Protocol S1

Impact of perturbation size on the prediction of fragile mechanisms. The impact of perturbation size upon the classification of mechanisms as robust or fragile was explored by constructing a family of parameters generated by randomly perturbing each nominal parameter by up to ± 2 -orders of magnitude. Overall State Sensitivity Coefficients (OSSC) were calculated for each mechanism in the model over the family of random parameter sets (N = 200). Comparison of OSSC values obtained for the small (up to $\pm 50\%$) versus large perturbation family (up to ± 2 -orders of magnitude) for the four treatment cases presented in the paper are shown in Fig. S1 (top). Several OSSC values are statistically different between the small and large perturbation families as determined by the Welch t-test, i.e., many OSSC values do not lie on the 45° line. However, the Spearman rank denoted by ρ and defined as:

$$\rho = 1 - \frac{6\sum_{i=1}^{P} d_i^2}{n\left(n^2 - 1\right)} \tag{1}$$

where d_i denotes the difference in the ordinal rank of parameter i between the small versus large perturbation cases and n denotes the number of pairs of values, shows an average of 96.6% agreement across treatment cases (see Fig. S1, bottom). Thus, while the numerical OSSC values are the not the same in the small versus large perturbation cases, the relative order of importance of the parameters is preserved. Therefore, the conclusions of this study would have been similar between small and large perturbations.



Figure S1: Sensitivity analysis results for large parameter perturbations. Top panel: Overall State Sensitivity Coefficients (OSSC) for the small versus large perturbation families for the different treatment cases considered. Bottom panel: Spearman rank correlation coefficients for the four treatment cases.

C and D \mathbf{F} cases А В Е fII 1700/2000 nM1700 nM $1400~\mathrm{nM}$ $1400~\mathrm{nM}$ $1400~\mathrm{nM}$ fVII $10 \ \mathrm{nM}$ FVIIa 0.5 nMTF $1 \ \mathrm{pM}$ TF-FVIIa 1.25 pM $5/10/50/500/5000~{\rm pM}$ 1.25 pM1.25 pM_ fV 20 nM20 nM20 nM20 nM20 nMfVIII 0.7 nM0.7 nM0.3 nM0.7 nM0.7 nMfIX 90 nM70 nM90 nM90 nM90 nMfΧ 135 nM170 nM170 nM 170 nM 170 nMTFPI 0/1/2.5/5 nM 0/2.5 nM2.5 nM3 nMATIII 0/3400 nM3000 nM _ \mathbf{PC} 65 nMTM0/1/10 nMPL (fM) 150150150150150PL binding sites on subendothelium (fM) 100 100100100 100fII/FIIa binding sites (nmol/fmol-active 505050200.5platelets) fV/FVa binding sites (nmol/fmol-active 5050502050platelets) 5×10^{-10} 5×10^{-10} 5×10^{-10} 5×10^{-10} 5×10^{-6} fVIII/FVIIIa binding sites (nmol/fmolactive platelets) fIX/FIXa binding sites (nmol/fmol-active $5 imes 10^{-8}$ 50505020platelets) Specific fIX/FIXa binding sites 50505020 5×10^{-8} (nmol/fmol-active platelets) fX/FXa binding sites (nmol/fmol-active 50502015050platelets)

Table S1: Initial conditions for the validation simulations. The initial concentration of surface binding sites

 were determined so that the validation simulations were consistent with experimental observations, all other

 parameters and initial conditions held constant.