

**A quantitative systems pharmacology model of plasma kallikrein-kinin system
dysregulation in hereditary angioedema**

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List of abbreviations

BDKR-B2	B2 receptor of bradykinin receptor family
BK	Bradykinin
C1-INH	C1-esterase inhibitor
C1q	Complement component 1
gC1q-R	Cofactors complement protein C1q
CK1	Cytokeratin 1
cHMWK	Cleaved high molecular weight kininogen
% cHMWK	Percentage of cleaved high molecular weight kininogen relative to the total of cleaved high molecular weight kininogen and high molecular weight kininogen
FXII	Factor XII
FXIIa	Activated factor XII
HAE	Hereditary angioedema
HMWK	High molecular weight kininogen
KAL	Kallikrein
k_{cat}	Turnover number
K_d	Binding affinity
KKS	Kallikrein-kinin system
K_m	Michaelis-Menten constant
K_{on}	Association constant
K_{off}	Dissociation constant
K_{syn}	Synthesis rate
PD	Pharmacodynamics
PK	Pharmacokinetics
PRCP	Prolylcarboxypeptidase
preKAL	Prekallikrein
Q2W	Every 2 weeks
Q4W	Every 4 weeks

QSP	Quantitative systems pharmacology
SC	Subcutaneous
uPAR	Urokinase plasminogen activating receptor
V_{medium}	Per endothelial cell-based vascular volume
V_{proximal}	Per endothelial cell-based proximal space volume

Method S1. Detailed description of model components

Factor XII (FXII) is an 80 kDa glycosylated protein consisting of a single polypeptide chain and circulates in plasma as a zymogen with a concentration of 30 µg/mL (375 nM) in healthy individuals. Upon contact with anionic surfaces, and in the presence of Zn^{2+} ions, FXII undergoes a conformational rearrangement leading to autoactivation or cleavage by kallikrein to generate FXIIa (the activated form of FXII).

Prekallikrein is an 86 kDa glycoprotein consisting of a single polypeptide chain that circulates in plasma as a zymogen at a median concentration of 31 µg/mL (365 nM) in healthy individuals [1], it is estimated that 75% of prekallikrein is bound to high molecular weight kininogen (HMWK) [2]. Prekallikrein binds to endothelial cells, platelets, and granulocytes in a Zn^{2+} -dependent interaction via the prekallikrein–HMWK complex [3]. It is cleaved by FXIIa to form kallikrein, the two-chain enzyme kallikrein. Prolylcarboxypeptidase (PRCP) has been identified as an endothelial cell activator of prekallikrein to kallikrein [4].

HMWK is a 120 kDa non-enzymatic glycoprotein with a plasma concentration of 80 µg/mL (670 nM) in healthy individuals. HMWK circulates in plasma both in a free or complexed form (with prekallikrein or kallikrein). The binding affinities of HMWK to prekallikrein and kallikrein are similar (K_d of 12 nM and 15 nM, respectively [5]).

Assembly of the kinin-kallikrein contact factor proteins on cell surfaces is mediated via urokinase plasminogen activating receptor (uPAR), and cofactors complement protein C1q (gC1q-R) and cytokeratin 1 (CK1). On the surface of endothelial cells, gC1q-R (with elevated levels of Zn^{+2} ions released from endothelial cells and activated platelets) is primarily responsible for assembly and activation of the FXII-HMWK-prekallikrein complex, as previously demonstrated using surface plasmon resonance measurements showing that HMWK

has the greatest affinity to gC1q-R (0.8 nM), followed by CK1 (15 nM) and then uPAR (2.3 μ M) [6]. Other studies suggested that FXII and HMWK compete with each other for binding to gC1q-R [7]. gC1q-R forms a multiprotein complex with uPAR and CK1, and the complex can bind to FXII [8]. In addition, the complex of uPAR with CK1 was shown to bind HMWK, resulting in prekallikrein activation at the endothelial cell surface [9].

The number of receptors, cofactors, and their complexes on endothelial cells has been reported. Among them, gC1q-R is the most abundant with over 1 million/cell [10], while uPAR (250,000/cell [11]) and CK1 (72,000/cell [9]) are expressed at lower levels. As the gC1q-R/CK1 complex preferentially binds HMWK, and FXII binds primarily to uPAR within the CK1-uPAR complex [12], it was assumed that CK1, with its lower expression, is the limiting factor for formation of the receptor complex in the activation of the surface KKS. The apparent affinity between FXII and HMWK to binding sites on human umbilical vein endothelial cells was reported to be 144 nM and 7–52 nM, respectively [7, 6].

Excessive bradykinin causes an increase in blood vessel permeability, which allows fluid to pass through the blood vessel walls, causing subcutaneous or submucosal swelling. The cleavage of HMWK by kallikrein produces a two-chain cleaved HMWK (cHMWK) and the bradykinin peptide. Bradykinin has a short half-life (less than 30 seconds in the blood of most species [13]) and strong affinity for the cell surface bradykinin B2 receptor (0.5 nM [14]). These properties make it challenging to obtain reliable measurements of bradykinin levels. However, cHMWK has been readily measured in clinical trials and serves as a valuable biomarker. Based on a saturation analysis of bradykinin binding to different human and rat bradykinin B2 receptors [14], we estimated the receptor number to be approximately 100,000/cell.

Table S1. Summary of clinical trials from which data were incorporated into the QSP model

Therapeutic	Phase	Study number	Title	Participants	Dosing	Reference
Lanadelumab	1a	NCT01923207	A Single Increasing Dose Study to Assess Safety and Tolerability of DX-2930 in Healthy Subjects	N = 32 healthy subjects	Single dose of placebo, 0.1, 0.3, 1.0, or 3.0 mg/kg lanadelumab	Chyung et al [15]
Lanadelumab	1b	NCT02093923	A Double-Blind, Multiple Ascending Dose Study to Assess Safety, Tolerability and Pharmacokinetics of DX-2930 in Hereditary Angioedema Participants	N = 37 patients with HAE type	Placebo or two doses of 30, 100, 300, 400 mg lanadelumab administered 14 days apart	Banerji et al [16]
Lanadelumab	3	NCT02586805	Efficacy and Safety Study of DX-2930 to Prevent Acute Angioedema Attacks in Patients With Type I and Type II HAE (HELP)	N = 125 patients with HAE type	Placebo or 150 mg Q4W, 300 mg Q4W, 300 mg Q2W lanadelumab for 26 weeks	Banerji et al [17]
Lanadelumab	3	NCT02741596	Long-term Safety and Efficacy Study of DX-2930 (SHP643) to Prevent Acute Angioedema Attacks in Patients With Type I and Type II HAE (HELP OLE)	N = 212 patients with HAE type	300 mg Q2W lanadelumab for up to 132 weeks	Banerji et al [18]
Cinryze®	3	NCT00289211	A Double-blind, Placebo-Controlled, Clinical Study to Investigate the Efficacy and Safety of Purified C1 Esterase Inhibitor (Human) for the Treatment of HAE in Acute Attacks	N = 68 patients with HAE	Placebo or 1000 U C1-INH (up to two injections)	Zuraw et al [19]
Cinryze®	3	NCT01005888	A Double-blind, Placebo-Controlled, Clinical Study to Investigate the Efficacy and Safety of Purified C1 Esterase Inhibitor (Human) as Prophylactic Treatment to Prevent HAE Attacks	N = 22 patients with HAE	Placebo or 1000 U C1-INH twice-weekly for 12 weeks	Zuraw et al [19]
Cinryze®	3	NCT00438815	Open-Label Safety/Efficacy Repeat Exposure Study of C1INH-nf (Human) in the Treatment of Acute HAE Attacks (CHANGE 2)	N = 113 patients with HAE	1000 U C1-INH as needed	Riedl et al [20]

Cinryze®	3	NCT00462709	Open-Label Use of C1INH-nf (Human) for the Prophylactic Treatment to Prevent HAE Attacks and as Treatment in Acute HAE Attacks (CHANGE 3)	N = 146 patients with HAE	1000 U C1-INH every 3 to 7 days	Shire [21]
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C1-INH C1 esterase inhibitor, *HAE* hereditary angioedema, *Q2W* every 2 weeks, *Q4W* every 4 weeks, *QSP* quantitative systems pharmacology

Table S2 List of model equations used to describe *ex vivo* KKS activation in the fluorogenic assay

$\frac{d(\text{preKAL})}{dt} = -\frac{k_{cat} * \text{FXIIa} * \text{preKAL}}{K_m + \text{preKAL}}$
$\frac{d(\text{KAL})}{dt} = \frac{k_{cat} * \text{FXIIa} * \text{preKAL}}{K_m + \text{preKAL}} - (\text{kon}_{\text{KAL}-\text{C1Inh}} * \text{KAL} * \text{C1Inh} - \text{koff}_{\text{KAL}-\text{C1Inh}} * \text{KAL}_{\text{C1Inh}}) - (\text{kon}_{\text{KAL}-\text{Lanadelumab}} * \text{KAL} * \text{Lanadelumab} - \text{koff}_{\text{KAL}-\text{Lanadelumab}} * \text{KAL}_{\text{Lanadelumab}})$
$\frac{d(\text{C1Inh})}{dt} = -(\text{kon}_{\text{KAL}-\text{C1Inh}} * \text{KAL} * \text{C1Inh} - \text{koff}_{\text{KAL}-\text{C1Inh}} * \text{KAL}_{\text{C1Inh}})$
$\frac{d(\text{KAL}_{\text{C1Inh}})}{dt} = (\text{kon}_{\text{KAL}-\text{C1Inh}} * \text{KAL} * \text{C1Inh} - \text{koff}_{\text{KAL}-\text{C1Inh}} * \text{KAL}_{\text{C1Inh}})$
$\frac{d(\text{Lanadelumab})}{dt} = -(\text{kon}_{\text{KAL}-\text{Lanadelumab}} * \text{KAL} * \text{Lanadelumab} - \text{koff}_{\text{KAL}-\text{Lanadelumab}} * \text{KAL}_{\text{Lanadelumab}})$
$\frac{d(\text{KAL}_{\text{Lanadelumab}})}{dt} = (\text{kon}_{\text{KAL}-\text{Lanadelumab}} * \text{KAL} * \text{Lanadelumab} - \text{koff}_{\text{KAL}-\text{Lanadelumab}} * \text{KAL}_{\text{Lanadelumab}})$

CI-INH: C1 esterase inhibitor, *FXII* factor XII, *KKS* kallikrein-kinin system, *KAL* kallikrein, k_{cat} turnover number, K_m Michaelis-

Menten constant, K_{off} dissociation constant, K_{on} association constant, *preKAL* prekallikrein

Table S3 Inputs and parameters used in the fluorogenic assay model

Input/Parameter	Explanation	Unit	Value used in model	Reference
preKAL	Basal prekallikrein level in plasma	nM	250–650	Madsen et al [1]
KAL	Basal kallikrein level in plasma	nM	0	Assumed negligible
C1Inh_normal	Basal plasma C1-INH level for normal population	nM	2400	Weidmann et al [22]
C1Inh_HAE	Basal plasma C1-INH level for HAE type 1 population	nM	600	Cicardi et al [23]
Kd_Lana_KAL	Binding affinity of lanadelumab to kallikrein	nM	0.12	Kenniston et al [24]
Kd_C1Inh_KAL	Binding affinity of C1-INH to kallikrein	nM	150	Harpel et al [25]
K _m	K _m for activation of prekallikrein by FXIIa	uM	1.8	Tankersley and Finlayson [26]
k _{cat}	k _{cat} for activation of prekallikrein by FXIIa	1/s	1.03	Tankersley and Finlayson [26]

C1-INH C1 esterase inhibitor, *FXIIa* activated factor XII, *HAE* hereditary angioedema, *KAL*

kallikrein, *k_{cat}* turnover number, *K_m* Michaelis-Menten constant, *preKAL* prekallikrein

Table S4 List of molecular species in the KKS model

Variable names used in model code	Unit	Description
In vascular space		
BK_degraded	nM	Degraded bradykinin concentration
BK_in_plasma	nM	Bradykinin concentration
C1Inh_degraded	nM	Degraded C1-INH concentration
C1Inh_FXIIa_degraded	nM	Degraded C1-INH_FXIIa complex concentration
C1Inh_FXIIa_in_plasma	nM	C1-INH_FXIIa concentration
C1Inh_in_plasma	nM	C1-INH concentration
C1Inh_KAL_degraded	nM	Degraded C1-INH_kallikrein complex concentration
C1Inh_KAL_HMWK_degraded	nM	Degraded C1-INH_kallikrein_HMWK complex concentration
C1Inh_KAL_HMWK_in_plasma	nM	C1-INH_kallikrein_HMWK complex concentration
C1Inh_KAL_in_plasma	nM	C1-INH_kallikrein complex concentration
FXII_degraded	nM	Degraded FXII concentration
FXII_in_plasma	nM	FXII concentration
FXIIa_degraded	nM	Degraded FXIIa concentration
FXIIa_in_plasma	nM	FXIIa concentration
HK2Chain_degraded	nM	Degraded cHMWK concentration
HK2Chain_in_plasma	nM	cHMWK concentration
HMWK_degraded	nM	Degraded HMWK concentration
HMWK_in_plasma	nM	HMWK concentration
KAL_degraded	nM	Degraded kallikrein concentration
KAL_HK2Chain_in_plasma	nM	kallikrein_cHMWK complex concentration

KAL_HMWK_in_plasma	nM	kallikrein_HMWK concentration
KAL_in_plasma	nM	kallikrein concentration
Lanadelumab_in_plasma	nM	Lanadelumab concentration
Lanadelumab_KAL_HMWK_in_plasma	nM	Lanadelumab_kallikrein_HMWK complex concentration
Lanadelumab_KAL_in_plasma	nM	Lanadelumab_kallikrein complex concentration
preKAL_degraded	nM	Degraded prekallikrein concentration
preKAL_HMWK_in_plasma	nM	prekallikrein_HMWK complex concentration
preKAL_in_plasma	nM	prekallikrein concentration
In proximal space		
BK	nM	Bradykinin concentration
BDKRB2	number/cell	Number of bradykinin B2 receptors
BDKRB2_degraded	number/cell	Number of degraded surface bradykinin B2 receptors
BK_BDKRB2	number/cell	Number of bradykinin_bradykinin B2 receptor complexes
BK_BDKRB2_degraded	number/cell	Number of degraded surface bradykinin_bradykinin B2 receptor complexes
C1Inh	nM	C1-INH concentration
C1Inh_FXIIa	nM	C1-INH_FXIIa concentration
C1Inh_FXIIa_gC1qR	number/cell	Number of C1-INH_FXIIa_gC1q receptor complexes
C1Inh_FXIIa_gC1qR_degraded	number/cell	Number of degraded surface C1-INH_FXIIa_gC1q receptor complexes
C1Inh_KAL_HMWK	nM	C1-INH_kallikrein_HMWK concentration

C1Inh_KAL_HMWK_gC1qR	number/cell	Number of C1-INH_kallikrein_HMWK_gC1q receptor complexes
C1Inh_KAL_HMWK_gC1qR_degraded	number/cell	Number of degraded surface C1-INH_kallikrein_HMWK_gC1q receptor complexes
FXII	nM	FXII concentration
FXII_gC1qR	number/cell	Number of FXII_gC1q receptor complexes
FXII_gC1qR_degraded	number/cell	Number of degraded surface FXII_gC1q receptor complexes
FXIIa	nM	FXIIa concentration
FXIIa_gC1qR	number/cell	Number of FXIIa_gC1q receptor complexes
FXIIa_gC1qR_degraded	number/cell	Number of degraded surface FXIIa_gC1q receptor complexes
gC1qR	number/cell	Number of gC1q receptors
gC1qR_degraded	number/cell	Number of degraded surface gC1q receptors
KAL_HK2Chain	nM	kallikrein_cHMWK concentration
KAL_HK2Chain_gC1qR	number/cell	Number of kallikrein_cHMWK_gC1q receptor complexes
KAL_HK2Chain_gC1qR_degraded	number/cell	Number of degraded surface kallikrein_cHMWK_gC1q receptor complexes
KAL_HMWK	nM	kallikrein_HMWK concentration
KAL_HMWK_gC1qR	number/cell	Number of kallikrein_HMWK_gC1q receptor complexes
KAL_HMWK_gC1qR_degraded	number/cell	Number of degraded surface kallikrein_HMWK_gC1q receptor complexes
Lanadelumab	nM	Lanadelumab concentration

Lanadelumab_KAL_HMWK	nM	Lanadelumab_kallikrein_HMWK concentration
Lanadelumab_KAL_HMWK_gC1qR	number/cell	Number of Lanadelumab_kallikreinL_HMWK_gC1q receptor complexes
Lanadelumab_KAL_HMWK_gC1qR_degraded	number/cell	Number of degraded surface Lanadelumab_kallikrein_HMWK_gC1q receptor complexes
preKAL_HMWK	nM	prekallikrein_HMWK concentration
preKAL_HMWK_gC1qR	number/cell	Number of prekallikrein_HMWK_gC1q receptor complexes
preKAL_HMWK_gC1qR_degraded	number/cell	Number of degraded surface prekallikrein_HMWK_gC1q receptor complexes

BDKR-B2 B2 receptor of bradykinin receptor family, *BK* bradykinin, *C1-INH* C1 esterase inhibitor, *C1q* complement component 1, *cHMWK* cleaved high molecular weight kininogen, *FXII* factor XII, *FXIIa* activated factor XII, *gC1q-R* cofactors complement protein C1q, *HK2* human kallikrein 2, *HMWK* high molecular weight kininogen, *KAL* kallikrein, *KKS* kallikrein-kinin system, *preKAL* prekallikrein

Table S5 List of governing mathematical equations of the KKS model

In vascular space	Equation number
$V_{medium} \frac{d\mathbf{BK_degraded}}{dt} = V_{medium} \cdot kdeg_{BK} \cdot \mathbf{BK_in_plasma}$	(E1)
$V_{medium} \frac{d\mathbf{BK_in_plasma}}{dt}$ $= -V_{medium} \cdot kdeg_{BK} \cdot \mathbf{BK_in_plasma} - (k12_{BK} \cdot \mathbf{BK_in_plasma} \cdot V_{medium}$ $- k21_{BK} \cdot \mathbf{BK} \cdot V_{proximal})$	(E2)
$V_{medium} \frac{d\mathbf{C1Inh_degraded}}{dt} = V_{medium} \cdot kdeg_{C1Inh} \cdot \mathbf{C1Inh_in_plasma}$	(E3)
$V_{medium} \frac{d\mathbf{C1Inh_FXIIa_degraded}}{dt}$ $= V_{medium} \cdot kdeg_{Bound_{C1Inh}} \cdot \mathbf{C1Inh_FXIIa_in_plasma}$	(E4)
$V_{medium} \frac{d\mathbf{C1Inh_FXIIa_in_plasma}}{dt}$ $= V_{medium} \cdot (kon_{C1Inh_FXIIa} \cdot \mathbf{C1Inh_in_plasma} \cdot \mathbf{FXIIa_in_plasma}$ $- koff_{C1Inh_FXIIa} \cdot \mathbf{C1Inh_FXIIa_in_plasma}) - V_{medium}$ $\cdot kdeg_{Bound_{C1In}} \cdot \mathbf{C1Inh_FXIIa_in_plasma} - (k12$ $\cdot \mathbf{C1Inh_FXIIa_in_plasma} \cdot V_{medium} - k21 \cdot \mathbf{C1Inh_FXIIa} \cdot V_{proximal})$	(E5)
$V_{medium} \frac{d\mathbf{C1Inh_in_plasma}}{dt}$ $= flux_{C1Inh_inj_nmol_per_hr} + V_{medium} \cdot ksyn_{C1Inh} - V_{medium} \cdot kdeg_{C1Inh}$ $\cdot \mathbf{C1Inh_in_plasma} - V_{medium} \cdot (kon_{C1Inh_KAL} \cdot \mathbf{C1Inh_in_plasma}$ $\cdot \mathbf{KAL_in_plasma} - koff_{C1Inh_KAL} \cdot \mathbf{C1Inh_KAL_in_plasma}) - V_{medium}$ $\cdot (kon_{C1Inh_KAL} \cdot \mathbf{C1Inh_in_plasma} \cdot \mathbf{KAL_HMWK_in_plasma}$ $- koff_{C1Inh_KAL} \cdot \mathbf{C1Inh_KAL_HMWK_in_plasma}) - V_{medium}$ $\cdot (kon_{C1Inh_FXIIa} \cdot \mathbf{C1Inh_in_plasma} \cdot \mathbf{FXIIa_in_plasma} - koff_{C1Inh_FXIIa}$ $\cdot \mathbf{C1Inh_FXIIa_in_plasma}) - (k12 \cdot \mathbf{C1Inh_in_plasma} \cdot V_{medium} - k21$ $\cdot \mathbf{C1Inh} \cdot V_{proximal})$	(E6)

$$V_{medium} \frac{dC1Inh_KAL_degraded}{dt} = V_{medium} \cdot kdeg_Bound_{C1Inh} \cdot C1Inh_KAL_in_plasma \quad (E7)$$

$$V_{medium} \frac{dC1Inh_KAL_HMWK_degraded}{dt} = V_{medium} \cdot kdeg_Bound_{C1Inh} \cdot C1Inh_KAL_HMWK_in_plasma \quad (E8)$$

$$V_{medium} \frac{dC1Inh_KAL_HMWK_in_plasma}{dt} = V_{medium} \cdot (kon_{C1Inh_KAL} \cdot C1Inh_in_plasma \cdot KAL_HMWK_in_plasma - koff_{C1Inh_KAL} \cdot C1Inh_KAL_HMWK_in_plasma) - V_{medium} \cdot kdeg_Bound_{C1Inh} \cdot C1Inh_KAL_HMWK_in_plasma - (k12 \cdot C1Inh_KAL_HMWK_in_plasma \cdot V_{medium} - k21 \cdot C1Inh_KAL_HMWK} \cdot V_{proximal}) \quad (E9)$$

$$V_{medium} \frac{dC1Inh_KAL_in_plasma}{dt} = V_{medium} \cdot (kon_{C1Inh_KAL} \cdot C1Inh_in_plasma \cdot KAL_in_plasma - koff_{C1Inh_KAL} \cdot C1Inh_KAL_in_plasma) - V_{medium} \cdot kdeg_Bound_{C1Inh} \cdot C1Inh_KAL_in_plasma \quad (E10)$$

$$V_{medium} \frac{dFXII_degraded}{dt} = V_{medium} \cdot kdeg_{FXII} \cdot FXII_in_plasma \quad (E11)$$

$$V_{medium} \frac{dFXII_in_plasma}{dt} = V_{medium} \cdot ksyn_{FXII} - V_{medium} \cdot kdeg_{FXII} \cdot FXII_in_plasma - (k12 \cdot FXII_in_plasma \cdot V_{medium} - k21 \cdot FXII \cdot V_{proximal}) \quad (E12)$$

$$V_{medium} \frac{dFXIIa_degraded}{dt} = V_{medium} \cdot kdeg_{FXIIa} \cdot FXIIa_in_plasma \quad (E13)$$

$$V_{medium} \frac{dFXIIa_in_plasma}{dt} = -V_{medium} \cdot kdeg_{FXIIa} \cdot FXIIa_in_plasma - V_{medium} \cdot (kon_{C1Inh_FXIIa} \cdot C1Inh_in_plasma \cdot FXIIa_in_plasma - koff_{C1Inh_FXIIa} \cdot C1Inh_FXIIa_in_plasma) - (k12 \cdot FXIIa_in_plasma \cdot V_{medium} - k21 \cdot FXIIa \cdot V_{proximal}) \quad (E14)$$

$$V_{medium} \frac{dHK2Chain_{degraded}}{dt} = V_{medium} \cdot kdeg_{cHMWK} \cdot HK2Chain_{in_plasma} \quad (E15)$$

$$V_{medium} \frac{dHK2Chain_{in_plasma}}{dt} = -V_{medium} \cdot (kon_{KAL_HK2Chain} \cdot KAL_{in_plasma} \cdot HK2Chain_{in_plasma} - koff_{KAL_HK2Chain} \cdot KAL_HK2Chain_{in_plasma}) - V_{medium} \cdot kdeg_{cHMWK} \cdot HK2Chain_{in_plasma} \quad (E16)$$

$$V_{medium} \frac{dHMWK_{degraded}}{dt} = V_{medium} \cdot kdeg_{HMWK} \cdot HMWK_{in_plasma} \quad (E17)$$

$$V_{medium} \frac{dHMWK_{in_plasma}}{dt} = V_{medium} \cdot ksyn_{HMWK} - V_{medium} \cdot kdeg_{HMWK} \cdot HMWK_{in_plasma} - V_{medium} \cdot (kon_{preKAL_HMWK} \cdot preKAL_{in_plasma} \cdot HMWK_{in_plasma} - koff_{preKAL_HMWK} \cdot preKAL_HMWK_{in_plasma}) - V_{medium} \cdot (kon_{KAL_HMWK} \cdot KAL_{in_plasma} \cdot HMWK_{in_plasma} - koff_{KAL_HMWK} \cdot KAL_HMWK_{in_plasma}) \quad (E18)$$

$$V_{medium} \frac{dKAL_{degraded}}{dt} = V_{medium} \cdot kdeg_{KAL} \cdot KAL_{in_plasma} \quad (E19)$$

$$V_{medium} \frac{dKAL_HK2Chain_{in_plasma}}{dt} = -(k12 \cdot KAL_HK2Chain_{in_plasma} \cdot V_{medium} - k21 \cdot KAL_HK2Chain \cdot V_{proximal}) + V_{medium} \cdot (kon_{KAL_HK2Chain} \cdot KAL_{in_plasma} \cdot HK2Chain_{in_plasma} - koff_{KAL_HK2Chain} \cdot KAL_HK2Chain_{in_plasma}) \quad (E20)$$

$$\begin{aligned}
V_{medium} \frac{dKAL_HMWK_in_plasma}{dt} &= V_{medium} \cdot (kon_{KAL_HMWK} \cdot KAL_in_plasma \cdot HMWK_in_plasma \\
&- koff_{KAL_HMWK} \cdot KAL_HMWK_in_plasma) - V_{medium} \cdot (kon_{C1In_KAL} \\
&\cdot C1Inh_in_plasma \cdot KAL_HMWK_in_plasma - koff_{C1Inh_KAL} \\
&\cdot C1Inh_KAL_HMWK_in_plasma) - V_{medium} \cdot (kon_{KAL_Lanadelumab} \\
&\cdot KAL_HMWK_in_plasma \cdot Lanadelumab_in_plasma \\
&- koff_{KAL_Lanadelumab} \cdot Lanadelumab_KAL_HMWK_in_plasma) - (k12 \\
&\cdot KAL_HMWK_in_plasma \cdot V_{medium} - k21 \cdot KAL_HMWK \cdot V_{proximal})
\end{aligned}
\tag{E21}$$

$$\begin{aligned}
V_{medium} \frac{dKAL_in_plasma}{dt} &= -V_{medium} \cdot kdeg_{KAL} \cdot KAL_in_plasma - V_{medium} \cdot (kon_{KAL_HMWK} \\
&\cdot KAL_in_plasma \cdot HMWK_in_plasma - koff_{KAL_HMWK} \\
&\cdot KAL_HMWK_in_plasma) - V_{medium} \cdot (kon_{C1In_KAL} \cdot C1Inh_in_plasma \\
&\cdot KAL_in_plasma - koff_{C1Inh_KAL} \cdot C1Inh_KAL_in_plasma) - V_{medium} \\
&\cdot (kon_{KAL_HK2Chain} \cdot KAL_in_plasma \cdot HK2Chain_in_plasma \\
&- koff_{KAL_HK2Chain} \cdot KAL_HK2Chain_in_plasma) - V_{medium} \\
&\cdot (kon_{KAL_Lanadelumab} \cdot KAL_in_plasma \cdot Lanadelumab_in_plasma \\
&- koff_{KAL_Lanadelumab} \cdot Lanadelumab_KAL_in_plasma)
\end{aligned}
\tag{E22}$$

$$\begin{aligned}
V_{medium} \frac{dLanadelumab_KAL_HMWK_in_plasma}{dt} &= V_{medium} \cdot (kon_{KAL_Lanadelumab} \cdot KAL_HMWK_in_plasma \\
&\cdot Lanadelumab_in_plasma - koff_{KAL_Lanadelumab} \\
&\cdot Lanadelumab_KAL_HMWK_in_plasma) - (k12 \\
&\cdot Lanadelumab_KAL_HMWK_in_plasma \cdot V_{medium} - k21 \\
&\cdot Lanadelumab_KAL_HMWK \cdot V_{proximal})
\end{aligned}
\tag{E23}$$

$$V_{medium} \frac{dLanadelumab_KAL_in_plasma}{dt} = V_{medium} \cdot (kon_{KAL_Lanadelumab} \cdot KAL_in_plasma \cdot Lanadelumab_in_plasma - koff_{KAL_Lanadelumab} \cdot Lanadelumab_KAL_in_plasma)$$

(E24)

$$V_{medium} \frac{dpreKAL_degraded}{dt} = V_{medium} \cdot kdeg_{preKAL} \cdot preKAL_in_plasma$$

(E25)

$$V_{medium} \frac{dpreKAL_HMWK_in_plasma}{dt} = V_{medium} \cdot (kon_{preKAL_HMWK} \cdot preKAL_in_plasma \cdot HMWK_in_plasma - koff_{preKAL_HMWK} \cdot preKAL_HMWK_in_plasma) - (k12 \cdot preKAL_HMWK_in_plasma \cdot V_{medium} - k21 \cdot preKAL_HMWK \cdot V_{proximal})$$

(E26)

$$V_{medium} \frac{dpreKAL_in_plasma}{dt} = V_{medium} \cdot ksyn_{preKAL} - V_{medium} \cdot kdeg_{preKAL} \cdot preKAL_in_plasma - V_{medium} \cdot (kon_{preKAL_HMWK} \cdot preKAL_in_plasma \cdot HMWK_in_plasma - koff_{preKAL_HMWK} \cdot preKAL_HMWK_in_plasma)$$

(E27)

In proximal space

$$V_{proximal} \frac{dBK}{dt} = (k12_{BK} \cdot BK_in_plasma \cdot V_{medium} - k21_{BK} \cdot BK \cdot V_{proximal}) + (kcat_{HMWK_cleavage} \cdot KAL_HMWK_gC1qR - (kon_{BK_BDK} \cdot BK \cdot BDKRB2 - koff_{BK_BDKRB2} \cdot BK_BDKRB2)) \cdot Num_to_Conc_converter \cdot V_{proximal}$$

(E28)

$$\frac{dBDKRB2}{dt} = ksyn_{BDKRB2} - kdeg_{BDKRB2} \cdot BDKRB2 - (kon_{BK_BDKRB2} \cdot BK \cdot BDKRB2 - koff_{BK_BDKRB2} \cdot BK_BDKRB2)$$

(E29)

$$\frac{dBDKRB2_degraded}{dt} = kdeg_{BDKRB2} \cdot BDKRB2$$

(E30)

$$\frac{dBK_BDKRB2}{dt} = (kon_{BK_BDKRB2} \cdot BK \cdot BDKRB2 - koff_{BK_BDKRB2} \cdot BK_BDKRB2) - kdeg_{BDKRB2} \cdot BK_BDKRB2 \quad (E31)$$

$$\frac{dBK_BDKRB2_degraded}{dt} = kdeg_{BDKRB2} \cdot BK_BDKRB2 \quad (E32)$$

$$\begin{aligned} \frac{dC1Inh_FXIIa_gC1qR}{dt} &= (kon_{C1Inh_FXIIa} \cdot C1Inh \cdot FXIIa_gC1qR - koff_{C1Inh_FXIIa} \\ &\cdot C1Inh_FXIIa_gC1qR) + (kon_{FXIIa_gC1qR} \cdot C1Inh_FXIIa \cdot gC1qR \\ &- koff_{FXIIa_gC1qR} \cdot C1Inh_FXIIa_gC1qR) - kdeg_{gC1qR} \\ &\cdot C1Inh_FXIIa_gC1qR \end{aligned} \quad (E33)$$

$$\frac{dC1Inh_FXIIa_gC1qR_degraded}{dt} = kdeg_{gC1qR} \cdot C1Inh_FXIIa_gC1qR \quad (E34)$$

$$\begin{aligned} \frac{dC1Inh_KAL_HMWK_gC1qR}{dt} &= (kon_{C1Inh_KAL} \cdot C1Inh \cdot KAL_HMWK_gC1qR - koff_{C1Inh_KAL} \\ &\cdot C1Inh_KAL_HMWK_gC1qR) + (kon_{HMWK_gC1} \cdot C1Inh_KAL_HMWK \\ &\cdot gC1qR - koff_{HMWK_gC1qR} \cdot C1Inh_KAL_HMWK_gC1qR) - kdeg_{gC1qR} \\ &\cdot C1Inh_KAL_HMWK_gC1qR \end{aligned} \quad (E35)$$

$$\frac{dC1Inh_KAL_HMWK_gC1qR_degraded}{dt} = kdeg_{gC1qR} \cdot C1Inh_KAL_HMWK_gC1qR \quad (E36)$$

$$\begin{aligned} \frac{dFXII_gC1qR}{dt} &= (kon_{FXII_gC1qR} \cdot FXII \cdot gC1qR - koff_{FXII_gC1qR} \cdot FXII_gC1qR) \\ &- Fold_increase_{FXII_AutoActivation} \cdot kcat_{FXII_AutoActivation} \cdot FXII_gC1qR \\ &- kcat_{FXII_AutoActivation} \cdot KAL_HMWK_gC1qR \cdot FXII_gC1qR \\ &\cdot Num_to_Conc_converter / (km_{FXII_AutoActivation} + FXII_gC1qR \\ &\cdot Num_to_Conc_converter) - kdeg_{gC1qR} \cdot FXII_gC1qR \end{aligned} \quad (E37)$$

$$\frac{dFXII_gC1qR_degraded}{dt} = kdeg_{gC1qR} \cdot FXII_gC1qR \quad (E38)$$

$$\begin{aligned}
\frac{dFXIIa_gC1qR}{dt} = & Fold_increase_{FXII_AutoActivation} \cdot kcat_{FXII_AutoActivation} \cdot FXII_gC1qR \\
& + kcat_{FXII_AutoActivation} \cdot KAL_HMWK_gC1qR \cdot FXII_gC1qR \\
& \cdot Num_to_Conc_converter / (km_{FXII_AutoActivation} + FXII_gC1qR \\
& \cdot Num_to_Conc_converter) + (kon_{FXIIa_gC1qR} \cdot FXIIa \cdot gC1qR \\
& - koff_{FXIIa_gC1qR} \cdot FXIIa_gC1qR) - (kon_{C1Inh_FXIIa} \cdot C1Inh \\
& \cdot FXIIa_gC1qR - koff_{C1Inh_FXIIa} \cdot C1Inh_FXIIa_gC1qR) - kdeg_{gC1qR} \\
& \cdot FXIIa_gC1qR
\end{aligned}
\tag{E39}$$

$$\frac{dFXIIa_gC1qR_degraded}{dt} = kdeg_{gC1qR} \cdot FXIIa_gC1qR
\tag{E40}$$

$$\begin{aligned}
\frac{dgC1qR}{dt} = & ksyn_{gC1qR} - kdeg_{gC1qR} \cdot gC1qR - (kon_{FXII_gC1qR} \cdot FXII \cdot gC1qR \\
& - koff_{FXII_gC1q} \cdot FXII_gC1qR) - (kon_{HMWK_gC1qR} \cdot preKAL_HMWK \\
& \cdot gC1qR - koff_{HMWK_gC1qR} \cdot preKAL_HMWK_gC1qR) - (kon_{FXIIa_gC1} \\
& \cdot FXIIa \cdot gC1qR - koff_{FXIIa_gC1qR} \cdot FXIIa_gC1qR) - (kon_{HMWK_gC1qR} \\
& \cdot KAL_HMWK \cdot gC1qR - koff_{HMWK_gC1q} \cdot KAL_HMWK_gC1qR) \\
& - (kon_{HK2Chain_gC1qR} \cdot KAL_HK2Chain \cdot gC1qR - koff_{HK2Chain_gC1qR} \\
& \cdot KAL_HK2Chain_gC1qR) - (kon_{FXIIa_gC1qR} \cdot C1Inh_FXIIa \cdot gC1qR \\
& - koff_{FXIIa_gC1qR} \cdot C1Inh_FXIIa_gC1qR) - (kon_{HMWK_gC1qR} \\
& \cdot C1Inh_KAL_HMWK \cdot gC1qR - koff_{HMWK_gC1qR} \\
& \cdot C1Inh_KAL_HMWK_gC1qR) - (kon_{HMWK_gC1qR} \\
& \cdot Lanadelumab_KAL_HMWK \cdot gC1qR - koff_{HMWK_gC1qR} \\
& \cdot Lanadelumab_KAL_HMWK_gC1qR)
\end{aligned}
\tag{E41}$$

$$\frac{dgC1qR_degraded}{dt} = kdeg_{gC1qR} \cdot gC1qR
\tag{E42}$$

$$\begin{aligned}
& \frac{dKAL_HK2Chain_gC1qR}{dt} \\
& = kcat_{HMWK_cleavage} \cdot KAL_HMWK_gC1qR + (kon_{HK2Chain_gC1qR} \\
& \quad \cdot KAL_HK2Chain \cdot gC1qR - koff_{HK2Chain_gC1qR} \\
& \quad \cdot KAL_HK2Chain_gC1qR) - kdeg_{gC1qR} \cdot KAL_HK2Chain \cdot gC1qR
\end{aligned}
\tag{E43}$$

$$\frac{dKAL_HK2Chain_gC1qR_degraded}{dt} = kdeg_{gC1qR} \cdot KAL_HK2Chain_gC1qR
\tag{E44}$$

$$\begin{aligned}
& \frac{dKAL_HMWK_gC1qR}{dt} \\
& = kcat_{preKAL_Activation} \cdot FXIIa_gC1qR \cdot preKAL_HMWK_gC1qR \\
& \quad \cdot Num_to_Conc_converter / (km_{preKAL_Activation} + preKAL_HMWK_gC1qR \\
& \quad \cdot Num_to_Conc_converter) - kcat_{HMWK_cleavage} \cdot KAL_HMWK_gC1qR \\
& \quad + (kon_{HMWK_gC1qR} \cdot KAL_HMWK \cdot gC1qR - koff_{HMWK_gC1qR} \\
& \quad \cdot KAL_HMWK_gC1qR) - (kon_{C1Inh_KAL} \cdot C1Inh \cdot KAL_HMWK_gC1qR \\
& \quad - koff_{C1Inh_KAL} \cdot C1Inh_KAL_HMWK_gC1qR) - kdeg_{gC1qR} \\
& \quad \cdot KAL_HMWK_gC1qR - (kon_{KAL_Lanadelumab} \cdot KAL_HMWK_gC1qR \\
& \quad \cdot Lanadelumab - koff_{KAL_Lanadelumab} \\
& \quad \cdot Lanadelumab_KAL_HMWK_gC1qR)
\end{aligned}
\tag{E45}$$

$$\frac{dKAL_HMWK_gC1qR_degraded}{dt} = kdeg_{gC1qR} \cdot KAL_HMWK_gC1qR
\tag{E46}$$

$$\begin{aligned}
& \frac{dLanadelumab_KAL_HMWK_gC1qR}{dt} \\
& = (kon_{KAL_Lanadelumab} \cdot KAL_HMWK_gC1qR \cdot Lanadelumab \\
& \quad - koff_{KAL_Lanadelumab} \cdot Lanadelumab_KAL_HMWK_gC1qR) \\
& \quad + (kon_{HMWK_gC1qR} \cdot Lanadelumab_KAL_HMWK \cdot gC1qR \\
& \quad - koff_{HMWK_gC1qR} \cdot Lanadelumab_KAL_HMWK_gC1qR) \\
& \quad - kdeg_{Lanadelumab_KAL_HMWK_gC1qR} \cdot Lanadelumab_KAL_HMWK_gC1qR
\end{aligned}
\tag{E47}$$

$$\frac{dLanadelumab_KAL_HMWK_gC1qR_degraded}{dt} = kdeg_{Lanadelumab_KAL_HMWK_gC1qR} \cdot Lanadelumab_KAL_HMWK_gC1qR \quad (E48)$$

$$\begin{aligned} \frac{dpreKAL_HMWK_gC1qR}{dt} = & (kon_{HMWK_gC1} \cdot preKAL_HMWK \cdot gC1qR - koff_{HMWK_gC1qR} \\ & \cdot preKAL_HMWK_gC1qR) - kcat_{preKAL_Activation} \cdot FXIIa_gC1qR \\ & \cdot preKAL_HMWK_gC1qR \cdot Num_to_Conc_converter / (km_{preKAL_Activation} \\ & + preKAL_HMWK_gC1qR \cdot Num_to_Conc_converter) - kdeg_{gC1qR} \\ & \cdot preKAL_HMWK_gC1qR \end{aligned} \quad (E49)$$

$$\frac{dpreKAL_HMWK_gC1qR_degraded}{dt} = kdeg_{gC1qR} \cdot preKAL_HMWK_gC1qR \quad (E50)$$

$$\begin{aligned} V_{proximal} \frac{dC1Inh}{dt} = & (k12 \cdot C1Inh_in_plasma \cdot V_{medium} - k21 \cdot C1Inh \cdot V_{proximal}) \\ & - (kon_{C1Inh_FXIIa} \cdot C1Inh \cdot FXIIa_gC1qR - koff_{C1Inh_FXIIa} \\ & \cdot C1Inh_FXIIa_gC1qR + kon_{C1Inh_KAL} \cdot C1Inh \cdot KAL_HMWK_gC1qR \\ & - koff_{C1Inh_KAL} \cdot C1Inh_KAL_HMWK_gC1qR) \cdot Num_to_Conc_converter \\ & \cdot V_{proximal} \end{aligned} \quad (E51)$$

$$\begin{aligned} V_{proximal} \frac{dC1Inh_FXIIa}{dt} = & (k12 \cdot C1Inh_FXIIa_in_plasma \cdot V_{medium} - k21 \cdot C1Inh_FXIIa \\ & \cdot V_{proximal}) - (kon_{FXIIa_gC1qR} \cdot C1Inh_FXIIa \cdot gC1qR - koff_{FXIIa_gC1qR} \\ & \cdot C1Inh_FXIIa_gC1qR) \cdot Num_to_Conc_converter \cdot V_{proximal} \end{aligned} \quad (E52)$$

$$\begin{aligned}
V_{proximal} \frac{dC1Inh_KAL_HMWK}{dt} &= (k12 \cdot C1Inh_KAL_HMWK_in_plasma \cdot V_{medium} - k21 \\
&\cdot C1Inh_KAL_HMWK \cdot V_{proximal}) - (kon_{HMWK_gC1qR} \cdot C1Inh_KAL_HMWK \\
&\cdot gC1qR - koff_{HMWK_gC1qR} \cdot C1Inh_KAL_HMWK_gC1qR) \\
&\cdot Num_to_Conc_converter \cdot V_{proximal}
\end{aligned}
\tag{E53}$$

$$\begin{aligned}
V_{proximal} \frac{dFXII}{dt} &= (k12 \cdot FXII_in_plasma \cdot V_{medium} - k21 \cdot FXII \cdot V_{proximal}) \\
&- (kon_{FXII_gC1} \cdot FXII \cdot gC1qR - koff_{FXII_gC1q} \cdot FXII_gC1qR) \\
&\cdot Num_to_Conc_converter \cdot V_{proximal}
\end{aligned}
\tag{E54}$$

$$\begin{aligned}
V_{proximal} \frac{dFXIIa}{dt} &= (k12 \cdot FXIIa_in_plasma \cdot V_{medium} - k21 \cdot FXIIa \cdot V_{proximal}) \\
&- (kon_{FXIIa_gC1qR} \cdot FXIIa \cdot gC1qR - koff_{FXIIa_gC1} \cdot FXIIa_gC1qR) \\
&\cdot Num_to_Conc_converter \cdot V_{proximal}
\end{aligned}
\tag{E55}$$

$$\begin{aligned}
V_{proximal} \frac{dKAL_HK2Chain}{dt} &= (k12 \cdot KAL_HK2Chain_in_plasma \cdot V_{medium} - k21 \cdot KAL_HK2Chain \\
&\cdot V_{proximal}) - (kon_{HK2Chain_gC1qR} \cdot KAL_HK2Chain \cdot gC1qR \\
&- koff_{HK2Chain_gC1qR} \cdot KAL_HK2Chain_gC1qR) \cdot Num_to_Conc_converter \\
&\cdot V_{proximal}
\end{aligned}
\tag{E56}$$

$$\begin{aligned}
V_{proximal} \frac{dKAL_HMWK}{dt} &= (k12 \cdot KAL_HMWK_in_plasma \cdot V_{medium} - k21 \cdot KAL_HMWK \\
&\cdot V_{proximal}) - (kon_{HMWK_gC1qR} \cdot KAL_HMWK \cdot gC1qR - koff_{HMWK_gC1qR} \\
&\cdot KAL_HMWK_gC1qR) \cdot Num_to_Conc_converter \cdot V_{proximal}
\end{aligned}
\tag{E57}$$

$$\begin{aligned}
V_{proximal} \frac{dLanadelumab_KAL_HMWK}{dt} &= (k_{12} \cdot Lanadelumab_KAL_HMWK_in_plasma \cdot V_{medium} - k_{21} \\
&\cdot Lanadelumab_KAL_HMWK \cdot V_{proximal}) - (kon_{HMWK_gC1qR} \\
&\cdot Lanadelumab_KAL_HMWK \cdot gC1qR - koff_{HMWK_gC1} \\
&\cdot Lanadelumab_KAL_HMWK_gC1qR) \cdot Num_to_Conc_converter \\
&\cdot V_{proximal}
\end{aligned}
\tag{E58}$$

$$\begin{aligned}
V_{proximal} \frac{dpreKAL_HMWK}{dt} &= (k_{12} \cdot preKAL_HMWK_in_plasma \cdot V_{medium} - k_{21} \cdot preKAL_HMWK \\
&\cdot V_{proximal}) - (kon_{HMWK_gC1qR} \cdot preKAL_HMWK \cdot gC1qR \\
&- koff_{HMWK_gC1qR} \cdot preKAL_HMWK_gC1qR) \cdot Num_to_Conc_converter \\
&\cdot V_{proximal}
\end{aligned}
\tag{E59}$$

BDKR-B2 B2 recepter of bradykinin receptor family, *BK* bradykinin, *C1-INH* C1 esterase inhibitor, *C1q* complement component 1, *FXII* factor XII, *FXIIa* activated factor XII, *gC1q-R* cofactors complement protein C1q, *HMWK* high molecular weight kininogen, *K_{off}* dissociation constant, *K_{on}* association constant, *KAL* kallikrein, *preKAL* prekallikrein, *V_{medium}* per endothelial cell-based vascular volume, *V_{proximal}* per endothelial cell-based proximal space volume

Table S6 List of model parameters

Parameter	Description	Unit	Value	Source
In vascular space				
Kd_KAL_HK2Chain	K _d for "KAL_in_plasma + HK2Chain_in_plasma ↔ KAL_HK2Chain_in_plasma"	nM	72	Kenniston et al [24]
Kd_KAL_HMWK	K _d for "KAL_in_plasma + HMWK_in_plasma ↔ KAL_HMWK_in_plasma"	nM	15	Bock et al [5]
Kd_preKAL_HMWK	K _d for "preKAL_in_plasma + HMWK_in_plasma ↔ preKAL_HMWK_in_plasma"	nM	12	Bock et al [5]
kdeg_BK	Degradation rate for bradykinin	1/h	55.452	Décarie et al (1996) [27]
kdeg_Bound_C1Inh	Degradation rate for bound C1-INH	1/h	13.863	Estimated from Jensen et al [28]
kdeg_C1Inh	Degradation rate for C1-INH	1/h	0.0165	Weidmann et al [22]
kdeg_cHMWK	Degradation rate for cHMWK	1/h	0.0619	Calibrated with guidance from typical range in proteins
kdeg_FXII	Degradation rate for FXII	1/h	0.0116	Weidmann et al [22]
kdeg_FXIIa	Degradation rate for FXIIa	1/h	8.318	Assumed same as kallikrein
kdeg_HMWK	Degradation rate for HMWK	1/h	0.0044	Weidmann et al [22]
kdeg_KAL	Degradation rate for kallikrein	1/h	8.318	Cumming et al [29]

kdeg_preKAL	Degradation rate for prekallikrein	1/h	0.0289	Calculated using a half-life of 24 hours (Labcorp.com)
koff_KAL_HK2Chain	k _{off} for kallikrein and cHMWK binding event	1/h	318.816	Calculated from K _d and k _{on} of the corresponding reaction
koff_KAL_HMWK	k _{off} for kallikrein and HMWK binding event	1/h	66.42	Calculated from K _d and k _{on} of the corresponding reaction
koff_preKAL_HMWK	k _{off} for prekallikrein & HMWK binding event	1/h	53.136	Calculated from K _d and k _{on} of the corresponding reaction
kon_KAL_HK2Chain	k _{on} for "KAL_in_plasma + HK2Chain_in_plasma ↔ KAL_HK2Chain_in_plasma"	1/(M×h)	4.428	Assumed 10× kon_HMWK_gC1qR, due to lower steric constraint
kon_KAL_HMWK	k _{on} for "KAL_in_plasma + HMWK_in_plasma ↔ KAL_HMWK_in_plasma"	1/(M×h)	4.428	Assumed 10× kon_HMWK_gC1qR, due to lower steric constraint
kon_preKAL_HMWK	k _{on} for "preKAL_in_plasma + HMWK_in_plasma ↔ preKAL_HMWK_in_plasma"	1/(M×h)	4.428	Assumed 10× kon_HMWK_gC1qR, due to lower steric constraint
ksyn_C1Inh	Synthesis rate for C1-INH	nM/h	11.883 HAE 39.608 Healthy	Calibrated with guidance from steady-state level data in Table S2

ksyn_FXII	Synthesis rate for FXII	nM/h	10.83	Calibrated with guidance from steady-state level data in Table S2
ksyn_HMWK	Synthesis rate for HMWK	nM/h	39.933	Calibrated with guidance from steady-state level data in Table S2
ksyn_preKAL	Synthesis rate for prekallikrein	nM/h	41.589	Calibrated with guidance from steady-state level data in Table S2
Vmedium	Per endothelial cell based plasma volume	L	1.23E-12	Estimated from Shah and Betts [30] and Bianconi et al [31]
In proximal space				
BDKRB2_per_Cell_SS	The number of bradykinin B2 receptors per cell at steady state	-	100,000	Estimated from Paquet et al [14]
gC1qR_per_Cell_SS	The number of gC1q receptors per cell at steady state	-	100,000	Mahdi et al [9], assumed limited by the CK1 cofactor
kcat_FXII_Activation	k _{cat} for FXII activation (substrate: FXII_gC1qR; enzyme: KAL_HMWK_gC1qR; product: FXIIa_gC1qR)	1/h	15	Calibrated with guidance from steady-state level data in Table S2

kcat_FXII_AutoActivation	k _{cat} for FXII autoactivation (substrate: FXII_gC1qR; enzyme: FXII_gC1qR; product: FXIIa_gC1qR)	1/h	0.0475	Calibrated with guidance from steady-state level data in Table S2
kcat_HMWK_cleavage	k _{cat} for cleavage of HMWK	1/h	394.7	Calibrated with guidance from steady-state level data in Table S2
kcat_preKAL_Activation	k _{cat} for prekallikrein activation (substrate: preKAL_HMWK_gC1qR; enzyme: FXIIa_gC1qR; product: KAL_HMWK_gC1qR)	1/h	18	Calibrated with guidance from steady-state level data in Table S2
Kd_BK_BDKRB2	k _d for "BK + BDKRB2 ↔ BK_BDKRB2"	nM	0.5	Paquet et al [14]
Kd_FXII_gC1qR	k _d for "FXII + gC1qR ↔ FXII_gC1qR"	nM	144	Reddigari et al [7]
Kd_FXIIa_gC1qR	k _d for "FXIIa + gC1qR ↔ FXIIa_gC1qR"	nM	144	Assumed same as Kd_FXII_gC1qR
Kd_HK2Chain_gC1qR	k _d for "KAL_HK2Chain + gC1qR ↔ KAL_HK2Chain_gC1qR"	nM	10.35	Assumed same as kon_HMWK_gC1qR
Kd_HMWK_gC1qR	k _d for "preKAL_HMWK + gC1qR ↔ preKAL_HMWK_gC1qR"	nM	10.35	Fernando et al [32]

kdeg_BDKRB2	Degradation rate for bradykinin B2 receptors	1/h	0.3466	Used typical degradation rate for receptors, as reported by Morelon and Dautry-Varsat [33]
kdeg_gC1qR	Degradation rate for gC1q receptor complex on endothelial cell surface	1/h	0.3466	Used typical degradation rate for receptors, as reported by Morelon and Dautry-Varsat [33]
kdeg_Lanadelumab_KAL_HMWK_gC1qR	Degradation rate for lanadelumab bound with KAL_HMWK_gC1qR receptor complex	1/h	0.3466	Assumed same as the degradation rate for unbound receptors
Km_FXII_Activation	K _m for FXII activation (substrate: FXII_gC1qR; enzyme: KAL_HMWK_gC1qR; product: FXIIa_gC1qR)	nM	510	Tankersley and Finlayson [26]
Km_FXII_AutoActivation	K _m for FXII autoactivation (substrate: FXII_gC1qR; enzyme: FXII_gC1qR; product: FXIIa_gC1qR)	nM	110	Bernardo et al [34]
Km_preKAL_Activation	K _m for prekallikrein activation (substrate: preKAL_HMWK_gC1qR; enzyme: FXIIa_gC1qR;	nM	91	Tankersley and Finlayson [26]

	product: KAL_HMWK_gC1qR)			
koff_BK_BDKRB2	k _{off} for bradykinin & bradykinin B2 receptor binding event	1/h	18	Calculated from K _d and k _{on} of the corresponding reaction
koff_FXII_gC1qR	k _{off} for FXII and gC1q receptor binding event	1/h	63.763	Calculated from K _d and k _{on} of the corresponding reaction
koff_FXIIa_gC1qR	k _{off} for FXIIa & gC1q receptor binding event	1/h	63.763	Calculated from K _d and k _{on} of the corresponding reaction
koff_HK2Chain_gC1qR	k _{off} for cHMWK & and gC1q receptor binding event	1/h	4.583	Calculated from K _d and k _{on} of the corresponding reaction
koff_HMWK_gC1qR	k _{off} for HMWK & gC1q receptor binding event	1/h	4.583	Calculated from K _d and k _{on} of the corresponding reaction
kon_BK_BDKRB2	k _{on} for "BK + BDKRB2 ↔ BK_BDKRB2"	1/(M×h)	36	Assumed fast binding for bradykinin due to its low molecular weight
kon_FXII_gC1qR	k _{on} for "FXII + gC1qR ↔ FXII_gC1qR"	1/(M×h)	0.4428	Assumed same as kon_HMWK_gC1qR
kon_FXIIa_gC1qR	k _{on} for "FXIIa + gC1qR ↔ FXIIa_gC1qR"	1/(M×h)	0.4428	Assumed same as kon_HMWK_gC1qR

kon_HK2Chain_gC1qR	k_{on} for "KAL_HK2Chain + gC1qR $\leftrightarrow$ KAL_HK2Chain_gC1qR"	1/(M×h)	0.4428	Assumed same as kon_HMWK_gC1qR
kon_HMWK_gC1qR	k_{on} for "preKAL_HMWK + gC1qR $\leftrightarrow$ preKAL_HMWK_gC1qR"	1/(M×h)	0.4428	Pixley et al [6]
ksyn_BDKRB2	Synthesis rate for bradykinin B2 receptors	number/ cell/h	34657.35 9	Estimated from steady-state level data from Paquet et al [14]
ksyn_gC1qR	Synthesis rate for gC1q receptor complex on endothelial cell surface	number/ cell/h	34657.35 9	Estimated from steady-state level data from Mahdi et al [9]
Vproximal	Proximal space volume near cell surface for each endothelial cell	L	8.00E-15	20 nm (height) × 400 μm^2 (surface area) [35, 36]

In vascular and proximal space

Kd_C1Inh_FXIIa	k_d for "C1Inh_in_plasma + FXIIa_in_plasma $\leftrightarrow$ C1Inh_FXIIa_in_plasma" and "C1Inh + FXIIa_gC1qR $\leftrightarrow$ C1Inh_FXIIa_gC1qR"	nM	1720	Estimated from Biacore data by Björkqvist et al [37]
Kd_C1Inh_KAL	k_d for "C1Inh_in_plasma + KAL_in_plasma $\leftrightarrow$ C1Inh_KAL_in_plasma", "C1Inh_in_plasma +	nM	150	Estimated from immunosorbent assay data by Harpel et al [25]

	KAL_HMWK_in_plasma ↔ C1Inh_KAL_HMWK_in_plasma", and "C1Inh + KAL_HMWK_gC1qR ↔ C1Inh_KAL_HMWK_gC1qR "			
Kd_KAL_Lanadelumab	k _d for "KAL_in_plasma + Lanadelumab_in_plasma ↔ Lanadelumab_KAL_in_plasma", "KAL_HMWK_in_plasma + Lanadelumab_in_plasma ↔ Lanadelumab_KAL_HMWK_in_plasma", and "KAL_HMWK_gC1qR + Lanadelumab ↔ Lanadelumab_KAL_HMWK_gC1qR"	nM	0.12	Shire measurement, Kenniston et al [24]
koff_C1Inh_FXIIa	k _{off} for "C1Inh_in_plasma + FXIIa_in_plasma ↔ C1Inh_FXIIa_in_plasma" and "C1Inh + FXIIa_gC1qR ↔ C1Inh_FXIIa_gC1qR"	1/h	229.104	Calculated from K _d and k _{on} of the corresponding reaction
koff_C1Inh_KAL	k _{off} for "C1Inh_in_plasma + KAL_in_plasma ↔ C1Inh_KAL_in_plasma",	1/h	9.18	Calculated from K _d and k _{on} of the corresponding reaction

	"C1Inh_in_plasma + KAL_HMWK_in_plasma ↔ C1Inh_KAL_HMWK_in_plasma", and "C1Inh + KAL_HMWK_gC1qR ↔ C1Inh_KAL_HMWK_gC1qR"			
koff_KAL_Lanadelumab	k _{off} for "KAL_in_plasma + Lanadelumab_in_plasma ↔ Lanadelumab_KAL_in_plasma", "KAL_HMWK_in_plasma + Lanadelumab_in_plasma ↔ Lanadelumab_KAL_HMWK_in_plasma", and "KAL_HMWK_gC1qR + Lanadelumab ↔ Lanadelumab_KAL_HMWK_gC1qR"	1/h	1.452	Calculated from K _d and k _{on} of the corresponding reaction
kon_C1Inh_FXIIa	k _{on} for "C1Inh_in_plasma + FXIIa_in_plasma ↔ C1Inh_FXIIa_in_plasma" and "C1Inh + FXIIa_gC1qR ↔ C1Inh_FXIIa_gC1qR"	1/(nM× h)	0.1332	Drouet et al [38]
kon_C1Inh_KAL	k _{on} for "C1Inh_in_plasma + KAL_in_plasma ↔	1/(nM× h)	0.0612	Drouet et al [38]

	C1Inh_KAL_in_plasma", "C1Inh_in_plasma + KAL_HMWK_in_plasma ↔ C1Inh_KAL_HMWK_in_plasma", and "C1Inh + KAL_HMWK_gC1qR ↔ C1Inh_KAL_HMWK_gC1qR"			
kon_KAL_Lanadelumab	k _{on} for "KAL_in_plasma + Lanadelumab_in_plasma ↔ Lanadelumab_KAL_in_plasma", "KAL_HMWK_in_plasma + Lanadelumab_in_plasma ↔ Lanadelumab_KAL_HMWK_in_plasma", and "KAL_HMWK_gC1qR + Lanadelumab ↔ Lanadelumab_KAL_HMWK_gC1qR"	1/(nM×h)	12.096	Shire measurement
Exchange between vascular and proximal space				
k12	Species exchange rate from plasma to proximal space	1/h	0.2341	Assumed based circulation time from Shah and Betts [30]

k21	Species exchange rate from proximal space to plasma	1/h	36	Assumed based circulation time from Shah and Betts [30]
k12_BK	Bradykinin exchange rate from plasma to proximal space	1/h	7.0244	Calibrated to match BK level in Table S2
k21_BK	Bradykinin exchange rate from proximal space to plasma	1/h	1080	Calibrated to match BK level in Table S2

BDKR-B2 B2 receptor of bradykinin receptor family, *BK* bradykinin, *C1-INH* C1 esterase inhibitor, *C1q* complement component 1, *cHMWK* cleaved high molecular weight kininogen, *CK1* cytokeratin 1, *FXII* factor XII, *FXIIa* activated factor XII, *gC1q-R* cofactors complement protein C1q, *HAE* hereditary angioedema, *HK2* human kallikrein 2, *HMWK* high molecular weight kininogen, *KAL* kallikrein, k_{cat} turnover number, K_d binding affinity, K_{off} dissociation constant, K_{on} association constant, K_{syn} synthesis rate, *preKAL* prekallikrein

Table S7 Model assumptions

1)	The CK1 cofactor with the lowest level of expression is the limiting number for formation of the gCq1-R/CK1/uPAR receptor complex in surface contact system activation. An apparent number of sites and affinity to different contact factors are used.
2)	The effects of Zn^{+2} dependency on binding affinities and the endothelial cell prekallikrein activator (PRCP) are not explicitly included in the model and their effects are assumed to be implicitly reflected in the model parameters.
3)	The exchange rate between the vascular space and the proximal space is assumed to be of the same order as the vascular volume circulation time (approximately 100 seconds [30]).
4)	Various triggers of HAE attacks (e.g. stress, physical trauma, a surgical or a dental procedure, infection, hormonal changes, mechanical pressure) are assumed to lead to a systematic perturbation that autoactivates the kinin-kallikrein pathway in the contact activation system.
5)	The rise in the pain is triggered by an acute attack and the duration of the attack is represented by the time period over which the level of FXII autoactivation remains elevated.
6)	The inhibitory activity of lanadelumab on kallikrein in plasma is the same as that of kallikrein bound to the surface, and the inhibitory activity of C1-INH on kallikrein and FXIIa in plasma is the same as on the surface.

C1-INH C1 esterase inhibitor, *CK1* cytokeratin 1, *C1q* complement component 1, *FXII* factor XII, *FXIIa* activated factor XII, *gC1q-R* cofactors complement protein C1q, *HAE* hereditary angioedema, *PRCP* prolylcarboxypeptidase, *uPAR* urokinase plasminogen activating receptor

Fig. S1 Overview of the process for development of the HAE QSP model. *C1-INH* C1 esterase inhibitor, *cHMWK* cleaved high molecular weight kininogen, *HAE* hereditary angioedema, *KKS* kallikrein-kinin system, *PD* pharmacodynamics, *PK* pharmacokinetics, *QSP* quantitative systems pharmacology

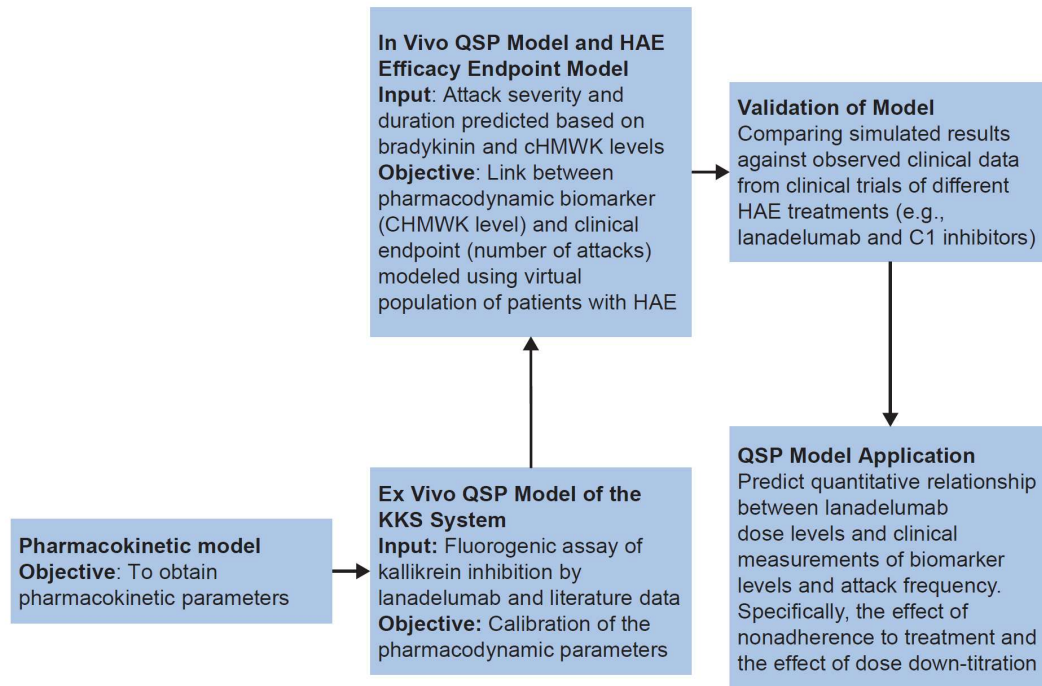


Fig. S2 Ex vivo assay measuring the inhibition of proteolytic activity of kallikrein by lanadelumab. The assay utilizes a synthetic peptide substrate (Pro-Phe-Arg-aminomethyl coumarin) that exhibits increased fluorescence upon proteolysis catalyzed by kallikrein. A key assumption is that the fluorescence readout can be directly related to the level of free kallikrein.

FXIIa activated factor XII, *SC* subcutaneous

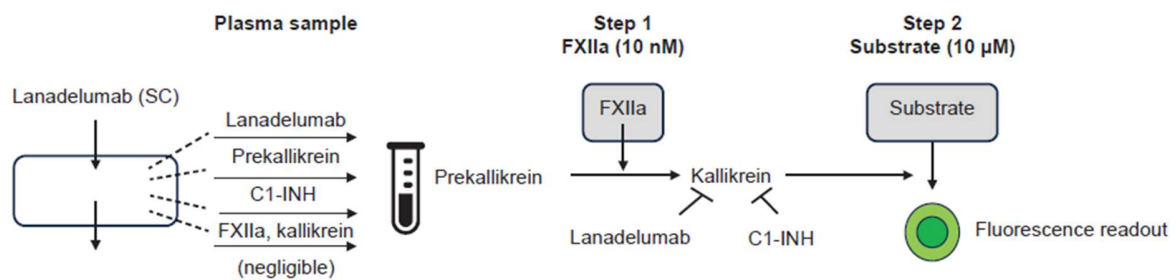


Fig. S3 Map of the KKS in HAE and pharmacodynamic interactions. The model reflects the biology of the KKS in HAE involving the essential contact factor proteins, such as FXII/FXIIa, prekallikrein/kallikrein and HMWK, which are activated on the endothelial cell surface to release the vasoactive peptide bradykinin. The pathway is a cascade of activation and cleavage reactions associated with these proteins and their complexes in plasma and bound to receptor complexes on the cell surface. *BDKR-B2* B2 receptor of bradykinin receptor family, *BK* bradykinin, *C1-INH* C1 esterase inhibitor, *FXII* factor XII, *FXIIa* activated factor XII, *HAE* hereditary angioedema, *HMWK* high molecular weight kininogen, *KAL* kallikrein, *mAb* monoclonal antibody, *KKS* kallikrein-kinin system, *preKAL* prekallikrein

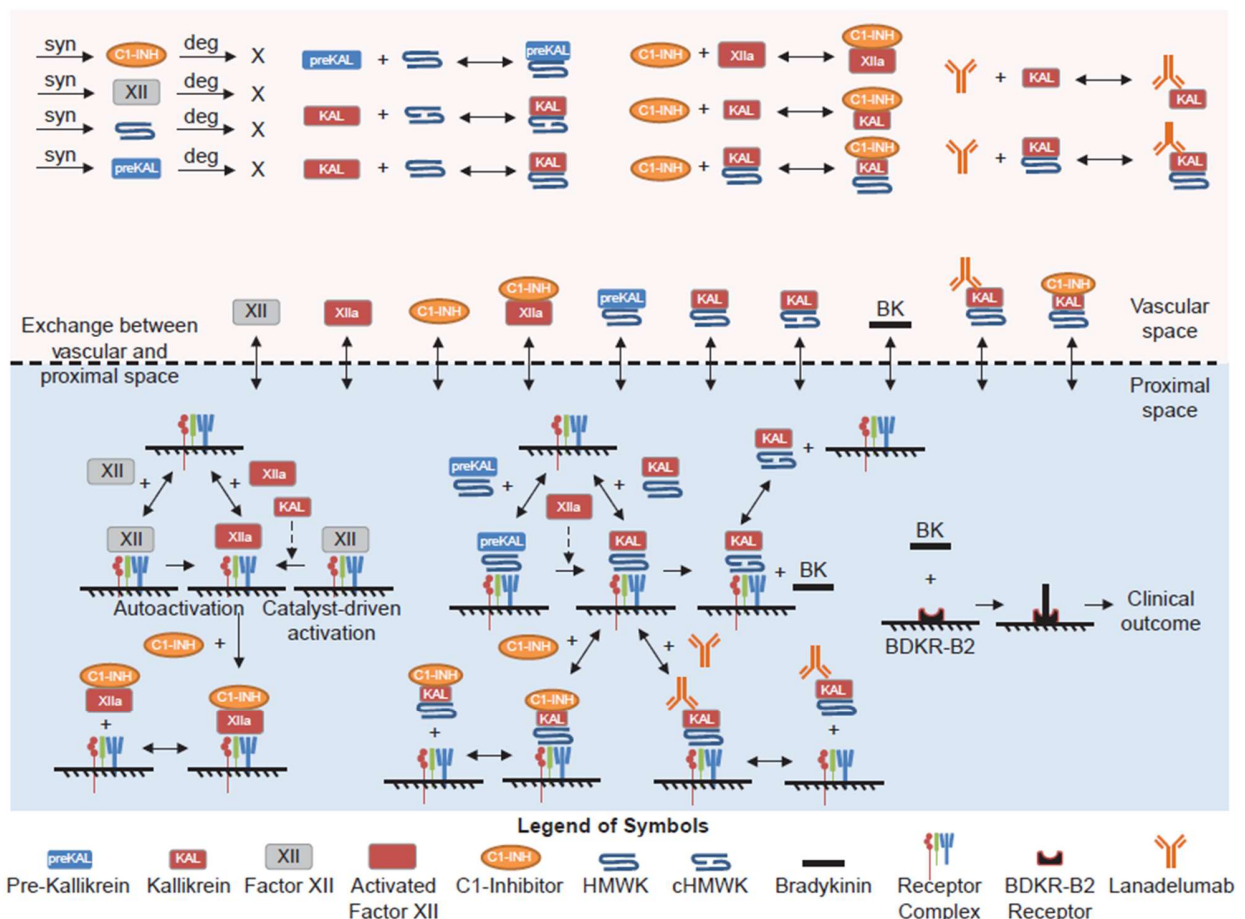


Fig. S4 Inhibition of plasma kallikrein by lanadelumab. Output from the simulation of lanadelumab treatment (shaded band) were compared to data from study phase 1 [15] (a) and phase 1b [16] (b) at different dose levels of lanadelumab. The prekallikrein range corresponds to the range reported by Madsen et al [1]. Negative % inhibition values arise when excess ex vivo KKS activation occurs in pre-dose samples during blood collection and processing to plasma, resulting in a lower enzymatic rate in the pre-dose sample [% inhibition = $(1 - \text{post-dose rate} / \text{pre-dose rate}) \times 100$]. *preKAL* prekallikrein

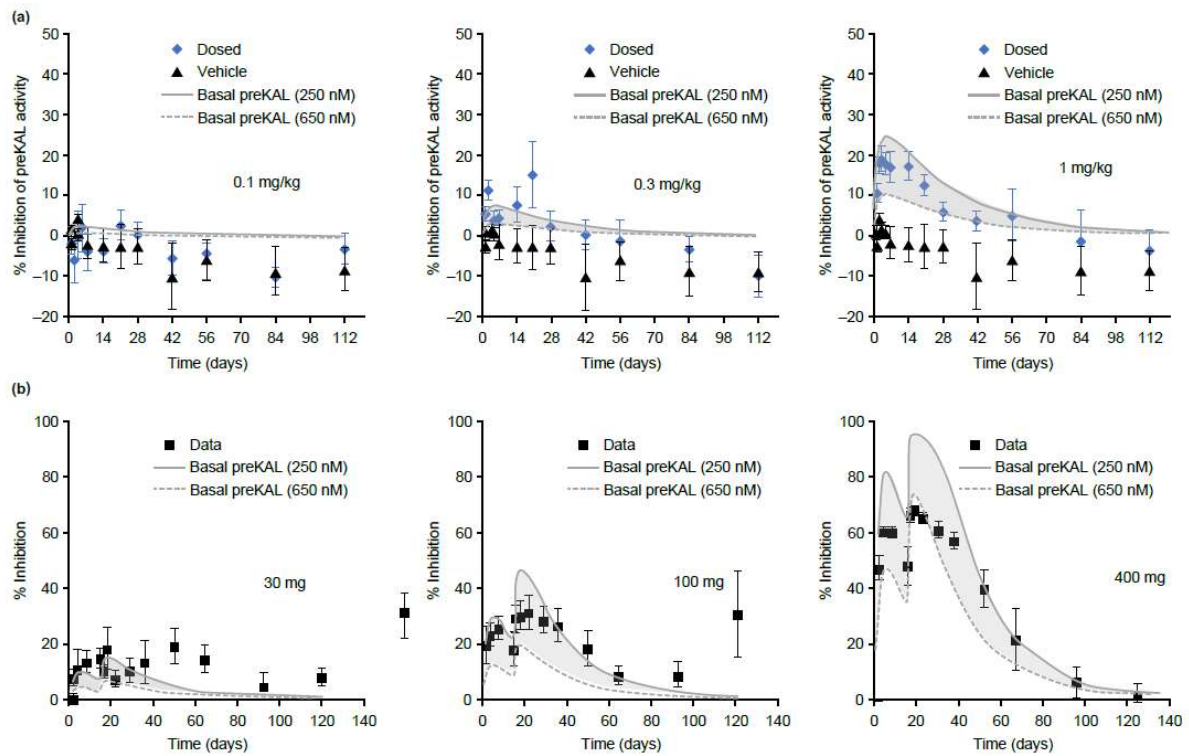


Fig. S5 Comparison of steady-state protein levels predicted by the KKS model to data from the literature. Error bars represent the range of levels reported in the literature. Free preKAL% refers to the percentage of prekallikrein in free form (i.e. not bound to HMWK) relative to total prekallikrein. %cHMWK is the percentage of cHMWK relative to the total of cHMWK + HMWK. *BK* bradykinin, *C1-INH* C1 esterase inhibitor, *cHMWK* cleaved high molecular weight kininogen, *HAE* hereditary angioedema, *HC* healthy controls, *HMWK* high molecular weight kininogen, *KKS* kallikrein-kinin system, *preKAL* prekallikrein

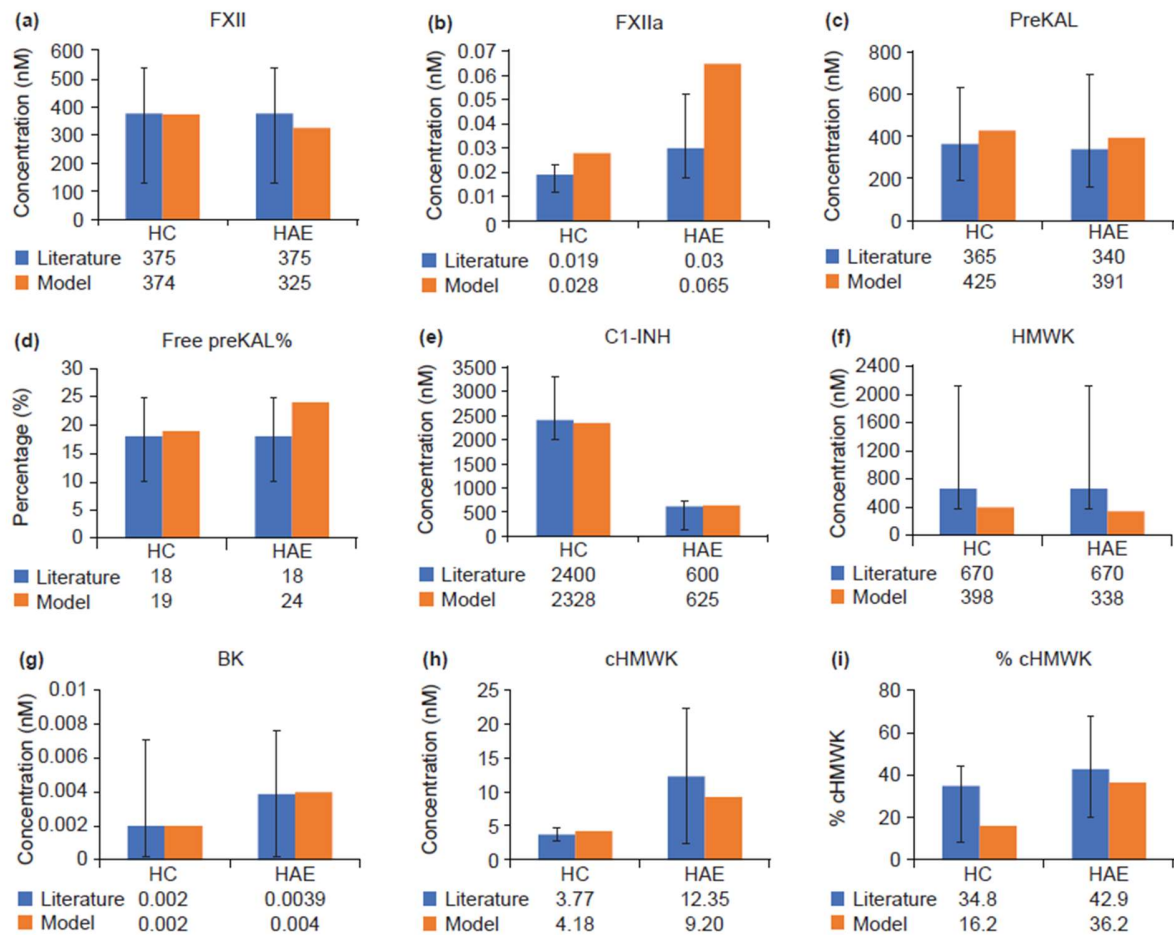


Fig. S6 Simulated PK profiles were compared with mean concentrations of functional C1-INH protein in HAE patients treated with (a) a single IV dose of 1000 IU C1-INH or (b) 1000 IU C1-INH twice/week for 12 weeks [19]. *C1-INH* C1 esterase inhibitor, *HAE* hereditary angioedema, *IV* intravenous, *PK* pharmacokinetics

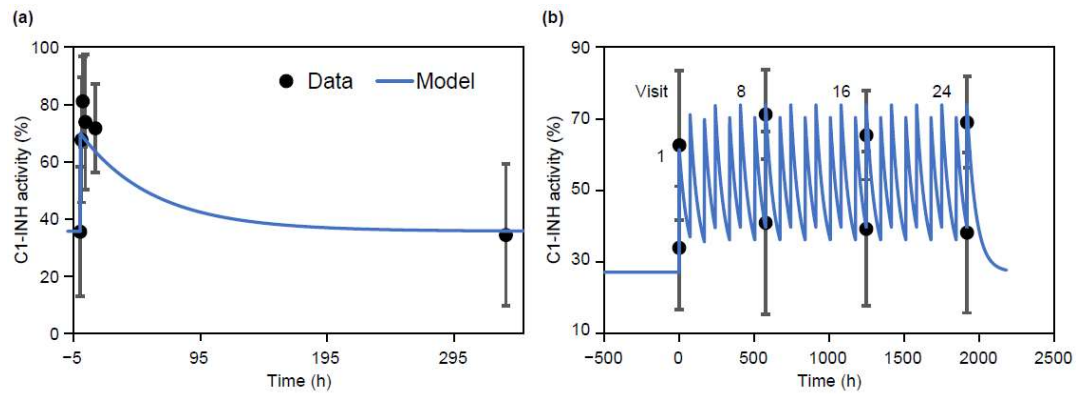
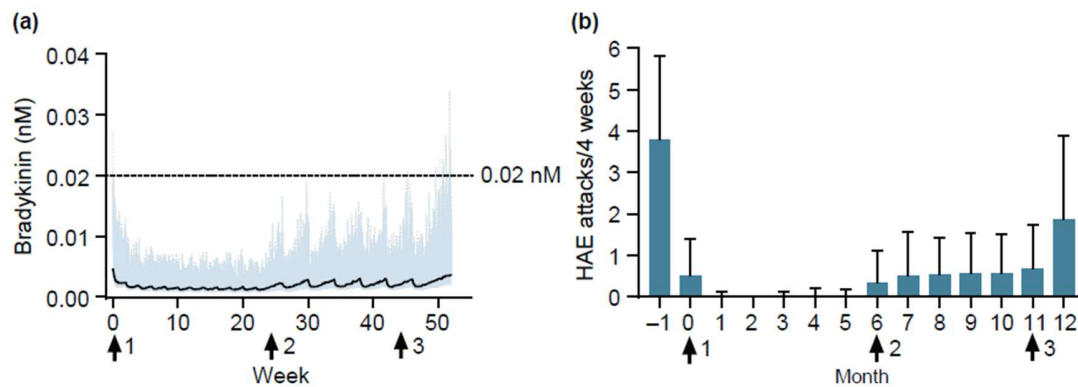


Fig. S7 Predicted impact of down-titration for patients with high baseline attack rate (6 attacks/month). (a) Bradykinin levels and (b) attack rate with lanadelumab Q2W dosing for 6 months followed by Q4W dosing for 6 months. Arrows indicate the following: (1) first 300 mg Q2W dose (week 0); (2) first 300 mg Q4W dose (week 24); and (3) last 300 mg Q4W dose (week 44). Bradykinin graph shows mean (black line) and 5% and 95% CI (shaded area). Dotted line shows bradykinin threshold (systemic concentration) for risk of attack (0.02 nM). Month -1 indicates baseline. *HAE* hereditary angioedema, *Q2W* every 2 weeks, *Q4W* every 4 weeks



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