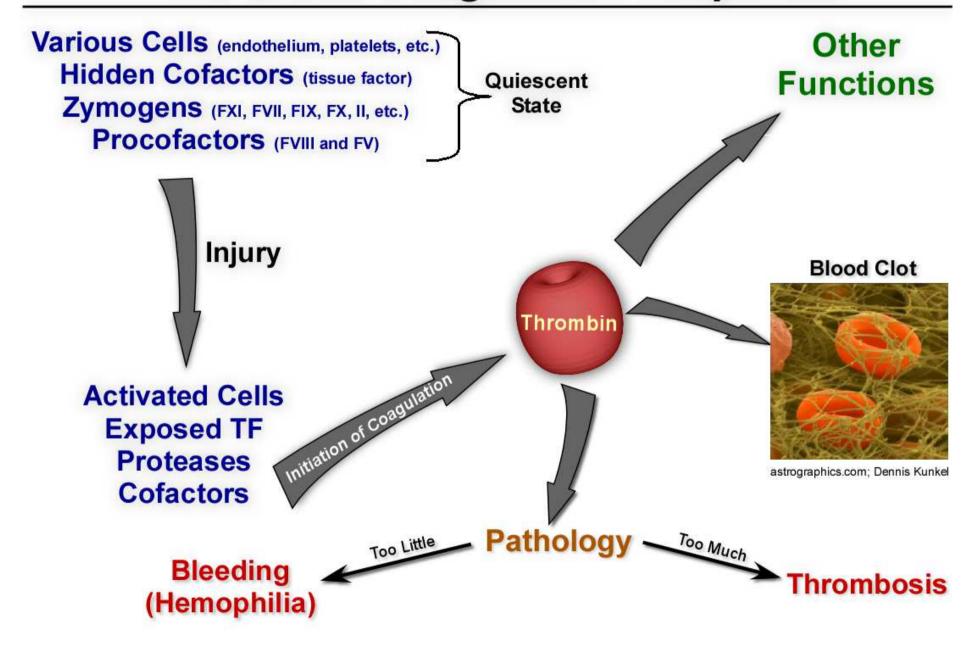
Activation and specificity of Thrombin

The Blood Coagulation Response:

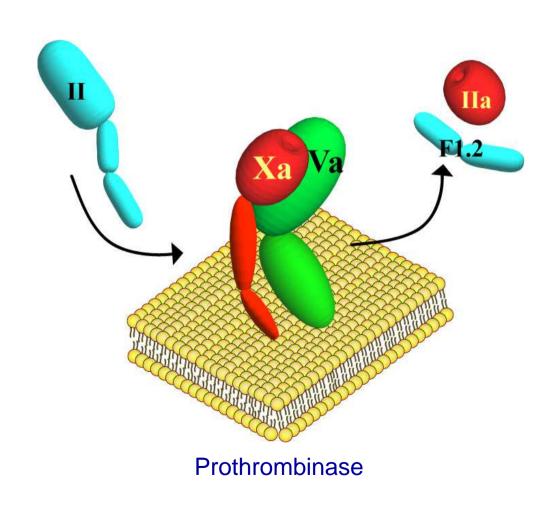


Summary

- Regulation of a Serine Protease: Thrombin
 - Zymogen Enzyme Substrate Specificity
- Staphylocoagulase
 - Bacteria know how a protease works (much more than we do...)
- Diagnostic applications
 - Imaging of Staphylococcus vegetations in the hearth

MECCANISMO ATTIVAZIONE

Prothrombin is activated to thrombin by two proteolytic cleavages

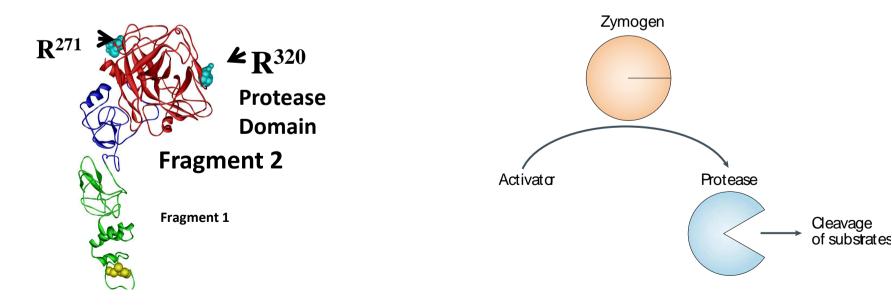


Thrombin is synthesized as a Zymogen: Prothrombin

Zymogen:

A proenzyme or inactive enzyme. It requires a biochemical change to reveal the active site for it to become an active enzyme.

Zymogens lack the structural attributes required for formation of the enzyme-substrate complex.



Serine Proteases: Conversion Pathway

- Cleavage between Arg^{15} - $Ile^{16} \rightarrow Exposure of new N-terminus$
- New N-terminus (IVGG) forms salt bridge with Asp¹⁹⁴
- N-terminal insertion leads to a conformational change in the "activation domain"

Active site (184-194)**N-terminus** (16-19)

Courtesy of W. Bode, Max Planck Institute of Biochemistry

SPECIFICITA SUBSTRATO

Trypsin and Thrombin have similar structures

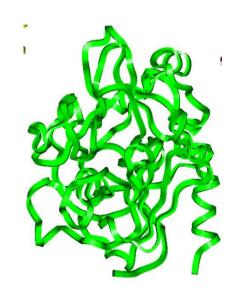
Trypsin

 Cleaves peptides on the Cterm of Lys and Arg amino acid residues

Thrombin

 Cleaves peptides at Arg (Pro, Arg, Ser/Ala/Gly/Thr, not acidic, Arg)





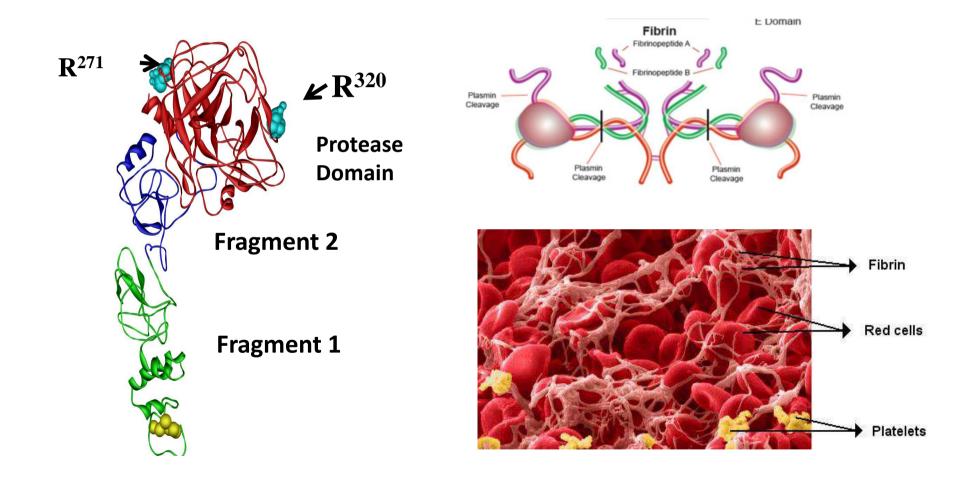
Thrombin cleaves different substrates

Thrombin cleaves after Arg residues

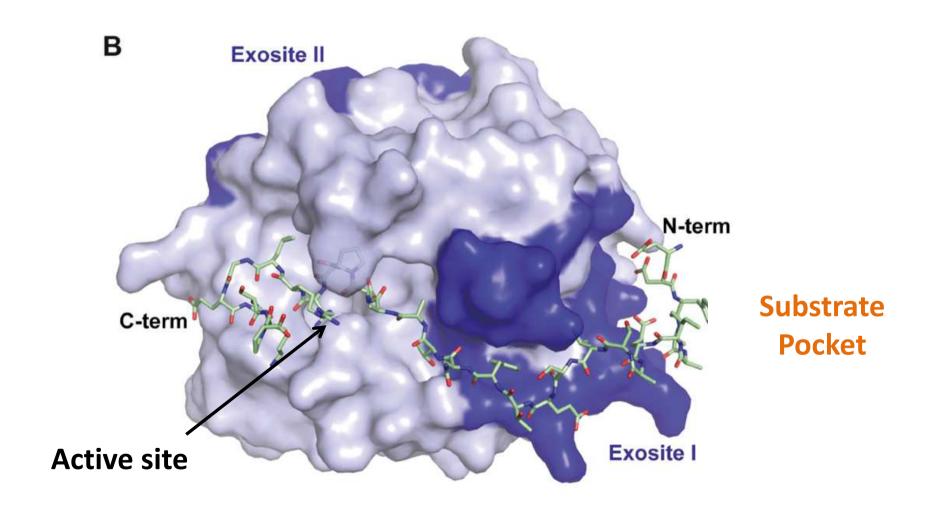
Cleavage Sites for Natural Thrombin Substrates

	P4	P3	P2	P1	P1 [′]	P2 [′]	P3 [′]
Fibrinogen (A)	Gly	Gly	Val	Arg	Gly	Pro	Arg
Fibrinogen (B)	Phe	Ser	Ala	Arg	Gly	His	Arg
FV (709)	Leu	Gly	Ile	Arg	Ser	Phe	Arg
FV (1018)	Leu	Ser	Pro	Arg	Thr	Phe	His
FV (1545)	Trp	Tyr	Leu	Arg	Ser	Asn	Asn
FVIII (372)	Ile	Gln	Ile	Arg	Ser	Val	Ala
FVIII (740)	Ile	Glu	Pro	Arg	Ser	Phe	Ser
FVIII (1689)	Gln	Ser	Pro	Arg	Ser	Phe	Gln
FXIII	Gly	Val	Pro	Arg	Gly	Val	Asn
PAR1	Leu	Asp	Pro	Arg	Ser	Phe	Leu
PAR4	Pro	Ala	Pro	Arg	Gly	Tyr	Pro
FXI	Ile	Lys	Pro	Arg	Ile	Val	Gly
PC	Val	Asp	Pro	Arg	Leu	Ile	Asp
ΓAFI	Val	Ser	Pro	Arg	Ala	Ser	Ala
AT	Ile	Ala	Gly	Arg	Ser	Leu	Asn

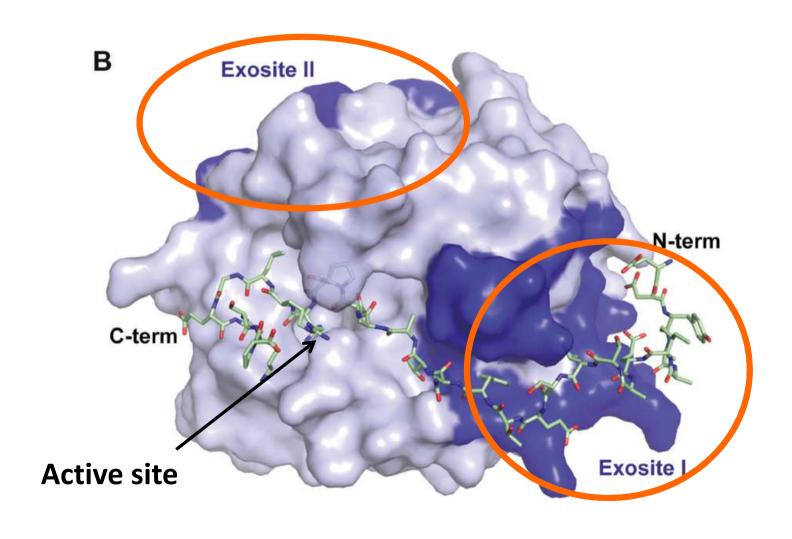
Thrombin cleavage of the plasma protein fibrinogen



Thrombin X-ray structure



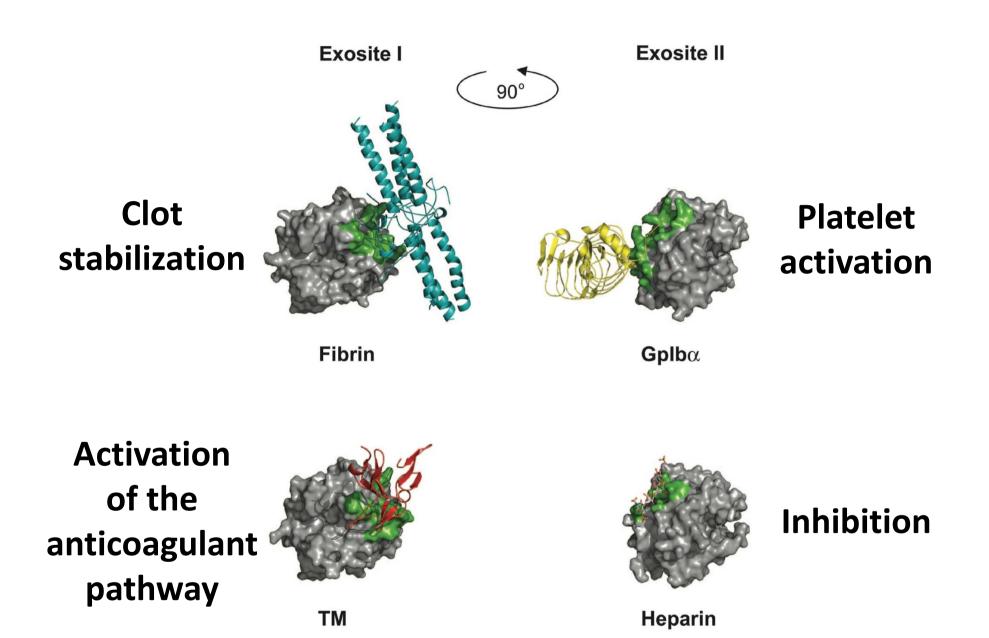
Thrombin X-ray structure



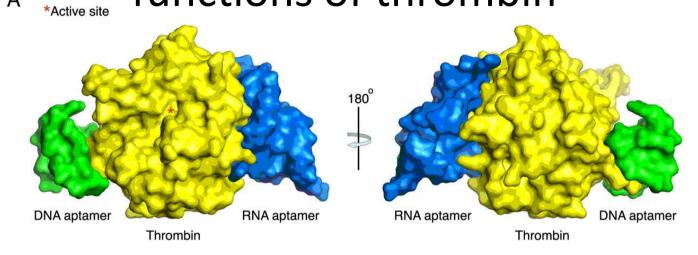
Exosite binding determines substrate specificity

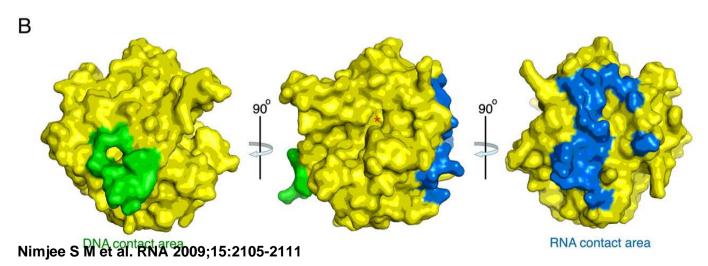
 Thrombin targets are restricted due to specific interactions between the protein substrate and residues outside the catalytic cleft termed Exosite

 Extended interactions at exosites drive substrate affinity and contribute to substrate specificity.



Exosites are good targets to inhibit specific functions of thrombin







be briefly stated as follows: The staphylococcus pyogenes aureus has a specific influence in causing coagulation of the blood. Bouillon cultures of the staphylococcus were much more potent than any one of the other organisms. The

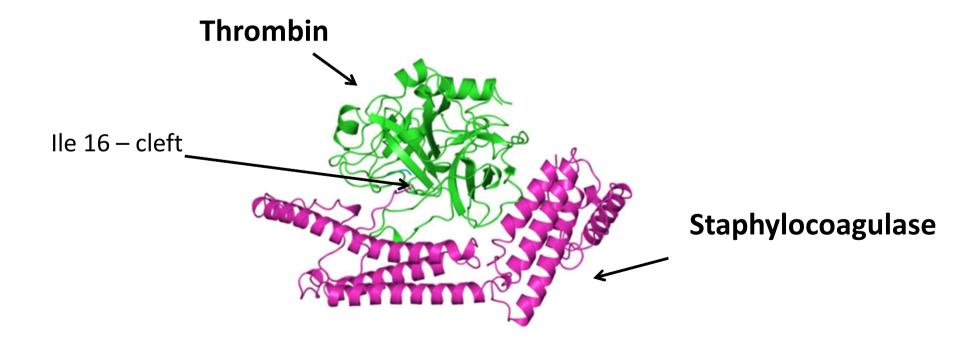
- Certain strains of Staphylococcus Aureus trigger coagulation (1903)
- Isolation of a bacterial agent that specifically activates thrombin: Staphylocoagulase (1970)
- SC does not cleave thrombin, No cleavage between Arg¹⁵Ile¹⁶

How is that possible???

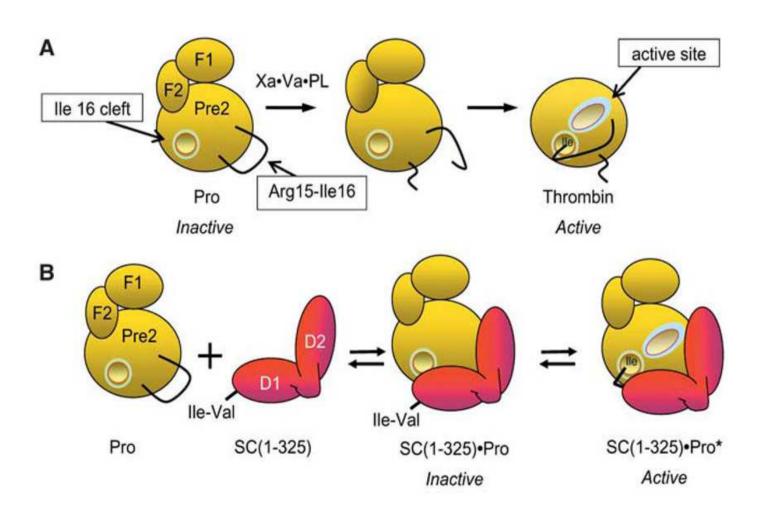
MECCANISMO ATTIVAZIONE BATTERICA

Staphylocoagulase (SC) X ray-structure

In 2003 crystal structure of (Pre2)Thrombinbound Staphylocoagulase was published (Friedrich, et al. *Nature*, 2003)

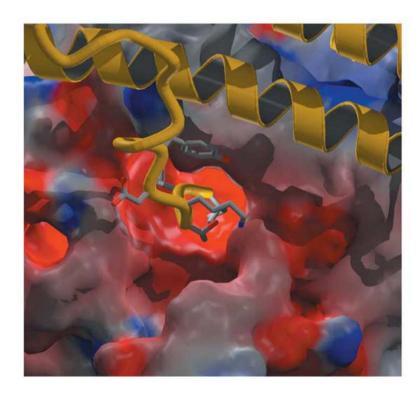


Non-Proteolytic Activation of Prothrombin by Staphylocoagulase support for the "Molecular Sexuality" Hypothesis



The observed insertion of the SC N-terminus into the Ile¹⁶ cleft of prethrombin 2, which triggers the activating conformational change, provided the first unambiguous structural evidence for the Molecular Sexuality mechanism of non-proteolytic zymogen activation.



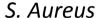


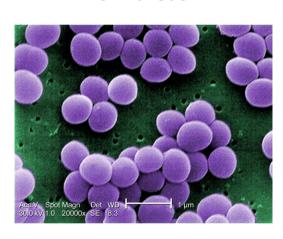
S. Aureus causes Endocarditis

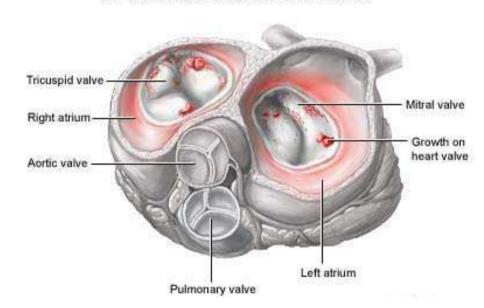
- Severe infection of the heart valves
- More than 50% of patients dies within days or weeks despite treatment
- Difficult diagnosis
 - new heart murmur, fever and the detection of circulating bacteria in blood cultures
- Coagulase-positive S. aureus causes 40–50% of neonatal endocarditis and 30–40% of endocarditis in adults

Acute bacterial endocarditis is characterized by vegetations on heart valves consisting of bacteria, platelets and fibrin

Infective endocarditis is an infection of the heart chambers or valves







- Growth and fortification of the vegetation by SC-induced fibrin deposition protects the bacteria in the vegetation from clearance by leukocytes and macrophages
- Heart valves are not easily accessible to the immune system

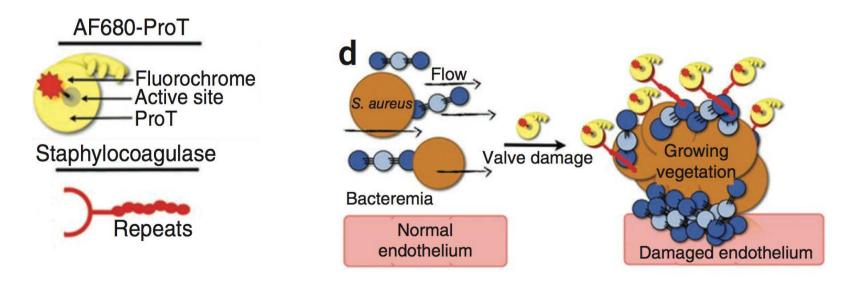


In vivo detection of Staphylococcus aureus endocarditis by targeting pathogen-specific prothrombin activation

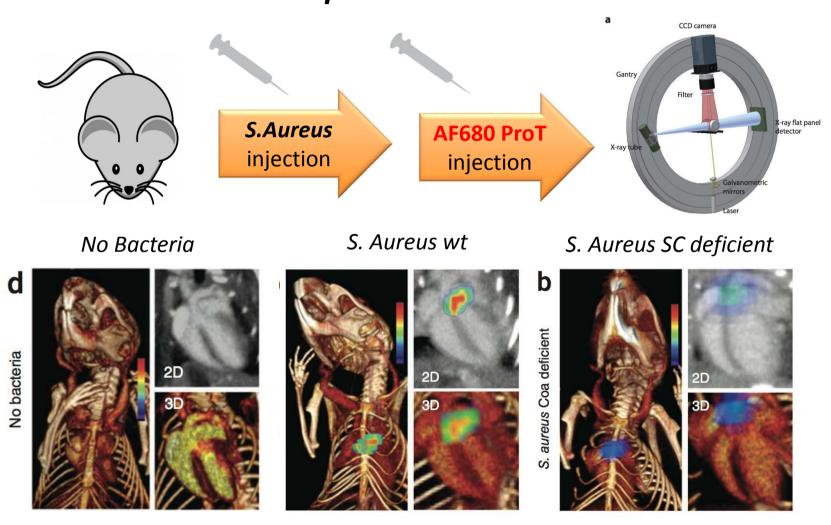
Peter Panizzi^{1,2,9}, Matthias Nahrendorf^{1,9}, Jose-Luiz Figueiredo¹, Jennifer Panizzi³, Brett Marinelli¹, Yoshiko Iwamoto¹, Edmund Keliher¹, Ashoka A Maddur⁴, Peter Waterman¹, Heather K Kroh⁴, Florian Leuschner¹, Elena Aikawa¹, Filip K Swirski¹, Mikael J Pittet¹, Tilman M Hackeng⁵, Pablo Fuentes-Prior⁶, Olaf Schneewind⁷, Paul E Bock⁴ & Ralph Weissleder^{1,8}

SC Prothrombin as a probe for *S. Aureus*

- SC binds prothrombin with high affinity and activates it through a conformation change
- SC-Prothrombin complex clots fibrinogen but is impervious to physiologic thrombin inhibitors.
- SC-Prothrombin is present in the vegetation
- Labeled Prothrombin can be used as a probe to detect bacterial vegetation in the heart



Fluorescent prothrombin co-localise with SC positive bacteria



Conclusion

- Zymogen activation requires conformational changes and maturation of the active site. This can be achieved even in the absence of canonical proteolysis.
- Exosite-Substrate interactions determine enzyme specificity.
- AF680ProT detects S.Aureus in vivo and can be used as a diagnostic tool to determine site, bacterial load and activity of the infection.

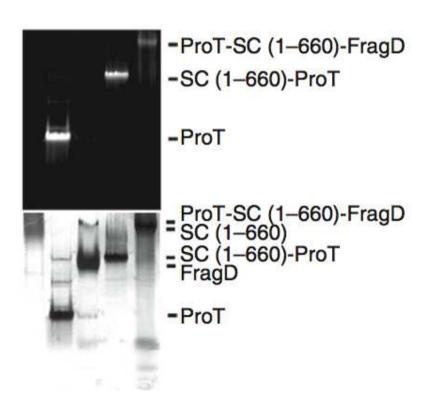
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Visualisation of *S. Aureus* in vivo using Near Infrared Imaging

The PROBE

AF680- Prothrombin



The DETECTOR

Fluorescence molecular tomography

- Computer Tomography

