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# Genetic disruption of the kindlin-3/β1-integrin interaction results in defective β1-integrin dependent platelet adhesion and granula secretion

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#### **Abbreviations**

ACD Blood coagulation buffer based on citrate-ions

ADP Adenosindiphosphat

C 807/ G 873 DNA base pair cytosine 807, guanine 873

CD Cluster of diffentiation CRP Collagen related peptide

CVX Convulxin
DAG Diacylglycerol
DCF Dichloroflourescein

DIC Differential interference contrast

DNA Desoxy- ribonucleic acid

dNTP Desoxynucleoside triphosphate

ECM extracellular matrix

EDTA Ethylenediaminetetraacetic acid

FAK Focal adhesion kinase

FcRy Fc Rezeptor- y

FERM Four- point-one, Ezrin, Radixin, Moesin Domain

FITC Flourescein isothiocyanate

GAPDH Glycerinaldehyd 3- phosphat- dehydrogenase

GEF Guanodin nucleotid-exchange-factors

GFOGER Amino acid sequence: Glycin-Phenylalanine-Pyrrolysine-

Glycine- Glutamic acid- Arginine

GPIba Glycoprotein Iba GPIX Glycoprotein IX GPV/ Glycoprotein V GPVI Glycoprotein VI

GTP Guanosin tri phosphate HCL Hydrogen chloride

HIS Histidine

HPLC High pressure liquid chromatography
Hpm Platelets , genotype: β1<sup>Hpm/-</sup> integrin

HRP Horse radish peroxidase HT  $\beta 1^{-/+}$  Heterozygous platelets HT Platelets , genotype:  $\beta 1+/-$  integrin

I- domain Inserted domain

ICAM-1 Inter- cellular adhesion molecule 1

Ig Imunoglobulin

ILK Integrin linked kinase IP3 Inositol- triphosphate

IPTG Isopropyl-beta-D-thiogalactopyranoside

ITAM Immunoreceptor tyrosine based activation motif

KO Knock out platelets β1<sup>-/-</sup> integrin

LAD III/ 3 Leucocyte adhesion deficiency type III / 3

LB medium Lysogeny broth

MAPK Mitogen activated protein kinase MIDAS Metal ion dependent adhesion site

MxCre Mx1- promoter driven Cre- Recombinase Gene

NFκB Nuclear factor κ B

NMD Nonsense mediated RNA decay

NPxY Amino acid sequence: Asparagine- Prolin- X – Tyrosine

NTA Nitrinoltriacetic acid

PBS Phosphate buffered saline PCR Polymerase chain reaction

PD Buffer Pulldown buffer

PDVF Polyvinylidene flouride

PE Phycoerythin

petB16 Expression plasmid for bacterial transfection

PFA Paraformaldehyde
PKC Protein kinase C
PLC2y Phospholipase C 2y

POLY I:C POLY I(losine):C(Cytidine)

PTB Phosphotyrosine- binding domain

Rap 1 Ras- related protein

RIAM Rap1–GTP-interacting adapter molecule

RNA Ribonucleic acid
RPM Rounds per minute
RT Room temperature
SDS Sodium dodecyl sulfate
SEM Standard error mean
SH2 domain Src homology domain 2

SILAC Stable isotope labeling with amino acids in cell culture

SOCE Store-operated Ca2+ entry

T 807 / A 873 DNA base pair Thymine 807, Adenine 873

TBST Tris buffered salins with TWEEN

TEMED Tetramethylethylendiamin

TFA Tri-flour-acid

TIRF Total interference reflection fluorescence

Tris Tris(hydroxymethyl)aminomethane

TT/AA  $\beta 1^{TTAA}$  mutated platelets

TT/AA Platelets carrying a β1<sup>TTAA/-</sup> integrin

U/ml Units/milliliter
U46619 Thromboxane- A
UV Ultra violet

VCAM-1 Vascular cell adhesion molecule 1

vWF von Willebrand factor

WT Platelets genotype β1<sup>+/+</sup> integrin (wild type)

Xg Multiple of g force αIlbβIlla α2bβ3a integrin αvβIll ανβ3 integrin

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#### 1. Introduction

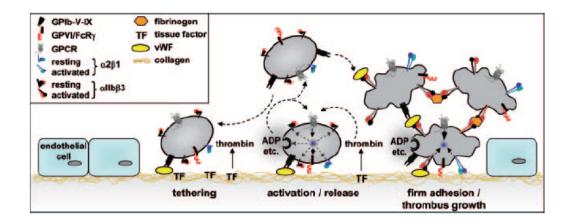
#### 1.1 Platelets in arterial thrombosis

Platelets represent the major cellular components of the hemostatic system. They are constantly sheared off from transendothelial proplatelets, which are protrusions of bone marrow megakaryocytes. This process is referred to as thrombopoesis (1). Once released into the blood, platelets patrol the vascular system in close proximity to the vessel wall in search for vascular injuries. When they get in contact with vessel wall lesions they become rapidly activated, which leads to platelet adhesion, aggregation and finally wound closure.

The molecular mechanisms that initiate primary hemostasis are well understood (2). Subendothelial lesions accompany with the exposure of various matrix components, which are recognized and bound by different platelet receptors. Platelets exhibit the GPIb-GPV-GPIX complex (around 25000 copies/cell) on their surface (3). In vivo and in vitro analyses have shown that under high shear conditions initial platelet adhesion to sites of vascular injuries or an activated endothelium is mediated by the interaction between GPIb( $\alpha$ ) on platelets and vWF, which is either bound to extracellular matrix (ECM) collagen or exposed by the activated endothelium (4). The interaction of GPIb with an activated endothelium was shown to promote leukocyte recruitment and atherosclerotic plaque formation in Apoprotein- E deficient mice (4).

However, during thrombus formation GPIb interactions are only transient and result in the deceleration and rolling of platelets, enabling other surface receptors to bind to the extracellular exposed matrices including collagen. Collagen belongs to the most thrombogenic matrix components of subendothelial lesions and is bound by GPVI and α2β1 integrin surface receptors. GPVI forms a non-covalently bound complex with FcRγ (5). Once the receptor is engaged to collagen it forms a GPVI homodimer which leads to tyrosine phosphorylation of the immunoreceptor tyrosine-based activation motif (ITAM) within the FcRγ by Src family kinases (6). GPVI signaling (9) triggers auto- and paracrine granula secretion and exposure of strong platelet activators such as Thromboxane- A2, ADP as well as a variety of procoagulant molecules including vWF, Fibrinogen, integrin αIIbβIIIA or P-selectine (7). Due to the release of these components, platelets contribute to a

prothrombotic and procoagulant environment at the lesion site (3). On the other hand, GPVI signals trigger the activation of surface receptors, such as  $\alpha IIb\beta 3$  and  $\beta 1$  integrins, which play fundamental roles in mediating platelet-matrix and platelet-platelet interactions (8) (*Supplementary Figure 1*).



Supplementary Figure 1: Platelet adhesion and aggregation on the ECM (from David Varga-Szabo et al. 2008(7)). The GPIb- vWF interaction mediates platelet tethering thereby enabling GPVI interaction with collagen. This triggers the shift of integrins to a high-affinity state and release of ADP and Thromboxane A2. In parallel, tissue factor (TF) locally triggers thrombin formation which also contributes to platelet activation.

#### 1.2 Integrins

Integrins represent the largest family of adhesion receptors which are exclusively expressed in metazoans. Integrins resemble heterodimeric transmembrane molecules consisting of non-covalently bound  $\alpha$  and  $\beta$  subunits (9). To date 18  $\beta$  and 8  $\alpha$  subunits are known, which form 24 different heterodimers. Both integrin subunits have a large extracellular domain consisting of  $\geq$  1600 aminoacids, whereas the intracellular tails (except for  $\beta$ 4 integrin) are relatively short with about 20- 50 aminoacids. The extracellular regions of integrin subunits consist of a stalk like amino acid chain attached via a bend "knee region" to a globular- head-domain, which facilitates a ligand recognition site. Some  $\alpha$ -subunits (i.e.  $\alpha$ 1,  $\alpha$ 2,  $\alpha$ L, and  $\alpha$ X,  $\alpha$ M ) expose a so-called I domain (inserted domain), which harbors a metal ion dependent adhesion site (MIDAS) coordinating the binding of an Mg<sup>2+</sup> ion. The  $\beta$ 1 integrin head domain consists of a  $\beta$ A- domain that also contain

several MIDAS. Therefore ligand binding to integrins such as  $\alpha 2\beta 1$  depends on extracelluar ion concentrations and can be induced by extracellular addition of ions such as Mn<sup>2+</sup>, irrespective of the intracellular activation mode (see below) (10).

#### 1.2.1 Integrin function

Integrins connect the intracellular cytoskeleton with the extracellular matrix. They act as bidirectional signaling machines which transmit mechanical forces via the plasma membrane in both directions. They are involved in various cellular processes such as cell spreading, migration, tumor metastasis, adhesion and cell survival (9, 11). Integrin derived signals maintain cell survival of most cell types (except for hematopoetic derived cells), as depriving integrin-extracellular matrix interactions leads to programmed cell death referred as "Anoikis", a mechanism which is bypassed in tumor cells (12). Most importantly,  $\beta$ 1 integrins are indispensable during embryonic development.  $\beta$ 1 integrin deficient mouse embryos develop normally until the blastocyst is formed. However already at this stage the inner cell mass fails to develop normally and dies due to a retarded growth shortly after the trophoblast had invaded the uterus (13).

The rising field of mechanotransduction addresses mechanisms by which physical stimuli are translated into a biological response on a molecular level. In this context, integrins were found to be major players with particular importance during the development of cardiovascular diseases (14-15). Atherosclerotic lesions develop preferentially at regions of so called turbulent flow within the vasculature. Turbulent flow patterns are found at the inner side of the aortic arch and at the outer side in carotid bifurcations. In these atheroprone regions, endothelial cell integrins are kept in an active state due to a continuously changing directionality of the blood flow and of locally exerted shear forces. Molecular studies found, that permanently activated integrins induce and maintain proinflammatory signaling pathways such as the activation of NFkB, a transcription factor that binds to kB response elements found in promoter regions of many inflammatory receptors such as ICAM-1 or VCAM-1 (16). The expression of these adhesion proteins leads to increased recruitment of inflammatory cells to the vessel wall, a key event in the development of arteriosclerosis. The importance for a differential  $\beta 1$  integrin

signaling in this process was convincingly demonstrated in vitro by Orr et al. (17). Endothelial cells seeded on fibronectin, a substrate for  $\alpha 5\beta 1$  integrin that accumulates at atheroprone regions within the vasculature, show an inflammatory activation state with sustained NF $\kappa$ B activation once they are exposed to shear stress. In contrast, cells seeded on collagen (the main ECM component of healthy vasculature), which anchor via  $\alpha 2\beta 1$  integrins, exhibit a silent, non- inflammatory phenotype under the same experimental settings. The authors suggested that atheroprotective behavior depends on the activation of p38–MAPK (17) downstream of  $\alpha 2\beta 1$  integrins.

So far it remains difficult to delineate signaling events emerging from activated integrins. Integrin outside-in signaling arises as soon as a specific ligand is bound and a multifaceted complex signaling platform, referred to as the adhesome (18) is formed. The adhesome involves more than 150 proteins that arrange in different signaling platforms with a certain local and temporal resolution depending on the surrounding microenvironment and activation state of the integrin (19).

# 1.2.2 Platelet integrins

Platelets express a series of integrins namely  $\alpha IIb\beta3$ ,  $\alpha\nu\beta3$ ,  $\alpha2\beta1$ ,  $\alpha5\beta1$ , and  $\alpha6\beta1$ , which have different ligand specificities. The  $\alpha IIb\beta3$  integrin (20) is the most abundant platelet integrin and binds to various ligands such as fibrinogen,  $\nu$ WF, and fibronectin. It is fundamental for platelet adhesion and aggregation and therefore represents the physiologically and therapeutically most relevant integrin. Clinical treatment with Abciximab, an integrin  $\alpha IIb\beta3$  inhibiting chimeric antibody, protects efficiently from arterial thrombosis, while increasing the risk of severe bleedings (21). Some patients suffering from Glanzmann thrombasthenia show mutations in  $\beta3$  or the  $\alpha II$  integrin subunit that cause either reduced receptor expression or severe functional defects of the integrin. These patients usually reveal a broad spectrum from minor to extensive, life threatening bleedings already at young ages (22). Similar observation can be seen in  $\beta3$  integrin deficient animals, which are protected from thrombus formation (23-24). The role of  $\alpha\nu\beta3$  in mediating binding to vitronectin and osteopontin remains still elusive in vivo, although both ECM proteins have been found in atherosclerotic plaques (25).

### 1.3 Role of platelet \$1 integrins

The biological significance of platelet  $\beta1$  integrins, namely  $\alpha2\beta1$  (collagen receptor),  $\alpha5\beta1$  (fibronectin receptor) and  $\alpha6\beta1$  (laminin receptor), in hemostasis is still controversially discussed. Studies using conditional  $\beta1$  knock-out mice suggested a marginal role during primary hemostasis in vivo (26) and platelet adhesion in a carotid ligation model (27). In contrast, studies on  $\alpha2\beta1$  deficient mice indicated a significant role in thrombus formation and thrombus stabilization in a ferric chlorid model (28). Furthermore these animals exhibit prolonged tail bleeding times (29) and delayed vessel occlusion after photochemical induced injury (30). During the last decades, most studies on  $\beta1$  integrins in platelets focused on collagen receptor  $\alpha2\beta1$ . For  $\alpha5\beta1$   $\alpha6\beta1$  so far only one study showed a role for both receptors on thrombus formation in a carotid artery model. However this was only indirectly by using a  $\alpha2\beta1$  knock-out mouse with an additional pharmacological blockade of  $\alpha11b\beta3$  integrins (27). A dedicated system to prove  $\alpha5\beta1$  or  $\alpha6\beta1$  integrin contribution to arterial thrombosis is still missing, as single receptor knock-out mice are not available.

In vitro flow chamber assays that quantified thrombus formation on fibrillar collagen or collagen containing plaque material showed as well diverse results. Experiments using whole blood form  $\beta1$  knock-out animals showed no differences in thrombus formation compared to wildtype blood (26). Penz et al. used flow chambers covered with homogenized atherosclerotic plaque material and human platelets which were incubated with  $\alpha2\beta1$  integrin blocking or control antibodies and found no effect on aggregate or thrombus formation (31) after  $\alpha2\beta1$  blockage. However enzymatic digest of collagen within the plaque material leads to reduced thrombus formation, to a similar extend as seen after GPVI inhibition by usage of a blocking antibody. In summary this date underlined the importance of GPVI as main collagen signaling receptor to induce thrombus formation.

Another study using blood from  $\alpha 2\beta 1$  or GPVI depleted animals showed an importance of either receptor on thrombus formation in flow chamber assays (29). Further, this group found prolonged tail bleeding times in GPVI or  $\alpha 2\beta 1$  receptor depleted mice. Pugh et al. used flow chambers covered with receptor specific substrate peptides or peptide mixtures, that recognize either GPVI (substrate: CRP),  $\alpha 2\beta 1$  (substrate: GFOGER) or GPIb (substrate: vWF) (32). Additionally, this

group performed all experiments under different shear conditions, as different molecular mechanism are thought to mediate platelet activation and adhesion at different shear rates. This study verified in an elegant way that at low shear rates of 100/s or 300/s both GPVI and  $\alpha2\beta1$  were required for thrombus formation. Whereas at a shear rate of 1000/s, GPIb and either of both collagen receptors was required and at even higher rate of 3000/s GPIb and GPVI were sufficient with only little contribution of  $\alpha2\beta1$ . Taken together, the in vitro data of above mentioned studies are rather conflicting and conclusive explanation for differences between the results are missing.

#### 1.4 Integrin activation and the role of talin and kindlin

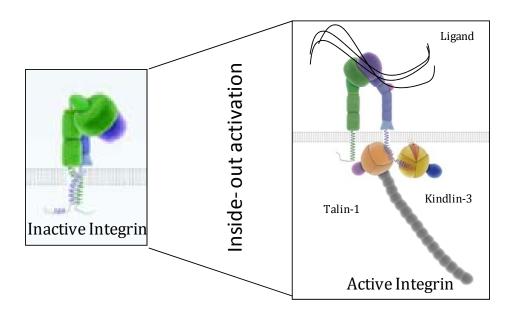
A hallmark of integrins is their ability to shift between an active and inactive conformation, which is controlled by intracellular signaling events. A tight control of integrin activity is a prerequisite for many cellular processes and is particularly important for platelets as defective or pathological activation of platelet integrins results in hemorrhages or thrombosis, respectively. Therefore integrins are found in an inactive conformation on resting platelets, but become rapidly activated when platelets get in contact to damaged vessel walls. It has been recently shown that integrin activation is under the control of two cytoplasmic protein families, talins (33) and the more recently described family of kindlin proteins (Supplementary figure 2).

Kindlins represent a novel family of three FERM domain proteins: kindlin-1 to 3. Historically, a mutation in kindlin-1 was identified as cause of Kindler syndrome, a rare genodermatosis characterized by an epithelial cell adhesion defect and a dermal presentation of poikiloderma and skin athrophy (34). While kindlin-1 and 2 are more widely expressed, kindlin-3 expression is restricted to the hematopoietic system. Recent in-vivo studies showed that absence of either kindlin-3 (35) or talin-1 (36-37) leads to a severe bleeding phenotype in mice due to impaired platelet integrin activation (8).

Talin and kindlin share a structurally conserved FERM domain (4.1-Protein (Four-point-one), Ezrin, Radixin, Moesin), which locates N-terminal and consist of four subdomains (F0-F3) in talin-1. In kindlin-3 it is C-terminal and comprises three subdomains (F1-F3) (33). The physical interaction to integrin cytoplasmic tails is

mediated via the F3 subdomains, which harbor a phosphotyrosine-binding domain (PTB). A structural feature of kindlin proteins is an inserted pleckstrin homology domain within the F2 domain, which mediates binding to phosphoinositides found at the inner side of the plasma membrane. Further, additional interaction sides exist and locate around the F3 subdomain in each protein and might regulate the differential recruitment of talin and kindlin to the integrin tail (see below) (33).

In-vitro pulldown experiments, using recombinant bacterial expressed cytoplasmic integrin tails revealed direct interactions of talin and kindlin with two NPxY motifs within the cytoplasmic domains of  $\beta1$ ,  $\beta2$  and  $\beta3$  integrins (35, 38). While the talin binding site was mapped to the membrane proximal NPxY motif, kindlins interact with the membrane distal NPxY motif. Further mutation analyses showed that the kindlin binding site extents more N-terminally and includes a double threonine motif and a serine/threonine motif within  $\beta1$  and  $\beta3$  integrins (35), respectively. This site is flanked by the two NPxY motifs and is not required for talin binding. It has been shown previously that disruption of the talin-1 binding sites within the  $\beta3$  integrin cytoplasmic domain results in an integrin  $\alpha$ IIb $\beta3$  activation defect and thrombosis resistance in mice (37). The molecular mechanism how kindlin-3 mediates its essential role during integrin activation in cooperation with talin is not known yet (34). Moreover, it was recently suggested that integrin activation can occur in the absence of kindlins and impaired integrin activity in kindlin mutant cells is due to so far unknown cellular functions of kindlin proteins (33, 39).



Supplementary figure 2: Integrin inside-out signalling (activation). Upon platelet activation, recruitment of talin-1 and kindlin-3 to the cytoplasmic tail induce a conformational shift within the integrin molecule which exposes a ligand binding site within the extracellular domain (adapted from Moser et al.(34))

#### 1.5 From platelet activation to integrin activation

As soon as platelets are activated, a machinery of signaling cascades is induced that finalize in integrin activation. Irrespective of the initial stimulus (almost) all signaling cascades merge at the step of phospholipase activation (6). PLC2y facilitates the enzymatic cleavage of plasma membrane phospholipids (Phosphatidyl 4,5 biphosphate), leading to the production of IP3 (Inositol-triphosphate) and DAG (Diacylglycerol). Both molecules act as 2<sup>nd</sup> messengers and trigger increased intracellular Ca<sup>2+</sup> levels through Ca<sup>2+</sup> release from endoplasmatic reticulum and subsequent opening of plasma membrane Ca<sup>2+</sup> channel Orail 1 (store-operated Ca2+ entry (SOCE)). Elevated calcium level and increased concentration of DAG activate CalDAG-GEF1 that functions as Guanosin nucleotide-exchange-factor for Rap 1 (Ras- related protein). Rap-1 is a small G-protein of the Ras family which once GTP loaded by CalDAG-GEF1 recruits the adapter protein RIAM (Rap1–GTP-interacting adapter molecule). Genetically depleted knock-out mice for CalDAG-GEF1 and Rap 1 exhibit severely

compromised platelet aggregation and thrombus formation (40-41). Beside CalDAG-GEF1, some PKC family members feature the ability to translate increased intracellular Ca<sup>2+</sup> and DAG levels into an even more sustained Rap1 activation (42).

Once established, the Rap-1-RIAM complex accompanies talin on its way to the integrin cytomplasmic tail at the plasma membrane. At the integrin tail, talin-1 destabilizes the interaction between the integrin  $\alpha$ - and  $\beta$ -subunit, which will lead to the separation of the subunits. The precise temporal and spatial resolution of talin and kindlin binding to the integrin cytoplasmic domain is still under investigation. Two models in which talin-1 and kindlin-3 bind either in a parallel or sequential manner are hypothesized (34).

At this stage integrins become structurally activated by a mechanism which transmits the cytoplasmic protein binding step (43) into a conformational change throughout the plasma membrane, to open up the extracellular binding site. A structural study on  $\beta 3$  integrins showed that binding of talin disrupts a stabilizing salt-bridge between  $\alpha$ IIb residue R995 and  $\beta 3$  integrin residue D723 (44). The physical separation induces a dissociation of both integrin subunits (45-46), that leads to a rearrangement of the transmembrane domain (33) as wells as switchblade like extension of the extracellular domain (47). A study, using recombinant expressed and artificially linked integrin dimers showed that cleaving an introduced C-terminal clasp between both integrin subunits will lead to separation of both cytoplasmic subunits by 14nm yielding an activated ligand binding form of  $\alpha 5\beta 1$  integrin receptors (48). However, the physical mechanism is still under investigation and might depend on more complex structural arrangements.

# 1.6 Clinical relevance of kindlin-3 in leucocyte adhesion deficiency type III

Recently, it was shown that mutations in the human kindlin-3 gene are responsible for Leukocyte adhesion deficiency type III (LAD-III) (54-55). LAD-III is a rare inherited immunodeficiency disease, which is caused by defective integrins on hematopoetic derived cells. The patients suffer from a severe bleeding phenotype and recurring bacterial infections, as all platelet and leukocyte integrins show

defective integrin inside-out signaling. Interestingly a similar phenotype is found in kindlin-3 knock out animals.(35)(49).

#### 1.7 Clinical relevance of $\alpha 2\beta 1$ integrins in arterial thrombosis

From a historical point of view, it was known that the α2β1 surface expression level in humans (50) vary at least around four fold. Kunicki et al. showed that increased surface expression correlates with increased adhesion to collagen, binding of fibronectin (via α5β1 integrin) and shorter lag times during collagen induced aggregation. Worth mentioning, as soon as the aggregation was initiated, a similar maximal aggregation was reached irrespective of the surface expression level. Further studies revealed that a polymorphism at base pair 807 and 873 within the  $\alpha$ 2 gene correlates with different surface expression levels. Thereby, the linked polymorphism of T 807 / A 873 has higher α2β1 expression levels compared to C 807/ G 873 (51). A first case control study by Moshfegh et al. (52) showed that patients carrying the T 807/ A 873 polymorphisms within the  $\alpha 2$  gene have a higher risk of myocardial infarction (or stroke) (53) compared to controls. In contrast, a meta-analysis including 6414 cases and 7732 control patients from 14 studies (1999-2003) with a myocardial infarction or coronary stenosis as primary endpoint, failed to show a significant correlation between the T807 polymorphism and the incidence of both endpoints (54). A similar results was found in a metaanalysis on stroke patients (55).

Another study focused on patients receiving coronary intervention and stent implantation, which leads to a severe vessel wall damage due to catheter intervention. The impact of T807 polymorphism in this clinical setting was analyzed in a large retrospective study (56). Von Beckenrath et al. observed 1797 patients after coronary artery intervention and stent implantation for the appearance of acute events such as death, myocardial infarction or urgent target vessel revascularization for a follow up period of 30 days. The investigators could not find any significant adverse outcome in patients carrying the T 807 polymorphism irrespective if patients were hetero- or homozygous for this polymorphism.

Today, it is general accepted that increased surface levels of  $\alpha 2\beta 1$  are no risk factor for arterial thrombotic events. However, all clinical studies were neither

designed to proof nor to reject the hypothesis that  $\beta1$  integrins contribute to arterial thrombotic disease. This is due to several limitations. First, to the best of our knowledge there are no patients reported who lack  $\beta1$  integrins in platelets. Second, evidence from clinical studies, in which the  $\alpha2\beta1$  integrin is inhibited, is missing as no  $\alpha2\beta1$  blocking agent is approved for clinical usage.

# 2. Aims of the study

First, we would like to gain more insides into the fundamental understanding of integrin-kindlin interactions. Kindlin-3 is required to activate integrins in platelets. However, so far it remains unclear if integrin inside-out-signaling requires a direct integrin-kindlin-3 interaction or if kindlin-3 acts apart from integrin tails via a so far unknown mechanism. To address this question we generated  $\beta 1^{TTAA/-}$  integrin mice, which carry a mutation within the kindlin-3 binding site. We decided to analyze platelets from  $\beta 1^{TTAA/-}$  integrin animals, as platelet are either switched on or off and have to be regulated within a fraction of a second.

Second, we would like to re-evaluate the role of  $\beta 1$  integrins during thrombus formation in vivo by facilitating newly generated  $\beta 1^{TTAA'}$  and  $\beta^{Hpm/}$  mice. The role of platelet  $\beta 1$  integrins during arterial thrombus formation in mice is still controversially discussed, ascribing  $\beta 1$  integrins a rather minor unimportant role. We would like to discriminate between the adhesive and the signaling functions of  $\beta 1$  integrins. Thereby we focus on collagen induced platelet responses.

#### 3. Materials and Methods

#### 3.1 Animal work

#### 3.1.1 Genetics

All mice were generated in the laboratory of Prof. Reinhard Fässler at the Max-Planck Institute of Biochemistry in Munich. All animal experiments were carried out in accordance to the German Animal protection law. All used strains and knock-out strategies are published elsewhere (see below). As β1 integrin null animals are not viable (13), we decided to use a conditional β1 integrin mouse (57). To allow blood cell specific gene deletion, the conditional β1 integrin mouse was crossed with an MxCre recombinase carrying mouse. Recombinase expression is regulated by the Mx-1 promoter, which is switched on in the presence of interferon and will remove the DNA between 2 introduced recombinase recognition sites (58). In our model, interferon production is triggered by an intraperitoneal application of poly I(Iosine):C(Cytidine) that forms a synthetic RNA double strand. Once applied, poly I:C is recognized by toll-like-receptors and triggers an immune response, usually found after viral infections (59). Hypomorph allele carrying mice were earlier generated in our lab and published before (60). The mutant BTTAA/- mouse was generated by Dr. Hannelore Meyer and Dr. Markus Moser and will be extensively characterized in another ongoing study. Briefly, the coding region of the integrin cytoplasmic tail localizes to the last both exons of the β1 integrin gene, exon 15 and 16. Thereby exon 16 codes the C- terminal 21 amino acids including both NPXY motifs and the double threonine 788/789 motif. A site directed mutagenesis within both threonine coding triplets replaced an adenosine-base pair against a guanine-base pair at the first position, thereby changing the aminoacid sequence from Thr 788/789 to Ala788/789. The precise cloning strategy will be published elsewhere, and is beyond the scope of this work. Interestingly, homozygous β1<sup>TTAA/TTAA</sup> and β1<sup>Hpm/Hpm</sup> animals are not viable as no homozygous animals were born ((60), unpublished). As one functional \( \beta 1 \) allele is sufficient to yield normal and healthy animals (13), we crossed heterozygous β1 TTAA/+ or β1Hpm/+ animals with β1<sup>flox/flox</sup> MxCre animals to yield β1 <sup>TTAA/flox</sup> MxCre or β1 <sup>Hpm/flox</sup> MxCre animals. Therefore, we were able to bypass embryonic lethality and to generate mice with

only one mutated β1 TTAA/- MxCre or β1 Hpm/- MxCre allele after MxCre induction by poly I:C injection.

#### 3.1.2 Genotyping of animals

For genotyping, DNA from the tail of 4 week old mice was isolated. At day one, 1 mm of the tail tip was cut and incubated over night in 500 µl lysis buffer. At day two, the incubation tube was centrifuged at maximal speed for 10 minutes. The supernatant was transferred into an equal volume of isopropanol. The tube was shaken until DNA precipitates were seen. Next the tube was centrifuged again at max. speed for another 10 minutes. The supernatant was carefully removed and the DNA pellet was washed with 2 ml 70% ethanol to remove remaining isopropanol. After a third centrifugation step at maximal speed for 10 minutes, the supernatant was discarded and the pellet was air dried and resuspended in 150 ul TE buffer.

Tail lysis buffer: 50 mM Tris (pH=8.0), 100 mM EDTA, 0.5% SDS, 1:100 Proteinase K (10 mg/ml proteinase K )

TE Buffer: 10 mM Tris HCl pH 8.0, 1 mM EDTA

reverse:

For genotyping the following PCR reaction mix (20 µl final) was prepared: 2 µl 10x PCR buffer (Metabion), 0.6 µl 50 mM MgCl<sub>2</sub> (Metabion), 10 mM dNTP (Sigma-Aldrich), 0.2 µl Primer forward, 0.2 µl Primer reverse, 0.3 µl Taq- polymerase (homemade by Dr. Moser).

## Genotyping PCR primer (Metabion):

MxCre Recombinase forward: 5'-TTC-GGA-TCA-TCA-GCT-ACA-CC-3' 5'-AAC-ATG-CTT-CAT-CGT-CGG-3' MxCre Recombinase reverse: 5'-GGG-TTG-CCC-TTC-CCT-CTA-G-3' β1integrin forward: β1integrin 5'-GTG-AAG-TAG-GTG-AAA-GGT-AAC-3' reverse:  $\beta 1^{TTAA}$ 5'-AGC-TGT-AAA-GGT-TCT-GTG-TGG-3' forward: B1<sup>TTAA</sup>

5'-CCA-AAA-CTA-CCC-TAC-TGT-GAC-3'

β1<sup>Hpm</sup> forward: 5'-TGC TCT CAG TAA TGT TTC ATA AC-3'

β1<sup>Hpm</sup> reverse: 5'-GTC CTA CTG GTC CCG AC-3'

## Following PCR program was used:

Step 1 95°C for 5 min

Step 2 95°C for 30 sec

Step 3 58°C for 30 sec

Step 4 72°C for 5 min

34 cycles of Step 2-4

Step 5 72°C for 5 min

Step 64 °C

To visualize PCR products, 5 µl of 6x loading dye were added to PCR reactions and then loaded on a 2% agarose gel (containing 0.01% ethidium bromide) and separated by applying a current of 100 V. To visualize PCR products, gels were placed under a UV-light source and pictures were taken using an Olympus camera system.

Loading dye 10x (50ml): 30% glycerol, 100 mg of Orange G (Sigma- Aldrich), water

#### 3.1.3 Generation of bone marrow chimeras

Femurs were isolated from adult mice and the bone marrow compartment was flushed out. Next, cells were rinsed through a cell strainer to remove cross-contaminant tissue pieces, counted and placed on ice. Next 4x10<sup>6</sup> cells were injected into the tail vein of two times lethally irradiated (7 Gray each) 8 weeks old recipient C57BL/6 mice. Three weeks after bone marrow transfer, mice were injected twice (day 0 and day 2) with poly I:C (GE Healthcare Life science) to induce MxCre mediated conditional knock-out. After two weeks platelets were isolated from isoflurane narcotized animals by retro-orbital plexus puncture and β1

integrin surface expression was analyzed by flow cytometry to check knock out efficiency.

#### 3.1.4 Carotid ligation model

This data set was generated with the help of Verena Barocke (Deutsches Herzzentrum München), who performed all experiments following the protocol described by Massberg et al. (61). Briefly, carotid arteries of anesthetized mice were dissected and ligated for 5 min by a small suture. To visualize single platelet adhesion, fluorescently DCF (Invitrogen Molecular Probes) labelled platelets (ca. 10% of platelets) from donor mice were infused and amount of adherent cells at the lesion quantified. As carotid ligation experiments were performed on different days for different groups, adherent platelet counts were expressed as percentage of adherent cells at 5 min time point in wildtype mice (which were taken as controls in every set of experiment).

#### 3.1.5 Bleeding time assay

Mice were anesthetized with isoflurane and placed on a 37°C heating pad. All experiments were performed in a steady state narcosis with constant gas flow, monitored by breathing frequency. Next, an 8 mm segment of the tail tip was cut off, so that arterial bleeding was visible. The bleeding tail was immediately placed into 37°C warm PBS. The time until the bleeding vanished was measured. All experiments were stopped latest after 15 min by cauterization to prevent death.

#### 3.2 Protein work

#### 3.2.1 Bacterial transformation and bacterial culture

1<sup>st</sup> day: 50 μl of competent BL 21 bacteria were incubated with 5ng of the designated DNA plasmid for 30 min on ice, to allow plasmid binding towards the cell surface. In a next step, the reaction tubes were transferred for 90 sec on a 42°C shaker. The heat shock facilitates plasmid translocation towards the inside of the cell. After 90 sec the tubes were immediately transferred on ice to seal the

bacterial membrane. 500  $\mu$ l of LB medium were added and bacteria were incubated at 37°C under shaking conditions for at least 60 min and then dispersed on an agar plate containing 50  $\mu$ g/ml of ampicillin (at 37°C).

2<sup>nd</sup> day: bacterial colonies were picked from each agar plate by the help of a pipette tip and transferred into 100 ml LB (50 µg/ml ampicillin) preculture and incubated at 37°C under shaking conditions. 3<sup>rd</sup> Day: 50 ml of preculture were transferred into 1 I LB (with 50 µg/ml ampicillin) medium and grown until the optical density at 600 nm reached 0.6. Next. **IPTG** (isopropyl-beta-Dthiogalactopyranoside) with a final concentration of 1mM was added to induce protein expression. After 5 h bacteria were harvested by centrifugation at 2500x q for 30 min at 4°C.

LB Medium (11): 25 g yeast extract, 50 g trypton, 50 g NaCl

#### 3.2.2 Integrin tail peptide production

The plasmid carrying the designated integrin tail sequence linked to 10 His residues in a Pet-16b cloning construct was kindly provided by David Calderwood. The following pulldown experiments used Nickel loaded agarose beads to bind the histidine tag. A disadvantage of highly charged nickel beads is the remaining unspecific background binding of proteins within the reaction batch that will contaminate the pulldown experiment. To circumvent this drawback we decided to redesign the integrin full length cytoplasmic tails to switch to a biotin (tag) streptavidin (magnetic bead) system. Therefore, we introduced a cysteine residue, three amino acids upstream of the cytoplasmic integrin tail coding region. The plasmid mutation work was done by a two step mutagenesis using a strategene quick mutagenesis kit following manufactures instructions (Primer: Step 1 forward: AGC GGC CAT ATC GAA GGT CGT CAT ATG TGT GGG CCC AAG CTT TTA ATG ATA ATT CAT, Step 1 reverse: ATG AAT TAT CAT TAA AAG CTT GGG CCC ACA CAT ATG ACG ACC TTC GAT ATG GCC GCT, Step 2 forward: ATC GAA GGT CGT CAT ATG CTG GAA GTT TGT GGG CCC AAG CTT TTA ATG ATA ATT CAT, Step 2 reverse: ATG AAT TAT CAT TAA AAG CTT GGG CCC ACA AAC TTC CAG CAT ATG ACG ACC TTC GAT ). Mutated plasmids were transfected into XL-1 bacteria and grown over night. Next day colonies were

picked, transferred into mini cultures and grown over night. The day after, a Quiagen mini-prep set (Quiagen was used to isolate DNA plasmid from bacterial cultures, before sending plasmids to the Max-Planck—core unit for sequencing.

#### 3.2.3 Protein purification

Bacteria pellets were washed in buffer A. As integrin tail peptides accumulate in bacterial inclusion bodies, bacteria were lysed with 8 M Urea containing buffer B on ice and additionally sonicated at maximal power for six times 15 sec each. Next, the lysate was centrifuged for another 30 min at 20000x g. Supernatants were incubated with 2ml of freshly prepared nickel loaded NTA-resin beads for 2 h on a rotator at room temperature. After washing with Buffer B, integrin tails were eluted by addition of 2 ml of elution buffer. In a next purification step proteins were loaded on a reversed phase HPLC (C-8 column, eluted by an acetonitril gradient in 0.08% TFA (tri-flour-acid) as carrier medium and aliquoted. Following a 2 hours vacuum centrifugation at 50 g at 37°C to evaporate the acetonitril, aliquots were lyophilized under high vacuum over night. The peptide powder was resuspended in steril PBS (ph 7.4). The biotinylation of the cysteine residue was performed as recommended by instructors guide (Pierce: EZ-Link NHS-SS-Biotin Kit). The reaction mixture was then purified by a size exclusion column (Vivaspin 3000) centrifugation step (30 min at 400g). The resulting peptide solution was kept under reducing conditions in the presence of β- mercaptoethanol at 4°C, until used.

Buffer A: 150 mM NaCl, 1 mM EDTA, 10 mM Tris, pH 8

Buffer B: 150 mM NaCl, 1 mM EDTA, 10 mM Tris, 8 M urea, pH 8

Elution buffer: 50 mM imidazole in buffer B

## 3.2.4 Protein concentration determination by Bradford assay

Protein concentration was determined, by the so called Bradford method, using Bradford reagent (Sigma-Aldrich). This method is based on the different light absorption of Coomassie blue dye as it binds to protein. Unprotonated Coomassie blue dye binds non-covalently to positively charged amino acid residues (e.g.,

arginin, lysine, histidine) and will increase light absorbtion at  $\lambda$  620nm in contrast to the protonated unbound form.

#### 3.2.5 UV spectroscopy to determine protein concentration

This method is based on the Beer–Lambert-law, which states that the amount of protein is proportional to the light it absorbs. Aromatic side chains containing amino acid such as tryptophan, tyrosine and phenylalanine absorb UV light at a wavelength of 280 nm. Each protein has a specific light extinction coefficient (L/mol x cm) which can be calculated (e.g. by Peptide property Calculator of the Northwestern University, USA). To calculate the concentration the Beer Lambert law:  $A=\epsilon cl$ 

aw. 71 coi

Is adapted as followed: c= A/ɛl

(A absorbance, ε extinction coefficient, I path length of the cuvette (cm))

#### 3.2.6 Pulldown assays

50 μg Biotin-fusion proteins were incubated with fresh platelet lysates in PD buffer for 4 h. Magnetic-Streptavidin labeled Dynalbeads (Invitrogen) were added for 30 min and then pulled out by using a magnet. Beads were washed three times with 0.5 ml PD buffer, boiled in 2x Laemmli buffer and subjected to immunoblotting. Tail loading was visualized by Coomassie staining.

PD buffer: 150 mM NaCl, 50 mM Tris, EDTA 1 mM, Triton-X 100 0.1%, Protease inhibitor tablets (Sigma-Aldrich) and phosphatase inhibitors P1 and P2 (Sigma-Aldrich), pH 7.4

2x Laemmli buffer: 4% SDS, 20% Glycin, 10%  $\beta\text{-Mercaptoethanol}$  , 0.004% bromphenol blue, 0.125 M Tris HCl

#### 3.2.7 Immunoblotting

SDS-polyacrylamide gel electrophoresis allows the separation of denatured proteins corresponding to its molecular weight. For SDS gel casting, electrophoresis and western blotting (of proteins on a PDVF membrane) a BIO-RAD system was used. First, a 10% resolving gel was casted (volume 10 ml: 4.0 ml water, 3.3 ml of 30% acrylamid mix, 2.5 ml 1.5 M Tris (pH 8.8), 0.1 ml 10% SDS, 0.1 ml 10% ammonium persulfate, 0.006 ml TEMED), second a stacking gel (volume 5 ml: 3.4 ml water, 0.86 ml of 30% acrylamid mix, 0.63 ml 1.0 M Tris (pH 6.8) 0.05 ml 10% SDS, 0.05 ml 10% ammonium persulfate, 0.006 ml TEMED) was prepared and a loading comb was inserted. Gels were placed into a SDS running buffer filled chamber. Samples containing a final volume of 1x Laemmli buffer were boiled for 5 minutes at 95°C, loaded into comb pockets and separated by a current of 40 mA. Next, the gel is placed on top of an isopropanol rinsed PDVF (polyvinylidene-difluoride, BIO-RAD) membrane, which is covered by 3 layers of whatman paper at the back and front side of the gel-membrane "sandwich". The "sandwich" is placed into a blotting chamber filled with blotting buffer and proteins were transferred onto the membrane by applying a current of 240 mA. Subsequently, membranes were blocked with 2.5 % milk powder dissolved in TBST buffer to saturate unspecific binding sites before primary antibodies were added over night for incubation. At the next day, the membrane was washed 3 times for 15 minutes with TBST, and corresponding secondary HRP (horse radish peroxidase) coupled antibodies were incubated for 1h at room temperature. After a second TBST wash (three times for 15 min) the membrane was incubated with a developing solution (Amersham: ECL- kit). The horse radish peroxidase generated chemi-luminescence signal was detected with a high sensitive photo film (Amersham Hyperfilm) and then developed in an appropriate film developing machine (Kodak).

Running buffer (11): Tris-base 3.03 g, glycine 14.4 g, SDS 1.0 g in H2O

Blotting buffer (11): glycine 28.8 g, Tris base 6.04 g, methanol 200 ml, ddH<sub>2</sub>O 1.6 L

TBST buffer (11):Tris 1.21 g, NaCl 8.76 g, Tween-20 0.5 ml

Following antibodies were used:

		Catalog	WB
Species	Vendor	number	concentration
rabbit	Biosource	#44-624G	1:1000
rabbit	Upstate	06-543, 21819	1:1000
goat	Santa Cruz	sc-7534	1:1000
rat	Chemicon	MAB1997	1:1000
rabbit	Moser et al.		1:5000
mouse	Calbiochem	CB1001	1:30000
mouse	Jackson	205-032-176	1:10000
goat	Jackson	115-035-174	1:10000
mouse	Jackson	211-032-171	1:10000
donkey	Jackson	712/035/150	1:10000
	rabbit rabbit goat rat rabbit mouse mouse goat mouse	rabbit Biosource rabbit Upstate goat Santa Cruz rat Chemicon rabbit Moser et al. mouse Calbiochem mouse Jackson goat Jackson mouse Jackson	Species Vendor number  rabbit Biosource #44-624G  rabbit Upstate 06-543, 21819  goat Santa Cruz sc-7534  rat Chemicon MAB1997  rabbit Moser et al.  mouse Calbiochem CB1001  mouse Jackson 205-032-176  goat Jackson 115-035-174  mouse Jackson 211-032-171

## 3.2.7.1 Stripping of western blot

To re-probe blotting membranes with another primary antibody, membranes were incubated with stripping buffer for 30 min at 55 °C. After washing twice with TBST for 15 min, membranes were re-blocked with milk-TBST solution and processed as described above.

Stripping solution: 2% SDS, 62.5 mM Tris pH 6.8, freshly added  $\beta$ -Mercaptoethanol (360 $\mu$ l in 50 ml)

#### 3.2.7.2 Coomassie staining

SDS gels were briefly rinsed in water before transferred into Coomassie staining solution. Gels were stained for 1 hour. Protein bands were visualized after placing the gel into destaining solution.

Coomassie staining solution: 50% methanol, 10% acetic acid, 40% water and 0.25% Coomassie blue R250.

Destaining solution: 10% acetic acid in water.

#### 3.2.8 Integrin phosphorylation analysis

Washed platelets were incubated in 100  $\mu$ l Tyrodes buffer (without phosphate) pH 7.4 for 30 min at 37 °C in the presence of 50 nM Calynculin A. Next, platelets were stimulated with 0.1 U/ml Thrombin. After 60 sec, an equal volume of 2x Laemmli buffer was added and the sample was immediately boiled for 5 min at 95°C. Boiled samples were subjected to SDS gel electrophoresis (8% gel) and Coomassie stained. The band corresponding to the molecular weight of  $\beta$ 1 integrin was cut out and sent to non-quantitative phospho-massspectrometry analysis at Max-Planck Institute of Biochemistry core unit in Martinsried.

#### 3.3 Platelet work

#### 3.3.1 Platelet isolation

Mice were anesthetized by isofluran inhalation and blood was drained from the retro-orbital plexus into 200 µl 20 U/ml Heparin- TBS, by using Heparin (Sigma-Aldrich) rinsed capillaries. Blood was centrifuged twice at 75x g and 100x g respectively, in a tabletop centrifuge (Eppendorf) for 10 min in soft acceleration/deceleration mode. Erythrocyte depleted serum was transferred into a new tube and then spun down by centrifuging for 5 min at 1500x g. After two washing steps with Tyrodes buffer ph 6.5, the platelet pellet was suspended in Tyrodes buffer pH 6.5 and the platelet concentration was determined using a Hemovet blood counter (Hemovet, Germany).

Tyrode's buffer (without phosphate): 137 mM NaCl, 2.8 mM KCl, 12 mM NaHCO<sub>3</sub>, 5.5 mM glucose and 10 mM Hepes, pH6.5 or 7.4,  $\pm$  1 mmMgCl<sub>2</sub>,  $\pm$ 1 mM CaCl<sub>2</sub>,  $\pm$ 1 mg/ml BSA

#### 3.3.2 Flow cytometry

Platelets were isolated and incubated with the desired fluorophore-conjugated antibodies for 15 min at room temperature and then directly analyzed on a FACScalibur flow cytometer (Becton Dickinson). For activation assays, platelets were stimulated for 15 min with the indicated agonists (Thrombin (Sigma- Aldrich), Convulxin (Alexis), ADP (Alexis), U46619 (Alexis)) and immediately analyzed.

#### Following antibodies were used:

Protein	Dye	Source	Clone	Vendor	Conc.
Integrin β1 (CD29)	PE	Hamster IgG	HMbeta1-1	BioLegend	1:200
Integrin β3 (CD61)	PE	Hamster IgG	2C9.G3	eBioscience	1:200
Integrin α2 chain					1:200
(GPIa)	FITC	Rat IgG2b	Sam.G4	Emfret	
GPVI	FITC	Rat IgG2a	JAQ1	Emfret	1:200
GPIX (CD42a)	FITC	Rat IgG1	Xia.B4	Emfret	1:200
CD62P (P-selectin)	FITC	Rat IgG1	Wug.E9	Emfret	1:200
Active integrin αIIbβ3	PE	Rat IgG2b	JON/A		1:200
IgG2a rat isotype ctrl	FITC	Rat IgG2a		NatuTec	1:200
IgG2b rat isotype					1:200
control	FITC	Rat IgG2b	TBE 15	EuroBiosciences	
IgG1 rat isotype					1:200
control	PE	Rat IgG1	R3-34	PharMingen	
IgG hamster isotype					1:200
con.	PE	Hamster IgG	HTK888	BioLegend	

#### 3.3.3 Platelet spreading

Glass slides were coated with 1 mg/ml fibrinogen (Sigma) or 25  $\mu$ g/ml acid soluble collagen I (Sigma- Aldrich, Collagen I from rat tail) overnight at 4°C. At day two, slides were rinsed with PBS and then blocked with 1% BSA (bovine serum albumin) in PBS for 1 h. 1.5 Mio platelets were suspended in 150  $\mu$ l Tyrodes buffer in the presence or absence of 0.75 mM Mn<sup>2+</sup>. Cells were stimulated with 0.1 U/ml, Thrombin (Sigma- Aldrich), immediately plated and the glass slides were incubated at 37°C. After 45 min unbound cells were washed off and DIC (differential interference contrast) microscopy was performed, using a 100x oil immersion lens on a Zeiss DIC microscope equipped with Metamorph- software. Acquired pictures were analyzed for platelet spreading size using ImageJ software and manual selection of spreading area.

PBS (1I) ph 7.4: NaCl<sub>2</sub> 8 g, KCl 0.2 g, Na<sub>2</sub>HPO4 1.44 g, KH<sub>2</sub>PO4 0.24 g, fill up with ddH<sub>2</sub>O (double distilled H2O)

#### 3.3.4 β1 integrin staining

WT cells were stimulated with 0.1 U/ml Thrombin and allowed to spread for 30 min on acid soluble collagen I coated glass slides. Next, cells were fixed for 15 min in 4% PFA in TBS. Then cells were permeabelized with 0.05% Triton X-100 TBS for 15 min. To block unspecific background binding, cells were incubated with 1% BSA in TBS for 1h. Primary antibody, anti-β1 integrin (Chemicon, Mab 1997, 1:500) in 1% BSA containing TBS, was incubated for 1h at RT (room temperature) and secondary, goat- anti- rat, C3 labeled antibody (Jackson) was incubated for another 30 min at RT in the dark. To visualize specific binding, isotype control staining (FITC Rat isotype control, eBioscience, 1:500) was performed in parallel. Pictures were acquired using as ZEISS TIRF microscope in DIC, epifluorescence or TIRF mode, equipped with MetaMorph software (Molecular devices).

#### 3.3.5 Aggregation assay

To determine platelet aggregation, light transmission was measured by adding 70  $\mu$ l of washed platelets (300000/ $\mu$ l) in 140  $\mu$ l Tyrodes buffer ph 7.4 in the presence or absence of fibrinogen (final concentration of 100  $\mu$ g/ml ) and Mn²+ (final concentration of 0.75 mM), respectively. Platelets were stimulated with 1ng/ml convulxin (Alexis) or 5  $\mu$ g/ml fibrillar collagen (Nycomed). Transmission was recorded on an Aggrolink (Aggrolink Germany, Probe and go) 2 channel aggregometer over 10 minutes and expressed as arbitrary units. Thereby, 100% transmission (unstimulated cells) corresponds to 0% aggregation. A decrease in light transmission, due to platelet shape change, was expressed as negative aggregation values (-x %).

#### 3.3.6 Fibrinogen secretion assay

Experiments were performed as described for aggregation experiments (see above) in the absence of fibrinogen and BSA. After 8 min of stirring, the reaction batch was spun down by centrifugation (14000x rpm table top centrifuge) for 30 sec and boiled in 1x Laemmli buffer. Lysates were subjected to immunoblotting for fibrinogen  $\gamma$ - chain. GAPDH was used as internal loading control to show equal loading. Decreases in cellular fibrinogen- $\gamma$  content were quantified by densitometry (Image J software).

#### 3.3.7 Evaluation of ATP release from dense granula

A luciferin-luciferase detection system (Promega) was used to quantify ATP release to monitor dense granule secretion. For these experiments, platelets were stimulated with 5  $\mu$ g/ml fibrillar collagen or 0.1 ng/ml convulxin in a lumino-aggregometer (Aggrolink, Germany) under stirring conditions in the presence of luciferase and its substrate luciferin. Emitted luminescence was recorded and maximal values were expressed as ratio of luminescence values from fibrillar collagen stimulate versus CVX stimulated platelets.

#### 3.4 Statistical analysis

Significance level are calculated by a two-tailed, unlinked student's T-test, and are indicated in the figures. Error bars represent standard error mean values.

#### 4. Results

#### 4.1 Integrin tail-peptide production and pulldown experiments

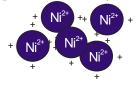
The kindlin-3 binding site at the β1 integrin cytoplasmic tail spans the region from the membrane distal NPXY motif to the more proximal located residues including a double threonine motif. For our earlier pulldown experiments we used full length integrin tail peptides (35, 38), labeled with a hexa- histidine tag, which were pulled down by nickel loaded agarose beads. A drawback of nickel beads is a high unspecific background binding which require harsh washing conditions. We wanted to decrease background binding by using streptavidin-coupled magnetic beads for the pulldown. We redesigned genetically our tails (Figure 1) so that a biotin molecule could be coupled via a disulfide bound to an introduced cysteine residue. To test the biological activity of freshly produced tails, pulldown experiments were performed, using Streptavidin coupled magnetic beads. We found significantly less kindlin-3 binding to β1<sup>TTAA</sup> integrin than to wild type tails without affecting talin binding. Coomassie staining of the low molecular weight fraction of the SDS gel showed equal amounts of tails per reaction. Furthermore mass spectrometry of marked (asterisk) bands verified a successful integrin peptide mutation and expression (Figure 2).

<u>β1 Integrin (wildtype) cytoplasmic tail (amino acid sequence)</u>

#### W- KLLMIHDRREFAKFEKEKMNAKWDTGENPIYKSAVTTVVNPKYEGK

Integrin cytoplasmic tail expressed in Pet-16b - HIS- Tag vector

MGHHHHHHHHHHSSGHIEGRHML EVLFQGP KLMIHDRREE......EGK



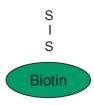
pulldown via Ni<sup>2+</sup>charged aggarose beads

# Redesinged Integrin cytoplasmic tail expressed in Pet-16b HIS Tag vector

Cysteine residue is introduced at position 23

5 Residues were deleted at Pos 24-28

MGHHHHHHHHHSSGHIEGRHM GP KLMIHDRREE.....EGK



pulldown via magnetic streptavidin beads

Figure 1: Amino-acid sequence of β1 integrin tail and bacterial expressed peptide sequence. Bacterial expression vector (Pet-16b) residues are in green, integrin coding sequence is shown in black, the deleted residues are in orange and newly introduced cysteine residue in red. Pulldown beads interaction sides are indicated.

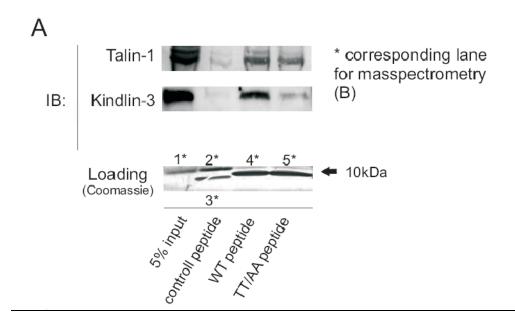


Figure 2: Pulldown experiment. Kindlin-3 binding to  $\beta 1^{TTAA}$  mutated cytoplasmic integrin tail is reduced. Recombinantly expressed cytoplasmic  $\beta 1$  integrin tails (WT vs. TT/AA) were incubated with WT platelet lysates. Pulldown eluates were separated on SDS gel and immuno-blotted for kindlin-3 and talin-1. Results are representative for 3 independent experiments.

# 4.2 Experimental design and generation of $\beta 1^{TTAA/fl}$ MxCre and $\beta 1^{Hpm/fl}$ MxCre mice

To analyze  $\beta1$  integrins lacking a functional kindlin binding site in vivo, we used mouse mutants in which the double threonine motif (TT788/789) within the  $\beta1$  cytoplasmic domain was replaced by two alanine residues (unpublished H. Meyer et al.). Mouse mutants that carry this mutation on both alleles ( $\beta1^{TTAA/TTAA}$ ) are not viable and their phenotype will be described elsewhere. Heterozygous mice ( $\beta1^{+/TTAA}$ ) are normal and were mated with conditional  $\beta1$  integrin mice ( $\beta1^{fl/fl}$ ), which additionally express an inducible cre-recombinase under the Mx1 promoter to obtain  $\beta1^{TTAA/fl}$  Mx1-Cre mice. Recombination and inactivation of floxed  $\beta1$  integrin yielded hematopoietic cells, with only the mutant  $\beta1^{TTAA}$  integrin expressed.

Furthermore, we included a  $\beta1$  integrin hypomorph mouse which expresses only minor amounts of (active) beta-1 integrins. As homozygous animals are also not viable, heterozygous ( $\beta1^{+/hpm}$ ) mice were bred and treated in parallel to mutant animals. Our experimental set up with all used mice strains is shown in Figure 3.

Λ	Mouse					
А		WT	HT	КО	TT/AA	Hpm
	$\beta_1$ Genotype	+/+	+/-	-/-	TT/AA//TT/AA	Hpm/Hpm
	Viability	+	+	*	- **	_***
	Generated bone marrow chimera C57/BI6 (12 weeks old donors)					
		WT	HT	КО	TT/AA	Hpm
	MxCre	-	+	+	+	+
	$\beta_1$ Genotype	+/+	+/fl	f /fI	TT/AA//fl	Hpm/fl

Figure 3: Experimental design. Table-A gives an overview of the subgroups used in this study. Used mice are listed by  $\beta1$  genotype (+ WT allele, - deleted  $\beta1$ integrin allele, TT/AA-mutated allele or Hpm-hypomorph allele) and viability (+ born alive, - not born; \* see text for further explanation). To bypass embryonic lethality, mice were bred into  $\beta1$  conditional background carrying a blood cell specific inducible MxCre system and utilized to generate bone marrow chimera.

#### 4.3 Platelet characterization

First, we isolated platelets from bone marrow chimera and asked whether the introduced ( $\beta1^{TTAA}$ ) mutation affects  $\beta1$  integrin surface expression. Indeed, upon cre-mediated deletion of the wild-type  $\beta1$  integrin allele only 18% of mutant  $\beta1^{TTAA}$  surface integrin was detected by flow cytometry in comparison to  $\beta1$  integrin levels on wild-type platelets (Figure 4A). Due to this reduction our experimental set up included inducible heterozygous ( $\beta1^{+/fl}$  MxCre) and homozygous ( $\beta1^{fl/fl}$  MxCre) conditional  $\beta1$  mice (62) (Figure 4A) as well as newly generated inducible  $\beta1$  hypomorph mice ( $\beta1^{Hpm/fl}$  MxCre) (unpublished, Figure 3) as internal controls. The latter expresses very low  $\beta1$  integrin level as a consequence of nonsense mediated RNA decay (NMD) (60). As expected, poly I:C induced Cre expression resulted in a complete absence and approximately 51% of wild-type  $\beta1$  integrin levels on platelets of  $\beta1^{fl/fl}$  MxCre and  $\beta1^{+/fl}$  MxCre mice, respectively. Platelets from  $\beta1^{Hpm/-}$  MxCre mice showed only 3,2%  $\beta1$  integrin expression level after Cremediated deletion of the floxed  $\beta1$  allele compared to wild-type platelets (Figure 4A).

Since the formation of functional heterodimeric integrin molecules occurs already in the endoplasmatic reticulum and only correctly formed heterodimers are transported to the plasma membrane, diminished expression of the  $\beta1$  integrin subunit caused a similar reduction in  $\alpha2$  integrin surface expression levels (Figure 4B) as well as  $\alpha5$  and  $\alpha6$  subunit expression (data not shown). In contrast, surface expression levels of  $\beta3$  integrin, GPVI and GPIX were similar between groups (Figure 4 D).

Western blot analysis from total platelet lysates also confirmed strongly reduced  $\beta 1^{TTAA}$  integrin expression levels indicating that the kindlin-binding site mutant  $\beta 1$  integrin has either reduced stability or expression. Expression of talin-1 and kindlin-3 was not changed in platelets of all genotypes (Figure 4C).

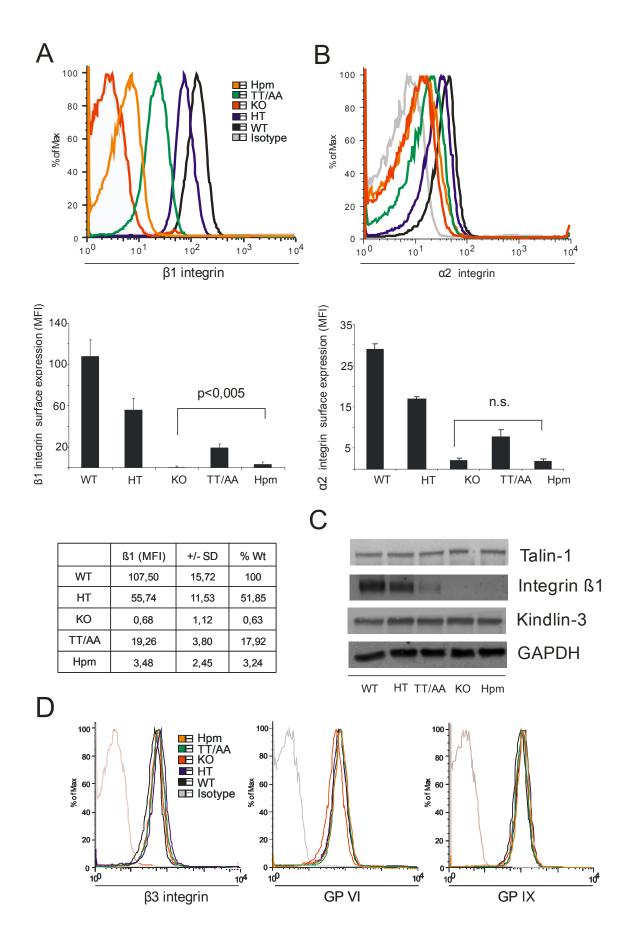


Figure 4: Characterization of platelets. Purified platelets (from  $n \ge 6$  chimeric animals per group) were analyzed by flow cytometry for  $\beta 1$  (A) and  $\alpha 2$  integrin (B) cell surface expression. (C) To determine total protein levels, platelets were lysed and immunoblotted for  $\beta 1$  integrin, talin-1, kindlin-3 and GAPDH as loading control. (D) No differences between groups were found for  $\beta 3$ , GPVI and GPIX cell surface expression.

# 4.4 Integrin activation assays

Next we wanted to test whether kindlin-binding deficient  $\beta 1^{TTAA}$  integrins can shift towards an active conformation. To this end, platelets from chimeric animals of all different genotypes were isolated and activated by different agonists in vitro. Active  $\beta 1$  integrins were measured by flow cytometry using the conformation specific 9EG7 antibody (63-64) that only binds to active  $\beta 1$  integrins. The acquired 9EG7 values were then corrected according to the  $\beta 1$  integrin surface expression levels between the groups. As shown in Figure 5A, treatment with 1ng/ml convulxin, 5 mmol thromboxane A analog U46619 or 0.1 U/ml thrombin, which all stimulate platelets via different receptor signaling pathways (6), fail to activate mutant  $\beta 1^{TTAA}$  integrins. In contrast  $\beta 1$  integrins on platelets from hypomorph and heterozygous mice became activated comparable to  $\beta 1$  integrins in wild-type platelets. In sharp contrast, activation of the  $\alpha 11b\beta 3$  integrin, as measured by conformation specific JON/A (active  $\alpha 11b\beta 3a$  integrin) antibody, was not affected in these cells (Figure 5B). These data indicate that a direct interaction between kindlin-3 and  $\beta 1$  integrin is required for  $\beta 1$  integrin inside-out activation in platelets.

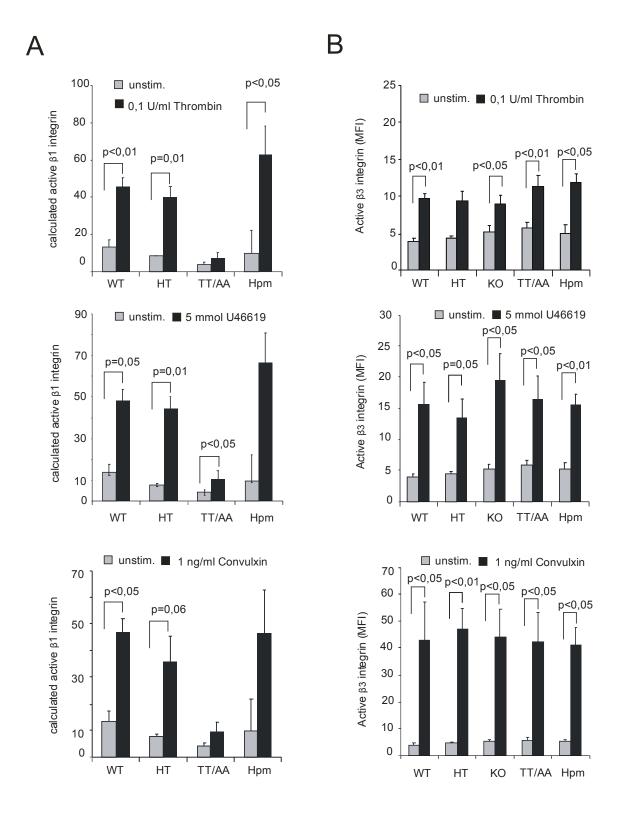


Figure 5:  $\beta$ 1 integrin activation is reduced in  $\beta$ 1<sup>TTAA/-</sup> platelets.  $\beta$ 3 integrin activation is unaffected. To determine  $\beta$ 1 integrin activation (A), platelets were stimulated with indicated agonists and 9EG7- FITC (conformational active  $\beta$ 1) binding was measured by flow cytometry. To adjust for different  $\beta$ 1 surface levels, acquired values were normalized

to 100% (WT)  $\beta$ 1 surface expression (see Figure 4). (B) Shown are MFI (mean fluorescence intensity) values for JON/A binding (conformational active  $\alpha$ IIb $\beta$ 3 integrin) after agonist stimulation. Bars represent mean values of independent experiments (n $\ge$ 3), error bars represent  $\pm$ SEM, significance levels are indicated. (Graphs are depicted at the following page)

#### 4.5 Platelet spreading

Being able to show that integrin inside-out activation is defect, we ask whether a direct integrin kindlin interaction is required for outside-in signaling. Integrin signaling triggers intracellular signals that re-organize the actin cytoskeleton during cell spreading (65). To further characterize the functional properties of \$1^TTAA integrins, platelet adhesion and spreading on soluble collagen or fibrinogen coated surfaces were analyzed. So far little is known about β1 integrin distribution in spreaded platelets on acid soluble collagen. In contrast, platelets spreading on fibrinogen form a typical pan-cake like shape thereby arranging an adhesion ring established of activated β3 integrins around the outer circumference (66). To check for β1 distribution we stained β1 integrins in WT platelets spread on collagen. TIRF microscopy (Total internal reflection microscopy) reveals indeed a β1 containing adhesive ring-like structure, although cells were unable to transform into a pancake-like structure most likely due to an overall decreased number of adhesions being formed. TIRF microscopy penetration depth is limited to several hundred nanometers (up to 400nm) and it might therefore fail to visualize additional intracellular pools of \$1 integrins. To exclude an additional integrin pool, epifluorescence microscopy was performed and showed a similar distribution pattern (Figure 6).

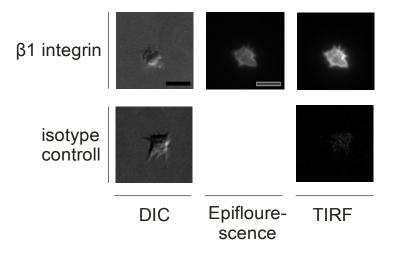


Figure 6:  $\beta$ 1-integrins form an adhesive ring at the outer circumference of spread platelets. Thrombin stimulated wild-type platelets were allowed to spread for 30 min on soluble collagen and afterwards fixed and stained for  $\beta$ 1 integrin. Pictures were acquired in DIC (Differential interference contrast), epifluorescence and TIRF microscopy mode, respectively. Images are representative for 3 independent experiments. Scale bar relates to 5  $\mu$ m.

Adhesion of washed platelets was performed in the presence of 0.1U/ml thrombin and with or without  $Mn^{2+}$ . Acidic soluble collagen I is bound by  $\alpha 2\beta 1$  integrins via a repeatedly found GFOGER motif (67). This motif is not recognized by GPVI, which requires a more complex binding motif present in fibrillar collagen (68). Hypomorphic and  $\beta 1$  deficient platelets showed a severe spreading defect on soluble collagen I, both in the presence or absence of  $Mn^{2+}$  (Figure 7A). These cells formed elongated filopodia and only occasionally small lamellipodia. Consistent with the defect in inside-out activation,  $\beta 1^{TTAA/-}$  platelets also failed to form lamellipodia and showed reduced spreading (Figure 7A). Notably,  $Mn^{2+}$  treatment rescued the spreading defect of  $\beta 1^{TTAA/-}$  platelets significantly. As control all cell types were seeded on fibrinogen a  $\alpha IIb\beta 3A$  integrin ligand and could spread and adhere normal. To further validate this hypothesis, we tested FAK phosphorylation as readout for integrin outside-in signaling (69). Tyrosine phosphorylation within the FAK molecule (e.g., Tyrosine 397) yield several docking sites for SH2 domain containing signaling molecules such as Src family kinases

thereby promoting cell spreading and adhesion (70-71). This time platelets were stimulated with fibrillar collagen to induce integrin inside out signaling and then seeded on soluble collagen in the presence or absence of  $Mn^{2+}$  for 30 minutes. Next, FAK phosphorylation was determined by Western blot analysis. Again, whereas  $\beta 1^{TTAAV-}$  mutant platelets revealed low pFAK levels similar to  $\beta 1^{-/-}$  knockout platelets,  $Mn^{2+}$  treatment resulted in an increase in pFAK levels in  $\beta 1^{TTAAV-}$  platelets but not in  $\beta 1^{-/-}$  knockout platelets (Figure 8).

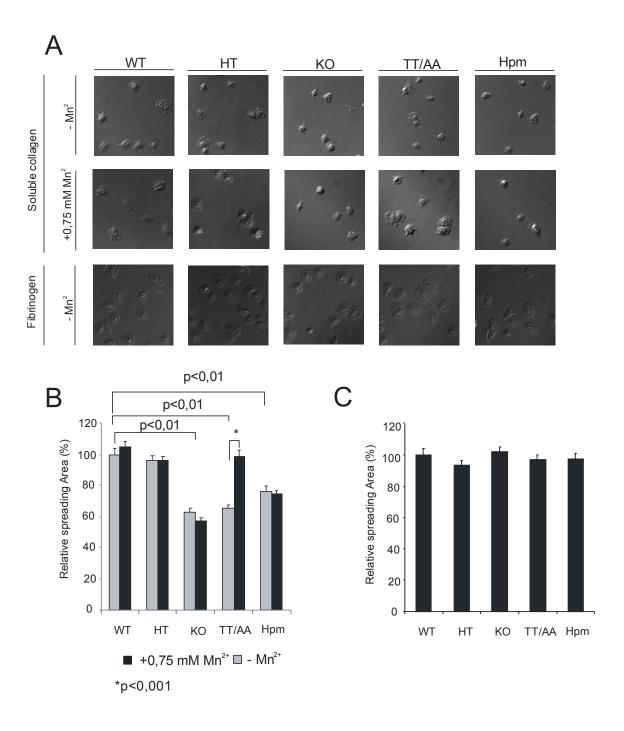


Figure 7: Spreading of thrombin stimulated platelets. Platelets ( $15x10^6$  cells/ $150 \mu$ I) were stimulated with 0.1 U/ml Thrombin and seeded on (A) acid soluble collagen I ( $25 \mu$ g/ml) or fibrinogen (1 mg/ml, no Mn<sup>2+</sup>), in the absence or the presence of 0,75mmol Mn<sup>2+</sup>. After 45 min, cells in suspension were carefully washed away, DIC pictures (100.6x) acquired and spreading size measured using Image J software. Cell size of at least 100 cells was measured and quantified as % of WT cell size of spread platelets on soluble collagen (B) or on fibrinogen (C), respectively. Pictures are representative for at least 3 independent

sets of experiments. Frame size correlates to 35x35 µm. Error bars represent SEM. Significance levels are indicated.

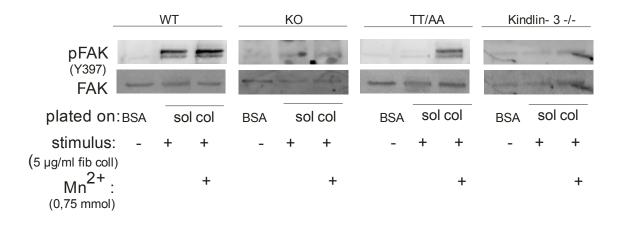


Figure 8:  $\beta 1^{TTAA/-}$  platelets show a defective outside-in signaling and FAK phosphorylation which can be rescued by addition of manganese. Notably, the presence of kindlin-3 is required for FAK phosphorylation. Platelets (15x10<sup>6</sup> cells/150  $\mu$ I) were stimulated with 5  $\mu$ g/ml fibrillar collagen and seeded on 25  $\mu$ g/ml acid soluble collagen I (sol col) in the absence or the presence of 0.75 mmol Mn<sup>2+</sup>. After 30min cells were immediately lysed in 1x laemmli buffer and subjected to immunoblotting for pFAK<sup>Y397</sup>.

#### 4.6 Tailbleeding assay

Being able to show a functional defect of  $\beta 1^{TTAA/-}$  platelets ex vivo we wanted to address the in vivo relevance of our observations. To exclude side effects due to Mx1-Cre mediated  $\beta 1$  integrin deletion in other organs (58, 72), all in vivo experiments were performed with bone marrow chimeras (Figure 3). Reduced  $\beta 1$  integrin expression levels or the expression of mutant  $\beta 1^{TTAA}$  integrins had no impact on thrombopoiesis as platelet counts were similar between all groups (Figure 9B). A former study suggested that deletion of  $\beta 1$  integrins in blood cells did not result in an obvious bleeding defect (26). However, recently designed small molecule inhibitors, targeting  $\alpha 2\beta 1$  integrin (73) conferred thrombosis protection in mice and supported previous findings by Sarrat et al. (29), which found prolonged tail bleeding times in  $\alpha 2$  integrin deficient mice. By using a more sensitive and easier controllable tail bleeding assay (discussed below), we measured a significantly prolonged tail bleeding time in  $\beta 1^{TTAA}$  mice, which was comparable to

 $\beta$ 1 deficient mice. Bleeding usually stops approximately after 8 to 9 min in both groups. Wild-type and heterozygous animals showed bleeding times between 2 to 3 min. Consistent with previous experiments, kindlin-3 deficient mice showed a severe bleeding phenotype due to an additional defect in  $\alpha$ IIb $\beta$ 3 integrin activation (35). Notably, hypomorphic mice expressing only ~3% of the normal  $\beta$ 1 integrin levels showed normal bleeding times, indicating that very low amounts of active  $\beta$ 1 integrins are already sufficient to trigger hemostasis (Figure 9A).

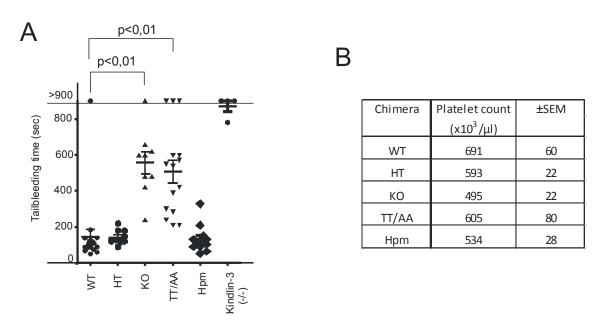


Figure 9:  $\beta 1^{-/-}$  and  $\beta 1^{TTAA/-}$  mice show prolonged bleeding times. (A) Tailtips of isoflurane narcotized animals were cut off. Tails were immediately placed into 37°C saline solution and time until bleeding stopped was measured (every symbol represents 1 animal). Tail bleeding exceeding 15 min was stopped by cauterization to prevent death. (B) Platelet counts were determined using Hemovet blood counter (WT (n=4), HT (n=3), KO (n=7), TT/AA (n=8), Hpm (n=8)). Mean bleeding times are indicated as bars, and error bars represent  $\pm$ SEM.

# 4.7 Carotid ligation assay

To analyze the role of  $\beta 1$  integrins in platelet adhesion to wounded vessel walls in more detail, we performed in vivo carotis ligation assays. In this model, single platelet adhesion to a severe vessel wound (61) is monitored over time. In contrast to an earlier study, we found a significant reduction in adherent  $\beta 1$  integrin deficient platelets at all time points (Figure 10). Beside  $\beta 1$  integrin deficient platelets,  $\beta 1^{Hpm/-}$  and  $\beta 1^{TTAA/-}$  platelets showed a similar reduction in adhesion to the wounded vessel wall. This indicated that 18%  $\beta 1^{TTAA}$  and 3% of  $\beta 1$  integrin surface expression are not sufficient to mediate initial platelet adhesion.

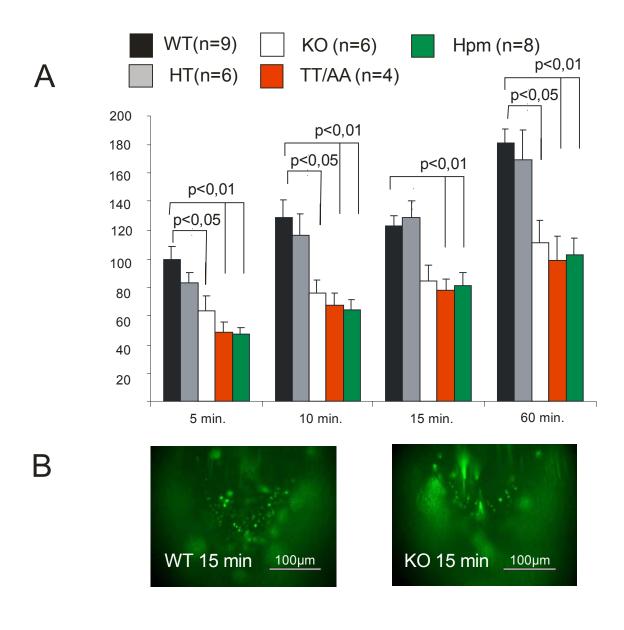


Figure 10: Single platelet adhesion analyzed in a carotid ligation model.  $\beta 1^{-/-}$ ,  $\beta 1^{TTAA/-}$ ,  $\beta 1^{Hpm/-}$  mice show reduced single platelet adhesion to a vessel wound. Carotid arteries were ligated for 5 min and DCF labeled adherent platelets were counted after 5, 10, 15 and 60 min (A). Significance levels and animal number (n) per group are indicated. Platelet adhesion is shown in % (y- axis of cells adherent at 5 min time point in WT animals). (B) Representative pictures of 15 min time point of WT and KO animal are shown.

## 4.8 Aggregation assays

As in vivo data accented a signaling instead of the adhesive function of β1 integrin in hemostasis, we decided to perform aggregation experiments using a light aggregometer. Convulxin a strong GPVI agonist induced similar aggregation in all tested platelet populations (Figure 11A, C). Stimulation with 5 µg/ml fibrillar collagen, which binds GPVI and α2β1 integrin leads to platelet aggregation in wildtype and heterozygous platelets. In strong contrast, β1 deficient platelets failed to aggregate at all and \(\beta1^{TTAA/-}\) mutant platelets showed significantly reduced aggregation. Hypomorph platelets showed an intermediate phenotype (Figure 11A, B). To show that defective \( \beta 1 \) integrin activation and subsequent signaling in β1<sup>TTAA/-</sup> platelets is responsible for the aggregation defect, β1<sup>TTAA/-</sup> mutant platelets and β1<sup>-/-</sup> platelets were incubated with fibrillar collagen in the presence of 0.75 mM Mn<sup>2+</sup>. Indeed, bypassing β1 integrin inside-out activation by Mn<sup>2+</sup> rescued the aggregation defect in β1<sup>TTAA/-</sup> mutant, but not in β1<sup>-/-</sup> knockout platelets (Figure 12), indicating that signals arising from activated \( \beta 1 \) integrin are crucial for platelet aggregation. Furthermore, addition of fibrinogen into the reaction tube rescued the aggregation effect of \$1<sup>+/TTAA</sup> and \$1<sup>-/-</sup> platelets, suggesting that defective secretion of fibrinogen loaded vesicles might be responsible for the aggregation defect (Figure 12).

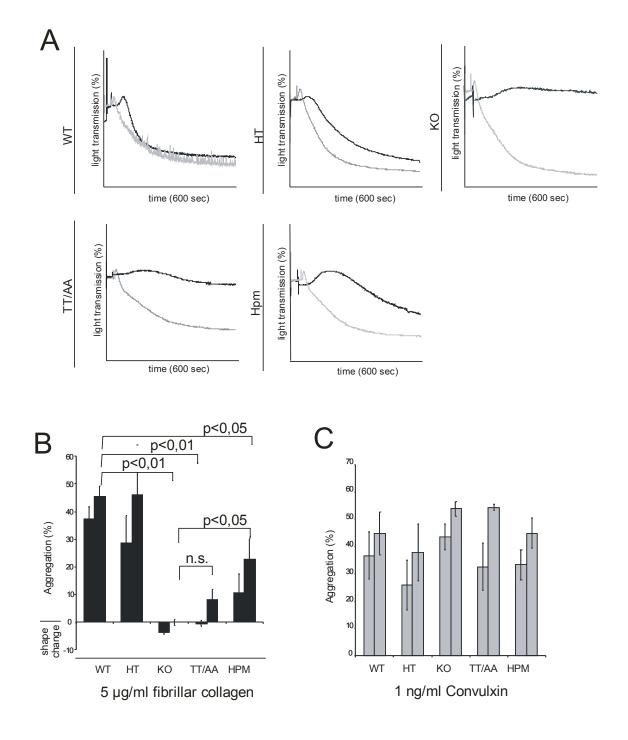


Figure 11:  $\beta$ 1 integrins are required for fibrillar collagen induced aggregation in vitro. (A) Platelets (1x10<sup>8</sup>) were stimulated with 1 ng/ml convulxin (gray graphs) or 5 µg/ml fibrillar collagen (black graphs) and aggregation was plotted for 600 sec N≥4 animals per group. Quantification of aggregation experiments after collagen (B) or convulxin stimulation (C) is shown. Two columns per group represent 300 sec and 600 sec time point. Significance levels are indicated. Error bars represent SEM.

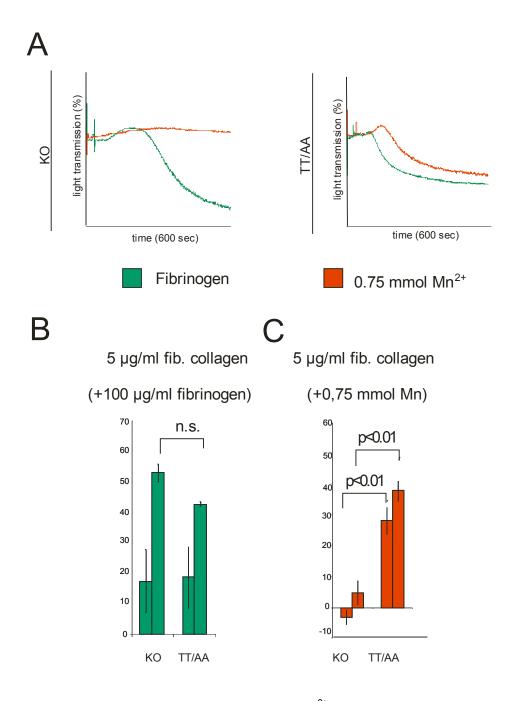


Figure 12: (A) Presence of fibrinogen or  $Mn^{2+}$  can rescue aggregation defect in  $\beta 1^{TTAA/-}$ . Aggregation of fibrillar collagen stimulated platelets, in the presence of 100  $\mu$ g/ml fibrinogen or  $Mn^{2+}$  is shown. Depicted graphs are representative for  $n \ge 3$  independent experiments per group. Quantification of aggregation experiments after collagen stimulation in the presence of fibrinogen (B) or manganese (C). Two columns per group represent 300 sec. and 600 sec. time points. Significance levels are indicated. Error bars represent SEM.

#### 4.9 Dense granula and α-granula secretion

In line with earlier studies on α2β1 deficient platelets (26, 74), we found a delayed maximal shape change in β1 knockout (t=144±23 sec. n=6) and β1<sup>TTAA/-</sup> platelets (t=143±34 sec, n=5) compared to wild-type (t=35±17sec, n=4) and heterozygous platelets (t=44±19, n=4). Hypomorph platelets showed again an intermediate time (t=94±23 sec, n=5). The initial shape change mainly depends on dense granule secretion, which releases different para- and endocrine agonists, such as ADP, ATP, GTP, GDP, serotonin, calcium, magnesium or 5-hydroxytryptamine that promote a positive feed forward loop. These mediators trigger an auto- and paracrine activation of platelets and subsequent aggregation and thrombus formation in vivo. To determine dense granule secretion we used a luciferase reporter system in a luminescence aggregometer. Platelets were stimulated either with 0.1 ng/ml convulxin or 5 µg/ml fibrillar collagen. The collagen induced luminescence signal value and the convulxin signal value are represented as ratio (Figure 14). Interestingly, integrin deficient and mutant β1<sup>TTAA/-</sup> platelets have a severe defect in dense granule secretion after fibrillar collagen stimulation. In parallel to dense granulas, platelets secrete numerous matrix molecules and other prothrombotic molecules stored in α- granules such as contain as fibringen, fibronectin, thrombospondin or vWF (75). Secretion from α-granule was indirectly determined by immunoblotting for cellular fibrinogen before and 8 min after agonist treatment (5 µg/ml fibrillar collagen or 1 ng/ml CVX), under stirring conditions in an aggreometer. Convulxin stimulation induced a strong fibrinogen release with a reduction of the intracellular fibrinogen pool in platelets of all genotypes. On the contrary, fibrillar collagen induced fibrinogen release only in wild-type and hypomorphic platelets. β1<sup>-/-</sup> knockout and β1<sup>TTAA/-</sup> mutant platelets showed almost no decrease of the cellular fibrinogen content (Figure 13A, B). To rule out that the observed differences are due to reduced overall fibrinogen level, we blotted for total fibrinogen level (Figure 13C).

Α

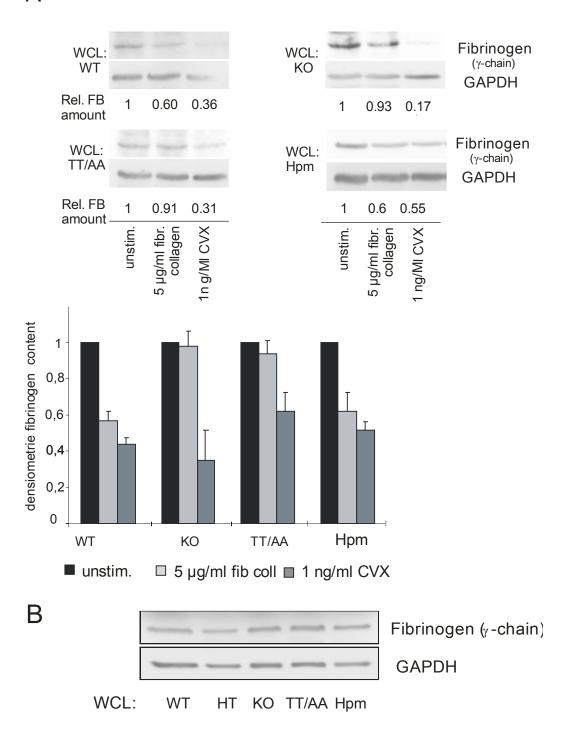


Figure 13:  $\beta$ 1 integrins are required for fibrinogen secretion after collagen stimulation. (A) Cells were stimulated for 8 min in an aggregometer with either 5  $\mu$ g/ml fibrillar collagen or 1 ng/ml convulxin (control). Next, cells were spun down and subjected to immunoblotting for fibrinogen ( $\gamma$  chain) and GAPDH as loading control. (B) No differences in cellular

fibrinogen loading in non-stimulated platelets is shown by immunoblotting of whole cell lysates for fibrinogen (γ-chain). GAPDH shows equal loading. Blots show one representative experiment.

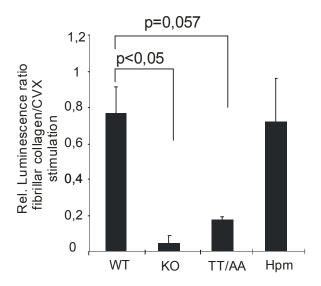


Figure 14: Dense granula secretion upon fibrillar collagen stimulation depends on  $\beta1$  integrins. Aggregation experiments (see above) were performed in the presence of a luciferin/luciferase reporter system to visualize ATP release from dense granula. Platelets were stimulated with either 5  $\mu$ g/ml fibrillar collagen or 1 $\mu$ g/ml convulxin. Detected luminescence (maximal signal strength) was expressed as ratio of fibrillar collagen stimulated to CVX stimulated cells. Bars represent mean values, error bars represents  $\pm$ SEM of at least 3 independent experiments ( $\mu$ ). Significance levels are indicated.

#### 4.10 Integrin tail phosphorylation

We showed that impaired kindlin-3 binding to double threonine mutated cytoplasmic tails has a physiological relevance. However, kindlin-3 binding might be also affect by phosphorylation within the tail. So far no phosphorylation of Thr 788 or 789 in platelets was reported. Hence, we wanted to know if both threonine residues can be phosphorylated in vivo and if so, can it be induced by platelet activation. Therefore, wild-type platelets were stimulated with 0.1 U/ml thrombin in the presence of 50 nmol Calynculin A and analyzed by non-quantitative mass spectrometry. We found peptides which correspond to β1 integrin protein with both

threonine residues individually phosphorylated, however, we failed to detect double threonine phosphorylated peptides. A kinase-data base search predicted PKC as a potential kinase (NetPhosK 1.0: sequence: WDTGENPIYKSAVTTVVNPKYEGK, Thr788: score 0.70/1; Thr789: score 0.76/1) (Figure 15).

No.	Detected	Peptide	Predicted Kinase
	sequences		(likelihood score)
1	-WDTGENPIYK-		
2	-SAVTTVVNPK-		
3	-SAV(ph)TTVVNF	PK-	PKC (0.70/1)
4	-SAVT(ph)TVVNF	PK-	PKC (0.76/1)

Figures 15: β1 integrins are phosphorylated in vivo, potentially by PKC family kinases. Lysates of thrombin stimulated platelets were separated on SDS gel, Coomassie stained and the corresponding bands of the approximate molecular weight of β1 integrin were sent to mass spectrometry to analyze phosphorylation state of threonines 788/789. Table summarizes detected phosphopeptides after 60 seconds of thrombin stimulation of two independent experiments (n=2). Phospho-data base search (NetPhosK 1.0: sequence: WDTGENPIYKSAVTTVVNPKYEGK) predicted PKC family kinases as potential kinases (likelihood score: Thr788: 0.70/1; Thr789: 0.76/1).

## 5. Discussion

# 5.1 β1<sup>TTAA</sup> integrins

Our former study showed that kindlin-3 is indispensable for integrin inside-out activation (35). We further showed that the kindlin binding site within the \( \beta 1 \) integrin cytoplasmic domain comprises a double threonine motif (TT788/789) which is located membrane proximal to the second NPxY motif, which is also required for kindlin binding. Since a recent study using recombinantly expressed αIIbβ3 integrin which was reconstituted as single heterodimers within membrane discs showed a conformational switch from an inactive towards an active conformation in a talin dependent manner but in the absence of kindlin (39), it was questioned whether integrin inside-out signaling requires an integrin-kindlin interaction. Therefore, it was suggested that the phenotypes that are caused by kindlin deficiencies in man and various animal models are due to so far unknown cellular defects. To clarify these points, we analyzed \$1 integrin function in platelets of mice, in which the double threonine motif was mutated to two alanine residues. Using this approach we can clearly show that the kindlin-3 binding motif within the β1 integrin cytoplasmic domain is indispensible for β1 integrin inside-out activation and subsequent downstream signaling. Our data therefore disprove that talin-1 is sufficient to activate integrins. A likely explanation for this contrariety might be given by the fact that Shattil et al. (33) used an artificial in vitro setting neglecting the complexity and variety of other protein interactions at the integrin tail found in a living cell. In contrast, our data come from in-vivo experiments that encountered all phyisological regulatory mechanisms.

# 5.1.1 β1<sup>TTAA</sup> integrin outside-in signaling

During our study we further showed that kindlin-3 is required for integrin outside-in signaling. Integrin downstream signaling leads to the phosphorylation of signaling molecules such as Src, Syk, SLP-76, phospholipases and FAK (7, 76). Kindlin-3 null and  $\beta 1^{TTAA/-}$  platelets fail to phosphorylate FAK upon collagen stimulation. Interestingly we found that addition of manganese that externally activates integrins, could rescue FAK phosphorylation and the spreading and aggregation defects in  $\beta 1^{TTAA/-}$  platelets but failed to rescue FAK phosphorylation in kindlin-3

null platelets. Although the precise mechanism remains to be described, we provide evidence that FAK phosphorylation and integrin outside-in signaling require the presence of kindlin-3 but not a direct interaction between kindlin-3 and the integrin.

#### 5.1.2 Integrin tail binding partner

A general limitation of our genetic animal model system remains that we cannot exclude that the β1<sup>TTAA</sup> mutation affects the binding of other integrin regulators, such as ILK, filamin A, or 14-3-3 protein, since their binding sites overlap with the kindlin binding site (77). However, these proteins are not involved in integrin inside-out signaling, but rather play a role in outside-in signaling. For example, ILK null platelets show only slightly decreased αllbβ3 integrin activation quantified by fibrinogen binding, whereas α-granula secretion and aggregation upon collagen stimulation were severely impaired (78). In our preliminary pulldown experiments we measured (data not shown) only a slight reduction of ILK binding to the β1<sup>TTAA</sup> cytoplasmic tail, arguing that threonines 788/789 are not essential for the ILK binding. However, future experiments should use a proteome wide approach to search for all known candidate interactors as well as unknown once. A quantitative mass spectrometry would offer a sophisticated and elegant way to address this question. The SILAC (Stable isotope labeling with amino acids in cell culture) approach uses cell lysates from mice which were fed with a diet that only contained a (13)C(6)-isotype substituted lysine (referred as "heavy lysate") (79) which can be distinguished from "non-heavy lysates" by mass spectrometry. Using lysates from SILAC mice versus non-SILAC mice will allow the calculation of protein binding ratios. A technical limitation of this approach arises from contaminations of unspecific bound proteins within pulldown eluates. Peptides used for pulldown experiments by us and others in earlier studies (35, 38) facilitated a hexa-histidine tag which was bound by nickel loaded agarose beads. Highly positively charged nickel beads exhibit a high degree of molecular stickiness and subsequent unspecific background protein binding. We therefore redesigned the integrin tails in a way that we could use a biotin streptavidin-bead system to pullout integrin tails. By this tool, we hope to decrease background binding within future mass spectrometry experiments.

#### 5.1.3 Threonine phosphorylation of the integrin tail

Another mechanism might as well contribute to reduced kindlin-3 binding to B1TTAA integrin. Replacing two potential phosphorylation sites could impair regulatory mechanism which control protein interactions at the tail. Protein phosphorylation is a fast and efficient way to rapidly change protein charges and binding affinities. Several so called phospho-switches by site directed phosphorylation at the integrin tail are described (77). Our data showed for the first time that both Thr 788 and Thr 789 residues are phosphorylated, although not simultaneously. However, phosphorylation was only found in the presence of the Serine/Threonine phosphatase inhibitor calinculyn A (80), arguing that only a minor pool of β1 integrins is phosphorylated. Experimental evidence for a threonine phospho-switch within the integrin tail comes from studies in T cells. Phosphorylation of Thr 758 (corresponding to the proximal Thr 788) in β2 integrin regulates Filamin-A and protein 14-3-3 binding to the cytoplasmic tail. Thereby, filamin-1 binds to the unphosphorylated β2 integrin subunit, but falls off from phospho Thr 758 allowing 14-3-3 binding (81), followed by increased cell spreading and migration (77). Another study showed that inhibition of Serine/Threonine phosphatases leads to a more sustained β3 integrin activation, likely through integrin phosphorylation (82). It is tempting to speculate that a similar phospho-switch mechanism regulates kindlin-3 binding to the tail and thereby inside-out activation of β1 integrins. A kinase data base search predicted PKC to be the potential kinase, responsible for Thr 788/789 phosphorylation although with a low likelihood score of around 0.7. In line with our hypothesis, a recent study showed that PKCα deficient platelets have defect β3 inside-out activation, granula secretion and reduced thrombus formation in vivo (83). Drawing another analogy to talin-1 reveals a similar phospho-switch mechanism for tail binding (34). Phosphorylation at the proximal NPxY motif of β3 integrins differentially regulates tail binding of talin-1 and tensin-1, an actin crosslinker protein. Interestingly talin binding is negatively regulated by phosphorylation and talin gets displaced from phospho-tyrosine residues enabling tensin to bind and to increase adhesion strength. In summary, our data support the hypothesis that kindlin-3 binding to the cytoplasmic tail is regulated by a threonine phospho-switch. Therefore, further experiments analyzing kindlin affinity to a differently phosphorylated tail are required to validate our hypothesis.

# 5.1.4 β1<sup>TTAA</sup> expression level

To our surprise, the introduction of the kindlin-binding site mutation into the  $\beta1$  integrin results in a strong reduction of  $\beta1$  integrin expression in platelets. We previously showed that kindlin-3 deficient platelets have reduced  $\beta1$  integrin surface levels (35) suggesting that an interaction between kindlin-3 and the cytoplasmic domain of the  $\beta1$  integrin is required for normal  $\beta1$  integrin expression. A similar observation was made in  $\beta3$  integrin mutant mice, which lack the last three c-terminal residues (RGT) (84). Although the RGT residues are directly adjacent to the membrane distal NPxY motif, it remains speculative whether kindlin binding is affected by this deletion. The molecular cause for these observations is still unclear and requires further investigations.

#### **5.2** β1 integrins in arterial thrombosis

Various groups analyzed platelet  $\beta1$  integrin contribution in hemostasis in animal models. However, the acquired data and conclusions remain inconsistent. Analyzing our two newly generated mouse models ( $\beta1^{TTAA/-}$  MxCre,  $\beta1^{Hpm/-}$  MxCre), we clearly show that  $\beta1$ -integrins facilitate an important signaling role in- and exvivo by regulating platelet granula secretion, aggregation and thrombus formation. Unexpectedly, we find that collagen signaling by GPVI is insufficient to activate platelets and requires an additional  $\beta1$  integrin mediated signal. Furthermore, we could show that  $\beta1$  integrins contribute to single platelet adhesion in vivo. A possible explanation for different results might arise from different experimental settings (discussed below) as well as different animal models, namely  $\beta1$  integrin depleted versus  $\alpha2\beta1$  depleted platelets, being used. Hence a careful review of experimental setups might give reasonable explanation.

Regarding clinical implications, our data underline the therapeutic potential of  $\beta 1$  integrin inhibition during treatment of acute arterial thrombosis. Although an obvious limitation might be that inhibitor molecules would have to reach 100% receptor blockade with a sufficient pharmacokinetics, as ~3% of intact integrin are sufficient to reach full platelet activation upon collagen stimulation.

#### 5.2.1 Bone marrow chimera

Bone marrow chimeras are an elegant way to avoid off-target conditional gene deletion in vivo. It was shown that poly I:C induced expression of Cre-recombinase efficiently deletes the conditional gene loci in the hematopoetic system and some other off-target tissues including kidney or liver cells (58). Preliminary experiments showed that poly I:C treated  $\beta 1^{-/-}$  MxCre animals have a severe and sustained liver failure with a strong inflammatory phenotype and a confined liver synthesis capacity (Petzold et al. unpublished). This phenotype is not seen in bone marrow chimeras carrying a  $\beta 1^{-/-}$  MxCre bone marrow in a C57/BL6 mouse background. As an altered liver synthesis might affect plasma proteins level and by this platelet granular content of internalized plasma proteins which get released upon platelet activation (85). In summary, generation of bone marrow chimeras is essential to prevent off-target effects.

#### 5.2.2 Carotid ligation assay and occlusive thrombus formation

Keeping in mind that  $\beta1$  integrin expression levels (e.g. approximately 3000  $\alpha2\beta1$  integrins per cell) are minor compared to  $\beta3$  integrin levels (e.g. approximately 60000 receptors per cells), it seems critical to find experimental conditions that are sensitive enough to uncover  $\beta1$  specific function. First, we analyzed single platelet adhesion in a carotid ligation model. Our results clearly show that  $\beta1$  integrins play a significant role in single platelet adhesion in vivo. The reason for the impaired adhesion of  $\beta1^{TTAA}$  mutant platelets is hard to interpret since we found an integrin activation defect and reduced  $\beta1^{TTAA}$  integrin surface expression. An appropriate control animal with around 18% functional integrin  $\beta1$  is needed to allow conclusions on the importance of  $\beta1^{TTAA}$  integrin in this process.

Our findings contradict the data generated by Gruner et al. (27), who found no adhesion defect in the same  $\beta 1^{-/-}$  MxCre animals and the same thrombosis model. A possible reason for the discrepancy might be that this group did not use bone-marrow chimeras (see above). Furthermore, they found occlusive thrombus formation in WT animals which was neither seen by us nor by Massberg et al. (61). Electron microscopic images from ligation areas by Massberg et al. revealed a well defined arterial lesion with endothelial cell denudation and matrix exposure after 5 min of vessel occlusion. In contrast, Gruner et al. used a ligation time of

only 30 sec without a precise description of their induced lesion. It is likely that other thrombosis inducing mechanisms (e.g. tissue factor) overweight the contribution of platelet-matrix interaction as primary stimulus.

Second, we wanted to analyse the role of  $\beta 1$  integrins in occlusive thrombus formation. Several models are applicable to induce an occlusive arterial thrombosis in mice (61, 86) and various were used to study \(\beta 1\)-integrin contribution. He et al. (30) found prolonged vessel occlusion times in a photochemical assay but no β1-integrin contribution in a pulmonary embolism model (86). The photochemical injury model is based on increased reactive oxygen production within the endothelium and potentially other less well defined mechanism (86). Hence a limitation of the model remains that it might be unsuitable to study ECM induced thrombosis. Kuijper used a Ferric-(III)-chloride model in integrin a2 null animals and found no difference in thrombus formation but decreased thrombus stability with frequently appearing embolic events in these animals (87). However, the underlying thrombosis inducing mechanism was shown to (mainly) rely on increased thrombin generation on the surfaces of ferric ion filled particles exposed on the endothelium and the exposed matrix shortly after placement of an Ferric chloride soaked paper on mesenteric arteries (88). This thrombosis model was shown to be insensitive to GPVI or β1 depletion and might therefore lead to an underestimation of β1 contribution in vivo. Trying to find a reasonable model to analyze (primarily) extra cellular matrix induced thrombosis, we decided to perform tailbleeding assays. Historically, bleeding assays were developed to screen for bleeding disorders in patients. For animal studies two different protocols were established in analogy to the clinical tests by Marx or Ivy. We decide to adapt the protocol by Marx to measure tail bleeding times. Therefore, tail tips of chimera animals were cut and placed into an isotonic 37°C warm water bath and time till bleeding stopped was measured. As a high degree of reproducibility and standardization is needed to reveal minor bleeding phenotypes, as expected for β1 integrin null animals, we set our wound 8 mm away from tail tips in 10-12 weeks old chimeras. Further, we choose an isofluran gas narcosis instead of intraperitoneal application of drugs (26, 89). This technique allowed a steady narcosis depth, monitored by breathing frequency, to prevent massive changes within circulatory parameters. Obviously, anatomical alteration of the tail vasculature between individual mice might lead to some variation in

bleeding behavior. Surprisingly, we found that as little as ~3% of functional  $\beta1$  integrins found in hypomorph mice are sufficient to yield normal bleeding and vessel occlusion times. In contrast  $\beta1$  integrin null mice or inactive integrin  $\beta1^{TT/AA}$  carrying mice showed prolonged bleeding times. Our results are in line with other studies using  $\alpha2\beta1$  depleted animals (29-30), but contradict two others (26, 89). The later used a bleeding assays adapted from clinical setting by Ivy, in which blood drops at the lesion were removed by whatman paper every 30 seconds until bleeding stopped. We tried to reproduce our data by this assay, however found big variations even inside the control group (preliminary data, not shown), as manipulating on the tail often caused disclosure of the lesion or sometimes an immediate bleeding arrest. In summary, our in vivo data show an important role of  $\beta1$  integrins for platelet adhesion and occlusive thrombus formation.

### 5.2.3 Aggregation and secretion

Trying to find a mechanism which might help to explain our tailbleeding data, we studied platelet aggregation. Earlier aggregation assays on β1 or α2β1 integrin deficient platelets were performed in the presence of 100 µg/ml fibrinogen and did not find any aggregation defect upon fibrillar collagen stimulation (26, 89). Interestingly, mouse fibringen plasma level in C57/BL6 mice range around 1.5 mg/ml and are therefore 15 times higher than used in above mentioned studies (90). As mimicking in vivo plasma conditions ex- vivo remains challenging, we performed our aggregation assays in the absence of fibrinogen or other plasma proteins and found an aggregation defect for β1 null and β1<sup>TTAA/-</sup> platelets. Moreover, addition of MnCl<sub>2</sub> to β1<sup>TTAA/-</sup> platelets rescued the aggregation defect and underlined that \( \beta \) derived signals are rather essential than just supportive as discussed above (91). Most surprisingly, we found that already minor amounts (~3%) of active β1 integrins are sufficient to trigger integrin outside-in signaling and subsequent aggregation. We further could show that the aggregation defect is due to an inhibited dense (ATP) and fibrinogen secretion from α-granula. So far, GPVI was thought to overtake all collagen signaling in vivo, leading to platelet activation, aggregation and granular secretion (26, 61-62, 92). However, earlier work from Chen et al. (91) suggested a two side two-step model in which α2β1 activation is downstream of GPVI and generates similar signals that merge into

GPVI downstream signaling pathways via Syk, SLP-76. These authors limited their hypothesis to conditions in which  $\alpha2\beta1$  signaling might compensate for insufficient GPVI signaling (e.g. in the presence of GPVI blocking reagents). We show that  $\beta1$  integrin signaling is required and indispensable for granula release and subsequent aggregation even in the presence of high concentrations (5µg/ml) of fibrillar collagen (93). Future studies have to address the precise molecular signaling pathways how  $\beta1$  integrin signaling affect granular secretion.

# 6. Summary

The affinity state of integrins is regulated by two cytoplasmic integrin binding proteins: talin and kindlin. Loss of either of them results in inactive integrins in various cell types. To test whether a direct interaction between kindlin and the integrin is required for integrin activation (integrin inside-out signaling) and β1 integrin mediated signaling (integrin outside-in signaling) we analyzed platelets from mice expressing a kindlin-binding deficient β1 integrin (β1<sup>TTAA</sup>). Our data clearly show that integrin inside-out signaling depends on a direct integrin-kindlin interaction as β1<sup>TTAA</sup> integrins remain inactive upon platelet activation by soluble agonists. On the contrary, exogenous activation of β1<sup>TTAAV-</sup> platelets by manganese results in normal cell spreading and FAK phosphorylation indicating that outside-in signaling does not depend on a direct interaction. Furthermore, mice expressing B1<sup>TTAA</sup> mutant integrins show prolonged bleeding times comparable to mice exhibiting a blood cell specific deletion of the \beta1 gene. Single platelet adhesion to injured vessel walls is strongly reduced in β1 null and β1<sup>TTAA</sup> platelets suggesting an important role of \( \beta 1 \) integrins for initial platelet adhesion. In line with this observation, we show that β1 integrin-mediated signals direct granula secretion, required for a pro-thrombogenic micro milieu favoring platelet adhesion and aggregation during the early phases of hemostasis. Most surprisingly, 3% of active β1 integrin are sufficient to rescue bleeding defects.

# 7. Zusammenfassung

Die Aktivität von Integrinen wird durch die beiden zytoplasmatischen Proteine Talin und Kindlin reguliert. Der Verlust eines von beiden führt zu inaktiven Integrinen in verschiedenen Zelltypen. Um zu prüfen, ob eine direkte Integrin-Kindlin Interaktion für die Integrinaktivierung als auch die anschließende Signalgenerierung benötigt wird, haben wir die Funktion von Mausthrombozyten analysiert, deren \( \beta 1 \) Integrine nicht mehr in der Lage sind, Kindlin zu binden. Unsere Daten zeigen, dass für die Integrinaktivierung eine direkte Kindlin-Integrin Interaktion notwendig ist, da mutierte β1<sup>TTAA</sup> Integrine nach Stimulation mit löslichen Agonisten inaktiv bleiben. Im Gegensatz dazu, findet sich nach externer Aktivierung der β1<sup>TTAA</sup> Integrine durch Mangan ein normales Zellspreading sowie eine normale FAK Phosphorylierung. Diese Beobachtung zeigt, dass Integrin vermittelte Signale keine direkte Interaktion von Kindlin mit dem Integrin benötigen. Weiterhin haben Mäuse, deren Thrombozyten β1<sup>TTAA</sup> Integrine exprimieren, verlängerte Blutungszeiten, die vergleichbar mit denen von ß1 Integrin (Blutzell-) depletierten Tieren sind. Die Anzahl adhärenter Thrombozyten an eine Gefäβläsion der Arteria carotis ist in β1-Integrin depletierten und β1<sup>TTAA</sup> exprimierenden Tieren reduziert, was die wichtige Rolle der β1 Integrine während der initialen Phase der Thrombozytenadhäsion hervorhebt. Weiterhin zeigen wir, dass β1 vermittelte Signale die Granulasekretion steuern, die zur Schaffung eines pro-thrombotischen Mikromilieus notwendig ist, um somit eine weitere Thrombozytenadhäsion und Aggregation während der frühen Phase der Blutstillung zu unterstützen. Überraschenderweise reichen bereits ~3% an aktiven β1 Integrinen auf den Thrombozyten, um eine normale Blutungszeit im Tiermodel zu erhalten.

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Focal adhesion kinase modulates activation of NF-[96]B by flow in endothelial cells. Petzold T, Orr AW, Hahn C, Jhaveri K, Parsons JT, Schwartz MA; Am J Physiol Cell Physiol. 2009

Genetic disruption of the Kindlin-3/ß1-integrin interaction results in defective platelet adhesion and granula secretion. Tobias Petzold, , Raphael Ruppert, Dharmendra Panday, Verena Barocke, Wolfgang Siess, Reinhard Fässler, Steffen Massberg and Markus Moser (script in preparation)