentrations: AKBA 20 μ M, wortmannin 200 nM and DNA-PKI 600 nM). Blots are representative of at least four experiments.

Akt phosphorylation by DNA-PK was tested in a radioactive ($[^{32}P]$ -ATP) in-vitro-kinase assay. Active DNA-PK (20 U) was incubated for 10 min at 30°C with BAs, CpG-ODNs (10 μ g/ml), $[^{32}P]$ -ATP (3 μ Ci) and 100 μ g DNA-PK peptide-substrate or 600 ng Akt-1 as substrate. After separation, the $[^{32}P]$ -phosphorylated proteins were exposed to a radioactivity-sensitive film. Phosphorylation of Akt (**fig. 433A**) and the DNA-PK peptide-substrate (**fig. 433B**) by DNA-PK is inhibited concentration-dependently by BAs.

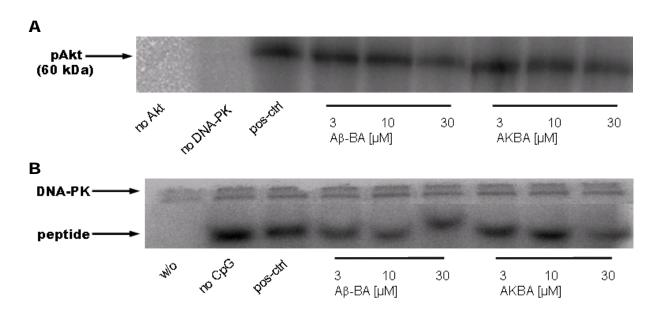


figure 433: Akt-phosphorylation is inhibited in-vitro. DNA-PK (20 U) was incubated for 10 min at 30°C with BAs, CpG-ODN (10 μ g/ml), [³²P]-ATP (3 μ Ci) and 100 μ g DNA-PK peptide substrate or 600 ng Akt-1 as substrate. Samples were separated as described in the *Methods* section and [³²P]-phosphorylated proteins were detected with a radioactivity-sensitive film. (A) In-vitro-kinase assay with recombinant Akt-1 and DNA-PK, (B) In-vitro-kinase assay with DNA-PK-peptide-substrate (1.75 kDa) and DNA-PK. Gel-loading was controlled with DNA-PK (upper band). Similar results were obtained in at least three additional experiments.

4.3.2 Akt-activity is directly inhibited by boswellic acids

In pulldown experiments with KBA-Seph and isolated recombinant DNA-PK and Akt, only Akt but not DNA-PK was detectable by immunoblotting in the precipitates (**fig. 434**). No Akt was detected in the sample derived from the negative control (Seph precipitates).

To determine the activity of Akt *in vitro*, purified enzyme was incubated with the test compounds and the specific substrate as described in the *Methods* section and fluorescence was quantified. BAs and ursolic acid (10 μ M each) potently inhibit Akt-activity. Staurosporine (1 μ M) was used as positive control; it inhibits the phosphorylation of substrate in an effective manner (**fig. 435A**). The 11-me-BAs are very potent inhibitors of Akt with an IC₅₀ of 300 nM for Aβ-BA and 1 μ M for β-BA whereas the 11-keto-BAs are less effective (AKBA 5 μ M; KBA 10 μ M). Akt inhibition by BAs is concentration-dependent (**fig. 435B**).

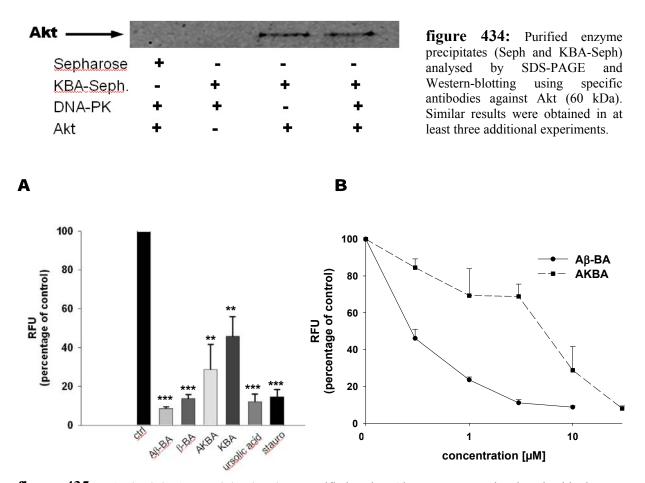


figure 435: BAs inhibit Akt activity in vitro. Purified active Akt enzyme was incubated with the test compounds at RT. After 15 min, eNOS was added as described in *Methods* section and fluorescence of secondary antibodies were detected and quantified with the Odyssey-system. (A) Akt inhibition by BAs, ursolic acid (each 10 μ M) and staurosporine (stauro 1 μ M) as positive control; (B) concentration-response curves of Akt inhibition by AKBA and Aβ-BA. Results are shown as mean + s.e.m, n = 5 and analysed by a one-way ANOVA followed by Bonferroni post-hoc test, **p<0.01, ***p<0.001.

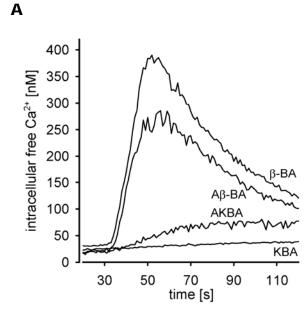
4.4 Modulation of signal transduction and functionality of platelets and monocytes by boswellic acids

4.4.1 Modulation of Ca²⁺ mobilization in washed human platelets by boswellic acids.

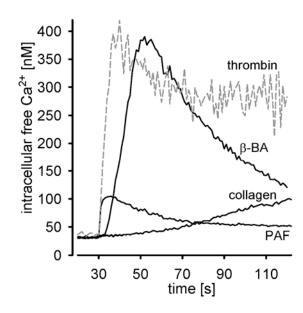
This study investigates, whether or not BAs, in particular the 11-keto-BAs, may have detrimental actions on agonist-evoked mobilization of intracellular free Ca²⁺. Thus, thrombin, collagen, platelet-activating factor (PAF) and the stable synthetic thromboxane analogue U-46619 were chosen as relevant platelet agonists ⁶⁹. Concentration-response studies in Fura-2-loaded washed human platelets revealed approximate EC₅₀ values of these agonists for Ca²⁺ mobilization as follows: 0.5 U/ml thrombin, 8 µg/ml collagen, 100 nM PAF, and 1 µM U-46619 (not shown). These concentrations were used for subsequent analysis of BA effects. Although ADP is known as potent physiological stimulus, it failed to mobilize Ca²⁺ in platelets under our assay conditions.

In the presence of extracellular Ca²⁺ (1 mM), BAs lacking the 11-keto moiety (Aβ-BA and β-BA, 10 mM each) induced a transient but robust elevation of $[Ca^{2+}]_i$ in washed platelets that peaked 18–30 s following exposure, whereas KBA was ineffective and AKBA caused only a weak and rather slow Ca²⁺ mobilization (fig. **441A** and **B**). β-BA was effective already at 3 μ M, though not yet significant (**fig. 441A**). At 10 μ M, the maximum elicited increase in $[Ca^{2+}]_i$ (381 ± 28 nM) was comparable to that obtained by thrombin (0.5 U/ml; 364 ± 34 nM), and exceeded the signal obtained by PAF (100 nM; 62 ± 5 nM, **fig. 441C**). However, thrombin- and PAF-induced Ca²⁺ mobilization was more rapid, peaking 5–10 s after exposure and (for thrombin) was more sustained. Collagen (8 μ g/ml) caused a slow and only moderate elevation of $[Ca^{2+}]_i$ (78 ± 7 nM, after 90 s).

 $[Ca^{2+}]_i$ was also measured in the absence of extracellular Ca^{2+} (not shown). BAs as well as thrombin evoked an internal Ca^{2+} release (no extracellular Ca^{2+}) with similar kinetics observed for the total Ca^{2+} response in the presence of extracellular Ca^{2+} , respectively. Nevertheless, in the absence of extracellular Ca^{2+} , elevation of $[Ca^{2+}]_i$ was reduced to about $37 \pm 14\%$ for thrombin and $28 \pm 17\%$ for β -BA, as compared to the total Ca^{2+} response.



С



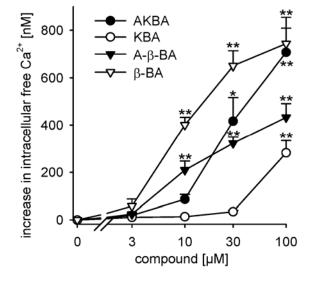


figure 441: BAs induce intracellular Ca²⁺ mobilization. To Fura-2-loaded platelets (10⁸ /ml PG buffer), 1 mM CaCl₂ was added 2 min prior stimulation, and $[Ca^{2+}]_i$ was determined. (A) Ca^{2+} mobilization in the presence of extracellular Ca^{2+} . BAs (10 mM, each) were added 30 s after the measurement was started. (B) Concentration-response curves of BAs in the presence of extracellular Ca^{2+} . The maximal increase in $[Ca^{2+}]_i$ obtained within 100 s of measurement is given. (C) Ca^{2+} mobilization induced by various agonists. The following agonists were used: β -BA (10 μ M), thrombin (0.5 U/ml), collagen (8 $\mu\text{g/ml}),$ and PAF (100 nM). Values are given as mean + s.e.m., n = 5; curves are representative of at least five experiments. One-way ANOVAs followed by Tukey HSD tests were applied to data related to unstimulated controls in (B), *p<0.05 or **p<0.01.

В

In agreement with previous studies, at a concentration of 10 μ M, AKBA (or KBA, not shown) caused only a slight and delayed (and transient) elevation of $[Ca^{2+}]_i$, whereas β -BA and A β -BA (10 μ M, each) led to substantial Ca²⁺ mobilization (**fig. 441A**). At higher concentrations (\geq 30 μ M), AKBA caused a slow yet continuous increase of $[Ca^{2+}]_i$ (not shown). Interestingly, pre-incubation of platelets for 15 min with AKBA (10 μ M) reduced Ca²⁺ mobilization induced by U-46619, PAF and collagen, but not when thrombin was used as agonist (**fig. 442A-D**).

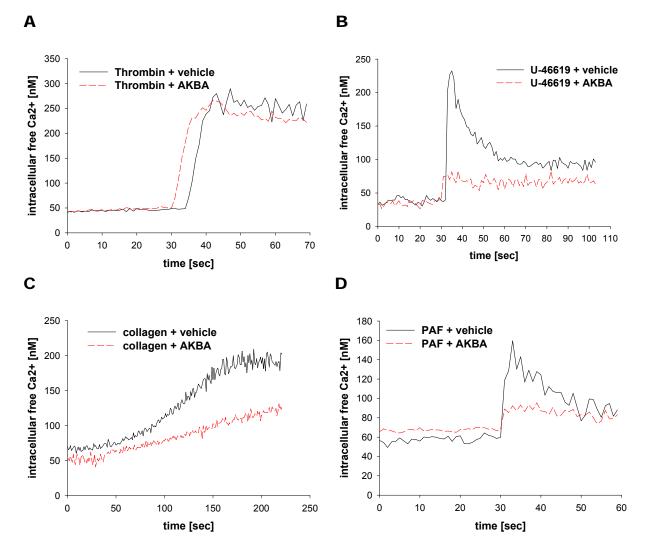


figure 442: Differential effects of boswellic acids on $[Ca^{2+}]_i$ in human washed platelets. Fura-2 loaded platelets (10⁸/ml PG buffer) were preincubated with 10 µM AKBA or with vehicle (DMSO) as indicated. After 15 min at 37°C the suspension was supplemented with CaCl₂ (1 mM) and the measurement of $[Ca^{2+}]_i$ was started. After 30 s, (A) 0.5 U/ml thrombin, (B) 1 µM U-46619, (C) 8 µg/ml collagen, or (D) 100 nM PAF were added. Curves are representative of at least four experiments.

Next, the efficacy of BAs (10 μ M, each) to prevent agonist-induced Ca²⁺ mobilization was compared. As shown in **fig. 443**, for platelets stimulated with collagen, PAF or U-46619, AKBA was most efficient to inhibit Ca²⁺ mobilization, followed by β -BA. A β -BA was somewhat less potent, whereas KBA even enhanced Ca²⁺ mobilization when compared to AKBA. Notably, all BAs essentially failed to prevent Ca²⁺ mobilization induced by thrombin (**fig. 443A**). IC₅₀ values of AKBA and β -BA for collagen-induced Ca²⁺ mobilization were 3 and 7 μ M, respectively, for PAF-induced Ca²⁺ mobilization 2 and 7 μ M and for U-46619-induced Ca²⁺ mobilization 5 and 8 μ M, respectively (**fig. 444A-D**). When thrombin was used as agonist, the IC₅₀ values for AKBA and β -BA were > 30 μ M. Pretreatment of platelets with AKBA also prevented the elevation of [Ca²⁺]_i induced by β -BA similar to that observed for collagen or U-46619.

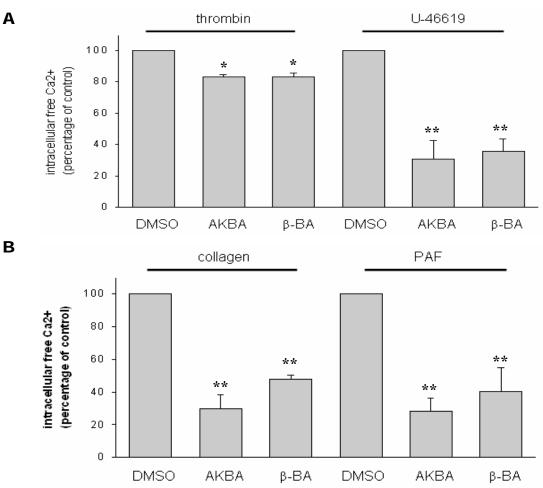


figure 443: Boswellic acids selectively suppress agonist-induced Ca^{2+} mobilization in human washed platelets. Fura-2 loaded platelets (10⁸/ml PG buffer) were supplemented with $CaCl_2$ (1 mM) and preincubated with the indicated BAs (10 μ M, each) or with vehicle (DMSO, negative control) as indicated. After 15 min at 37°C, the measurement of $[Ca^{2+}]_i$ was started and after another 30 sec, (A) 0.5 U/ml thrombin, 1 μ M U-46619, (B) 8 μ g/ml collagen, or 100 nM PAF, were added. The maximal increase in $[Ca^{2+}]_i$ determined within 100 sec of measurement is expressed as percentage of control (DMSO). Values are given as mean + s.e.m., n = 4-6 One-way ANOVAs followed by Tukey HSD tests were applied to data related to unstimulated controls in figs A to D, p<0.05 (*) or <0.01 (**)

Since β -BA ($\geq 3 \mu$ M), and to a minor degree also AKBA ($\geq 10 \mu$ M), caused a transient elevation of $[Ca^{2+}]_i$ returning to baseline after about 5 to 7 min ⁶⁷, it appeared possible that such an unspecific increase in $[Ca^{2+}]_i$ leading to desensitized platelets could be the reason for the subsequent failure of Ca^{2+} mobilization upon addition of other agonists. Accordingly, U-46619 which causes a transient Ca^{2+} mobilization similar to that observed for 11-methylene-BAs, was first added to platelets and after 15 min, platelets were stimulated with either collagen or thrombin. In contrast to 11-methylene-BAs, preincubation with U-46619 failed to substantially suppress elevation of $[Ca^{2+}]_i$ evoked by either collagen or thrombin, although a slight reduction of the signals was detected. Taken together, depending on the structure (e.g. the presence of the 11-keto- and 3-O-acetyl group), BAs differentially modulate Ca^{2+} mobilization in platelets which is further influenced by the nature of the platelet agonist.

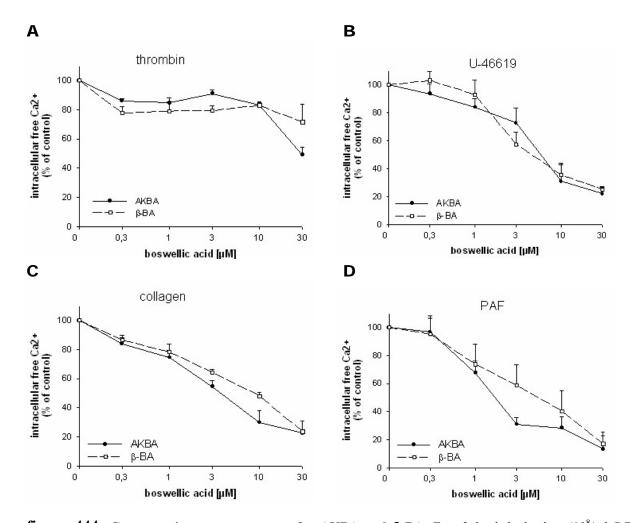


figure 444: Concentration-response curves for AKBA and β -BA. Fura-2 loaded platelets (10⁸/ml PG buffer, supplemented with 1 mM CaCl₂) were preincubated with the indicated concentrations of AKBA or β -BA and after 15 min at 37°C, the measurement of [Ca²⁺]_i was started. After 30 s, (A) 0.5 U/ml thrombin, (B) 1 μ M U-46619, (C) 8 μ g/ml collagen, or (D) 100nM PAF were added and the maximal increase in [Ca²⁺]_i was determined within 100 s, expressed as percentage of control (DMSO). Values are given as mean + s.e.m., n = 4-5.

4.4.2 Src family kinases are involved in β-BA-induced Ca²⁺ mobilization

The role of Src family kinases in β -BA-induced Ca²⁺ mobilization was assessed using the selective Src family kinase inhibitors PP2 (and its inactive analogue PP3) ¹⁴² and SU6656 ¹⁴³. PP2 (3 μ M) blunted the Ca²⁺ response initiated by β -BA (92 ± 2% inhibition, n = 7, **fig. 445**), whereas the inactive analogue PP3 (3 μ M) was hardly effective (89 ± 8% residual activity, n = 4, **fig. 445**). In addition, the structurally unrelated Src kinase inhibitor SU6656 (10 μ M) also abolished the β -BA signal (93 ± 1% inhibition, n=4, **fig. 445**). In sharp contrast, no such inhibitory effects of PP2 on Ca²⁺ signals induced by thrombin, PAF, or AKBA were apparent (not shown).

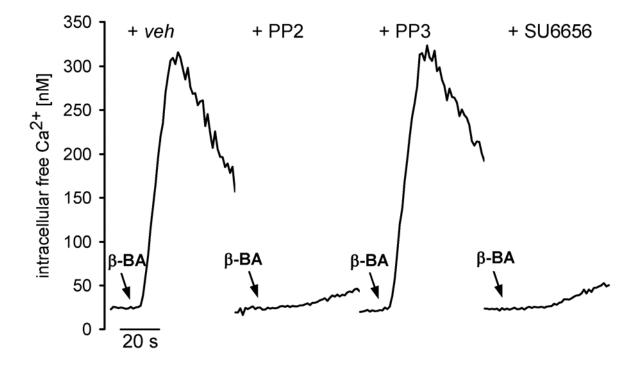


figure 445: Effects of Src family kinase inhibitors. Fura-2-loaded platelets (10^8 /ml PG buffer) were preincubated with PP2 (3 μ M), PP3 (3 μ M), SU6656 ($10 \ \mu$ M), or vehicle (DMSO) for 15 min. CaCl₂ (1mM) and β -BA ($10 \ \mu$ M) were added, and [Ca²⁺]_i was determined. Curves are representative of at least four experiments.

4.4.3 BAs and modulation of thrombin generation and expression of activation markers

β-BA was tested for its ability to generate thrombin from PRP, expressed as the endogen thrombin potential (ETP). β-BA (10 μM) potently stimulated thrombin generation, whereas AKBA (10 μM) was inactive (**fig. 446A**, **left panel**). Although collagen was only moderately superior to β-BA in the peak thrombin generation velocity, there was again a delayed onset of the β-BA effect, visualized by the kinetic progression of the ETP (**fig. 446B**). In the absence of Ca²⁺, neither stimulus induced a marked increase in the ETP over time although analysis of the ETP-AUCs revealed a slight stimulatory effect of β-BA (10 μM) as compared to DMSO and collagen that both were inactive (**fig. 446A, right panel**).

Finally, the expression of the activation markers PAC-1 (the activated GPIIb/IIIa-receptor for fibrinogen) and CD62, which indicates the release of platelet alpha-granules, were assessed. Incubations were carried out in (I) whole blood (containing 3.13% citrate), (II) recalcified PRP, and (III) washed platelets in Ca²⁺-containing PGC buffer, for 2 or 15 min. Neither β -BA (30 or 100 μ M) nor AKBA (30 μ M) led to a significant expression of CD62 and PAC-1 under all experimental settings (I–III), whereas TRAP (used as positive control) was a strong activator (**fig. 446C**).

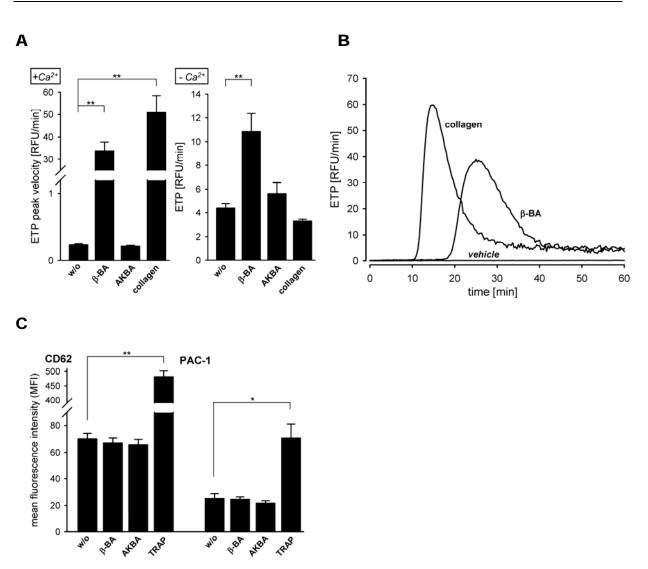


figure 446:Thrombin generation and activation marker expression. (A) Thrombin generation was assessed in recalcified PRP (given as ETP peak velocity, left bar chart), or citrate-chelated PRP (given as ETP-AUC, right bar chart). PRP and buffer containing the indicated stimuli were added to each well of a 96-well microtitre plate. β -BA (10 μ M), AKBA (10 μ M), collagen (2 μ g/ml, final concentrations each), and vehicle (DMSO) were tested for their ability to induce thrombin generation. Data are expressed as mean + s.e.m., n = 4 (β -BA, AKBA, collagen) or n=8 (vehicle). One-way ANOVA and Tukey HSD tests were performed, **p<0.01. (B) Representative original traces of the ETP kinetic progression. Cells in recalcified PRP were stimulated as described above. (C) Expression of the platelet activation markers CD62 and PAC-1. Flow cytometry in recalcified PRP was performed as described in the *Methods* section. Expression of CD62 (left bar chart) and PAC-1 (right bar chart) after stimulation with vehicle (DMSO), β -BA (30 μ M), AKBA (30 μ M), or TRAP (10 μ M) is given. The percentage of CD62-positive cells (%) as well as their mean channel fluorescence intensity (MFI) was determined (left diagram). Right, a histogram of PAC1-FITC against cell events was generated and MFI of total platelet population was recorded, n = 4. One-way ANOVA and Tukey HSD tests were performed, **p<0.05 or **p<0.01.

4.4.4 β-BA induces aracidonic acid release in human platelets

An elevation of $[Ca^{2+}]_i$ and/or activation of members of the MAPK family are considered important for the liberation of AA by the cPLA₂¹⁴⁴. In platelets AA is the substrate of COX 1 and platelet-type 12-LO. AA-metabolism by COX and 12-LO leads to prostaglandins, 12-HETE and other signal molecules in atherosclerosis and inflammation. On the other hand, AA is an important component of the lipid bilayer responsible for the fluidity of the cell membrane. Incubation of [³H]-AA-labelled platelets with β-BA caused a concentrationdependent increase in the amounts of [³H]-AA released into the medium. At about 50 μ M, β-BA was equipotent to 2 U/ml thrombin (**fig.447**).

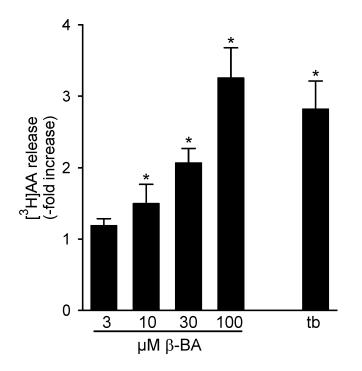


figure 447: β -BA induces the release of AA. Platelets were labelled with [³H]-AA for 2 h. CaCl₂ (1 mM) was added to the cells (10⁸ in 1ml PG buffer), and after 2.5 min, cells were stimulated with the indicated concentrations of β -BA or 2 U/ml thrombin (tb). [³H]-AA released into the medium was measured after 5 min at 37°C. Data are expressed as increase over unstimulated cells, values are given as mean + s.e.m., n = 5. Statistical analysis (directed t-tests for correlated samples) was applied to original data prior to normalisation, *p<0.05.

4.4.5 AKBA attenuates agonist-induced elevation of [Ca²⁺]_i in MM6

The attempt was made to investigate whether or not AKBA could also prevent agonistinduced elevations of $[Ca^{2+}]_i$ in monocytes. Agents that elevate $[Ca^{2+}]_i$ involving PLC/IP3 signalling (e.g. PAF, fMLP and *m*-3M3FBS) but also stimuli that raise $[Ca^{2+}]_i$ independent of the PLC/IP3 pathway like ionomycin were added to MM6 cells that received BAs, 20 s prior to agonist addition. As shown in **fig. 448**, AKBA, but not Aβ–BA, potently inhibited the subsequent Ca²⁺ mobilization induced by PAF or fMLP as well as by the direct PLC activator *m*-3M3FBS ¹⁴⁵. The IC₅₀ value for AKBA was in the range of 10 – 30 µM, depending on the stimulus. In contrast, initial elevation of $[Ca^{2+}]_i$ induced by the ER/SR Ca²⁺ ATPase inhibitor thapsigargin (TG) or by the Ca²⁺-ionophore ionomycin were not affected (not shown).

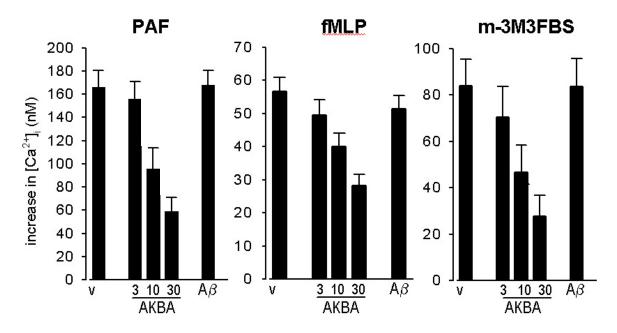


figure 448: AKBA antagonizes agonist-induced Ca²⁺ mobilization. Fura-2 loaded MM6 cells $(3x10^6/ml PG buffer, supplemented with 1 mM CaCl₂) were treated with vehicle (v), AKBA (3, 10, 30 µM), or Aβ-BA (Aβ, 30 µM) followed by the addition of PAF (0.1 µM), fMLP (0.1 µM), or$ *m*-3M3FBS (50 µM) after 20 s as indicated. The amplitude of the agonist-induced elevation of [Ca²⁺]_i was determined. Values are given as mean + s.e.m., n = 4 - 5, and compared to the positive controls.

4.4.6 AKBA attenuates Ca²⁺ mobilization from intracellular stores in monocytic cells.

Next, the test was carried out to investigate wether or not AKBA may also affect the PAFinduced release of Ca^{2+} from intracellular stores, a process that is typically PLC/IP3dependent. MM6 cells were resuspended in Ca^{2+} -free buffer containing 1 mM EDTA and treated with AKBA or Aβ-BA, followed by the addition of PAF after another 20 s. Neither AKBA nor Aβ–BA exhibited an effect on basal $[Ca^{2+}]_i$ in resting cells under these conditions. However, AKBA reduced the release of Ca^{2+} from internal stores elicited by PAF (**fig. 449B** left panel), although slightly higher concentrations of AKBA were required as compared to those needed to suppress total Ca^{2+} mobilization in the presence of extracellular Ca^2 panel) was also partially antagonised by AKBA, implying that PLC inhibition may not be the sole mechanism by which AKBA affects $[Ca^{2+}]_i$, since TG-mediated Ca^{2+} mobilization circumvents the PLC/IP3 route.

А

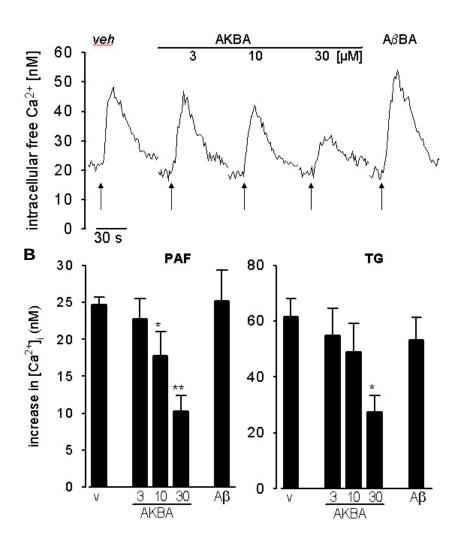


figure 449: Effects of **BAs on Ca²⁺ release from** internal stores. MM6 cells were prepared as described in fig. 448, except that 1 mM EDTA was added instead of 1 mM Ca²⁺. (A) Original $[Ca^{2+}]_i$ recordings of samples stimulated by μM) PAF (0.1 after preincubation with vehicle (veh), AKBA (3, 10, 30 μ M), or Aβ-BA (30 μ M) for 20 s. Curves are representative of 3 - 4 independent determinations. (B) Cells were treated with vehicle (v), AKBA (3, 10, 30 μM), or Aβ-BA (Aβ, 30 μ M) followed by the addition of PAF (0.1 µM, left panel), or thapsigargin (TG, 0.1 µM, right panel). The amplitude of the stimulus-induced elevation of $[Ca^{2+}]_i$ was determined. Values are given as mean + s.e.m., n = 4, p<0.05 (*) or <0.01 (**).

4.4.7 Effects of boswellic acids on TNFα-release in MM6 cells

Macrophages and monocytes are considered to be the main source of TNF α ¹⁴⁶, which as a prototypical proinflammatory cytokine, plays a key role not only in chronic inflammatory diseases but also in innate immunity ¹⁴⁷. To investigate whether Aβ-BA or AKBA are able to affect the TNF α generation in fMLP- and in LPS-stimulated MM6 cells (**fig. 4410**). TNF α is synthesized as a precursor, which is processed and released from the membrane ¹⁴⁸, implying that regulation can occur at any of those steps. Unstimulated cells did not release detectable amounts of TNF α . AKBA and Aβ-BA (10 µM each) slightly inhibit fMLP- or LPS-induced TNF α -release.

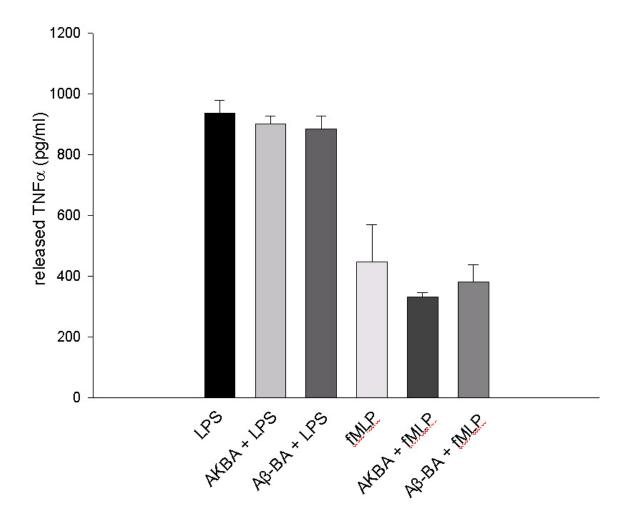


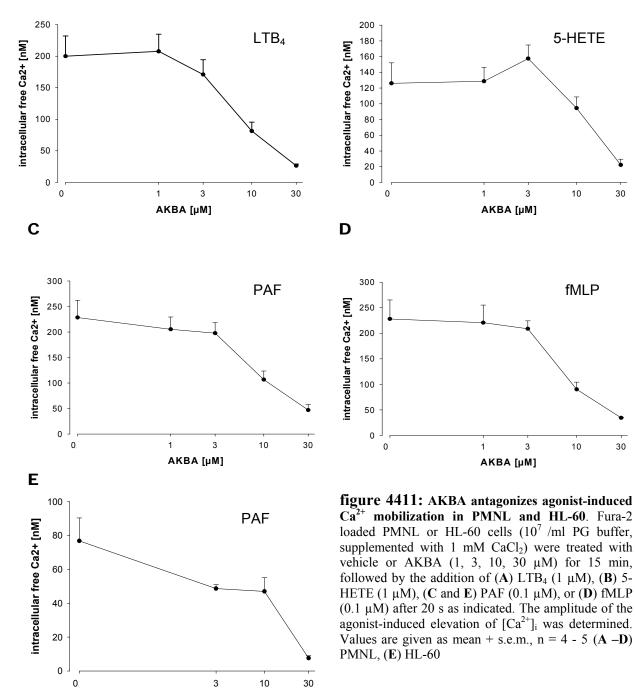
figure 4410: Effects of BAs on TNF α -release in MM6 cells. 3 x 10⁶ MM6 cells were preincubated with AKBA or A β -BA (10 μ M each) for 1 h and then stimulated with either LPS 50 ng/ml or fMLP 1 μ M. Supernatants were tested by ELISA for TNF α as described in the methods section. Results are given as mean + s.e.m., n = 3.

4.4.8 Suppression of agonist-evoked Ca²⁺ mobilization in PMNL and HL-60 cells by boswellic acids

The influence of BAs on Ca^{2+} -influx in agonist stimulated PMNL and HL-60 was investigated. The following agents were used: PAF, LTB₄, 5-HETE and fMLP for PMNL (**fig. 4411A-D**) and PAF for the leukemic cell line HL-60 (**fig. 4411E**). Preincubation with AKBA results in a concentration-dependent inhibition of agonist-evoked Ca^{2+} -influx (**fig. 4411**), the IC₅₀ values were determined at 5 to 15 μ M. AKBA is the most potent boswellic acid, 11-methylene BAs are less effective and KBA is even inactive (not shown).



В



AKBA [µM]

4.4.9 Suppression of agonist-evoked aggregation of washed human platelets by boswellic acids

Rapid and pronounced elevation of $[Ca^{2+}]_i$ in platelets is a determinant for platelet aggregation in response to various stimuli ^{69, 149}. Recently, β-BA (10 to 30 µM) strongly elevates $[Ca^{2+}]_i$ in platelets also causes platelet aggregation, although moderately and in a delayed manner, whereas in contrast AKBA was barely active even at high concentrations (30 µM). Since AKBA potently prevented the elevation of Ca^{2+} in platelets stimulated by collagen and U-46619 at rather low effective concentrations ($\leq 3 \mu$ M), it seemed reasonable that keto-BAs could inhibit aggregation induced by these agonists in a similar manner. First, the capacity of selected agonists and BAs themselves were analyzed for their ability to induce aggregation of washed platelets. As shown in **fig. 4412A**, collagen, thrombin and U-46619 caused marked aggregation of platelets within seconds or a few minutes. Differential effects for the BAs were observed: Whereas both 11-methylene BAs at 30 µM efficiently induced platelet aggregation, the 11-keto-BAs AKBA as well as KBA (30 µM, each) failed in this respect (**fig. 4412B**).

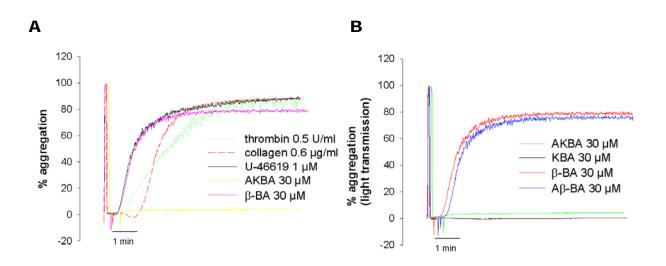
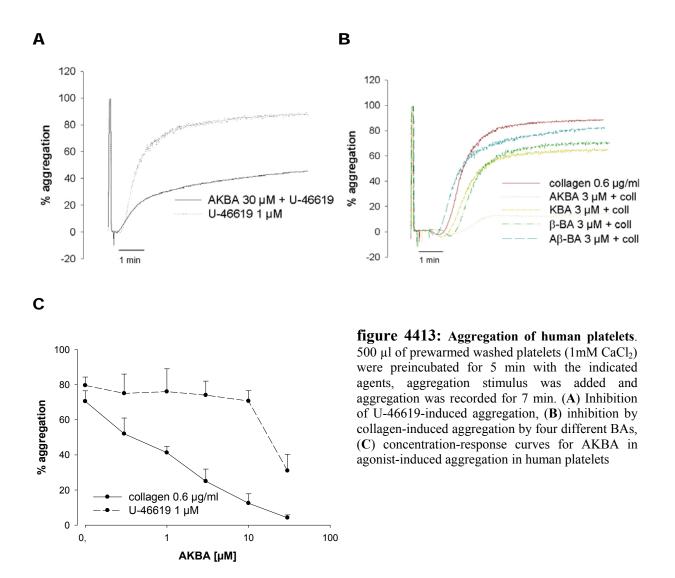


figure 4412: Aggregation of human platelets. To 500µl of prewarmed recalcified washed platelets (1mM CaCl₂) the indicated agents were added and aggregation was recorded for 7 min. (A) Aggregation induced by various stimuli; (B) BAs induced aggregation.

Next, washed platelets were preincubated with BAs (3 μ M, each) for 15 min and subsequently stimulated with collagen, thrombin, and U-46619 and aggregation was analyzed. Among the four BAs, only AKBA potently suppressed collagen-induced aggregation (**fig. 4413B**). In contrast, thrombin-evoked aggregation was not prevented by AKBA (up to 30 μ M) or any other BA (not shown). This is of interest since U-46619-induced aggregation AKBA was hardly efficient and KBA did not inhibit the effect of U-46619 (**fig. 4413A**). Detailed concentration-response studies showed that AKBA prevented platelet aggregation induced by collagen with an IC₅₀ value of 1.1 μ M, whereas for U-46619 the IC₅₀ value was approximated at 25 μ M (**fig. 4413C**). Despite the robust inhibition of collagen-induced aggregation by AKBA, only a moderate effect was measured with KBA (IC₅₀ = 10 μ M). Apparently, the 3-*O*-acetyl group and the 11-keto moiety present in AKBA render this β -configurated BA a potent inhibitor of collagen-evoked platelet aggregation, implying that defined structure-activity relationships exist excluding unspecific effects of AKBA.



4.5 Celecoxib

4.5.1 Suppression of 5-LO product formation in human whole blood

To investigate the effects of COX-2-selective inhibitors on eicosanoid formation, a human whole blood assay using Ca²⁺-ionophore A23187 as stimulus was applied, equipped with a highly sensitive LC/MS-MS methodology capable of selectively detecting a broad number of eicosanoids ¹⁴⁰. Celecoxib but no other COX-2 inhibitor (up to 100 μ M) concentration-dependently inhibited the formation of the 5-LO products LTB₄ and 5-HETE with IC₅₀ values \approx 30.9 and 42.3 μ M, respectively (**fig. 451 A** and **B**). Zileuton, a well-known iron-ligand inhibitor of 5-LO ¹⁵⁰ used as positive control, caused significant reduction of LTB₄ and 5-HETE already at 1 μ M. All COX inhibitors reduced PGE₂ formation with diclofenac and celecoxib showing the highest efficacy (**fig. 451 C**). Interestingly, celecoxib failed to significantly suppress the levels of 12(S)-HETE (**fig. 451D**), and the formation of 15(S)-HETE (**fig. 451E**) was only slightly decreased (IC₅₀ > 100 μ M). Therefore, besides inhibition of PGE₂ synthesis, celecoxib has the unique property among COX inhibitors tested to suppress the formation of 5-LO products¹⁵¹. This work was done in collaboration with Dr. T.J. Maier.

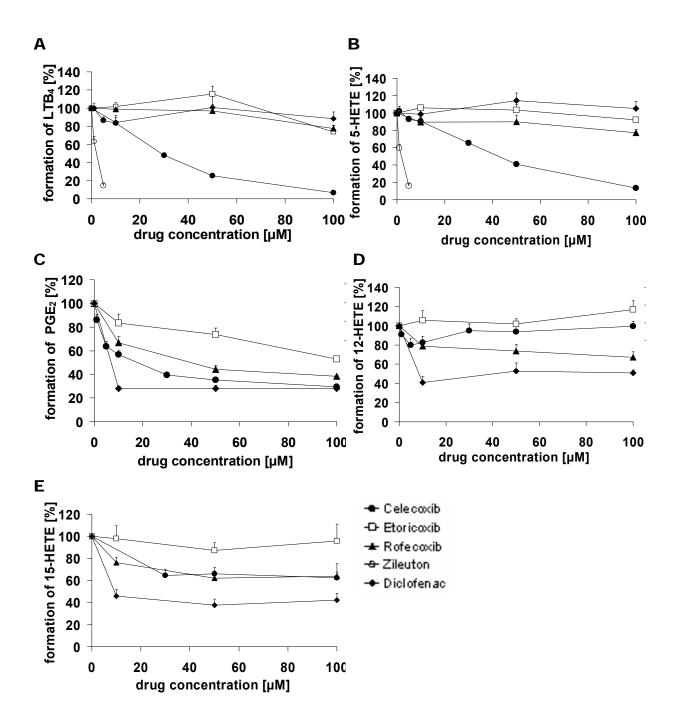


figure 451: Effects of COX inhibitors on eicosanoid formation in human whole blood. The experimental procedure is described in *Methods*. The eicosanoids in the plasma supernatant analyzed by LC/MS-MS were LTB₄ (**A**), 5-HETE (**B**), PGE₂ (**C**), 12(S)-HETE (**D**) and 15(S)-HETE (**E**). For LTB₄, 5-HETE and 12(S)-HETE data are given as mean + s.e.m. of 8 (celecoxib and zileuton; n = 14-26 values), 4 (etoricoxib, rofecoxib; n = 11-13 values) and 2 (diclofenac; n = 6 values) independent experiments. For PGE₂ and 15-HETE data are given as mean + s.e.m. of \geq 4 (celecoxib, n = 12-15 values) or \geq 2 (rofecoxib, etoricoxib, diclofenac, n = 5-6 values) independent experiments.

4.5.2 Inhibition of 5-LO product formation in activated human PMNL

The efficacy of celecoxib was assessed in A23187-stimulated human PMNL, a frequently used model for evaluation of 5-LO inhibitors ¹⁵². Pretreatment (10 min) of PMNL with celecoxib caused a concentration-dependent inhibition of 5-LO product formation with an IC₅₀ \approx 8.4 µM (**fig. 452A**). To exclude effects of celecoxib on the availability of endogenous AA as substrate and consequently circumvent cPLA₂ activity, exogenous AA (2, 10, or 20 µM) was provided. No alteration in the efficacy of celecoxib on 5-LO product formation was observed. Data are shown in **fig. 452A** for incubations with 20 µM AA (IC₅₀ \approx 7.0 µM) and without AA. In agreement with the results obtained from whole blood assays, other coxibs (i.e. etoricoxib and rofecoxib) as well as the unselective COX-inhibitor diclofenac (up to 30 µM, each), showed no or only weak inhibition of 5-LO activity in A23187-activated PMNL, irrespective of the presence of exogenous AA (**fig. 452B**). BWA4C, a potent 5-LO inhibitor, structurally related to zileuton, was used as positive control. In summary, only celecoxib among various COX inhibitors suppresses 5-LO product synthesis in activated intact human PMNL.

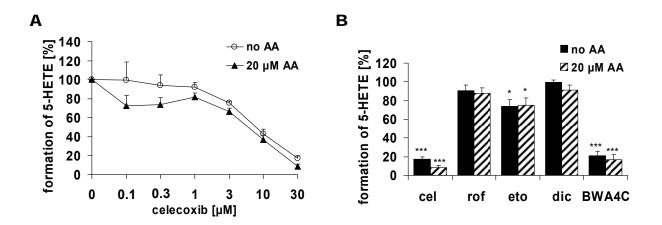


figure 452: (A) Inhibition of 5-LO product formation in intact PMNL by celecoxib. 5-LO activity in absence or presence of AA (20 μ M) was assayed as described in *Methods* and expressed as percentage compared to vehicle control (100 %). Results are mean + s.e.m., n = 6. (B) 5-LO-inhibitory activity of celecoxib, rofecoxib, etoricoxib, diclofenac and BWA4C in intact PMNL in the absence (black bars) or presence (striped bars) of 20 μ M AA as compared to vehicle control (100 %). For experimental procedures see *Methods*. Results are mean + s.e.m., n = 5. ***, $p \le 0.001$; *, $p \le 0.05$. cel, celecoxib; eto, etoricoxib; rof, rofecoxib; diclofenac; AA, arachidonic acid

4.5.3 Inhibition of 5-LO activity of in cell-free assays

5-LO is a tightly regulated enzyme and suppression of cellular product formation by any compound does not unequivocally indicate a direct interference with 5-LO catalysis. Thus, cellular regulatory components or mechanisms such as FLAP, mitogen-activated protein kinases (MAPKs), Ca²⁺ mobilization, nuclear membrane association ⁷⁹ and interaction with CLP¹⁵³ might theoretically be targeted by celecoxib. To demonstrate a direct action on 5-LO, the effects of celecoxib on 5-LO product synthesis were analyzed in cell free assays. Celecoxib concentration-dependently inhibited 5-LO product formation in homogenates supplemented with 20 μ M AA, though less potent (IC₅₀ \approx 23.4 μ M, fig. 453A) as compared to intact cells (IC₅₀ \approx 8.4 μ M, fig. 451A). Such a loss of efficacy in cell-free assays was observed also for nonredox-type 5-LO inhibitors, ¹⁵⁴ but addition of thiols (to reduce the lipid hydroperoxide levels via glutathione (GSH) peroxidases) restored potent inhibition. However, addition of GSH had no significant effect on the efficacy of celecoxib (fig. 453A, $IC_{50} \approx 20.5$ μ M). In accordance with the results obtained from the whole blood assay, celecoxib (up to 30 μ M) did not suppress the formation of 12(S)-HETE or 15(S)-HETE in homogenates of human platelets or eosinophils, respectively (fig. 453B). Finally, celecoxib concentrationdependently inhibited the activity of partially (ATP-affinity) purified human recombinant 5-LO with an IC₅₀ \approx 30.9 μ M (fig. 453C). Rofecoxib, etoricoxib and diclofenac (up to 100 μ M) caused no or only modest inhibition of 5-LO activity in the cell free assays (fig. 453D and E). Taken together, celecoxib can thus be said to be a direct inhibitor of 5-LO.

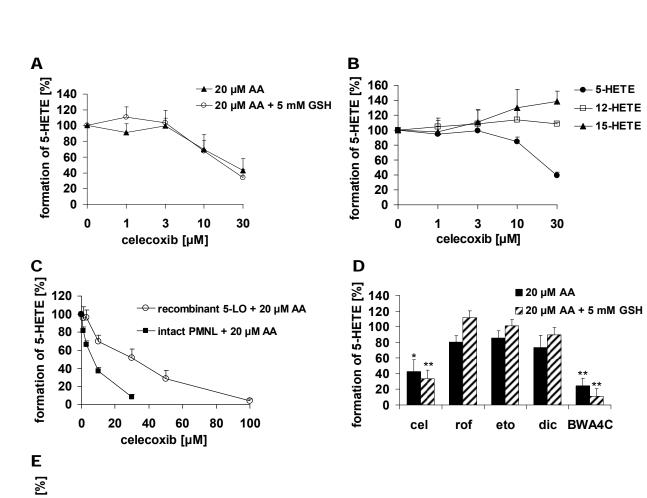


figure 453: Effect of celecoxib, etoricoxib, rofecoxib, diclofenac and BWA4C on 5-LO activity in cellfree assays. 5-LO activity is expressed as percentage compared to vehicle control (100 %). Respective 5-LO activities were determined as described in *Methods*. (A) Inhibition of 5-LO product formation in broken cell homogenates by celecoxib in presence or absence of 5 mM GSH. Results are given as mean + s.e.m., n = 4. (B) Determination of 5-, 12- and 15-LO product formation in broken cell homogenates in the presence of increasing concentrations of celecoxib, 20 μ M AA and 25 mM GSH. Results are given as mean + s.e.m., n = 3. (C) Inhibition of partially purified human recombinant 5-LO by celecoxib as compared to intact PMNL. Results are given as mean + s.e.m., n \geq 5. (D) Comparison of the 5-LO-inhibitory activity of celecoxib, rofecoxib, etoricoxib, diclofenac and BWA4C in PMNL homogenates in the absence (black bars) or presence (striped bars) of 5 mM GSH as compared to vehicle control (100 %). Results are given as mean + s.e.m., n = 4. (E) Effect of celecoxib, rofecoxib, etoricoxib, diclofenac and BWA4C on recombinant 5-LO activity as compared to vehicle control (100 %). Results are given as mean + s.e.m., n = 3. ***, $p \leq 0.001$; **, $p \leq 0.01$; *, $p \leq 0.05$. cel, celecoxib; eto, etoricoxib; rof, rofecoxib; dic, diclofenac.

formation of 5-HETE

rof

eto

cel

dic BWA4C

5 Discussion

5.1 Cathepsin G is a target of boswellic acids

Our findings provide strong evidence that human CatG is a high-affinity target of BAs. By means of a protein-fishing approach, besides several other proteins, a 25.8 kDa protein from lysates of human PMNL was found to selectively bind to an affinity matrix composed of KBA, linked covalently to insoluble EAH Sepharose 4B. MALDI-TOF-MS analysis combined with immunological detection revealed that this 25.8 kDa BA-binding protein is CatG. All four β -configurated BAs inhibited the activity of human CatG in vitro. With IC₅₀ values of 0.6 and 0.7 μ M, respectively, AKBA and β -BA were most potent. The proteolysis of related serine proteases was less efficiently inhibited, confirming the selectivity of BAs for CatG. Moreover, BAs potently inhibited CatG-mediated cellular responses such as chemoinvasion of PMNL, and Ca²⁺ mobilization in platelets evoked by fMLP-stimulated PMNL. Hence, BAs not only bind CatG but also functionally interact with this protease. Along these lines, AKBA with a high average docking score docked into the active site of CatG, where the proteolysis of the substrate takes place. Since CatG plays a role in inflammatory diseases such as asthma, psoriasis and rheumatoid arthritis $^{28, 155}$ for which B. spec extracts have proven to possess therapeutic benefit ⁵, the high-affinity BA-CatG interaction may provide a molecular basis for these beneficial effects and might be of pharmacological relevance.

BAs, in particular AKBA, have been proposed as the main active principles of *B. spec* preparations that are traditionally used to cure chronic inflammatory disorders ^{4, 5}. Initial attempts to elucidate the molecular mechanisms responsible for the pharmacological actions of *B. spec* extracts identified 5-LO as a reasonable target of AKBA. Thus, AKBA as well as *B. spec* extracts reduced the formation of LTs in activated leukocytes ^{8, 65, 156, 157} and AKBA inhibited 5-LO activity in cell-free assays ^{65, 101, 156} or isolated 5-LO enzyme ¹⁵⁶. Aside from AKBA, other BAs were either hardly active or not active at all ⁶⁵. Since LTs play pivotal roles in inflammatory reactions ⁷⁹ and because *B. spec* extracts show clinical effects similar to typical 5-LO inhibitors, it appeared reasonable that intervention of BAs with 5-LO and thus with LT formation, is the molecular basis underlying the anti-inflammatory effectiveness of *B. spec* extracts observed *in vivo* ⁵.

Besides 5-LO, AKBA was shown to interfere with several other enzymes or signalling events that contribute to inflammation including HLE 10 , p12-LO 9 , the NF κ B pathway 12 ,

intracellular Ca²⁺ levels and MAPKs activities ¹⁷. However, the required concentrations in the respective experimental systems frequently appeared rather high (> 10 μ M) ^{10, 18, 101, 157}, whereas the plasma concentrations of AKBA obtained after oral application of standardised *B. spec* extracts are in the submicromolar range ^{26, 27}. The most affine AKBA target was proposed to be 5-LO, but strikingly different potencies (IC₅₀ = 1.5 up to 50 μ M) were reported ^{8, 65, 101, 104}. These discrepancies seemingly depend on the different experimental settings (intact cells or cell-free assays, different species and cell type, AA concentration etc.). Thus, AKBA was most efficient to inhibit 5-LO in Ca²⁺ ionophore-stimulated PMNL from rat (IC₅₀ = 1.5) ⁸, whereas for direct interference with isolated human 5-LO, the IC₅₀ was determined at 16 μ M ¹⁵⁶, a phenomenon that needs to be resolved. Taken together, the pharmacological relevance of most of the proposed target interactions and effects of AKBA observed *in vitro* and the pharmacological relevance *in vivo* are a matter of debate.

In fact, isolated CatG binds to KBA-Seph, but not to Seph without ligand, in the same manner as CatG in PMNL lysates. Most of the other identified proteins that selectively bind to KBA-Seph are directly or indirectly involved in inflammatory processes, immune responses or cancer formation ^{30, 53, 63, 134, 158}.

The successful identification of CatG as target of BAs by means of the fishing approach using the affinity resin is the result of the combination of several fortunate circumstances. First, CatG is a rather abundant protein in PMNL. Second, the relatively high affinity of CatG to KBA-Seph enabled binding under the stringent conditions (1% Triton X-100) and the ability to endure the thorough and long-lasting washing (overnight) of the precipitates. Third, KBA is a highly rigid molecule with only few conformations favourable for binding to CatG. Finally, the C3-OH moiety used to link KBA to the resin via an ester bound is not detrimental for binding, in fact esterified BAs (i.e. AKBA or A β -BA) are virtually equipotent or even more potent than those with a free OH group (i.e. KBA and β -BA).

In collaboration with HLE, CatG is largely responsible for the destruction of bacteria by neutrophils ²⁸. Both proteases as well as proteinase-3 are not only involved in defence against bacterial infections, these enzymes are also important players in ulcerative colitis and rheumatoid arthritis ¹⁵⁹⁻¹⁶² where BS extracts are beneficial ^{49, 163}. Moreover, this work demonstrates that CatG activity can be reduced by oral administration of BS extracts in healthy volunteers and patients with ulcerative colitis in remission (*ex vivo*). Our findings might prove useful for the development of new drugs for treatment of bowel diseases.

5.2 DNA-PK and Akt are targeted by BAs

The ability of apoptotic stimuli to induce cell death is counteracted by the activity of antiapoptotic proteins. The PI3K/Akt pathway downstream of RTKs plays an important role in cell survival. It was shown before that PKCξ acts as an antiapoptotic protein during TNFαinduced cell death ¹⁶⁴ and that there is a cross-talk between PKCE and Akt. PKCE acts upstream of Akt/PKB to exert its antiapoptotic function 165 . Activation of Akt by PKC ξ is mediated by DNA-PK, a member of the PIKK subfamily of protein kinases, and depletion of DNA-PK reversed the antiapoptotic function of PKC ξ during TNF α -induced apoptosis ¹⁶⁵. The observation that inhibition of DNA-PK can reverse antiapoptotic signalling by Akt establishes a new role for DNA-PK in the extrinsic cell death pathway. Multiple signalling pathways that exist in certain cell types might regulate/influence the final outcome of cell death or survival. Cells that are resistant to TNF α contain constitutively active Akt ¹⁶⁶. Activation of Akt is an early event following binding of $TNF\alpha$ to its cell surface receptors. Complete activation of Akt requires phosphorylation at Thr308 and Ser473 by PDK1 and PDK2, respectively¹⁶⁷. Since phosphorylation of Akt at Ser473 is mediated by PDK2 and since DNA-PK has been recently identified as PDK2⁵⁵, the recruitment of DNA-PK during DNA damage-induced apoptosis is reasonable. In fact, DNA-PK is activated in response to DNA damage ¹⁶⁸ and autophosphorylation of DNA-PK has been shown to inhibit DNA-PK activity ¹⁶⁹. These results suggest that DNA-PK may also play a critical role in receptorinitiated cell survival via activation of Akt/PKB. The mechanism of how DNA-PK localizes to the membrane is not clear, but DNA-PK has been reported to be associated with epidermal growth factor receptor ⁵⁷.

It was shown, that activation of Akt via DNA-PK inhibits TNFα-induced apoptosis in breast cancer cells ¹⁶⁵. BAs inhibit LPS-mediated TNFα induction in monocytes ¹⁰⁰. The cross-talk between multiple signalling pathways is an important determinant of cell survival and cell death, but is currently not well understood. The involvement of DNA-PK during DNA damage induced apoptosis is well known and a new role for DNA-PK during receptor-initiated apoptosis was established ⁵⁵. DNA-PK is activated upon DNA damage by UV irradiation, as is Akt ⁵⁸. Induction of apoptosis by cisplatin was explained by a decrease in DNA-PK activity through proteolytic degradation of DNA-PK ⁶⁰. Interestingly, Akt activity and Ser473 phosphorylation are also inhibited by cisplatin treatment ^{61, 170}. Mouse and human cells deficient in DNA-PK are hypersensitive to ionizing radiation and to radiomimetic drugs ¹⁷¹⁻¹⁷³; a similar phenotype can be observed in Akt1/PKB knock-out mice ¹⁷⁴ and with mouse

embryo fibroblasts derived from such mice ¹⁷⁵. It was often reported that BAs induce apoptosis in several cancer cell lines in vitro and reduce tumour growth in vivo ^{13, 14, 16, 18} Modulation of DNA-PK and inhibition of Akt by BAs may be responsible for induction of apoptosis in several tumour cell lines ^{13 14 16}.

Akt is an upstream kinase of the IKK pathway which is indirectly inhibited by boswellic acids ^{12, 18}. Akt is bound selectively by immobilized KBA (KBA-Seph). Beneath NF κ B pathway (IKK) and MAPK pathway, Akt influences several other pathways like cell growth, cell death, cardiovascular homoeostasis (eNOS), JNK and glycogen synthesis. Akt is inhibited potently (A β -BA IC₅₀ = 300 nM) and directly in vitro but also in vivo by BAs. Phosphorylation of Akt on Ser473 was reduced by BAs and phosphorylation of the direct downstream-target Bad by Akt was inhibited (Bad phosphorylation in LNCaP). These results provide strong evidence that Akt is a high affinity target of BAs. The fact that also Akt phosphorylation by DNA-PK is inhibited *in vitro* and *in vivo* implies that BAs may inhibit the interaction of Akt and DNA-PK, which results in the inactivation of their activity.

The DNA-PK pathway plays a crucial role in controlling transcription, cell cycle progression, and apoptosis ¹⁷¹. Similarly, Akt is also implicated in the regulation of many different cellular processes ^{53, 54}. It was shown that BAs induce apoptosis in cancer and cancer cell lines^{13, 14, 18}. These anticarcinogenic effects may be influenced by inhibition of DNA-PK and Akt. Hyperproliferation and overactivation of RTKs in cancers lead to invasiveness and metastasis of tumours. BAs block the growth signal on an early stage of the pathway. Because Akt is potently inhibited by Aβ-BA (IC₅₀ = 300 nM), *in vivo* relevance could be implicated; plasma levels up to 6.4 μ M β-BA / 4.9 μ M Aβ-BA have been achieved. In conclusion, the RTK/PI3K pathway is directly modulated by BAs, DNA-PK and Akt are the main targets.

Beneath Akt and DNA-PK, other BA-binding proteins are involved in cancer development and immune response ^{30, 53, 63, 134, 158}. Farnesyl pyrophosphate synthetase (FPPs) is the target of the bisphosphonates, which are effective in the treatment of osteoporosis. Though FPPs is not influenced by BAs *in vitro*, there could still be a certain relevance *in vivo* ¹⁷⁶. Proteinase-3 is a human protease involved in tissue degradation and penetration of macrophages. Proteinase-3 is very similar in sequence and structure to CatG and HLE, but inhibition by BAs is weak (IC₅₀ ~ 30 μ M). Moreover, other proteins found by pulldown experiments, e.g. cathelicidin antimicrobial peptide (hCAP18), UNC-112 and prohibitin are important key mediators in inflammation and the autocrine immune response, but the whole context still remains unclear ¹⁵⁸.

5.3 Modulation of signal transduction and functionality of platelets and monocytes by boswellic acids

The current opinion regarding the molecular mechanisms underlying the anti-inflammatory properties of BS extracts or BAs, favours 5-LO as the most relevant target ⁵, but AKBA interferes with several other relevant enzymes or signalling events that contribute to inflammation as described above. In this work, additional anti-inflammatory implications of AKBA, that is, suppression of agonist-induced Ca²⁺ mobilization in platelets and platelet aggregation, are implicated. These inhibitory actions may contribute to the anti-inflammatory effectiveness of AKBA-containing extracts of B. *spec*. observed *in vivo* as well. Moreover, inhibition of platelet aggregation suggests novel beneficial properties of AKBA in cardiovascular diseases such as atherosclerosis, heart attack and stroke that are characterized by aggregating platelets and thrombotic states ¹⁷⁷. However, besides AKBA, BS extracts also contain 11-methylene-BAs that activate platelets and when BS extracts were analysed, the overall effects on platelets resembled those of 11-methylene-BAs.

As mentioned before, there is a discrepancy between plasma levels of BAs and the concentrations required for inhibition of proposed targets. Along these lines, it was previously shown that in platelets, AKBA at concentrations \geq 30 μ M caused moderate elevation of $[Ca^{2+}]_i$ and activation of MAPKs, accompanied by the release of AA ⁶⁷. Furthermore, our group showed before that relatively high concentrations of AKBA (10 up to 100 µM) stimulated the mobilization of Ca²⁺ and the activation of MAPK in PMNL ⁶⁶, coupled to the formation of reactive oxygen species (ROS), the release of AA and its metabolism to leukotrienes, similarly as observed for the chemoattractants fMLP or PAF ¹⁰⁴. Such actions of AKBA suggest a pro-inflammatory rather than an anti-inflammatory potential. However, these events were evoked only at quite high concentrations of AKBA ($\geq 10 \ \mu M$) ^{17, 66, 67, 104} that we consider not pharmacologically relevant in vivo. Intriguingly, in the present study, effective concentrations of AKBA (\leq 3 µM) significantly antagonised Ca²⁺ mobilization evoked by the platelet agonist collagen and suppressed agonist-induced platelet aggregation. Note that AKBA itself is unable to activate platelets at such low concentrations ⁶⁷. AKBA was somewhat less efficient to suppress Ca^{2+} mobilization and platelet aggregation induced by U-46619, and when thrombin was used to evoke these responses, AKBA even failed. Presumably, AKBA targets a component pivotal in the signal transduction of collagen (and U-46619), which apparently is dispensable for thrombin-induced pathways leading to Ca^{2+} mobilization and platelet aggregation. It should be noted that the signal transduction pathways

of collagen and U-46619 or thrombin are quite distinct and, at least in part, utilize different types of signalling molecules. U-46619 and thrombin act on GPCRs and signals via $G_{i/q}$ proteins and PLC to rapidly release Ca^{2+} from IP₃-sensitive stores ¹⁷⁸, whereas collagen binds integrin $\alpha_2\beta_1$ and glycoprotein VI and slowly allows Ca^{2+} entry via a Src family kinase/PLC γ -mediated pathway ¹⁷⁹. For β -BA, Src family kinases and the PLC/IP3 pathway seem to be involved in Ca^{2+} mobilization, and β -BA causes activation of ERK2 and the PI3K/Akt route. Moreover, β -BA induces the release of AA, a pronounced generation of thrombin, and Ca^{2+} dependent platelet aggregation. In contrast, AKBA-induced Ca^{2+} mobilization is not connected to Src family kinases and PLC/IP3 signalling, and AKBA failed to induce phosphorylation of Akt and ERK2, as well as functional platelet responses. Detailed analysis of the interference of AKBA with these signalling pathways is needed to reveal the molecular modes of action.

Among the BAs tested for induction of Ca^{2+} mobilization, β -BA is the most potent analogue. At 10 μ M, the effectiveness of β -BA exceeded that of PAF or collagen, and was comparable with that of the potent platelet agonist thrombin. Such β -BA concentrations are in the range of β -BA levels in human plasma (10.1 μ M), determined after oral application of 4 x 786 mg BS extract/day within 10 days²⁷. The 3-O-acetyl group slightly hampers (receptor-)activation and the 11-keto moiety significantly decreases the potency and also alters the signalling routes in platelets. In sharp contrast to platelets, only 11-keto BAs, but not 11-me-BAs, caused stimulation of PMNL ^{66, 104}. Possibly, PMNL and platelets selectively express closely related but not identical receptors specific for AKBA or 11-me-BAs, respectively. Important receptors for soluble agonists known to regulate $[Ca^{2+}]_i$ in platelets are the purinergic P_2X_1 and P_2Y_{12} receptors, the TXA₂ receptor, the PAF receptor, the 5-HT_{2A} receptor, and the PAR-1 and -4^{180} . Whether β -BA acts at one (or more) of these receptors is unknown. However, antagonists of thrombin (argatroban), PAF (WEB 2086), and ADP (NF449 and MRS2179) did not affect β -BA-induced Ca²⁺ mobilization. Thrombin is the most potent platelet agonist acting via PAR-1 and -4^{181, 182} and the influence of 2-APB on cellular Ca²⁺ influx systems has been reported ¹⁸³. Interestingly, the proximal routes mediating PLC/IP3-dependent Ca²⁺ mobilization appear to be different for β-BA and thrombin (or PAF). Thus, Src family kinase inhibitors abolished the β -BA-induced response, but not the responses elicited by thrombin or PAF. PLC γ is the most abundant PLC isoform in platelets ¹⁸⁴ and is an operative element in Ca²⁺ mobilization in response to activation by adhesion receptors ¹⁸⁵. Whereas, soluble ligands such as thrombin, ADP, PAF, or TXA₂ act via GPCRs to stimulate PLC_β isoenzymes,

the PLC γ isoforms are regulated through phosphorylation by Src family kinases ¹⁸⁵. In analogy to agonists that act via adhesion receptors but unlike thrombin, β -BA may utilise the Src family kinases/PLC γ pathway to induce Ca²⁺ mobilization. Another difference between β -BA- and thrombin-mediated Ca²⁺-mobilization is the significant delay of the response to β -BA as compared to the rapid effect of thrombin. Aside from acting as a direct ligand at a certain (adhesion) receptor, β -BA may possibly first induce the generation of an endogenous agonist that in turn causes PLC- γ /IP3-coupled Ca²⁺ mobilization via (adhesion) receptors. Attempts to unravel a putative autocrine mode of action are in progress in our laboratory.

Typical platelet agonists such as thrombin, collagen, or TXA₂ activate PI3K and its downstream effector Akt, important mediators of agonist-induced platelet activation ¹⁸⁶, as well as p38 MAPK and ERKs ^{71, 72, 187}. The MAPK are a point of convergence of complex signalling networks, regulating cell proliferation and differentiation ⁷². In platelets, the functions of MAPK are mainly uncharacterised and the signal transduction steps are poorly understood. All BAs tested activated p38 MAPK with similar efficacy, but only β-BA (and Aβ-BA) rapidly and significantly activated ERK2. Also, β-BA, but not AKBA, evoked Akt phosphorylation, and in analogy to thrombin, the PI3K and/or the PLC/Ca²⁺ pathwav are involved. Therefore, the receptor for BAs mediating p38 MAPK activation might differ from those transmitting signals to activate ERK2 and Akt. The latter (11-me-BA specific) receptor may also mediate increases in $[Ca^{2+}]_i$, generation of thrombin, release of AA and aggregation, since AKBA and KBA failed to elicit these events. Investigation of the platelet functions elicited by β-BA provided controversial results. As a rule, the distinct responses of activated platelets depend on the strength (potency) of the agonist, and these responses can be arranged in an activation sequence: (1) aggregation, (2) granule secretion, (3) AA liberation, and (4) acid hydrolase secretion 188 . The magnitude of Ca²⁺ mobilization is an important parameter for the induction of these responses. In fact, β -BA (10–30 μ M) substantially elevated [Ca²⁺]_i and potently induced thrombin generation, being equipotent in this respect with collagen at 2 mg/ml in a model utilising native platelets. Also, β -BA potently evoked the liberation of free AA from washed platelets, although at concentrations slightly higher than those required for Ca²⁺ mobilization, probably due to the presence of fatty acid-free albumin that may bind BAs. In general, liberation of free AA is a response distal of aggregation and degranulation, and its induction normally requires potent agonist-activating platelets with high strength. Surprisingly, however, the efficacy of β -BA was greatly reduced by the induction of aggregation. In contrast to collagen, the response of β -BA was strictly dependent on the

presence of extracellular Ca^{2+} . Therefore, β -BA may facilitate aggregation by other factors rather than being a full agonist. Moreover, β -BA failed to induce degranulation and fibrinogen receptor activation (CD62, PAC-1 expression). Together, despite the pronounced elevation of $[Ca^{2+}]_{i}$, only select functional platelet responses were observed after stimulation with β -BA. Along these lines it was found that platelets in polycythaemia vera exhibit decreased aggregation after stimulation with PAF, although an equal increase in $[Ca^{2+}]_i$ was seen as compared to platelets from healthy donors ¹⁸⁹. Furthermore, a patient was described with defective platelet aggregation in response to Ca^{2+} ionophore A23187, despite normal increases in $[Ca^{2+}]_i^{190}$. Hence, elevation of $[Ca^{2+}]_i$ in platelets is one important signalling step for eliciting various platelet responses, but must not necessarily lead to the induction of all Ca^{2+} -dependent platelet functions. It is conceivable that β -BA on one hand is a platelet agonist that potently induces central signalling pathways (Ca²⁺ mobilization, MAPK/Akt phosphorylation) and select responses such as thrombin generation and AA release, but on the other hand lacks the stimulation of certain signalling components or executing molecules particularly important for a rapid aggregation, degranulation, and fibrinogen receptor activation. At the present time, the findings cannot be readily related to the antiinflammatory properties of BS extracts, observed in animal models or in studies with human subjects ⁶. Nevertheless, due to its high effectiveness and the importance of the signalling molecules and the select platelet functions induced, the receptor(s) mediating the actions of β -BA in platelets warrant further elucidation. Effective concentrations of β -BA (10 μ M) are in range of β -BA levels in human plasma (see above), but platelet agonism does not appear to be relevant in vivo (see above).

Besides the agonism of 11-me-BAs to Ca^{2+} -influx in human platelets, particularly the keto-BAs antagonize potently agonist-induced increases in $[Ca^{2+}]_i$ in monocytes, PMNL and HL-60-cells. These findings indicate that keto-BAs especially AKBA desensitize blood cells to endogenous inflammation-stimuli. Based on present findings and previous studies, the 3-*O*-acetyl group and the 11-keto moiety strongly determine the pharmacodynamics in the particular assay system, and AKBA is frequently the most effective or the exclusive active analogue ^{9, 17, 65, 66}. In this work, AKBA suppressed Ca²⁺ mobilization induced by collagen or U-46619 with almost equal potencies (IC₅₀ = 3 and 5 μ M), implying the interference with a common type of target or pathway distal of Ca²⁺ release. Besides AKBA, also β-BA reduced U-46619- and collagen-induced Ca²⁺ mobilization, though less efficient (IC₅₀ = 7 and 8 μ M, respectively). Accordingly, one would expect that KBA, containing an 11-keto moiety or Aβ-BA, possessing a 3-*O*-acetyl-group should be more potent than β-BA. However, KBA caused no suppression at all but rather enhanced the signals induced by collagen, U-46619 and also by thrombin, and Aβ-BA was less effective than AKBA or β-BA. It is possible that AKBA and β-BA act at different targets both leading to suppression of Ca²⁺ mobilization.

The finding that β -BA (and A β -BA) suppressed agonist-induced Ca²⁺ mobilization was actually quite surprising, since these 11-methylene-BAs themselves cause a transient increase of $[Ca^{2+}]_i$ in platelets (this study and ⁶⁷). Such transient increase of $[Ca^{2+}]_i$ could desensitize platelets by unspecific actions (e.g. depletion of intracellular Ca²⁺ stores) and thus be the reason for the subsequent failure of Ca²⁺ release evoked by a second agonist. However, elicitation of Ca²⁺ mobilization by U-46619 and subsequent stimulation of platelets with collagen or thrombin did not substantially prevent the effects of the agonists as in the case of BAs.

At the moment it is not possible to provide a molecular basis explaining the puzzling effects of the various BAs on the Ca²⁺ homeostasis in platelets. Nonetheless, since elevation of $[Ca^{2+}]_i$ is a determinant for aggregation of washed platelets ^{69, 149}, it is reasonable to speculate that inhibition of collagen-induced aggregation by AKBA could be the result of suppressed Ca^{2+} mobilization. On the other hand, lower concentrations of AKBA were sufficient to suppress collagen-evoked aggregation ($IC_{50} = 1.1 \ \mu$ M) as compared to those required to inhibit Ca^{2+} mobilization ($IC_{50} = 3 \ \mu$ M). Moreover, AKBA was about equipotent for suppression of Ca^{2+} mobilization in response to U-46619 as compared to collagen, but U-46619-evoked aggregation was hardly affected by AKBA ($IC_{50} = 25 \ \mu$ M). Therefore, the inhibitory effects of ABKA on agonist-induced Ca^{2+} mobilization and on aggregation also might be separated. However, it is possible that already a minimal impairment of $[Ca^{2+}]_i$ as in the case of collagen is sufficient to substantially affect aggregation. In contrast to AKBA, β -BA at higher concentrations caused aggregation itself⁶⁷. Once again, this indicates that AKBA interacts with other molecules required for aggregation, and thus concrete structureactivity relationships may exist in this respect.

In summary, AKBA (and to a minor extent also β -BA) acts as an efficient blocker of select agonist-induced Ca²⁺ mobilization, and AKBA but no other BA, potently inhibits collagenevoked aggregation of platelets. Importantly, the concentrations of AKBA needed to block aggregation are in the range of plasma levels of AKBA detected *in vivo* after oral administration of standardized doses of B. *spec* extracts. For patients taking such BS preparations, the amounts of AKBA reached in the blood may be sufficient to antagonise collagen-mediated pathophysiological events, particularly platelet aggregation.

5.4 Celecoxib is an inhibitor of 5-LO

Celecoxib, a well-tolerated antirheumatic drug, reduces familial adenomatous polyp formation⁹². This effect could not merely be explained by COX-2 inhibition. The COX-2-selective drug is, as shown here, also an inhibitor of 5-LO both *in vitro* and *in vivo*. Celecoxib inhibited 5-LO product synthesis in different cellular models using A23187 as agonist, that is, in human whole blood and human isolated PMNL. A direct interaction between 5-LO and celecoxib is visualized in cell-free assays, where celecoxib blocked the activity of isolated human recombinant 5-LO or 5-LO in homogenates from human PMNL. Together, the data strongly suggest that pharmacologically relevant doses of celecoxib affect 5-LO activity in leukocytes and may thus contribute to the unique pharmacology of this drug.

5-LO contains a non-heme iron in the active site coordinated by three His and a C-terminal Ile residue that cycles between the ferrous and the ferric state, being essential for catalysis ¹⁹¹. Most pharmacological 5-LO inhibitors target this iron by retaining the ferrous state and/or by chelation, whereas nonredox-type inhibitors interfere with fatty acid binding cleft(s) ¹⁹¹. Based on the structure of celecoxib, an iron-chelating- or a redox-related mechanism for inhibition can be excluded and a typical pattern of a nonredox-type 5-LO inhibitor is not evident. Besides direct inhibition of 5-LO, interference with FLAP (by e.g. MK-886) blocks cellular 5-LO product formation ¹⁹². Although the cellular environment influences inhibition of 5-LO product synthesis by celecoxib, FLAP is apparently not involved in 5-LO inhibition in intact cells.

A number of studies were conducted to identify non-COX-2 targets of celecoxib responsible for its pharmacological profile. The pharmacological effects include a distinctive chemopreventive and tumour-regressive efficacy at higher celecoxib doses as well as a favourable gastrointestinal tolerability as compared to classical NSAIDs. Various COX-independent mechanisms predominantly in the higher μ M range were considered to explain the unique pharmacology of celecoxib such as inhibition of 3-phosphoinositide-dependent kinase-1 (PDK-1, IC₅₀ = 48 μ M)⁹⁷ or endoplasmatic reticulum Ca²⁺-ATPase (IC₅₀ \approx 35 μ M)⁹⁵ in human prostate cancer cells, degradation of the oncogenic survival factor beta-catenin in human colon carcinoma cells at 60-100 μ M⁹⁸ or inhibition of adenylyl-cyclases (IC₅₀ = 375 μ M)⁹⁶. However, due to the strong discrepancy between the plasma concentrations of celecoxib after a single intake of 800 mg (c_{max} \approx 8 μ M)⁹⁹ and those concentrations required to affect non-COX-2 targets (\geq 30 μ M, *in vitro* assays) the relevance of these findings is discussed controversially. Inhibition of 5-LO product synthesis occurs at comparably low celecoxib concentrations (IC₅₀ $\approx 8 \ \mu$ M, cellular assay) similar to the plasma levels in humans after intake of high celecoxib doses ⁹⁹. The findings are supported by a study of Chen et al. demonstrating that high doses of celecoxib in chow (1000 ppm) reduced the content of LTB₄ by approx. 50 % in oesophageal adenocarcinoma xenografts of rats, whereas lower doses were almost ineffective ¹⁹³. Also, a 50% reduction of LTB₄ and 5-HETE formation in A23187-stimulated human whole blood by 100 μ M celecoxib was described using different experimental conditions as compared to the present study ¹⁹⁴. In contrast, Mao et al. showed that administration of celecoxib 400 mg twice daily to active smokers increased the production of LTB₄ in broncho alveolar lavage fluids (BAL) by around 36 % ¹⁹⁵, indicating that modulation of 5-LO product synthesis by celecoxib may depend on the tissue type and its state of health.

Distinctive inhibition of tumour growth as a pharmacological effect arising from 5-LO inhibition was reported for human prostate, oesophageal, pancreatic, lung and colorectal cancer ¹⁹⁶. Interestingly, all these tumour types were susceptible to celecoxib treatment ¹⁹⁷ providing a possible link between the antiproliferative potency of celecoxib and inhibition of 5-LO. Indeed, the celecoxib-mediated antiproliferative effects are mechanistically similar to those observed during 5-LO inhibitor-triggered cell death. For instance, both 5-LO inhibitors and celecoxib attenuated the growth of human carcinoma cells by releasing cytochrome c from mitochondria, by activating caspase-3 and -9 followed by cleavage of poly(ADP-ribose)polymerase (PARP), and by inducing the cell cycle inhibitor p21^{kip1 94, 198, 199}. Moreover, inhibition of 5-LO was described to implicate downregulation of LTD₄-mediated beta-catenin-signalling ²⁰⁰, and this pathway is suppressed also by celecoxib ⁹⁸. Finally, LTB₄ was reported to activate Akt by PI3-kinase/PDK-1-mediated phosphorylation ²⁰¹, suggesting that inhibition of LTB₄ formation by celecoxib may contribute to the downregulation of cellular Akt activity as described for this drug few years ago ⁹⁷. Taken together, celecoxib may exert its antiproliferative effects via inhibition of 5-LO product synthesis.

Dual inhibitors of COXs and the 5-LO pathway, such as licofelone, constitute a valuable alternative to classical NSAIDs since they cause less severe gastrointestinal (GI) side effects than non selective COX-inhibitors alone ²⁰². Recent studies showed that the gastroprotective effects of licofelone can substantially be attributed to an inhibition of LT-mediated leukocyte adhesion to intestinal blood vessels and subsequent mucosal inflammation ²⁰³. Together, since celecoxib combines COX-2 selectivity with 5-LO-inhibitory activity, it may counteract such ulcerogenic effects by reducing LTB₄ levels which is highly chemotactic for leukocytes and plays an important role in the development of NSAID-induced gastrointestinal lesions ²⁰⁴.

Interestingly, it has been shown that celecoxib has a favourable gastrointestinal tolerability profile superior to that of unselective NSAIDs ²⁰⁵⁻²⁰⁷, perhaps partly due to its 5-LO-inhibitory activity.

In conclusion, this work shows that celecoxib inhibits human 5-LO in cell-free assays, in cellular systems and in human whole blood. These findings may contribute to a better understanding of the unique pharmacological profile of celecoxib among coxibes, characterized by a distinctive antiproliferative efficacy and a favourable gastrointestinal tolerability.

6 Summary

Extracts of Boswellia serrata, also known as Indian frankincense, have been used to treat inflammatory diseases in the Indian ayurvedic medicine or Chinese traditional medicine (TCM) for over 3000 years, but the molecular mechanisms of the anti-inflammatory effects are still not well understood. It is obvious that the boswellic acids, the major compounds in the extracts, are responsible for the efficacy. This work employed a protein fishing technique to identify putative targets of boswellic acids at different stages within the inflammatory cascade. For fishing experiments, boswellic acids were immobilized to sepharose and incubated with cell lysates. After washing and boiling, fished proteins were separated by SDS-PAGE and analysed by MALDI-TOF-MS. CatG, DNA-PK and the protein kinase Akt were identified by protein pulldowns with immobilised BAs and characterised as selective and important targets for BAs with an IC₅₀ in the range of physiologically achievable plasma levels up to 5 μ M. In addition, the influence on several signal transductions by BAs was tested. Calcium influx, arachidonic acid release, platelet aggregation and TNF α -release were assayed to reveal further pharmacological effects of BAs.

Celecoxib is a well-known selective COX-2 inhibitor that is in clinical use. In this work, it is demonstrated that celecoxib is also a highly potent direct 5-LO inhibitor. Celecoxib is used in arthritis and its gastro-intestinal side effects are reduced compared to non-selective NSAIDs. In patients with a familiar disposition to polyp forming, celecoxib reduced polyps and the incidence of colon cancer. Because of lowered leukotriene levels in patients under celecoxib therapy it was plausible to test whether celecoxib interferes with 5-LO. Here it is shown that the activity of 5-LO is inhibited in PMNL and cell-free assays with IC₅₀ of 8 μ M in intact cells, 20 μ M with supplemented arachidonic acid and 30 μ M in cell-free systems. Thus, celecoxib is a dual inhibitor of COX-2 and 5-LO. Since 2006, celecoxib has been approved as an orphan drug for the treatment of familial adenomatous polyposis. Aside from this indication, it could be useful for treatment of asthma and other diseases where 5-LO is implicated.

7 Zusammenfassung

Extrakte des indischen Weihrauchs werden seit mehr als 3000 Jahren in der Volksmedizin zur Behandlung chronisch entzündlicher Erkrankungen sowie in der ayurvedischen Medizin in Indien und in der Traditionellen Chinesischen Medizin (TCM) verwendet. Pharmakologisch wirksame Inhaltsstoffe sind vor allem Boswelliasäuren, die den Hauptbestandteil der Säurefraktion des Harzes ausmachen. Bislang sind die molekularen Mechanismen nur unvollständig aufgeklärt. Diese Arbeit befasst sich mit der Auffindung und Charakterisierung von Zielstrukturen, die innerhalb der Entzündungskaskade von Boswelliasäuren moduliert werden. Zur Identifizierung wurden 11-Keto-Boswelliasäure sowie ß-Boswelliasäure an Sepharose-Beads gebunden und damit immobilisiert. Die immobilisierten Boswelliasäuren wurden zusammen mit Zelllysaten aus verschiedenen Zelltypen (PMNL, Thrombozyten, MM6, LNCaP, MCF-7, HL-60 und RBL-1) inkubiert. Die Beads wurden von dem Überstand getrennt, gewaschen und mittels SDS-PAGE aufgetrennt. Als Vergleichsprobe diente ungebundene Sepharose, die unter gleichen Bedingungen behandelt wurde. Banden, die nur in den Präzipitaten der BA-Sepharosen auftraten, wurden mittels MALDI-TOF-MS analysiert. Bei diesem als "Fischen" bezeichneten Verfahren wurden selektiv folgende Proteine als mögliche Interaktionspartner von BAs gefunden: CatG, Rap1b, DNA-PK, Akt, Proteinase-3, Prohibitin, UNC-112, cathelicidin antimicrobial peptide (hCAP18), farnesylpyrophosphate synthase (FPPs), VAT-1 und ATP-synthase.

Die meisten dieser Proteine sind wichtige Mediatoren in der Zelltransduktion, im Immunsystem oder spielen eine Rolle im Entzündungsgeschehen sowie bei der Apoptose. Der Einfluss der BAs auf PKC und FPPs ist noch unklar, da *ex vivo* und *in vitro* Experimente zwar negativ ausfielen, Untersuchungen in klinischen Studien aber auf eine Beeinflussung schließen lassen¹⁷⁶. Rap1b wird durch BAs in seiner Aktivität gehemmt, hier sind noch weitere Studien zur genaueren Charakterisierung nötig. CatG, DNA-PK und die Proteinkinase Akt konnten als Zielstrukturen von BAs mit hoher Affinität (IC₅₀-Werte < 5 µM) identifiziert werden, wobei die erreichbaren Plasmakonzentrationen mancher BAs pharmakologische Relevanz implizieren. Andere in der Literatur beschriebene Interaktionspartner benötigen für eine Hemmung sehr hohe Konzentrationen an BAs (z.B. 5-LO: 15-30 µM), so dass eine pharmakologische Relevanz *in vivo* bei der 5-LO und der HLE fraglich erscheint. Die proteolytische Aktivität von CatG wird konzentrationsabhängig, kompetitiv und reversibel mit IC₅₀-Werten von 0,6 µM (AKBA), 0,8 µM (β-BA), 1,1µM (Aβ-BA) und 3,7 µM (KBA) gehemmt. Den BAs ähnliche Triterpene wie Amyrin oder Ursolsäure konnten die Aktivität von CatG bis zu einer Konzentration von 30 μM nicht signifikant supprimmieren. Auch die Aktivität anderer Serinproteasen wurde zum Teil sehr potent inhibiert, so konnte für Chymotrypsin ein IC₅₀-Wert von 4,8 μM (Aβ-BA) ermittelt werden, die anderen BAs waren weniger stark wirksam. Insgesamt ist die Effektivität bei CatG aber am höchsten. Nicht nur die proteolytische Aktivität von CatG wird in vitro inhibiert, auch die CatG-induzierte Migration von PMNL durch Matrigel wird mit einem IC₅₀ von 2,9 μM (Aβ-BA) gehemmt. In unserem kleinen pharmakokinetischen Test (Phase I) konnte gezeigt werden, dass bei einem Plasmaspiegel von 6 μg/ml Aβ-BA die proteolytische Aktivität von CatG nur noch ca. 75% des Ausgangswertes beträgt.

DNA-PK und Akt sind Kinasen in der Signaltransduktion von Rezeptor-Tyrosinkinasen, deren Liganden Wachstumsfaktoren sind. Durch zellbasierte und zellfreie Experimente wurde gezeigt, dass BAs die Aktivität von DNA-PK und Akt hemmen und somit Wachstumssignale inhibieren. Die Phosphorylierung von Akt wird intrazellulär und im zellfreien System durch BAs stark reduziert, und die Aktivität von Akt wird extrazellulär durch BAs konzentrationsabhängig im nanomolaren Bereich inhibiert.

Weiterhin wurde der Einfluss BAs auf verschiedene Elemente von der Signaltransduktionskette untersucht. Hierbei stellen vor allem Ca²⁺-Einstrom, Beeinflussung des MAP-Kinaseweges, Arachidonsäurefreisetzung, Thrombozytenaggregation und TNFa-Freisetzung interessante Ansatzpunkte dar. Die gefundenen Ergebnisse beleuchten die pharmakologischen Wirkungen der Weihrauchextrakte ein wenig besser, vor allem die sehr guten Ergebnisse bei chronisch entzündlichen Darmerkrankungen könnten nun aufgrund der sehr potenten Hemmung von Cathepsin G erklärbar sein.

Mittels sepharose-gebundener Boswelliasäuren konnten Zielstrukturen entdeckt werden, die selektiv und hochaffin an die BA-Sepharosen binden. Diese Strukturen sind größtenteils in wichtige Prozesse der zellulären Signalkaskade involviert. Durch die Identifizierung dieser hochaffinen Targets ist es möglich, die zelluläre Wirkung der BAs genauer aufzuklären und die BA-Grundstruktur zur Entwicklung neuer selektiver Arzneistoffe gegen chronisch entzündliche Erkrankungen und Krebs zu nutzen.

Des Weiteren wurde gezeigt, daß Celecoxib, ein gut charakterisierter, selektiver COX-2-Inhibitor, als hoch potenter 5-LO Inhibitor fungiert. Der Wirkstoff wird zur Therapie von Arthrosen und Rheumatoider Arthritis angewendet und kam als erster selektiver COX-2-Hemmer auf dem Markt. Die gastrointestinalen Nebenwirkungen sind gegenüber den unselektiven NSAR deutlich reduziert, zusätzlich konnte durch klinische Studien gezeigt werden, dass die Gefahr für kardiovaskuläre Ereignisse im Vergleich zu den anderen Coxiben – Rofecoxib (Vioxx[®]) wurde deshalb vom Markt genommen – stark erniedrigt ist und auf dem Niveau der klassischen Antirheumatika wie Ibuprofen oder Diclofenac liegt.

Durch Zufall wurde entdeckt, dass neben der bisher bekannten Wirkung Celecoxib bei Patienten mit familiärer Prädisposition die Polypentstehung im Darm stoppen kann. Diese Darmpolypen sind eine Vorstufe von Darmkrebs. Dieser Effekt konnte weder mit klassischen NSAR noch mit den anderen Coxiben erzielt werden, weshalb neben der COX-2 noch weitere Zielstrukturen beeinflusst werden mussten. Bei weiteren Studien konnte mit 800 mg/d Celecoxib bei einigen Patienten eine Remission von Darmtumorvorstufen erreicht werden, unzureichend mit einer COX-Hemmung erklärbar ist. Durch Untersuchungen an Patientenvollblut konnten verminderte Level an Leukotrienen unter Therapie gezeigt werden, die auf eine Beeinflussung des Leukotrienstoffwechsels hindeuten. Als interessante Zielstrukturen wurden FLAP und die 5-LO ausgewählt und bearbeitet. Schon schnell stellte sich heraus, dass FLAP nicht gehemmt wird (Untersuchungen von Dr. Maier und M. Hörnig). Die Aktivität der 5-LO wird sowohl in PMNL als auch in zellfreien Systemen inhibiert und zwar mit folgenden IC₅₀-Werten: 8 μ M in intakten PMNL, 20 μ M nach Zugabe von 20 μ M Arachidonsäure und 30 μ M im zellfreien System.

Seit 2006 ist Celecoxib in Europa als Orphan Drug zur Therapie der adenomatösen familiären Polyposis zugelassen. Weitere Anwendungen bei Erkrankungen mit hoher Beteiligung der 5-LO wie zum Beispiel Asthma bronchiale, entzündliche Erkrankungen oder bestimmte Krebsarten sind sinnvoll. Die duale Hemmung von 5-LO und COX eröffnet neue Behandlungsmöglichkeiten, so dass neben verstärkter antientzündlicher Wirkung die Nebenwirkungsrate gesenkt werden kann. Celecoxib ist zur Zeit der einzige zugelassene Arzneistoff, der über diesen dualen Wirkmechanismus verfügt.

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9 Presentations

9.1 Poster presentation

Tausch L, Poeckel D, Altmann A, Werz O Studies on the anti-inflammatory mode of action of boswellic acids in human platelets DPhG PhD student meeting, Leipzig, 2005

Tausch L, Siemoneit U, Poeckel D, Kather N, Franke L, Schneider G, Holtmeier W, Beckhaus, T, Karas M, Jauch J, Werz O Identification of targets and molecular modes of action of boswellic acids DPhG Weihrauch Symposium 2008

9.2 Oral presentations

Tausch L, Werz O

Protein pulldown with immobilized boswellic acids and characterisation of fished targets. Plenary lecture

Summerschool of Pharmceutical Institute, University of Tuebingen, Arnsburg/Oberjoch, September 2006

Tausch L, Werz O Target identification of boswellic acids Seminar of the Institute of Pharmaceutical Chemistry, Frankfurt, February 2007

Tausch L Short oral presentations at work group seminars of AK Steinhilber

10 Publications

- Poeckel D, Tausch L, Kather N, Jauch J, Werz O. Boswellic acids stimulate arachidonic acid release and 12-lipoxygenase activity in human platelets independent of Ca²⁺ and differentially interact with platelet-type 12lipoxygenase. *Mol Pharmacol. 2006 Sep;70(3):1071-8..*
- Poeckel D, Tausch L, George S, Jauch J, Werz O.
 3-O-acetyl-11-keto-boswellic acid decreases basal intracellular Ca²⁺ levels and inhibits agonist-induced Ca²⁺ mobilization and mitogen-activated protein kinase activation in human monocytic cells. *J Pharmacol Exp Ther. 2006 Jan;316(1):224-32.*
- Michaelis M, Suhan T, Michaelis UR, Beek K, Rothweiler F, Tausch L, Werz O, Eikel D, Zornig M, Nau H, Fleming I, Doerr HW, Cinatl J Jr. Valproic acid induces extracellular signal-regulated kinase 1/2 activation and inhibits apoptosis in endothelial cells. *Cell Death Differ. 2006 Mar;13(3):446-53.*
- Poeckel D, Tausch L, Altmann A, Feisst C, Klinkhardt U, Graff J, Harder S, Werz O. Induction of central signalling pathways and select functional effects in human platelets by beta-boswellic acid. *Br J Pharmacol. 2005 Oct;146(4):514-24.*
- Maier TJ, Tausch L, Hörnig M, Coste O, Schmidt R, Angioni C, Metzner J, Grösch S, Steinhilber D; Werz O, Geisslinger G Celecoxib is a 5-LO inhibitor *Biochem Pharmacol;, accepted2008*
- Poeckel D, Greiner C, Verhoff M, Rau O, Tausch L, Hörnig C, Steinhilber D, Schubert-Zsilavecz M, Werz O Carnosic acid and carnosol potently inhibit human 5-lipoxygenase and suppress proinflammatory responses of stimulated human polymorphonuclear leukocytes *Biochem Pharmacol; accepted2008*
- 7. Tausch L, Kather N, Franke L, Schneider G, Beckhaus T, Karas M, Chen J, Jauch J, Werz O.
 Identification of human cathepsin G as a functional target of boswellic acids from the anti-inflammatory remedy frankincense.
 Nature Chemical Biology, presubmission accepted

- Tausch L, Kather N, Jauch J, Werz O. Molecular interference of boswellic acids with Akt/protein kinase B and DNAdependent protein kinase. *Manuscript*
- Tausch L, Siemoneit U, Jauch J, Werz O.
 3-O-Acetyl-11-keto-β-boswellic acid potently antagonises select agonist-induced Ca²⁺ mobilisation and aggregation of human platelets. Manuscript
- 10. Kather N, Tausch L, Poeckel D, Werz O, Herdtweck E, Jauch J Immobilization of Boswellic acids at EAH Sepharose[®] for "target fishing" *Manuscript*

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