

UNIVERSITY OF BIRMINGHAM

University of Birmingham Research at Birmingham

Modelling and simulation of flow and agglomeration in deep veins valves using discrete multi physics

Ariane, M; Wen, W; Vigolo, D; Brill, A; Nash, F G B; Barigou, M; Alexiadis, A

DOI:

10.1016/j.compbiomed.2017.07.020

License

Creative Commons: Attribution-NonCommercial-NoDerivs (CC BY-NC-ND)

Document Version
Peer reviewed version

Citation for published version (Harvard):

Ariane, M, Wen, W, Vigolo, D, Brill, A, Nash, FGB, Barigou, M & Alexiadis, A 2017, 'Modelling and simulation of flow and agglomeration in deep veins valves using discrete multi physics', *Computers in Biology and Medicine*, vol. 89, pp. 96-103. https://doi.org/10.1016/j.compbiomed.2017.07.020

Link to publication on Research at Birmingham portal

General rights

Unless a licence is specified above, all rights (including copyright and moral rights) in this document are retained by the authors and/or the copyright holders. The express permission of the copyright holder must be obtained for any use of this material other than for purposes permitted by law.

- Users may freely distribute the URL that is used to identify this publication.
- Users may download and/or print one copy of the publication from the University of Birmingham research portal for the purpose of private study or non-commercial research.
- User may use extracts from the document in line with the concept of 'fair dealing' under the Copyright, Designs and Patents Act 1988 (?)
- Users may not further distribute the material nor use it for the purposes of commercial gain.

Where a licence is displayed above, please note the terms and conditions of the licence govern your use of this document.

When citing, please reference the published version.

Take down policy

While the University of Birmingham exercises care and attention in making items available there are rare occasions when an item has been uploaded in error or has been deemed to be commercially or otherwise sensitive.

If you believe that this is the case for this document, please contact UBIRA@lists.bham.ac.uk providing details and we will remove access to the work immediately and investigate.

Accepted Manuscript

Modelling and simulation of flow and agglomeration in deep veins valves using discrete multi physics

M. Ariane, W. Wen, D. Vigolo, A. Brill, F.G.B. Nash, M. Barigou, A. Alexiadis

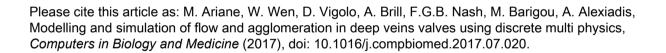
PII: S0010-4825(17)30244-5

DOI: 10.1016/j.compbiomed.2017.07.020

Reference: CBM 2731

To appear in: Computers in Biology and Medicine

Received Date: 8 May 2017
Revised Date: 10 July 2017
Accepted Date: 28 July 2017



This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.



- 1 Modelling and simulation of flow and agglomeration in deep veins valves using
- 2 Discrete Multi Physics.
- 3 M. Ariane^{a,*}, W. Wen^a, D. Vigolo^a, A. Brill^b, F. G.B Nash^b, M. Barigou^a, A.
- 4 Alexiadis^a,*
- 5 ^a School of Chemical Engineering, University of Birmingham, Birmingham, United
- 6 Kingdom
- 7 b Institute of Cardiovascular Sciences, University of Birmingham, Birmingham, United
- 8 Kingdom

9

10 **Abstract**

- 11 The hemodynamics in flexible deep veins valves is modelled by means of discrete 12 multi-physics and an agglomeration algorithm is implemented to account for blood accrual in the flow. Computer simulations of a number of valves typologies are carried 13 14 out. The results show that the rigidity and the length of the valve leaflets play a crucial 15 role on both mechanical stress and stagnation in the flow. Rigid and short membranes 16 may be inefficient in preventing blood reflux, but reduce the volume of stagnant blood potentially lowering the chances of thrombosis. Additionally, we also show that in 17 18 venous valves, cell agglomeration is driven by stagnation rather than mechanical stress.
- Keywords: Discrete Multi-Physics, Smoothed Particle Hydrodynamics, biological
 venous valve, Clot, Deep Venous Thrombosis.

1. Introduction

23	Deep venous thrombosis (DVT) is a dangerous and painful condition in which blood
24	thrombi form in deep veins. Such thrombi contain blood cells (including red blood cells
25	and platelets) within a mesh of coagulated protein which is predominantly fibrin. If one
26	of these aggregates detaches from the vein, it can reach the lungs resulting in a life-
27	threatening complication known as pulmonary embolism (PE). In the UK alone, DVT
28	and PE (designated together as venous thromboembolism, VTE) cause an estimated
29	25,000 deaths annually which exceeds the number of deaths from breast cancer, AIDS
30	and road traffic accidents combined (Hunt 2009).
31	One of the factors exacerbating DVT is prolonged immobility (e.g. bed ridden after
32	surgery, limb paralysis and long-haul flights), where the insufficient efficacy of the
33	muscle pump, which normally assists blood flow through the leg veins, leads to sluggish
34	flow. Stasis and low flow states are classically associated with a high probability of
35	thrombus formation (Reitsma et al., 2012). Because of this, it is likely that specific flow
36	patterns in veins, especially around the valve flaps, can play a fundamental role in the
37	formation of thrombi (Bovill and van der Vliet, 2011). Since hydrodynamics is affected
38	by the valve characteristics, the valve characteristics can also affect thrombus
39	agglomeration but the actual mechanisms remain unclear.
10	Moreover, once a person has developed DVT and has been successfully treated, he or
1 1	she is likely to develop other thrombus in the future, suggesting that the person's
12	specific valve geometry or flexibility may also contribute to the solid blood formation.

44	By providing hydrodynamic information of the blood flow around the valve, computer
45	simulations can improve our understanding of the link between fluid dynamics and
46	DVT. Computational blood dynamics has been widely and successfully used for cardiac
47	valves (e.g. De Hart et al., 2000; Fenlon and David, 2001; Van Loon et al., 2004;
48	Buxton and Clarke, 2006; Watton et al., 2007; Van Loon 2010; Astorino et al., 2012;
49	Espino et al., 2012; Bahraseman et al., 2014; Al-Azawy et al., 2015; Borazjani 2015;
50	Halevi et al., 2015; Kamensky et al., 2015; Marom 2015; Miandehi et al., 2015; Bavo et
51	al., 2016), but with a few exceptions so far little attention has been given to venous
52	valves (e.g. Wijeratne and Hoo 2008; Keijsers et al., 2015). In the majority of the
53	venous valve simulations, the valve is fixed and the complex interaction between the
54	flow and the moving leaflets is lost. Recently, flexible structures were investigated with
55	the Fluid Structure Interaction (FSI) method (Simao et al., 2016). Nevertheless, analyses
56	remain often limited to few cycles or implemented with one symmetrical leaflet. One of
57	the reasons is related to the intrinsic difficulty of the FSI to simulate the leaflets' contact
58	at the end of the closing phase. A few studies (e.g. Van Loon et al. 2010, Kamensky et
59	al. 2015) used additional contact algorithms to account for the mechanical contact of
60	leaflets. However, the implementation of these algorithms is complex and the frequency
61	of the re-meshing close to the contact point remains an issue.
62	In this work, we use the Discrete Multi-Physics (DMP) approach developed in
63	Alexiadis (2015) to model both the fluid dynamics and the flexible leaflets. This
64	approach was previously used for cardiac valves (Ariane et al., 2017). In this previous
65	paper, based on both dimensionless analysis and direct numerical simulations, we have
66	shown that size and rigidity of the leaflets, together with inlet velocity, are key
67	parameters and have determined which factors most affect the hydrodynamics around

68	the valve. For comparison, we focus on these parameters in this study, to specific flow
69	patterns and stress profiles in a venous valve system using the DMP approach.
70	Other advantages of the approach proposed here is that, contrary to FSI simulation, it
71	can account for a complete valve closure without the use of a stabilisation algorithm.
72	Finally, using DMP gives the possibility of introducing an agglomeration algorithm that
73	transforms a portion of the liquid into a solid. Other studies (e.g. Simao et al., 2016)
74	simulated cell aggregates by tweaking the viscosity of the liquid, but with the method
75	proposed here we can form actual solid structures within the liquid phase. By
76	introducing the agglomeration algorithm, we identify among the regions where thrombi
77	are most likely to form and which have the highest growth probability.
78	

79 **2. Modelling**

80

2.1. Modelling approach

81 The DMP modelling technique used in this work is based on the so-called discrete 82 multi-hybrid system (DMHS). In the DMHS model, the liquid is represented by 83 Smoothed Particle Hydrodynamics (SPH) particles (Monaghan 1994; Morris et al., 84 1997; Liu and Liu 2003), while the solid structure is divided into many notional 85 particles linked by computational springs (to model the elastic modulus of the solid), 86 computational hinges (to model the flexural modulus) and computational dashpots (to 87 model the viscous material behaviour). Mathematically, this is similar to the treatment 88 of molecular bonds used in Molecular Dynamic (MD) simulations. In the original paper 89 where the DMHS was first proposed (Alexiadis 2014), this part of the model was 90 referred to as Coarse Grained Molecular Dynamics (CGMD) to highlight its MD origin. 91 Here we prefer the term Mass-Spring Model (MSM) as the scales involved are 92 macroscopic. Readers can refer to Appendix A for details and to (Ariane et al., 2017) for application of this methodology to biological valves. 93

2.2. Geometry

94

In this study, we use a 2D schematic representation of the leg venous valve (Wijeratne and Hoo 2008) as illustrated in Fig. 1. The channel radius is Z = 0.004 m, the radius of the valve chamber is R = 0.007 m and its length is Y = 0.04 m. Three different lengths L of the membrane are studied: long (0.0256 m), medium (0.0175 m) and short (0.01 m). In order to distinguish different parts of the geometry, we refer to the region between

100	the two leaflets as the 'opening region' and to the two regions between the wall and the
101	leaflet as 'sinus regions'. Fig. 1 shows the location of the opening region and one of the
102	two sinus regions (the other is symmetric and located above the upper leaflet).

Fig. 1. Illustration of the venous valve 2D geometry and particle representation.

The leaflets are represented by (solid) MSM particles joined together by springs and hinges (fig. 1) as discussed in Appendix A. SPH particles are used for the fluid and stationary (solid) particles for the walls. Three layers of particles are used for the channel and two for each leaflet.

There are two types of parameters required for the simulations: model parameters and simulation parameters. The first group consists of internal parameters used by the SPH and MSM solvers (Table 2); the second refers to the operative conditions detailed

2.3 Simulation conditions

103

111

112

below.

The Young's modulus E and the flexural modulus F of the membrane are the results of 113 114 the MSM particles joined together by numerical springs and hinges. The relation 115 between the spring (k_b) and hinge (k_a) constants and the actual Young's modulus and 116 the flexural modulus is given in Ariane et al. (2017). A viscous coefficient is added to 117 the MSM springs to confer viscoelastic properties to the membrane as in a Kelvin-118 Voigt material. 119 Periodic boundary conditions are used at the inlet/outlet and we implement the same 120 pulsatile flow (purely oscillatory) used by (Wijeratne and Hoo 2008) and imposing to 121 each liquid particle the acceleration g as shown in equation 1

$$g = g_0 \sin(2\pi f t), \qquad (1)$$

123

124

125

126

127

128

129

130

131

132

133

134

135

136

137

138

139

140

141

142

143

144

with amplitude g_{θ} (given in Table 1), time t and oscillation frequency f = 1/T (with T the period oscillation). We use equation 1 as a simple means of forcing alternating flow in the valve, but the real oscillation is not sinusoidal and the frequency is not constant. We chose T = 4 s, which is long enough to ensure full closure. Here we limit the total time of calculation with 4 full cycles (opening and closing) which correspond to 16 s. Previous work (Wijeratne and Hoo 2008) used T = 3 s and simulated only one full cycle. When the muscles contract, the blood within the veins is compressed and the valve opens; when the muscles dilate, the valve closes preventing backward flow. The blood velocity depends on the force of the muscle contraction and, in general, it is related to the level of physical activity of a specific person. In order to account for three levels of physical activity, we take into account three values of g_0 (0.1 m s⁻², 0.25 m s⁻² and 0.4 m s⁻²), which result in three different flows with maximum velocities in the inlet channel of 0.03 m s⁻¹ (low physical activity), 0.07 m s⁻¹ (intermediate case), 0.13 m s⁻¹ (high level of physical activity) respectively. The low velocity is from (Simao et al., 2016), the intermediate velocity is from (Wijeratne and Hoo 2008) and we include the third highest velocity to account for high levels of physical activity. In all cases, the flow is laminar. The length and the flexibility of the membrane vary from person to person (Mühlberger et al., 2008; Moore et al., 2011). In order to investigate a variety of individual variations, we consider three membrane lengths (0.0256 m, 0.0175 m, and 0.01 m). The longest length is from (Wijeratne and Hoo 2008) and the shortest length is chosen as the minimum size allowing a complete closure of the leaflets. Regarding the flexibility and

the stiffness, in our previous paper (Ariane et al. 2017), the literature review for the aortic valve has shown that the membrane has three dynamic regimes based on the membrane stiffness (Bavo et al. 2016; Ledesma et al. 2014; De Hart et Al. 2000; Van Loon et al. 2006). In the simulation, we vary the stiffness of the valve according to these regimes (see Table 1).

Table 1. List of simulations with fluid velocities and membrane parameters.

Variation of the	e membrane le	ength and the velocity with $k_a = 0.01J$
Length of the membrane L [m]	V [m s ⁻¹]	Designation
Short	0.03	L0.01/V0.03/k _a 0.01
L= 0.01 m	0.07	L0.01/V0.07/k _a 0.01
	0.13	L0.01/V0.13/k _a 0.01.
Medium L= 0.0175 m	0.03	L0.0175/V0.03/k _a 0.01
	0.07	L0.0175/V0.07/k _a 0.01
	0.13	L0.0175/V0.13/k _a 0.01.
Long L= 0.0256 m	0.03	L0.0256/V0.03/k _a 0.01
	0.07	L0.0256/V0.07/k _a 0.01
	0.13	L0.0256/V0.13/k _a 0.01
Variation of the me	embrane flexib	pility with $L = 0.0256$ m and $V = 0.07$ m s ⁻¹
Angular coeffici	ent k _a [J]	Designation
0.0001		L0.0256/V0.07/k _a 0.0001
0.002		L0.0256/V0.07/k _a 0.002
0.005		L0.0256/V0.07/k _a 0.005
0.02		L0.0256/V0.07/k _a 0.02
0.05		L0.0256/V0.07/k _a 0.05

2.4 Agglomeration algorithm

In order to understand the agglomeration dynamics, we introduce in the model the
agglomeration algorithm developed in Ariane et al. (2017) (a brief overview of the
method can be found in Appendix B). At this stage, our focus is to understand if
hydrodynamics alone favours agglomeration at different locations. The actual
biochemical process of thrombus formation is an extremely complex phenomenon (e.g.
Panteleev et al., 2015) and it is beyond the scope of this article.
Specific particle points are used as agglomeration seeds. The algorithm every N time-
steps checks all the fluid particles at a distance $R_{\rm MAX}$ from the seeds and, with a certain
probability P , transforms some of these particles into solid agglomerate-particles.
In the simulations, the values of N , R_{MAX} and P are given in Table 2. These values do
not correspond to the real time-scale of agglomeration but were chosen in order to
accelerate agglomeration and to observe significant growth in few cycles. Since our
goal is to determine where agglomeration is more likely, this 'accrual acceleration' does
not affect the validity of the results, as long as the timescale of agglomeration is longer
than the timescale of the flow.

Table 2. Model parameters used in the simulations.

SPH (eq.s A.5–A.7)	
Parameter	Value
Number of SPH wall particles (3 layers)	5360
Number of SPH valve particles (2 layers)	(1) 1026, (2) 702, (3) 402
Number of SPH fluid particles	(1) 89592, (2) 89726, (3) 90012
Mass of each particle (fluid)	1.05·10 ⁻⁵ kg
Mass of each particle (wall and valve)	2·10 ⁻⁵ kg
Initial distance among particles Δr	1·10 ⁻⁴ m
Smoothing length h	2.5·10 ⁻⁴ m
Artificial sound speed c_0	10 m s ⁻¹
Density ρ_0	1056 kg m ⁻³
Time step Δt	10 ⁻⁷ s
CGMD (eq.s A.10–A.11)	42
Parameter	Value
Angular coefficient k_a	See Table 1
Hookian coefficient k_b	$1 \cdot 10^6 \text{ J m}^{-2}$
Viscous damping coefficient k_{ν}	0.01 kg s ⁻¹
Equilibrium distance r_0	1·10 ⁻⁴ m
Equilibrium angle θ_0	π/2 rad
BOUNDARIES (eq. A.15)	· ·
Constant K	$4.10^{-4} \mathrm{J}$
Repulsive radius r*	1·10 ⁴ m
SOLID FORMATION (Section Formation of	solid aggregates)
Number of time step for solid formation N	$0.5 \cdot 10^6 \text{ s}$
R _{max}	2.5·10 ⁻⁴ m
Agglomeration probability P	50 %
Max bonds per solid particle	4
(a) long, (b) medium, (c) short membrane	

170

3. Results and discussion

172

173

3.1 Stress and residence time

174	According to (Simao et Al., 2016), causes of DVT among young people remain
175	unknown in most of half of the cases. When the origin of DVT is known, thrombus
176	initiation is often associated with blood coagulability, changes in the vessel wall or
177	immobility (Esmon 2009). In the case of immobility, low velocity and high residence
178	time are the most causes of blood aggregation (Menichini et Al. 2016, Bovill et Al.
179	2011). Although, shear stress can also play a role in platelet aggregation and activation.
180	In fact, components of the coagulation cascade and platelets can be activated in all shear
181	stresses, just mechanisms will be different. For instance, at low shear stress, platelets
182	adhere to fibrinogen, whereas at high shear stress to von Willebrand factor (Ikeda et al.,
183	1991). Likewise, abnormal shear stress distribution can initiate and accelerate the
184	formation of thrombi (Hou et Al., 2015). In the arterial setting, high stress could be an
185	activator of platelet aggregation (Zhang et al., 2002). However, arterial and venous
186	thrombi are structurally different. In arteries, high shear may induce platelet activation
187	and formation of what is sometimes called white thrombi with few red cells in it. In
188	veins, the thrombus is red with many red cells trapped in coagulated proteins. In this
189	case, coagulation seems to be the dominant process, suggesting a slower, time
190	dependant, accrual.
191	The following discussion focuses on both mechanical stress and residence time, to
192	account for the two factors that are most generally related to thrombus formation
193	(Zhang et al., 2002).

194	3.1.1 Mechanical stress
195	Our DVT calculations show that shear stress is high only in the opening region and
196	almost negligible everywhere else (Fig. 2a). Clinical experience, however, indicates that
197	thrombi do not form in the opening region, where shear stress is high, but rather in the
198	sinus region (Bovill and van der Vliet 2011) where it is at its lowest (Ju et al., 2016).
199	These observations suggest that total mechanical stress (Ttot), rather than shear stress
200	(Tshear) should be investigated and our results show that, in this case, Ttot is high or
201	both sides of the membrane (Fig. 2b).
202	In a fluid, the total mechanical stress (or total mechanical force) is the sum of shear
203	stress (or viscous forces), inlet pressure (or pressure force, Ptot) and gravity (or Body
204	force, ignored here). Fig. 2c shows the pressure profile and indicates that the higher
205	total mechanical stress on the membrane cannot be justified by pressure alone. When a
206	solid body moves in the fluid (acceleration or deceleration), it generates additional
207	forces (virtual mass force) that simultaneously move the volume of the surrounding
208	fluid. These inertial forces so-called added forces explain the higher Ttot on the
209	membrane in the sinus region.
210	Fig. 2. Shear stress (a), total mechanical stress (b), pressure (c), and velocity magnitude
211	(d) for L0.0256/V0.03/k _a 0.01
212	3.1.2. Residence time in the sinus
213	In Lagrangian approach, displacement is used as a proxy for residence time and the
214	following discussion is based on this parameter instead of residence time.

215	As shown in Fig. 2d, the velocity in the sinus region is low compared to that of the
216	opening region; as a consequence, the residence time of fluid particles in this region is
217	higher. Fig. 3 illustrates this point. We highlight the particles initially in the sinus in
218	blue and we track their position during the simulation. At the end of four cycles, a
219	fraction of the particles has left the sinus, while the rest remains confined in this region.
220	In Fig. 3, the particles are coloured according to their displacement, defined as the
221	distance travelled by each particle during the simulation. Blue particles do not move
222	very much and are substantially stagnant; red particles have higher velocity and show
223	higher displacement.
224	Fig. 3. Simulation snapshots illustrating the fluid motion of the particles initially in the
225	sinus at different times (beginning of each new cycle): for L0.0256/V0.03/ k_a 0.01;
226	particles coloured according to their displacement.
227	In the sinus, we can identify two areas of high fluid displacement. The first (called
228	'mixing region' in Fig. 3) corresponds to the recirculation region created by the
229	backflow (Fig. 4). The second (called 'compression region' in Fig. 3) is below the
230	membrane and corresponds to the part of the fluid displaced by the oscillating
231	movement of the membrane. There is a fundamental difference between the two
232	regions. While in the mixing region the fluid particles are actually moved out of the
233	sinus by the backflow, in the compression region the particle only oscillates around the
234	same point due to the alternate motion of the membrane.
235	Fig. 4. Velocity profile in the sinus area (vectors) for L0.0256/V0.03/k _a 0.01.
236	Despite the fact that both regions show high displacement, the actual residence time is
237	high only in the compression region.

238	Displacement alone, therefore, is not enough to distinguish between regions of low and
239	high residence time. In order to account for this, in the next section, we introduce the
240	time-averaged displacement as a more accurate proxy for the residence time.

3.2. Parametric study

In this section, we investigate how Ttot and displacement are affected by (i) membrane flexibility, (ii) leaflet length and (iii) level of physical activity (fluid velocity). According to our simulations, all of these three parameters are particularly significant for the performance of the valve. Comparing the effect of these parameters among different setups, however, is not straightforward because it changes in space and time. To compare results with respect to the same reference point, we identify the fluid particle in the sinus region with the highest mechanical stress and for every setup we measure stress and displacement at this location. In this way, we carry out all our measurements at the same relative position. However, as indicated in Fig. 5a and 5b, displacement and Ttot also change with time. To account for this, in the case of displacement (Fig. 5a), we use the time-average instead of the instantaneous displacement. In the case of Ttot (Fig. 5b), we use the maximal rather than the average stress because agglomeration is more affected by the peak of the stress rather than its average.

Fig. 5. Time evolution of the local fluid displacement (a) and total mechanical stress magnitude (b) for the particle of maximal stress (L0.0256/V0.03/k_a0.01).

- We can also quantify how both these parameters oscillate with time by calculating their standard deviation; in the subsequent Fig.s, the error bars indicate the standard deviation.

 3.2.1. Effect of membrane flexibility

 The flexibility of the membrane depends on its flexural modulus. In Ariane et al. (2017)
- we showed that the flexural modulus is mostly affected by the k_a , for this reason, in this section we focus on how time-averaged displacement and maximal stress vary with this parameter.

 In Fig. 6, both average displacement and Ttot decrease as the membrane flexibility
- 267 increases to a value $k_a = 0.02$ J because mechanical deformation is lower for rigid membranes. However, the displacement for very rigid membranes ($k_a = 0.05 \text{ J}$) 268 increases. The reason for this can be understood by comparing Fig. 7a and 7b: at k_a = 269 270 0.05 J, the leaflets maintain a straight profile during the closure phase (Fig. 7a), while at $k_a = 0.02$ J they bend under the flow (Fig. 7b). When the leaflets bend, they partially 271 272 shield the sinus region from the backflow and reduce the velocity (and therefore the 273 displacement). Conversely, very flexible membranes ($k_a < 0.005$ J) highly deform and 274 fluctuate under the flow (Fig. 7c). This explains the higher standard deviation in Fig. 6 and the irregular profile of Fig. 6b for $k_a < 0.005$ J. 275
- Fig. 6. Time-averaged displacement (a) and total mechanical stress (b) versus k_a (valve flexibility) for cases: L = 0.0256 m, V = 0.07 m s⁻¹ and k_a from 0.0001J to 0.05J.
- Fig. 7. Simulation snapshots illustrating the fluid motion of the particles initially in the sinus for long valve, $V = 0.07 \text{ m s}^{-1}$ and three flexibilities: (a) $k_a = 0.05 \text{ J}$, (b) $k_a = 0.02 \text{ J}$, and (c) $k_a = 0.0001 \text{ J}$, particles coloured according to their displacement.

281	3.2.2. Effect of membrane length and inlet velocity
282	Fig. 8 shows the effect of the inlet velocity on the displacement and stress for three
283	membrane sizes. For medium or long membranes, as expected, higher velocities are
284	associated with higher stress and displacement. The short membrane, however, behaves
285	differently.
286	Fig. 8. Evolution of displacement (a) and total mechanical stress magnitude (b) with
287	the maximum inlet velocity.
288	Contrary to the medium and long membrane (see Fig. 2), in short membranes, the
289	highest stress (see Fig. 9) is located at the tip rather than the middle of the valve. At the
290	tip, the motion of the particles depends on the hydrodynamics at the opening region
291	rather than that at the sinus region and, therefore, they are easily transported away by
292	the flow and the displacement increases significantly.
293	Fig. 9. Total mechanical stress (a), velocity magnitude (b), vector velocity (c), and
294	displacement (d) in the short valve case for L0.01/V0.07/ $k_{\rm a}0.01$.
295	3.3. Agglomeration
296	The main physical parameters that affect agglomeration are the residence time and
297	mechanical stress. The simulations highlight two key locations: one at the sinus side of
298	the membrane, where stress is the highest (point P_1 in Fig. 10), and the other at the
299	valve/wall connection where the residence time is the highest (point P ₂ in Fig. 10).
300	The higher mechanical stress at P ₁ pushes particles closer, increasing the number of
301	particles inside R_{MAX} ; however, because the velocity is higher, these particles remain

inside $R_{\rm MAX}$ only for a short time. At P_2 , the opposite happens: the mechanical stress is lower, but also, because the velocity is low (and, therefore, the residence time is high) and the particles remain inside $R_{\rm MAX}$ for longer.

High stresses and high velocities, therefore, have opposite effects on agglomeration. Fig. 10 shows faster growth at P_2 , suggesting that residence time may be more

Fig. 10. Solid aggregates in the sinus region at two different times for $L0.0175/V0.07/k_a0.01$.

important for aggregation propagation in the venous valve than mechanical stress.

Conclusions

This article presents a discrete multi-physics model for both blood dynamics and leaflets mechanics of a leg venous valve. In the simulations, we focused on mechanical stress and flow stagnation (high residence time) in the sinus region because these two factors have been linked to the onset of blood solid formation. The model is subsequently coupled with an agglomeration algorithm to account for the formation and propagation of solid aggregates in the flow.

The results show that the flexibility and the length of the membrane play a crucial role in both stress and flow stagnation. Rigid membranes do not close completely and, therefore, they may be inefficient in preventing blood reflux. However, they also allow for a larger flow exchange between the sinus region and the central flow reducing stagnation and, potentially, lowering the chances of thrombosis. Similarly, short membranes reduce the volume of the sinus region, which also decreases stagnation.

323	We also focused on the issue in venous valves and whether it is mechanical stress or
324	stagnation that favours cell agglomeration which may lead to thrombosis.
325	In order to compare the role of these two factors, we identified the location in the sinus
326	with the highest stress and that with the highest stagnation. We placed an agglomeration
327	seed in each of these two locations and implemented our agglomeration algorithm.
328	The growth of the agglomerate at the point of maximum stagnation was considerably
329	higher than that at the point of maximal stress. This implies that, in the case of the
330	venous valve, stagnation can be more important than mechanical stress in thrombus
331	formation and propagation.
332	This result, combined with the fact that membrane flexibility and length determine the
333	level of stagnation in the sinus, highlights the potential for personalised diagnostics in
334	the fight against deep venous thrombosis. In principle, length and stiffness could be
335	evaluated in clinical setting using existing diagnostic methods. Currently, they are not
336	evaluated, but based on our results, if they were added, in the future, to the toolkit of
337	physicians they could, potentially, help predicting the likelihood of DVT.
338	These data, in fact could be introduced into our discrete multi physics model to predict,
339	for that particular valve, the location of maximum stagnation and provide information
340	that, potentially, could be converted into a probability of thrombus formation for a
341	specific individual.
342	Acknowledgements
343	This work was supported by the Engineering and Physical Sciences Research Council

344

(EPSRC) grant number: EP/N033698/1.

Supporting Information

345

362

363

346	A Appendix
347	B Appendix
348	References
349	Al-Azawy, M. G., A. Turan and A. Revell (2015). "Assessment of turbulence models
350	for pulsatile flow inside a heart pump." Computer Methods in Biomechanics and
351	Biomedical Engineering: 1-15.
352	Alexiadis, A. (2014). "A smoothed particle hydrodynamics and coarse-grained
353	molecular dynamics hybrid technique for modelling elastic particles and breakable
354	capsules under various flow conditions." International Journal for Numerical
355	Methods in Engineering 100 (10): 713-719.
356	Alexiadis, A. (2015). "The Discrete Multi-Hybrid System for the Simulation of Solid-
357	Liquid Flows." Plos One 10(5).
358	Ariane, M., M. H. Allouche, M. Bussone, F. Giacosa, F. Bernard, M. Barigou and A.
359	Alexiadis (2017). "Discrete multi-physics: A mesh-free model of blood flow in
360	flexible biological valve including solid aggregate formation." Plos One 12(4).
361	Astorino, M., J. Hamers, S. C. Shadden and JF. Gerbeau (2012). "A robust and

364 Bahraseman, H. G., K. Hassani, A. Khosravi, M. Navidbakhsh, D. M. Espino, N.

Numerical Methods in Biomedical Engineering **28**(9): 937-959.

efficient valve model based on resistive immersed surfaces." International Journal for

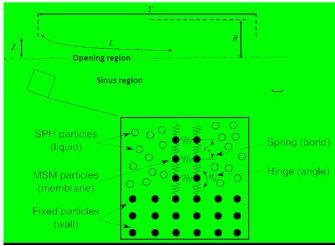
- Fatouraee and D. Kazemi-Saleh (2014). "Combining numerical and clinical methods
- to assess a ortic valve hemodynamics during exercise." Perfusion-Uk **29**(4): 340-350.

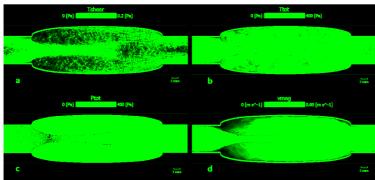
- Bavo, A. M., G. Rocatello, F. Iannaccone, J. Degroote, J. Vierendeels and P. Segers
- 368 (2016). "Fluid-Structure Interaction Simulation of Prosthetic Aortic Valves:
- 369 Comparison between Immersed Boundary and Arbitrary Lagrangian-Eulerian
- Techniques for the Mesh Representation." Plos One **11**(4).
- 371 Borazjani, I. (2015). "A Review of Fluid-Structure Interaction Simulations of Prosthetic
- 372 Heart Valves." **25**(1-2): 75-93.
- 373 Bovill, E. G. and A. van der Vliet (2011). Venous Valvular Stasis-Associated Hypoxia
- and Thrombosis: What Is the Link? Annual Review of Physiology, Vol 73. D. Julius
- and D. E. Clapham. **73**: 527-545.
- 376 Buxton, G. A. and N. Clarke (2006). "Computational phlebology: The simulation of a
- vein valve." Journal of Biological Physics **32**(6): 507-521.
- De Hart, J., G. W. M. Peters, P. J. G. Schreurs and F. P. T. Baaijens (2000). "A two-
- dimensional fluid-structure interaction model of the aortic value." Journal of
- 380 Biomechanics **33**(9): 1079-1088.
- 381 Esmon, C. T. (2009). "Basic mechanisms and pathogenesis of venous thrombosis.".
- 382 Blood Reviews 23(5): 225-229.
- Espino, D. M., D. E. T. Shepherd and D. W. L. Hukins (2012). "Evaluation of a
- transient, simultaneous, arbitrary Lagrange-Euler based multi-physics method for
- simulating the mitral heart valve." Computer Methods in Biomechanics and
- 386 Biomedical Engineering **17**(4): 450-458.
- Fenlon, A. J. and T. David (2001). "Numerical models for the simulation of flexible
- artificial heart valves: part I--computational methods." Computer methods in
- biomechanics and biomedical engineering **4**(4): 323-339.

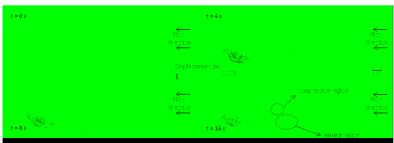
- 390 Halevi, R., A. Hamdan, G. Marom, M. Mega, E. Raanani and R. Haj-Ali (2015).
- 391 "Progressive aortic valve calcification: Three-dimensional visualization and
- biomechanical analysis." Journal of Biomechanics **48**(3): 489-497.
- 393 Hou, X. Y., Sun, X., Shi, Y. T., Zhang, K. L. and Yao, J. T. (2015). "Simulation of the
- formation mechanism of coronary thrombosis based on dem-cfd coupling." 2015 8th
- 395 International Conference on Biomedical Engineering and Informatics (Bmei): 24-28.
- 396 Hunt, B. J. (2009). "The prevention of hospital-acquired venous thromboembolism in
- the United Kingdom." British Journal of Haematology **144**(5): 642-652.
- 398 Ikeda, Y., Handa, Kawano, M., Kamata, T., Murata, M., Araki, Y., Anbo, H., Kawai,
- 399 Y., Watanabe, K., Itagaki, I., Sakai, K., and Ruggeri, Z. M. (1991). "The role of
- 400 vonwillebrand-factor and fibrinogen in platelet-aggregation under varying shear-
- stress." Journal of Clinical Investigation 87(4): 1234-1240.
- Ju, L. N., Y. F. Chen, L. Z. Xue, X. P. Du and C. Zhu (2016). "Cooperative unfolding of
- distinctive mechanoreceptor domains transduces force into signals." eLife 5.
- Kamensky, D., M.-C. Hsu, D. Schillinger, J. A. Evans, A. Aggarwal, Y. Bazilevs, M. S.
- Sacks and T. J. R. Hughes (2015). "An immersogeometric variational framework for
- fluid-structure interaction: Application to bioprosthetic heart valves." Computer
- 407 Methods in Applied Mechanics and Engineering **284**: 1005-1053.
- 408 Keijsers, J. M. T., C. A. D. Leguy, W. Huberts, A. J. Narracott, J. Rittweger and F. N.
- van de Vosse (2015). "A 1D pulse wave propagation model of the hemodynamics of
- 410 calf muscle pump function." International Journal for Numerical Methods in
- 411 Biomedical Engineering **31**(7).
- 412 Liu, G. R. and M. B. Liu, Eds. (2003). Smoothed Particle Hyrodynamics: a meshfree
- 413 method. Singapore, World Scientific Publishing Co. Pte. Ltd.

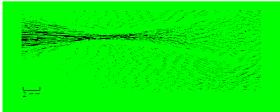
- 414 Marom, G. (2015). "Numerical Methods for Fluid-Structure Interaction Models of
- Aortic Valves." Archives of Computational Methods in Engineering **22**(4): 595-620.
- 416 Menichini, C., Cheng, Z., Gibbs, R. G. J. and Xu, X. Y. (2016). "Predicting false lumen
- 417 thrombosis in patient-specific models of aortic dissection." Journal of the Royal
- 418 Society Interface 13(124).
- 419 Miandehi, E. E., M. H. Aazami, H. Niazmand, Y. Mesri, A. Deyranlou and S. Eslami
- 420 (2015). "Clinical simulation of aortic valve: a narrative review." Studies in health
- technology and informatics **210**: 612-616.
- 422 Monaghan, J. J. (1994). "Simulating Free Surface Flows with SPH." Journal of
- 423 Computational Physics **110**(2): 399-406.
- 424 Moore, H. M., M. Gohel and A. H. Davies (2011). "Number and location of venous
- valves within the popliteal and femoral veins a review of the literature." Journal of
- 426 Anatomy **219**(4): 439-443.
- 427 Morris, J. P., P. J. Fox and Y. Zhu (1997). "Modeling Low Reynolds Number
- Incompressible Flows Using SPH." Journal of Computational Physics 136(1): 214-
- 429 226.
- 430 Mühlberger, D., L. Morandini and E. Brenner (2008). "An anatomical study of femoral
- vein valves near the saphenofemoral junction." Journal of Vascular Surgery **48**(4):
- 432 994-999.
- Panteleev, M. A., N. M. Dashkevich and F. I. Ataullakhanov (2015). "Hemostasis and
- 434 thrombosis beyond biochemistry: roles of geometry, flow and diffusion."
- 435 Thrombosis Research **136**(4): 699-711.

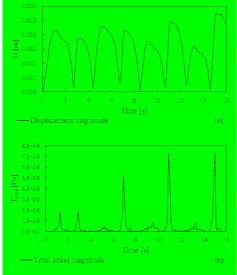
436 Reitsma, P. H., H. Wersteeg and S. Middeldorp (2012). "Mechanistic View of Risk 437 Factors for Venous Thromboembolism." Arteriosclerosis Thrombosis and Vascular 438 Biology **32**(3): 563-568. 439 Simao, M., J. M. Ferreira, J. Mora-Rodriguez and H. M. Ramos (2016). "Identification 440 of DVT diseases using numerical simulations." Medical & Biological Engineering & 441 Computing **54**(10): 1591-1609. 442 van Loon, R. (2010). "Towards computational modelling of aortic stenosis." 443 International Journal for Numerical Methods in Biomedical Engineering 26(3-4): 444 405-420. van Loon, R., P. D. Anderson, J. de Hart and F. P. T. Baaijens (2004). "A combined 445 446 fictitious domain/adaptive meshing method for fluid-structure interaction in heart valves." International Journal for Numerical Methods in Fluids 46(5): 533-544. 447 448 Watton, P. N., X. Y. Luo, X. Wang, G. M. Bernacca, P. Molloy and D. J. Wheatley 449 (2007). "Dynamic modelling of prosthetic chorded mitral valves using the immersed 450 boundary method." Journal of Biomechanics 40(3): 613-626. 451 Wijeratne, N. S. and K. A. Hoo (2008). Numerical studies on the hemodynamics in the 452 human vein and venous valve. 2008 American Control Conference, Vols 1-12. New 453 York, Ieee: 147-152. Zhang, J.-n., A. L. Bergeron, Q. Yu, C. Sun, L. V. McIntire, J. A. López and J.-f. Dong 454 (2002). "Platelet Aggregation and Activation under Complex Patterns of Shear 455 456 Stress." Thrombosis and Haemostasis 88(11): 817-821. 457

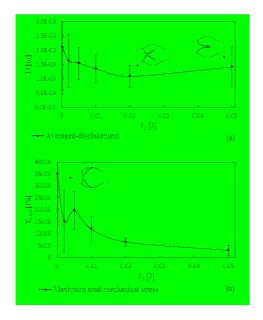


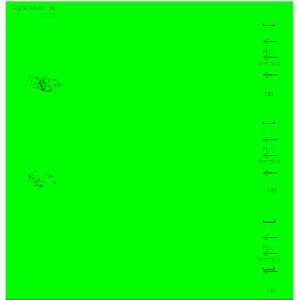


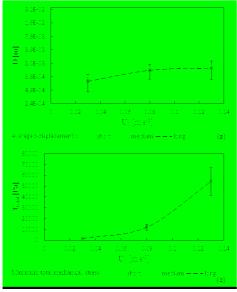


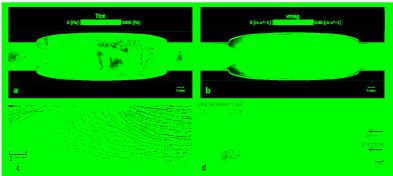














Highlights:

- Development of a discrete multi-physics model for both the blood dynamics and the leaflets mechanics in a leg venous valve.
- The model accounts for the hydrodynamics, the valve deformation with contact closure, and the solid aggregation at the same time.
- The key role of the flexibility and the length of the valve in both stress and flow stagnation are investigated.
- In venous valve, stagnation can be more important than stress in thrombus formation and propagation.