

# Antithrombotic drugs

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## Chapter Eight

# Antithrombotic drugs

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## 1. Summary

The haemostatic-thrombotic system (HTS) can be inhibited at dozens of different sites by hundreds of different drugs. The ideal antithrombotic is an inhibitor that downregulates the HTS without causing bleeding or other side-effects, and that can be given orally in a fixed dose. These are pharmacological criteria. Indeed, the three current therapies (heparins, oral anticoagulant therapy and aspirin) act by completely different mechanisms. It is not the mechanism of inhibition which is important but the pharmacological properties of the inhibitor; no compound has been found that inhibits the HTS and does not show an antithrombotic effect.

The HTS is a non-linear system containing a number of nested positive and negative feedback loops. At the present state of knowledge it is impossible to predict the effect of inhibition of a single reaction on the response of the complete system. For this reason one cannot predict the antithrombotic potency of a compound from its biochemical properties. It is shown that clotting tests, measurement of platelet aggregation/adhesion or tests of individual functions within the system are only of limited use. Antithrombotics therefore should be evaluated by their action on a representative function test of the complete HTS. Such a test is not routinely available at the moment. Two promising possibilities are flow chambers using non-anticoagulated blood and thrombin generation in blood or platelet rich plasma (PRP).

## 2. Background

Thrombosis is caused by the same mechanism that causes haemostasis. Administering antithrombotic drugs therefore requires careful manipulation of the balance between thrombosis and bleeding. For the safe use of

antithrombotic drugs, we need a probe for the functional status of the HTS. Such a test is not routinely available at this moment. Bleeding times are hopelessly inaccurate [1] and clotting times are to be considered surrogate variables of limited use. There is no general test to indicate hypercoagulability and thrombosis risk, or to monitor the effect of antithrombotic treatment. Current management of thrombosis is like managing diabetes by the taste of the urine, or hypertension with only a pulse to feel.

This situation has an immediate influence on the search for new antithrombotic drugs. There is no easy, valid laboratory test to indicate therapeutic efficacy or optimal dosage. The research goal is defined in terms like 'doubling the activated partial thromboplastin time (aPTT)' or 'inhibition of platelet aggregation' that have, at the best, a semiquantitative relation to haemostasis and thrombosis. Further testing is in animal thrombosis models and, eventually, clinical trials.

In this article we will defend the position that it is unimportant how the function of the HTS is inhibited as long as the inhibition fulfils certain pharmacological criteria. Consequently, we will not dwell in detail on individual mechanisms of inhibition or on inhibitor design. Rather we will stress the importance of assessing HTS function as a whole.

### 3. Medical need

In recent years it has been firmly established that virtually every arterial occlusive event (such as myocardial or cerebral infarction) is due to the formation of a local thrombus on a damaged internal surface of the vessel. Thrombin plays an all-important role in this process [2-6]. A damaged wall exposes triggers for thrombosis, such as tissue factor (TF) present in the smooth muscle cells of neointima [7], in the adventitia and in the plaque [8-10], as well as the platelet-activating collagen surface [11]. Non-occlusive arterial thrombosis may lead to downstream occlusion if the thrombus is carried with the bloodstream. In strategic positions, such as the carotid arteries, this may have serious consequences. Arterial thrombosis is not only the consequence of atherosclerosis; through microthrombus formation it also plays a role in the development of arterial wall degeneration. Furthermore, venous thrombosis and its sequelae, ranging from sudden death due to pulmonary embolism to chronic leg-ulcers, is not a negligible source of illness. At this moment, in the Western World, arterial and venous thrombosis and atherosclerosis together are responsible for well over 50% of all mortality and serious morbidity. In a few decades this will be the case worldwide [12]. Good antithrombotic drugs, therefore, are a necessity.

#### 3.1 Existing treatment

##### 3.1.1 Heparins

Heparins, either unfractionated (UFH) or of the low molecular weight type (LMWH) [14], reduce the incidence of thrombosis by about 50%. They have the drawback of parenteral administration. They may cause antithrombin (AT) deficiency, thrombocytopenia, bleeding and a few more rare side-effects. The methods for laboratory control of heparin dosage, notably the aPTT, are inadequate. The question of whether adapted doses of heparin would do better than current practice remains unsolved because control methods to guide the adaptation are not available. Prolonged heparin treatment, in particular with UFH, is not favoured.

### 3.1.2 Oral anticoagulation

Oral anticoagulation (OAC) is maintained with coumarin congeners, i.e., anti-vitamin K drugs. They inhibit the oxidation reduction cycle of vitamin K in the liver, required for the normal  $\gamma$ -carboxylation of the clotting Factors II, VII, IX and X, protein C and protein S. There are large variations in individual need and many drug interactions (e.g., with sleeping pills and alcohol). Therefore, regular control of the level of anticoagulation is necessary. This can be readily carried out *via* determination of the thromboplastin time and expressing the result as an international normalised ratio (INR).

### 3.1.3 Aspirin

Aspirin is the only antiplatelet drug in general use. The effect is dose-independent between 40 and 1000 mg/day. There are few complications seen. The protective effect in venous thrombosis is not significant and in arterial thrombosis it is modest (10 - 20%). Ticlopidine and clopidogrel (Iscover<sup>TM</sup>, Plavix<sup>TM</sup>) are alternative possibilities. Abciximab, the antibody against the fibrinogen receptor of plasma, is used in acute coronary infarction.

## 4. Therapeutic class review

Heparins are glycosaminoglycans with molecular weights ranging from 1500 to 100,000 Da [14]. Those that contain a specific pentasaccharide sequence bind to AT and catalyse the irreversible inhibition of activated clotting factors by this plasma protein. The inhibitory capacity of AT towards thrombin is roughly three times that towards Factor Xa; that towards IXa and XIa is smaller and probably without practical significance. The relative activity against the different factors is not changed by heparin, with one notable exception: heparins smaller than 5400 Da will not cause inactivation of thrombin, but do catalyse that of Factor Xa. Inhibition of Factor Xa is an inefficient way to inhibit thrombin formation. LMWHs contain a fair amount of molecules of < 5400 Da, whereas UFH does not; therefore LMWHs have a higher anti-Xa activity than UFH. The difference is small in the presence of Ca<sup>++</sup> ions (i.e., *in vivo*) but appears much larger in the usual *in vitro* test, where Ca<sup>++</sup> is absent [15]. A high anti-Xa activity has been the guideline in searching for efficient LMWHs [14]. Despite the inefficiency of Factor Xa inhibition this has led to useful products, because preparations with a high anti-Xa activity also contain a large proportion of material in the 5400 - 9000 Da range. Such material is an efficient anticoagulant because of its persistent AT activity, together with a high bioavailability after sc. injection [16,17]. It has also been surmised that the very large molecular weight components present in UFH, but not in LMWH, have specific haemorrhagic properties [18]. In retrospect, the anti-Xa activity must be considered largely an artefact; LMWHs are good drugs developed on the basis of a mistaken concept.

AT only partially inhibits clot-bound thrombin, if at all. Heparin co-factor II (HCII) [19] is capable of inhibiting this thrombin, but only in the presence of relatively high concentrations of acid polysaccharides (dermatan sulfate, pentosan polyphosphate, lactobionic acid, etc.) [20-24]. Heparin can also activate HCII, but at doses much higher than the normal therapeutic and preventive ones.

### 4.1 Direct thrombin inhibitors

The prototype of direct thrombin inhibitors is hirudin [25-28], which has proven its value in experimental thrombosis as well as in clinical trials. Indeed direct thrombin inhibition is a feasible route to obtain antithrombotic action [6,29-33]. Modern techniques of

combinatorial chemistry can construct large peptide derivative libraries and facilitate the search for specific inhibitors [34,35]. There is also continuous interest in natural inhibitors from haematophagous animals and other sources [36-40], CVS1123 [41], inogatran (HA-31427) [42-44], melagatran (H-31968) [42-47] and napsagatran (Ro-46-6240) [48], and a large variety of others [49-65]. Some have been tested in the clinical situation. Most of them are administered parenterally, but oral administration is feasible [66]. The overall impression is that, like hirudin, they can replace heparin, without showing obvious advantages except their use in AT deficiency and their ability to inactivate clot-bound thrombin [44,49,67,68].

There is an essential difference between irreversible thrombin inhibitors (such as hirudin) and reversible ones (such as argatroban [Novastan™]). The irreversible ones bind with 1:1 stoichiometry to thrombin. If drug administration is stopped, the inactivated thrombin molecules remain inhibited. Serious bleeding problems are to be expected as soon as the concentration of an irreversible inhibitor approaches the level of prothrombin in plasma (~2µM), so that all the thrombin that can be formed is immediately inhibited. Reversible inhibitors bind a fraction of the available thrombin; the proportion inhibited is dependent upon the inhibition constant ( $K_i$  = the concentration at which half-maximal inhibition is observed) and upon the concentration of the inhibitor. If the concentration of the inhibitor decreases, the inhibition is relieved. Reversible inhibitors probably act because they diminish thrombin concentration at the site of thrombosis, where thrombin formation is triggered. They will enter into competition with the natural substrates of thrombin (procoagulants: e.g., fibrinogen; Factors V, VIII, XI; platelets, as well as anticoagulants: e.g., thrombomodulin). All reversibly inhibited thrombin molecules will be eventually released and taken over by natural, irreversible inhibitors (e.g., AT). So, irreversible inhibitors arrest thrombin action and reversible ones dampen it. Both mechanisms seem to lead to antithrombotic activities. It is unknown to the authors whether there are fundamental differences in haemorrhagic properties.

#### 4.2 Direct inhibitors of other clotting enzymes

A large number of natural and synthetic inhibitors of Factor Xa has been described [35,69-86]. The original idea behind their development was that it is more efficient to inhibit thrombin production ('closing the tap') than thrombin itself ('mopping the floor'). This view, based upon the early concepts of the clotting cascade as a linear mechanism, became untenable as soon as it was realised how important thrombin-mediated feedback reactions are and how important Factor Xa is for the inhibition of the TF Factor VIIa complex (TF-VIIa) *via* tissue factor pathway inhibitor (TFPI). Also, Factor Xa has to be inhibited by 90% in order to inhibit prothrombin conversion by about 50%.

Inhibition of the first step in thrombin generation, i.e., at the level of Factor VIIa and TF, is also possible [87-89]. An interesting concept is the inhibition of TF by the administration of recombinant, active site-inhibited Factor VIIa [90-92], or by peptides that mimic part of this molecule [93].

All VIIa and Xa inhibitors tested appear capable of inhibiting experimental thrombosis (e.g., [94-96]). They have not yet been used in the clinic, and it is not established whether they show specific advantages. However, an important finding has recently been made that shows that circulating TF plays a role in the formation of the thrombus [170]. A competitive VIIa inhibitor could inhibit this thrombus formation. Theoretically, this approach towards thrombus prevention would have a broad therapeutic range since the amount of circulating

TF is much easier to inhibit than the local TF exposure after vessel damage, thus minimising bleeding risk.

The main result from research on direct inhibitors as a group is that any inhibitor of the clotting system, independent of the enzyme that it attacks or of the mode of inhibition is antithrombotic, in any thrombosis model.

### 4.3 Platelet inhibitors

Many excellent reviews on platelet inhibitors have been published [e.g., 97,98]. Aspirin, in all probability, acts by irreversibly inhibiting cyclooxygenase and, thereby, platelet thromboxane A<sub>2</sub> (TXA<sub>2</sub>) synthesis [99]. Other non-steroidal anti-inflammatory drugs (NSAIDs) and sulfapyrazone inhibit the same enzyme reversibly. Dipyridamole increases intraplatelet cyclic AMP (cAMP) [99-101], as do prostaglandin E1 and prostacyclin (which, due to its instability, is of limited therapeutic use) [102,103]. Stable analogues of the latter are iloprost (SH-401) [104,105] and ciprostone (U-61431F) [106]. Thromboxane synthase inhibitors and inhibitors of the thromboxane receptor are of doubtful efficiency [107]. Ketanserin, one of the group of serotonin receptor antagonists seems to do no better [108,109].

Inhibitors of the ADP receptor, such as ticlopidine [110-113] or clopidogrel [114], are better than aspirin and may be combined with it. Inhibitors of the platelet fibrinogen receptor glycoprotein (GP) IIb/IIIa have shown promising results [115-119].

Platelet inhibitors have been developed as inhibitors of adhesion or aggregation. Recently, it appeared that all those tested hitherto also inhibit thrombin generation in PRP. Aspirin, for example, causes an inhibition of about 10%, roughly equal to the reduction of arterial thrombosis that is seen in large clinical trials. Prostacyclin analogues also inhibit thrombin generation in PRP [120] (unpublished observations). Inhibition of GP IIb/IIIa, the fibrinogen receptor of the platelet membrane, reduces thrombin generation by about 50% [121,122]. Administration of these inhibitors produces the same defect as is congenitally present in Glanzmann's thrombasthenia, a disease in which a diminution of thrombin generation is observed [121,122]. Also, inhibition of von Willebrand factor (vWF), GP Ib and the vWF receptor strongly inhibits thrombin generation in PRP [123,124]. Thrombin generation in PRP is evidently also dependent upon the function of the plasmatic clotting factors and upon the interactions between plasma and platelets. This suggests that thrombin generation in PRP and/or blood might be a good screening technique for antithrombotics, independent of whether they act *via* the platelet or as anticoagulants.

## 5. Current research goals

The ultimate research goal is an antithrombotic that decreases the incidence of thrombosis (of any type) by more than 50%, that can be administered orally, that is immediately active, that can be given in a standard dosage without requiring laboratory control and that does not cause bleeding when overdosed. None of the available antithrombotics fulfil these criteria.

The complicated HTS lends itself to inhibition at a large number of sites, each of them a potential target for pharmacological action. The three established therapies have completely different modes of action. Multiple other modes of inhibition have been shown to work in experimental thrombosis. This proves that the desired goal can be attained in many different ways. The question of what specific target in the HTS should be attacked in

order to obtain the ideal antithrombotic is often posed. It is mostly answered *a priori* by theoretical considerations that need to be corroborated *a posteriori* by animal thrombosis experiments. The result, as stated, is that any inhibition entails antithrombotic activity. Advanced knowledge of the subsystems of the HTS may, in the long run, give the answer. This goal is not within easy reach however. At this moment, it would be more useful to conceive of a screening test that represents total HTS function, and that is simpler and less expensive than animal thrombosis models.

### 5.1 Laboratory models for the function of the HTS

The ideal model, like real thrombosis, should respond in a similar way to clinically effective concentrations of all antithrombotics independent of the molecular focus of their effect. This means that clotting times in PRP (and *a fortiori* inhibition tests of individual clotting factors), platelet aggregation or adhesion tests carried out in anticoagulated PRP, are not reasonable options. In view of their widespread use, we will nevertheless discuss clotting times in more detail.

### 5.2 Clotting times

Blood (plasma) clots when more than 95% of the prothrombin is still to be converted [120]. Clotting times do not reflect the amount of thrombin that is still to be formed at the moment of clotting. Yet it is the amount of thrombin that determines the extent of the thrombotic reaction. Therefore, a clotting time will not reflect directly the relevant parameter. This is also seen from the fact that a different type of clotting time is required for estimating the effect of different types of anticoagulants. The thromboplastin time or 'prothrombin time' (PT), for example, indicates the depth of oral anticoagulation but is not sensitive to the effect of heparin. The aPTT is particularly sensitive to variations in the concentration and/or activity of Factor VIII and UFH, but not to LMWH and activators of HCII. Doubling the aPTT is thought to represent adequate heparin administration. This is based on very limited studies in animals [125] and man [126,127]. There is no relation between recurrent thromboembolism and the aPTT response [128]. The mechanism by which direct thrombin or Factor Xa inhibitors prolong the aPTT is significantly different from that of heparin [129]. Nevertheless 'doubling the aPTT' is also used as an indicator for therapeutic action with any type of clotting inhibitor. We conclude that clotting tests are not a valid general indicator of antithrombotic action. This does not mean that they cannot be used in special cases: the thromboplastin time is a good indicator of oral anticoagulation [130] and the Ecarin time is useful in assessing the level of direct thrombin inhibitors [131,132].

### 5.3 Platelet tests

Platelet aggregation and flow-chamber or other tests that are carried out on washed platelets or on PRP in which the coagulation mechanism has been inhibited, will not reflect interactions between platelets and the clotting system, and therefore are of limited use [133-135]. Experiments in which non-anticoagulated blood is passed through a flow chamber containing a thrombogenic surface are not subject to this drawback. A model system that tries to mimic the concert of thrombotic triggers and flow parameters has been developed [136-140]. A flow chamber is directly coupled to the antecubital vein of a volunteer, and flow of non-anticoagulated blood is metered by a peristaltic pump. The flow chamber contains a piece of the media of a pig aorta, cultured endothelial cells, smooth muscle cells or collagen-coated surfaces and human atherosclerotic plaques. After the experiment, the exposed material will be stained and examined for platelet and fibrin deposition. Drugs can be administered and tested for antithrombotic properties. This

model for thrombosis has proven to be a workable system that closely mimics physiologic reality under standardised conditions.

#### 5.4 The thrombin generation curve

The thrombin generation curve (TGC), one of the oldest tools of the clotting trade, has lately become the subject of renewed interest [15,74,120,122-124,141-44]. It contains all the relevant information on the physiological function of the coagulation system. The area under the curve (the endogenous thrombin potential (ETP) [142]) represents the total enzymatic action that thrombin can develop during its lifetime in plasma. In blood and in PRP, the ETP can be measured from a TGC obtained by subsampling. If a substrate with the right kinetic properties is added to plasma, thrombin generation can be monitored continuously as the first derivative of the product-time curve. At this moment, only chromogenic substrates have been described for this use so that the method is restricted to defibrinated PRP.

We demonstrated that higher than normal ETP is seen in all blood-based thrombosis-prone states tested, either congenital (e.g., deficiency of AT, protein S or protein C, factor V Leiden, prothrombin 20210A) or acquired (e.g., from the use of contraceptive pills). In coronary infarction, months after the acute phase as well as in patients presenting with a deep vein thrombosis, the ETP is increased. In the haemophilias (deficiency of Factors VII, IX and XI) the ETP is significantly lower than normal [145].

OAC treatment (INR = 1.5 - 3.0) decreases the ETP to 20 - 40% of normal. Heparin administration in the therapeutic range has the same effect. The ETP also accurately reflects the effect of mixed OAC and heparin treatment. Platelet aggregation inhibitors diminish thrombin generation in PRP. In fact, we encountered no anticoagulant that would not decrease the ETP in PRP, and no antiplatelet agent that would not inhibit the ETP in PRP. This made us think that the ETP in PRP might be a useful model for antithrombotic action.

## 6. Scientific rationale

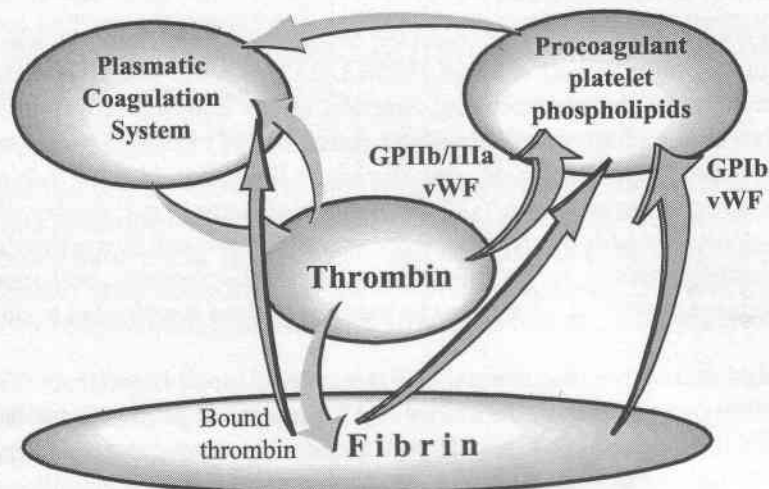
The HTS is traditionally seen as the result of the sequential action of primary and secondary haemostasis. The first, reflected in the bleeding time, is considered to be a platelet function and its overshoot is the arterial, platelet-rich thrombus. The second is seen as an activity of the plasmatic clotting mechanism; the plasma clotting time is considered the relevant test and venous thrombosis the pathological derailment.

The separation has been artificially stressed because platelet adhesion and aggregation are studied in anticoagulated blood, and clotting in platelet-free plasma. Under these experimental circumstances the co-operation between platelets and coagulation is artificially prevented. The impressive increase of our knowledge of the details of the clotting mechanism and platelet function tends to favour further subdivision. This is reflected in drug research. Not only is a sharp distinction made between anticoagulants and platelet inhibitors, but often a more restricted target is defined (e.g., Factor VIIa or G-proteins) and screening is for an action on that target rather than on global functioning of the HTS.

It is becoming increasingly clear that, *in vivo*, the HTS functions by the concerted action of platelets and plasma, rather than as a consequence of an adhesion-aggregation function of the platelets and a clotting function of the plasma (Figure 1). Thrombin is the most potent platelet activator, and platelet activation is essential for thrombin generation.



**Figure 1:** Outlines of the interactions between the coagulation system and the platelet glycoprotein (GP) IIb/IIIa and GPIb receptors (from [123]). VWF: von Willebrand factor.



Anticoagulants inhibit arterial as well as venous thrombosis [146,147]. Platelet inhibitors inhibit thrombin generation in PRP, etc.

This is not the place to discuss the mechanisms in detail. We refer to books and review articles for this purpose. Here we only sketch outlines, together with some recent findings that may have an important bearing on drug development.

In the plasmatic coagulation system, Factor Xa is formed by the action of the TF-VIIa. Factor Xa binds to TFPI, and the TFPI-Xa complex inhibits the TF-VIIa complex. Thrombin activates Factors V, VIII and XI and so accelerates its own generation, but it also binds to thrombomodulin and so starts the protein C mechanism that breaks down Factors V and VIII, thus indirectly inhibiting further thrombin generation. An important fraction (~ 30%) of all thrombin formed in clotting plasma is bound to the fibrin clot. Clot-bound thrombin does retain its thrombotic properties; it can clot fibrinogen, activate Factors V, VIII and XI and platelets [123,148,149]. It is not inhibited by AT and, upon fibrinolysis, it may reappear in circulation.

Four different thrombin receptors have been described on the platelet membrane. Thrombin action causes GPIIb/IIIa receptors in the platelet membrane to bind fibrinogen, which causes platelet aggregation. GPIIb/IIIa activation also leads to the exposure of procoagulant phospholipids in a vWF-dependent reaction [123,124]. These phospholipids are required for the proper activation of Factor X and prothrombin. Lately, the picture has been complicated by the discovery that fibrin, previously thought to be the inert end product of coagulation, plays an active role itself. It binds and activates vWF, which in turn activates platelets *via* GPIb, and provokes the exposure of procoagulant phospholipids *via* an alternative pathway.

Thus, the co-operation between platelets and the coagulation system is central to the HTS, and the mechanism shows an abundance of positive and negative (often nested) feedback loops. This necessarily is a non-linear system [150]. In the most simple terms this means that the dose-response relationships become highly unpredictable ('chaotic'). Furthermore, thrombin generation reactions often go to completion, so that classical initial rate kinetics is

of limited applicability. The literature abounds with studies that conjecture on antithrombotic effects of enzyme inhibition on the basis of oversimplified models. Extensive experience in enzyme kinetics tells us that our knowledge of the system (and of non-linear systems in general) is simply insufficient to predict such effects. We may be glad if we can understand certain phenomena *a posteriori*.

An example is the much-debated issue of inhibition of Factor Xa. The original idea was that it was more efficient to inhibit the prothrombin-converting enzyme (Factor Xa) than the product (thrombin) [151,152]. This seemed a very logical idea until it was fully recognised that, under physiological circumstances, Factor Xa requires thrombin-activated Factor V to increase its turnover number [154], and needs adsorption onto procoagulant phospholipid to decrease its  $K_m$  below the plasma concentration of prothrombin [153]. The amount of prothrombin-converting complex that can be maximally formed is not higher than the scarcest component. In PRP, with adequate amounts of phospholipid added, it is easily seen that the rate-limiting component is not Factor Xa. The reasons are:

- the plasma concentration of Factor X is approx. 120 nM, whilst that of Factor V is approx. 20 nM, which may rise to 40 nM if the platelets release their Factor V [155]
- the serum concentration of Factor X is considerable ('stable factor'), whereas that of Factor V is near to zero ('labile factor')
- decrease of the Factor Va concentration *via* the APC system efficiently limits prothrombin conversion.

In PRP, the rate limiting component is neither Factor Va nor Factor Xa, but the exposure of procoagulant phospholipids by the platelets [120]. Still closer to real life, if the procoagulant phospholipid surface is much larger than the molecules involved, as in the case of platelets sticking to a wounded surface, the reaction is diffusion-limited and therefore no longer dependent even upon the concentration of the prothrombinase complex on the surface. Only the concentration of prothrombin in the plasma counts [156-158]. Therefore, there are many arguments to show that, in clotting plasma, Factor Xa is available in excess and has to be inhibited by 90% in order to inhibit prothrombinase activity by 50%.

Still, it is often argued that one can expect from direct Factor Xa inhibition an efficacy:safety ratio superior to that achieved by inhibiting thrombin once it is formed. There is no rational theoretical basis for this statement as soon as one realises the importance of thrombin-mediated feedback reactions, and the excess of Factor Xa present in clotting plasma. This does not mean to say that Factor Xa inhibition may not be the mechanism *via* which good antithrombotic efficiency can be obtained. If it is pushed far enough it may certainly cause inhibition of prothrombin conversion. It is only meant as an illustration that predictions on the basis of available theory are precarious. At a previous state of knowledge, one would expect a benefit of Factor Xa inhibition. At present, one would rather predict a narrow therapeutic margin (*viz.* that between 90 - 100% inhibition of Factor X). It is the very essence of this article to show that even sophisticated theoretical arguments cannot predict the *in vivo* behaviour of an antithrombotic, but that a physiological test of the overall function of the HTS is needed instead. Nevertheless, it is a rewarding operation to refute theoretical considerations that are no longer up-to-date, but that tend to survive in pharmaceutical circles.

The proof of the pudding will remain in the eating. It has been shown that thrombin generation can be effectively inhibited by the injection of the pentasaccharide with anti-Factor Xa

activity [159]. This pentasaccharide (Sanofi [SP/90107]; Organon [Org31540]) is now in Phase II clinical trials. DX9065a (Daiichi) is in Phase I clinical trial and TFPI SC59735 is in Phase II. If there are any specific advantages of Factor Xa inhibition they will be seen in such studies.

## 7. Potential development issues

### 7.1 Catalysts of physiological inhibition

#### 7.1.1 Purer preparations of the active fractions within natural heparins

Heparins, as they are presently obtained from natural sources, are highly heterogeneous [14]. Apart from the anticoagulant molecules, there is 50 - 80% of other material that is not necessarily without biological action. Also, it has been postulated on good grounds that material with a molecular weight lower than 5400 is much less active than higher molecular weight material [160-162], whereas material larger than about 9000 has an extremely low bioavailability [16]. In addition, it might be more haemorrhagic than lower molecular weight material [18].

#### 7.1.2 Synthetic heparins

It has been an important achievement of Petitou (within the Choay group) to synthesise the pentasaccharide that is responsible for high affinity AT binding in natural heparins. Later research in the Sanofi and Organon teams has led to variants with higher activity [163,164]. These materials have anti-Xa activity only. High doses can be given, however, so that thrombin generation can be inhibited in this way. In view of the much greater efficiency of AT action than of anti-Xa activity, it is to be expected that heparins in which a long inert glycosaminoglycan chain is attached to the active pentasaccharide would make a pure and very active heparin.

#### 7.1.3 Activators of heparin co-Factor II

Dermatan sulfate and pentosan polyphosphate are long since known as antithrombotics [20,164,165]. These drugs have the drawback that concentrations above the micromolar range have to be administered, which at a molecular weight of 10,000 to 100,000 requires administration of large amounts of therapeutic material. The development of pure, low molecular weight material should be considered. There is only 1  $\mu\text{M}$  HCII in plasma. This is lower than the level of prothrombin (2  $\mu\text{M}$ ). Optimal activation of all HCII will not exhaust the thrombin-forming capacity of the plasma. Unlike heparin, which acts on 2.4  $\mu\text{M}$  AT, activators of HCII cannot, therefore, be overdosed to cause bleeding.

A special class of substances, the thioxylosides, is capable of altering synthetic reactions in the body of glycosaminoglycans that interact with HCII. Administration of these drugs raises the level of glycosaminoglycans in the plasma, and in this way activate HCII [166,167].

### 7.2 Direct inhibitors

Enormous progress has been made in our understanding of the stereochemistry of clotting enzymes [168]. This, together with modern techniques of organic synthesis and screening of inhibitory activity [34], has provided us with a number of highly specific inhibitors of individual clotting enzymes, and no doubt more are to come. Sanderson and Naylor-Olsen [169] recently published an excellent review on thrombin inhibitor design, to which we

refer you for details. The hope is that from this line of research a compound will be identified that can be administered orally, and that has pharmacokinetic properties which allow standard dosage without control. Evidently, one of the problems is the large amount of serine proteases present in the intestinal tract.

It will be clear from the above discussion that, in our opinion, there is no predictable relation between the biochemical mode of action of these inhibitors and the pharmacological efficiency. Elg *et al.* compared the kinetic constants of eight different direct thrombin inhibitors and saw that, despite a 50,000-fold difference in  $K_i$  value, hirudin and melagatran had similar antithrombotic effects at around 0.14  $\mu\text{M}$  plasma concentration [129]. These authors did report a relation between  $k_{\text{on}}$  and the slope of the dose-response curve of thrombus inhibition. In theory, a steeper slope indicates a narrower therapeutic interval. To judge inhibitors on this criterion does not seem realistic, however, as long as we are not certain that the pharmacokinetics allow maintenance of stable plasma levels and, more importantly, as long as the dose-response of haemorrhage is not known.

### 7.3 Platelet inhibitors

At this moment, there are no inhibitors in clinical use that attack *via* GP Ib/IX, the receptor absent in Bernard-Soulier's disease. Antibodies against this inhibitor also strongly (~50%) inhibit thrombin generation. A combination of antibodies against both the GP Ib and the GP IIb/IIIa/IIIa receptors will almost completely abolish thrombin generation in PRP. vWF has been shown to be a necessary co-factor for both receptors; consequently, in the absence of this factor thrombin generation is almost nil, independent of the availability of Factor VIII. This suggests that interfering with these interactions can be an efficient means of antithrombotic action.

We saw that fibrin can play an active role in the development of platelet procoagulant activity, and hence in thrombin generation and further thrombus growth. It may be interesting to try and find inhibitors of the (vWF-mediated) platelet-fibrin interaction, which might lead to early pacification of a growing thrombus.

## 8. Editorial analysis

In the search for new and better antithrombotics, the open options are as big as the number of vulnerable sites in the HTS (i.e., every enzyme and every receptor involved) multiplied by the number of compounds that can inhibit each site). It is hard to believe that the ideal antithrombotic is not there to be found. The question is how to find it. The present strategy is to first define a subsystem of interest within the HTS; this clotting factor, that membrane receptor. Then a test is set up for that subsystem. Promising compounds are selected on the basis of their performance in that test. Such compounds are then tested in animal thrombosis models and may make it to the phases of investigation in humans.

The problem lies in the choice of the subsystem to be tested. Clotting times and platelet aggregation/adhesion assays are only of limited use because they do not reflect interactions between platelets and the clotting system, recently shown to be extremely important. More specific tests on individual clotting factors or platelet functions are even less fit for the task. The risk is that we find inhibitors of an irrelevant test, and pass by good antithrombotics that do not inhibit this test. The example of LMWHs is a good illustration. These drugs were developed upon the criterion of Factor Xa inhibition, which later was shown to be of no relevance. Later research showed that the performance of these drugs is due to

thrombin-inhibition together with high bioavailability and, probably, absence of large molecular weight molecules that induce bleeding.

Therefore, there is an obvious need for a representative, functional test of the HTS. This test should respond quantitatively in the same manner to any antithrombotic given at a dosage within its therapeutic window. It should, for example, be equally sensitive to doses of antiplatelet drugs and anticoagulants of comparable efficacy.

There are two tests in which the interaction between platelets and plasma is not artificially disrupted: thrombin generation in PRP (or whole blood), and flow chamber experiments in which non-anticoagulated human blood is drawn over a thrombogenic surface. Both tests require much development. In our opinion, progress in technical development of this type of test would be a major step forward in the search for the ideal anticoagulant.

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